

bcb	Procedure codes related to the CMS Inpatient Only List	have been removed, however the inpatient admission care requires an authorization.
	Acupuncture	
Code	Code Description	Comments
20560	Needle insertion without injection; 1 or 2 muscle	
20561	Needle insertion without injection; 3 or more muscles	
97810	Acupuncture, one or more needles, without electrical stimulation, initial 15 min	
97811	Acupuncture, 1 or more needles; without electrical stimulation, each additional 15 minutes of personal one-on- one contact with patient, with insertion of needle	
97813	Acupuncture, one or more needles, with electrical stimulation, initial 15 min	
97814	Each additional 15 minutes of personal one-on-one contact with the patient, with insertion of needle	
97014	Lach additional 13 minutes of personal one-on-one contact with the patient, with insertion of needle	
	Behavioral Health	
Code	Code Description	Comments
G0410	Group psychotherapy other than of a multiple-family group, in a partial hospitalization setting, approx 45 to 50 mins	
	Cardiac Echo	
Code	Code Description	Comments
93321	Doppler echocardiography, pulsed wave &/or continuous wave w/spectral display; follow-up or limited study	
93350	Echocardiography, transthoracic real time, rest & stress; interpret & report	
93352	Use of echocardiographic contrast agent during stress echocardiography	
C8928	TEE w or wo contrast foll by w contrast real-time w image doc (2D) w or wo M-mode rec during rest and c/v stress test	
	Diagnostic Testing	
Code	Code Description	Comments
20560	Needle insertion without injection; 1 or 2 muscle	
20561	Needle insertion without injection; 3 or more muscles	
92517	Vestibular evoked myogenic potential (VEMP) testing, with interpretation and report; cervical (cVEMP)	
92519	Vestibular evoked myogenic potential (VEMP) testing, with interpretation and report; cervical (cVEMP) and ocular (oVEMP)	
93321	Doppler echocardiography, pulsed wave &/or continuous wave w/spectral display; follow-up or limited study	
93350	Echocardiography, transthoracic real time, rest & stress; interpret & report	
93351	Echocardiography, transthoracic, real-time w/image documentation (2D), includes M-mode recording, when performed, during rest & cardiovascular stress test using treadmill, bic	
93352	Use of echocardiographic contrast agent during stress echocardiography	
97810	Acupuncture, one or more needles, without electrical stimulation, initial 15 min	
97811	Acupuncture, 1 or more needles; without electrical stimulation, each additional 15 minutes of personal one-on- one contact with patient, with insertion of needle	
97813	Acupuncture, one or more needles, with electrical stimulation, initial 15 min	
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Updated: 07/01/2025

Chattanooga, Tennessee 37402
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97814	Acupuncture, 1 or more needles; with electrical stimulation, each additional 15 minutes of personal one-on-one		
	contact with patient, with insertion of needle		
C8928	TEE w or wo contrast foll by w contrast real-time w image doc (2D) w or wo M-mode rec during rest and c/v stress		
C6926	test		

Durable Medical Equipment				
Code Description		Comments		
4.4000	Supply allowance for non-adjunctive, non-implanted continuous glucose monitor, includes all supplies &			
A4239	accessories, 1 month supply = 1 unit of service			
A6023	A6023 Collagen Dressing, sterile, size more than 48 sq. in., each			
E0118	Crutch, substitute, lower leg platform, with or without wheels, each			

A6023	Collagen Dressing, sterile, size more than 48 sq. in., each			
E0118	Crutch, substitute, lower leg platform, with or without wheels, each			
E0144	PRICE MOD REQ Walker, enclosed, 4 sided framed, rigid or folding, wheeled with posterior seat			
E0147	PRICE MOD REQ Walker, heavy duty, multiple braking system, variable wheel resistance			
E0149	PRICE MOD REQ Walker, heavy duty, wheeled, rigid or folding, any type			
E0158	PRICE MOD REQ Leg extensions for walker, per set of 4			
E0159	PRICE MOD REQ Brake attachment for wheeled walker, replacement, each			
E0165	PRICE MOD REQ Commode chair, mobile or stationary, with detachable arms			
E0181	PRICE MOD REQ Powered pressure reducing mattress overlay/pad, alternating with pump, heavy duty			
E0188	PRICE MOD REQ Synthetic sheepskin pad			
E0191	PRICE MOD REQ Heel or elbow protector, each			
E0193	PRICE MOD REQ Powered air flotation bed			
E0194	AIR FLUIDIZED BED			
E0210	PRICE MOD REQ Electric heat pad, standard			
E0218	Fluid circulating cold pad with pump, any type			
E0249	PRICE MOD REQ Pad for water circulating heat unit, for replacement only			
E0250	PRICE MOD REQ Hospital bed, fixed height, with any type side rails, with mattress			
E0255	PRICE MOD REQ Hospital bed, variable height, hi/lo, with any type side rails, with mattress			
E0260	PRICE MOD REQ Hospital bed, semielectric, with any type side rails, with mattress			
E0261	PRICE MOD REQ Hospital bed, semielectric, with any type side rails, without mattress			
E0271	PRICE MOD REQ Mattress, innerspring			
E0277	PRICE MOD REQ Powered pressurereducing air mattress			
50000	PRICE MOD REQ Hospital bed, heavy duty, extra wide, with wt. capacity > 350 lb., but < = 600 lb., with any type			
E0303	side rails, with mattress			
E0305	PRICE MOD REQ Bed side rails, half length			
E0310	PRICE MOD REQ Bed side rails, full length			
F0.404	PRICE MOD REQ Portable gaseous oxygen system, rental; includes portable container, regulator, flowmeter,			
E0431	humidifier, cannula or mask, & tubing			
F0404	PRICE MOD REQ Portable liq. oxygen system, rental; includes portable container, supply reservoir, humidifier,			
E0434	flowmeter, refill adaptor, contents gauge, cannula or mask, & tub			
F0.400	PRICE MOD REQ Stationary liq. oxygen system, rental; includes container, contents, regulator, flowmeter,			
E0439	humidifier, nebulizer, cannula or mask, & tubing			
E0441	Stationary oxygen contents, gaseous, 1 month's supply = 1 unit			



PRICE MOD REO IV pole

E0776

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505	stmedicare.com  Procedure codes related to the CMS Inpatient Only Lis	st have been removed, however the inpatient admission care requires an authorization.
E0781	PRICE MOD REQ Ambulatory infusion pump, single or multiple channels, electric or battery operated, with	
EU/01	administrative equipment, worn by patient	
E0784	PRICE MOD REQ External ambulatory infusion pump, insulin	
E0912	PRICE MOD REQ Trapeze bar, heavy duty, for patient wt. capacity > 250 lb., free standing, complete with grab ba	г
E0935	PRICE MOD REQ Continuous passive motion exercise device for use on knee only	
E0940	PRICE MOD REQ Trapeze bar, free standing, complete with grab bar	
E0944	PRICE MOD REQ Pelvic belt/harness/boot	
E0955	PRICE MOD REQ Wheelchair accessory, headrest, cushioned, any type, including fixed mounting hardware, eac	ի
E0956	PRICE MOD REQ Wheelchair accessory, lateral trunk or hip support, any type, including fixed mounting hardware, each	
E0957	PRICE MOD REQ Wheelchair accessory, medial thigh support, any type, including fixed mounting hardware, each	
E0960	PRICE MOD REQ Wheelchair accessory, shoulder harness/straps or chest strap, including any type mounting hardware	
E0966	PRICE MOD REQ Manual wheelchair accessory, headrest extension, each	
E0971	PRICE MOD REQ Manual wheelchair accessory, antitipping device, each	
E0973	PRICE MOD REQ Wheelchair accessory, adjustable height, detachable armrest, complete assembly, each	
E0986	PRICE MOD REQ Manual wheelchair accessory, pushrim activated power assist system	
E0990	PRICE MOD REQ Wheelchair accessory, elevating leg rest, complete assembly, each	
E0995	PRICE MOD REQ Wheelchair accessory, calf rest/pad, replacement only, each	
E1002	PRICE MOD REQ Wheelchair accessory, power seating system, tilt only	
E1004	PRICE MOD REQ Wheelchair accessory, power seating system, recline only, with mechanical shear reduction	
E1007	PRICE MOD REQ Wheelchair accessory, power seating system, combination tilt & recline, with mechanical shear reduction	er
E1012	PRICE MOD REQ Wheelchair accessory, addition to power seating system, center mount power elevating leg	
E1012	rest/platform, complete system, any type, each	
E1039	PRICE MOD REQ Transport chair, adult size, heavy duty, patient wt. capacity > 300 lbs.	
E1161	PRICE MOD REQ Manual adult size wheelchair; includes tilt in space	
E1225	PRICE MOD REQ Wheelchair accessory, manual semireclining back, each	
E1226	PRICE MOD REQ Wheelchair accessory, manual fully reclining back, each	
E1390	PRICE MOD REQ Oxygen concentrator, single delivery port, capable of delivering 85% or > oxygen concentration at prescribed flow rate	
E1392	PRICE MOD REQ Portable oxygen concentrator, rental	
E1399	DME (Durable medical equipment) - miscellaneous	
E2102	Adjunctive, non-implanted continuous glucose monitor or receiver	
E2103	Non-adjunctive, non-implanted continuous glucose monitor or receiver	
E2202	PRICE MOD REQ Manual wheelchair accessory, nonstandard seat frame, width 24 - 27 in.	
E2203	PRICE MOD REQ Manual wheelchair accessory, nonstandard seat frame, depth 20 - < 22 in.	
E2205	PRICE MOD REQ Manual wheelchair accessory, handrim without projections, any type, replacement only, each	



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capacity 301 - 450 lb.

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Updated: 07/01/2025

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K0848	PRICE MOD REQ Power wheelchair, group 3 standard, sling/solid seat/back, patient wt. capacity up to &	
K0040	including 300 lb.	
K0849	PRICE MOD REQ Power wheelchair, group 3 standard, captain's chair, patient wt. capacity up to & including 300	
K0849	lb.	
K0861	PRICE MOD REQ Power wheelchair, group 3 standard, multiple power option, sling/solid seat/back, patient wt.	
K0801	capacity up to & including 300 lb.	
K0862	PRICE MOD REQ Power wheelchair, group 3 heavy duty, multiple power option, sling/solid seat/back, patient wt.	
KU862	capacity 301 - 450 lb.	
Q0506		
Q0300	Battery, lithium-ion, for use with electric or electric/pneumatic ventricular assist device, replacement only	
V2625	Enlargement of ocular prosthesis	
V2626	Reduction of ocular prosthesis	
V2632	Posterior chamber intraocular lens	

#### **Parenteral and Enteral Nutrition** Code **Code Description** Comments Enteral feeding supply kit; syringe fed, per day, includes but not limited to feeding/flushing syringe, B4034 Prior Authorization for BlueCare Plus only administration set tubing, dressings, tape Enteral feeding supply kit; pump fed, per day, includes but not limited to feeding/flushing syringe, administration B4035 Prior Authorization for BlueCare Plus only set tubing, dressings, tape Enteral feeding supply kit; gravity fed, per day, includes but not limited to feeding/flushing syringe, B4036 administration set tubing, dressings, tape B4087 Gastrostomy/jejunostomy tube, standard, any material, any type, each B4088 Gastrostomy/jejunostomy tube, low-profile, any material, any type, each Enteral formula, nutritionally complete with intact nutrients, includes proteins, fats, carbohydrates, vitamins & B4150 Prior Authorization for BlueCare Plus only minerals, may include fiber, administered Enteral formula, nutritionally complete, calorically dense with intact nutrients, includes proteins, fats, B4152 Prior Authorization for BlueCare Plus only carbohydrates, vitamins & minerals, may include fiber, administered Enteral formula, nutritionally complete, hydrolyzed proteins, includes fats, carbohydrates, vitamins & minerals, B4153 may include fiber, administered through enteral feeding tube Enteral formula, nutritionally complete, for special metabolic needs, excludes inherited disease of metabolism, B4154 includes altered composition of proteins, fats, carbohydrates Enteral formula, nutritionally incomplete/modular nutrients, includes specific nutrients, carbohydrates, B4155 Prior Authorization for BlueCare Plus only proteins/amino acids, fat or combination, administered through enteral B4185 Parenteral nutrition solution, not otherwise specified, 10 g lipids Parenteral nutrition solution: compounded amino acid & carbohydrates with electrolytes, trace elements, & B4189 Prior Authorization for BlueCare Plus only vitamins, including preparation, any strength, 10 to 51 g of protein, Parenteral nutrition solution: compounded amino acid & carbohydrates with electrolytes, trace elements, & B4193 vitamins, including preparation, any strength, 52 to 73 g of protein Parenteral nutrition solution; compounded amino acid & carbohydrates with electrolytes, trace elements & B4197 vitamins, including preparation, any strength, 74 to 100 g of protein B4220 Parenteral nutrition supply kit; premix, per day Prior Authorization for BlueCare Plus only



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B4224 Parenteral nutrition administration kit, per day		
B9002 Enteral nutrition infusion pump, any type		Prior Authorization for BlueCare Plus only
B9998 NOC for enteral supplies		Prior Authorization for BlueCare Plus only

#### **Gene Therapy**

Code	Code Description	Comments
0540T Chimeric antigen receptor T-cell (CAR-T) therapy; CAR-T cell administration, autologous		
C9301	Obecabtagene autoleucel, up to 410 million cd19 car-positive viable t cells, including leukapheresis and dose	
	preparation procedures, per therapeutic dose	

Genetic Testing	
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Antigen 1 genotyping, ITGB3 , gene analysis, common variant, HPA- 1a/b  Antigen 2 genotyping, GP1BA , gene analysis, common variant, HPA- 2a/b  Antigen 3 genotyping, ITGA2B , gene analysis, common variant, HPA- 3a/b  Antigen 4 genotyping, ITGB3 , gene analysis, common variant, HPA- 4a/b  Antigen 5 genotyping, ITGA2 , gene analysis, common variant  Antigen 6 genotyping, ITGB3 , gene analysis, common variant, HPA- 6a/b	
Antigen 3 genotyping, ITGA2B , gene analysis, common variant, HPA- 3a/b Antigen 4 genotyping, ITGB3 , gene analysis, common variant, HPA- 4a/b Antigen 5 genotyping, ITGA2 , gene analysis, common variant Antigen 6 genotyping, ITGB3 , gene analysis, common variant, HPA- 6a/b	
Antigen 4 genotyping, ITGB3 , gene analysis, common variant, HPA- 4a/b Antigen 5 genotyping, ITGA2 , gene analysis, common variant Antigen 6 genotyping, ITGB3 , gene analysis, common variant, HPA- 6a/b	
Antigen 5 genotyping, ITGA2 , gene analysis, common variant Antigen 6 genotyping, ITGB3 , gene analysis, common variant, HPA- 6a/b	
Antigen 6 genotyping, ITGB3 , gene analysis, common variant, HPA- 6a/b	
Antigen 9 genotyping, ITGA2B , gene analysis, common variant, HPA- 9a/b	
Antigen 15 genotyping, CD109 , gene analysis, common variant, HPA-15a/b	
variants	
variants	
n) (eg, Duchenne/Becker muscular dystrophy) deletion analysis, and duplication analysis, if	
gene analysis; full sequence analysis & full duplication/deletion analysis	
gene analysis; full sequence analysis	
gene analysis; full duplication/deletion analysis	
alysis; full sequence analysis	
alysis; full duplication/deletion analysis	
alysis; full duplication/deletion analysis	
L1;14)) (eg, mantle cell lymphoma) translocation analysis, major breakpoint, qualitative and performed	
alysis, variants in kinase domain	
ysis; eval to detect abnormal alleles	
ysis; characterization of alleles	
s; full gene sequence	
s; known familial variant	
alysis; full gene sequence	
alysis; targeted sequence analysis	
ysis, eval to detect abnormal alleles	
alysis, eval to detect abnormal alleles	
alysis, eval to detect abnormal alleles	
a y:	lysis; targeted sequence analysis sis, eval to detect abnormal alleles



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81180	ATXN3 gene analysis, eval to detect abnormal alleles		
81181	ATXN7 gene analysis, eval to detect abnormal alleles		
81182	ATXN8OS gene analysis, eval to detect abnormal alleles		
81183	ATXN10 gene analysis, eval to detect abnormal alleles		
81184	CACNA1A gene analysis; eval to detect abnormal alleles		
81185	CACNA1A gene analysis; full gene sequence		
81186	CACNA1A gene analysis; known familial variant		
81187	CNBP gene analysis, eval to detect abnormal alleles		
81188	CSTB gene analysis; eval to detect abnormal alleles		
81189	CSTB gene analysis; full gene sequence		
81190	CSTB gene analysis; known familial variant		
81191	NTRK1 (neurotrophic receptor tyrosine kinase 1) (eg, solid tumors) translocation analysis		
81192	NTRK2 (neurotrophic receptor tyrosine kinase 2) (eg, solid tumors) translocation analysis		
81193	NTRK3 (neurotrophic receptor tyrosine kinase 3) (eg, solid tumors) translocation analysis		
81194	NTRK (neurotrophic-tropomyosin receptor tyrosine kinase 1, 2, and 3) (eg, solid tumors) translocation analysis	_	
81195	Cytogenomic (genome-wide) analysis, hematologic malignancy, structural variants and copy number variants,		
	optical genome mapping (OGM)	T	
81200	ASPA gene analysis, common variants		
81201	APC (adenomatous polyposis coli) gene analysis; full gene sequence		
81202	APC (adenomatous polyposis coli) gene analysis; known familial variants		
81203	APC (adenomatous polyposis coli) gene analysis; duplication/deletion variants		
81204	AR gene analysis; characterization of alleles		
81205	BCKDHB gene analysis, common variants		
81206	BCR/ABL1 translocation analysis; major breakpoint, qualitative or quantitative		
81207	BCR/ABL1 translocation analysis; minor breakpoint, qualitative or quantitative		
81208	BCR/ABL1 translocation analysis; other breakpoint, qualitative or quantitative		
81209	BLM gene analysis, 2281del6ins7 variant		
81210	BRAF , gene analysis, V600 variant		
81212	BRCA1, BRCA2 gene analysis; 185delAG, 5385insC, 6174delT variants		
81215	BRCA1 gene analysis; known familial variant		
81216	BRCA2 gene analysis; full sequence analysis		
81217	BRCA2 gene analysis; known familial variant		
81218	CEBPA , gene analysis, full gene sequence		
81219	CALR , gene analysis, common variants in exon 9		
81220	CFTR gene analysis; common variants		
81221	CFTR gene analysis; known familial variants		
81222	CFTR gene analysis; duplication/deletion variants		
81223	CFTR gene analysis; full gene sequence		
81224	CFTR gene analysis; intron 8 poly-T analysis		
81225	CYP2C19, gene analysis, common variants		
81226	CYP2D6 cytochrome P450, family 2, subfamily D, polypeptide 6 eg, drug metabolism, gene analysis, common		
01220	variants		



81269

HBA1/HBA2, gene analysis; duplication/deletion variants

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. 81227 CYP2C9, gene analysis, common variants 81228 Cytogenomic constitutional microarray analysis 81229 Cytogenomic constitutional microarray analysis 81230 CYP3A4, gene analysis, common variant 81231 CYP3A5, gene analysis, common variants 81232 DPYD, gene analysis, common variant 81233 BTK gene analysis, common variants 81234 DMPK gene analysis; eval to detect abnormal alleles 81235 EGFR (epidermal growth factor receptor) gene analysis, common variants 81236 EZH2 gene analysis, full gene sequence 81237 EZH2 gene analysis, common variant 81238 F9, full gene sequence 81239 DMPK gene analysis; characterization of alleles 81240 F2 gene analysis, 20210G>A variant 81241 F5 gene analysis, Leiden variant 81242 FANCC gene analysis, common variant 81243 FMR1 gene analysis; evaluation to detect abnormal alleles 81244 FMR1 gene analysis; characterization of alleles 81245 FLT3, gene analysis; internal tandem duplication variants 81246 FLT3 , gene analysis; tyrosine kinase domain variants 81247 G6PD, gene analysis; common variant 81248 G6PD, gene analysis; known familial variant 81249 G6PD, gene analysis; full gene sequence 81250 G6PC gene analysis, common variants 81251 GBA gene analysis, common variants 81252 GJB2 (gap junction protein, beta 2, 26kDa; connexin 26) gene analysis; full gene sequence 81253 GJB2 (gap junction protein, beta 2, 26kDa; connexin 26) gene analysis; known familiar variants 81254 GJB2 (gap junction protein, beta 6, 30kDa; connexin 30) gene analysis; common variants 81255 HEXA gene analysis, common variants 81256 HFE gene analysis, common variants 81257 HBA1/HBA2, gene analysis; common deletions or variant 81258 HBA1/HBA2, gene analysis; known familial variant 81259 HBA1/HBA2, gene analysis; full gene sequence 81260 IKBKAP gene analysis, common variants IGH@ gene rearrangement analysis to detect abnormal clonal population(s); amplified methodology 81261 81262 IGH@ gene rearrangement analysis to detect abnormal clonal population(s); direct probe methodology 81263 IGH@, variable region somatic mutation analysis 81264 IGK@ gene rearrangement analysis, evaluation to detect abnormal clonal population(s) 81265 Comparative analysis using Short Tandem Repeat markers; patient & comparative specimen 81266 Comparative analysis using Short Tandem Repeat markers; each add'l specimen 81267 Chimerism analysis, post transplantation specimen 81268 Chimerism analysis, post transplantation specimen, each cell type



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MGMT promoter methylation analysis

MCOLN1 gene analysis, common variants

MLH1 gene analysis; full sequence analysis

MLH1 gene analysis; known familial variants

MSH2 gene analysis; full sequence analysis
MSH2 gene analysis; known familial variants

MSH6 gene analysis; full sequence analysis

MSH6 gene analysis; known familial variants

MECP2 gene analysis; full sequence analysis

MECP2 gene analysis; known familial variants

MYD88 gene analysis, p.Leu265Pro variant

NUDT15 gene analysis, common variant

PALB2 gene analysis; full gene sequence

NPM1 gene analysis, exon 12 variants

PALB2 gene analysis; known familial variant

MECP2 gene analysis; duplication/deletion variants

PIK3CA gene analysis, targeted sequence analysis

NRAS, gene analysis, variants in exon 2 & exon 3

PABPN1 gene analysis, eval to detect abnormal alleles

normal tissue, if performed

MLH1 gene analysis; duplication/deletion variants

MSH2 gene analysis; duplication/deletion variants

MSH6 gene analysis; duplication/deletion variants

Microsatellite instability analysis markers for mismatch repair deficiency, includes comparison of neoplastic &

MTHFR gene analysis, common variants

MLH1 gene analysis; promoter methylation analysis
FXN gene analysis; known familial variant

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bcbs	stmedicare.com	Procedure codes related to the CMS Inpatient Only List I	have been removed, however the inpatient admission care requires an authorization.	
81313	PCA3/KLK3 prostate	cancer antigen 3 non-protein coding kallikrein-related peptidase 3 prostate specific antigen		
61313	ratio			
81314	PDGFRA , gene analysis, targeted sequence analysis			
81315	PML/RARalpha, translocation analysis; common breakpoints, qualitative or quantitative			
81316	PML/RARalpha, trans	slocation analysis; single breakpoint, qualitative or quantitative		
81317	PMS2 gene analysis;	full sequence analysis		
81318	PMS2 gene analysis;	known familial variants		
81319	PMS2 gene analysis;	duplication/deletion variants		
81320	PLCG2 gene analysis	s, common variants		
81321	PTEN (phosphatase	and tensin homolog) gene analysis; full sequence analysis		
81322	PTEN (phosphatase	and tensin homolog) gene analysis; known familial variant		
81323	PTEN (phosphatase	and tensin homolog) gene analysis; duplication/deletion variant		
81324	PMP22 (peripheral m	nyelin protein 22) gene analysis; duplication/deletion analysis		
81325	PMP22 (peripheral m	nyelin protein 22) gene analysis; full sequence analysis		
81326	PMP22 (peripheral m	nyelin protein 22) gene analysis; known familial variant		
81327	SEPT9 promoter met	hylation analysis		
81328	SLCO1B1, gene ana	lysis, common variant - RN Review >= 12/13/2018		
81329	SMN1 gene analysis;	dosage/deletion analysis, includes SMN2 analysis, if performed		
81330	SMPD1 gene analysis	s, common variants		
81331	SNRPN/UBE3A, metl	hylation analysis		
81332	SERPINA-1, gene and	alysis, common variants		
81333	TGFBI gene analysis,	common variants		
81334	RUNX1, gene analys	is, targeted sequence analysis		
81335	TPMT, gene analysis	, common variants		
81336	SMN1 gene analysis;	full gene sequence		
81337	SMN1 gene analysis;	known familial sequence variant		
81338		ogene, thrombopoietin receptor) (eg, myeloproliferative disorder) gene analysis; common W515K, W515L, W515R)		
		ogene, thrombopoietin receptor) (eg, myeloproliferative disorder) gene analysis; sequence		
81339	analysis, exon 10	-6,,,,,,		
81340		gement analysis to detect abnormal clonal population(s); using amplification methodology		
81341	TRB@, gene rearrang	gement analysis to detect abnormal clonal population(s); using direct probe methodology		
81342	TRG@, gene rearrang	gement analysis, evaluation to detect abnormal clonal population(s)		
81343	PPP2R2B gene analy	rsis, eval to detect abnormal alleles		
81344	TBP gene analysis, e	val to detect abnormal alleles		
81345	TERT gene analysis, t	targeted sequence analysis		
81346	TYMS , gene analysis	s, common variant		
81347	٠,, ٠	or [3b] subunit B1) (eg, myelodysplastic syndrome/acute myeloid leukemia) gene analysis, g, A672T, E622D, L833F, R625C, R625L)		
01240	SRSF2 (serine and ar	rginine-rich splicing factor 2) (eg, myelodysplastic syndrome, acute myeloid leukemia) gene		
81348	analysis, common va	ariants (eg, P95H, P95L)		



genes, including CYP2C19, CYP2D6, and CYP2D6 duplication/deletion analysis

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Festal chromosomal aneuploidity genomic sequence analysis panel, circulating cell-free PMIn in maternal blood		Procedure codes related to the CMS inpatient Only List	thave been removed, however the inpatient admission care requires an authorization.	
Blodd, must include analysis of chromosomes 13, 18, and 21	81420			
Hereditary breast cancer-related disorders; genomic sequence analysis panet, must include sequencing of at least 10 genes, always including BRCA1, BRCA2, CDH1, MLH1, MSH2, MSH4 Hereditary breast cancer-related disorders; a capital profession analysis panet, must include analyses for SRCA1, BRCA2, MLH1, MSH2, ASTK11  81434 Hereditary breast cancer-related disorders; a capital profession analysis panet, must include sequencing of at least 19 genes, including ABCA4, CNCA1, CRB1, FVS, PDEA8, PDE68, PRPS13, PRPH2, RD  81435 Hereditary colon cancer disorders; genomic sequence analysis panet, must include sequencing of at least 10 genes, including APC, BMPR1A, CDH1, MLH1, MSH2, MSH6, MUTHY, PTIN, S  81436 Hereditary colon cancer disorders; updated involved temper analysis panet, must include analysis of at least 10 genes, including APC, BMPR1A, CDH1, MLH1, MSH2, MSH6, MUTHY, PTIN, S  81437 Hereditary resurrence disorders; updated involved temper analysis panet, must include analysis of at least 5 genes, including MCH1, MSH2, PCPAM, SMAD4, STK11 Hereditary resurrence disorders; updated involved temper analysis panet, must include sequencing of at least 5 genes, including MCH1, MSH2, PCPAM, SMAD4, STK11 Hereditary resurrence disorders; updated penes analysis panet, must include sequencing of at least 5 genes, including postary part analysis panet, must include sequencing of at least 5 genes analysis of at least 10 genes, including BCS11, Cloridary, CoO2, COVID, DCIUO, MPV17, OPA1, PDSS2, PO  Nuclear encoded mitochrondriat genes, genomic sequence panet, must include sequencing of at least 12 genes, including BCS11, Cloridary, CoO2, COVID, DCIUO, MPV17, OPA1, PDSS2, PO  Nuclear encoded mitochrondriat genes genomic sequence analysis panet, must include sequencing of at least 15 genes analysis panet, part part part part part part part part				
191432 least 10 genes, always including BRCA1, BRCA2, CDH1, MLH1, MSH2, MSH 191433 least 10 genes, always including BRCA1, BRCA2, CDH1, MLH1, MSH2, MSH 191434 letretiary intend storeters is deplication/deletion analysis panel, must include analyses for BRCA1, BRCA2, MLH1, MSH2, & STK11 191434 letretiary intend storeters, genomic sequence analysis panel, must include sequencing of at least 15 genes, including ARCA, CNOA1, CRS1, ER3, PDESA, PDESA, PRPH2, RDH 191435 letretiary colon cancer disorders, genomic sequence analysis panel, must include sequencing of at least 10 genes, including MSLA, BMPH2, CDH1, MH1, MSH2, PCBAR, MUTH, MSH2, PCBAR, MADA, & STK11 191437 Hereditary neuroendocrine tumor disorders 191438 Hereditary neuroendocrine tumor disorders 191439 Letreditary neuroendocrine tumor disorders 191439 Letreditary cardiomyopathy, genomic sequence analysis panel, must include sequencing of at least 15 genes, including MSH1, CDTC2, COQ2, COXID, GUICK, MPT1, OPA1, PDSS2, PO 1914340 Nuclear encoded mition-nondrial genes, genomic sequence analysis of at least 100 genes, including MSH1, CDTC2, COQ2, COXID, GUICK, MPT1, OPA1, PDSS2, PO 1914441 Nuclear encoded mition-nondrial genes, genomic sequence analysis panel, must include analysis of at least 100 genes, including MSH1, CDTC2, COQ2, COXID, GUICK, MPT1, OPA1, PDSS2, PO 1914440 Nuclear encoded mition-nondrial genes, genomic sequence analysis panel, must include sequencing of at least 100 genes, including BSH2, CBL, HRSS, RMSS, MAP2XI, MAP2X2, NMSS, PPN11, RAP1, 191445 Nuclear encoded mition-nondrial genes, genomic sequence analysis panel, must include sequencing of at least 15 genes, including BSH2, CBL, HRSS, RMSS, MAP2XI, MAP2X2, NMSS, PPN11, RAP1, 191445 Nuclear encoded mition-nondrial genes, genomic sequence analysis panel, storetial genes, interrogation	81422			
least 10 genes, always including BRCA1, BRCA2, CDH1, MLH1, MSH2, MSH  81434 BRCA1, BRCA2, MLH1, MSH2, & SIK11  81436 BRCA1, BRCA2, MLH1, MSH2, & SIK11  81437 BRCA2, MLH1, MSH2, & SIK11  81438 BRCA4, CNCA1, CRB1, PSY, PDEA, PDEB, PRPS1, PRPH2, RD  81439 BRCA4, BRCA4, CNCA1, CRB1, PSY, PDEA, PDEB, PRPS1, PRPH2, RD  81439 BRCA4, CNCA1, CRB1, PSY, PDEA, PDEB, PRPS1, PRPH2, RD  81439 BRCA4, CNCA1, CRB1, PSY, PDEA, PDEB, PRPS1, PRPH2, RD  81439 BRCA4, CNCA1, CRB1, PSY, PDEA, PDEB, PRPS1, PRPH2, RD  81439 BRCA4, CNCA1, CRB1, PSY, PDEA, PDEB, PRPS1, PRPH2, RD  81439 BRCA4, CNCA1, CRB1, PSY, PDEA, PDEB, PRPS1, PRPH2, RD  81439 BRCA4, CNCA1, CRB1, PSY, PDEA, PDEB, PRPS1, PRPH2, RD  81439 BRCA4, CNCA1, C	81432	Hereditary breast cancer-related disorders; genomic sequence analysis panel, must include sequencing of at		
88CA1_BRCA2_MH11_MSH2_EX_SIST11 81434 Hereditary retinal disorders, genomic sequence analysis panel, must include sequencing of at least 15 genes, including ABCA4_CNGA1_CRB1_EYS_P.DEAR_PDEB8_RPF31_RPRH2_RD 81438 Hereditary colon cancer disorders; genomic sequence analysis panel, must include sequencing of at least 10 genes, including RACA4_CNGA1_CRB1_EYS_P.DEAR_PDEB8_RPF31_RPRH2_RD 81438 Hereditary colon cancer disorders; depulication/deletion gene analysis panel, must include analysis of at least 5 genes, including MRH1_MSH2_EPCAB_NAMA4_RS_INTER_PRH2_RD 81438 Hereditary neuroendocrine tumor disorders 81438 Hereditary neuroendocrine tumor disorders 81439 Hereditary neuroendocrine tumor disorders 81439 Hereditary neuroendocrine tumor disorders 81440 Nuclear encoded mitochondrial genes, genomic sequence panel, must include sequencing of at least 10 genes, including MRDS_INTER_PRH2_RD 81440 Nuclear encoded mitochondrial genes, genomic sequence panel, must include analysis of at least 100 genes, including MRDS_INTER_PRH2_RD 81441 Blacktara maemia, Shwachman-Diamond syndrome, GATA2 deficiency syndrome, congenital amagekanyocytic 81442 Noona spectrum disorders, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF_CRB_HMS_KRAS_MAPZK_I_MAPZK_RMS_FPRH1_I_MR1_ 81443 Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 12 genes of including BRAF_CRB_HMS_KRAS_MAPZK_I_MAPZK_RMS_FPRH1_I_MR1_ 81444 Side of the sequence analysis panel, sequence analysis panel, must include sequencing of at least 15 genes of the sequence analysis panel, sequence analysis panel, must include sequencing of at least 15 genes of the sequence analysis panel, sequence analysis panel, must include sequencing of at least 15 genes of the sequence analysis panel, sequence analysis panel, must include sequencing of at least 15 genes of the sequence analysis panel, solid organ reoplasm of sequence analysis, and the sequence analysis panel, solid organ	01402	least 10 genes, always including BRCA1, BRCA2, CDH1, MLH1, MSH2, MSH		
8RCA1, BRCA2, HLH,I, MSH2, & STK11  61434 including ABCA4, CNGA1, CRBI, FYS, DEBA, PDEB, PREP13, PRPH2, RD  61435 deneritary extend disorders, genomic sequence analysis panel, must include sequencing of at least 10 genes, including APC, BMPR1A, CDH1, MLH1, MSH2, MSH6, MUTYH, PTEN, S  61436 deneritary coion cancer disorders; duplication/deletion gene analysis panel, must include analysis of at least 5 genes, including HLH, MSH2, EVAN, SMADA, 4 STK11  61437 Hereditary neuroendocrine tumor disorders  61438 dereditary neuroendocrine tumor disorders  61439 Hereditary neuroendocrine tumor disorders  61440 Nuclear encoded mitochondrial genes, genomic sequence panel, must include sequencing of at least 10 genes, including MSD31, C10erd, C002, C0X10, DGUOX, MPV17, DPA1, POSS2, PO  61441 Stackfan anemia, Swacchman-Danoma Syndrome, GATA, Sedencinery syndrome, congenitar amegakanyocytic  61442 Noonan spectrum disorders, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF, CBL, HRAS, KRAS, MAPZK1, MAPZK2, NRAS, PIPN11, RAF1,  61443 Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF, CBL, HRAS, RARS, MAPZK1, MAPZK2, NRAS, PIPN11, RAF1,  61445 Solid organ neoplasm, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF, CBL, HRAS, SRAS, MAPZK1, MAPZK2, NRAS, PIPN11, RAF1,  61449 Sequence analysis panel, solid organ neoplasm, 5-50 genes, interrogation for sequencing of at least 12 genes, proper proper inherited an endorse special research proper inherited an endorse special research proper inherited research	81/33			
hereditary colon cancer disorders; genomic sequence analysis panel, must include sequencing of at least 10 genes, including APC, BMPRIA, CDH, MLH, MSI2, EMSA, MUTH, PTEN, S  Hereditary colon cancer disorders; deplication (deletion gene analysis panel, must include analysis of at least 5 genes, including MLH, MSI2, EPCAN, SMADA, & STISTI  Hereditary neuroendocrine tumor disorders  Hereditary cardiomyopathy, genomic sequence analysis panel, must include sequencing of at least 5 cardiomyopathy-related genes  Nuclear encoded mitochondrial genes, genomic sequence panel, must include analysis of at least 100 genes, including BCS11, C100rt2, COQ2, COX10, DGUOK, MPV17, OPA1, PDSS2, PO  Bischan analmais, Shwachman-Diomond syndrome, CAR2 deficiency syndrome, congenital american analmais, analmais analmais, analmais analmais, analmais analmais, analmais analmais, analmais analmais, analmais	01400	BRCA1, BRCA2, MLH1, MSH2, & STK11		
including ABCM, CNGAI, CRBI, EYS, PDE68, PRP63, PRP612, BPH2, RD fereditary colon cancer disorders; genomic sequence analysis panel, must include sequencing of at least 10 genes, including APC, BMPRIA, CDHJ, MLHJ, MSH2, MSH6, MUTH, PTEN, S fereditary colon cancer disorders; supplication/desilenting open analysis panel, must include analysis of at least 5 genes, including MLHJ, MSH2, EPCAM, SMAD4, & STK11 fereditary neuroendocrine tumor disorders fereditary cardiomyopathy, genomic sequence analysis panel, must include analysis of at least 5 cardiomyopathy, genomic sequence analysis panel, must include analysis of at least 10 genes, including MSH2, CD (100, 100, 100, 100, 100, 100, 100, 100	81/3/	Hereditary retinal disorders, genomic sequence analysis panel, must include sequencing of at least 15 genes,		
81439 genes, including APC, BMPRIA CDHI, MLHI, MSH2, MSH6, MUTYH, PTEN, S  81438 Hereditary coton cancer disorders, duplication/deletion gene analysis panel, must include analysis of at least 5  81437 Hereditary neuroendocrine tumor disorders  81438 Hereditary neuroendocrine tumor disorders  81439 Hereditary neuroendocrine tumor disorders  81440 Hereditary cardiomyopathy, genomic sequence analysis panel, must include sequencing of at least 5  81440 Cardiomyopathy-related genes  81440 Nuclear encoded mitochnodrial genes, genomic sequence panel, must include analysis of at least 100 genes, including BCSIL, C10ortz, COQ2, COX10, DOUGK, MPV17, OPAI, PDSS2, PO  81441 Blackfan anemia, Shwachman-Diamond syndrome, GATA2 deficiency syndrome, congenital amegakanyocytic  81442 Noonan spectrum disorders, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BBAF, GEN, HASS, KRAS, AMAZYAI, MAPZAY, NAPZAY, N	01404	including ABCA4, CNGA1, CRB1, EYS, PDE6A, PDE6B, PRPF31, PRPH2, RD		
genes, including APC, BMPRIA, CDH1, MLH1, MSH2, KSH6, MUTH, PTEN, S  ### recrifiatry colon cancer disorders, subjuiciation/deleting gene analysis panel, must include analysis of at least 5 genes, including MLH1, MSH2, EPCAM, SMADA, & STK11  ### recrifiatry neuroendocrine tumor disorders  ### recrifiatry cardiomyopathy, genomic sequence analysis panel, must include sequencing of at least 5 exaction/myopathy-related genes  ### Nuclear encoded mitochondrial genes, genomic sequence panel, must include analysis of at least 100 genes, including BCS1L, C10ortz, COQ2, COX10, DOUDK, MPV17, OPA1, PDSS2, PO  ### Nuclear encoded mitochondrial genes, genomic sequence panel, must include analysis of at least 100 genes, including BCS1L, C10ortz, COQ2, COX10, DOUDK, MPV17, OPA1, PDSS2, PO  ### Nuclear encoded mitochondrial genes, genomic sequence, panel, must include sequencing of at least 120 genes, including BCS1L, C10ortz, COQ2, COX10, DOUDK, MPV17, OPA1, PDSS2, PO  ### Noonan spectrum disorders, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF, CBL, HRAS, KRAS, MAPZLA, MAPZZL, VARSZLA, VARSZLA, MAPZLA, MAPZL	81/35	Hereditary colon cancer disorders; genomic sequence analysis panel, must include sequencing of at least 10		
81437 Hereditary neuroendocrine tumor disorders 81438 Hereditary neuroendocrine tumor disorders 81439 Hereditary neuroendocrine tumor disorders 81439 Hereditary varionendocrine tumor disorders 81439 Hereditary varionendocrine tumor disorders 81439 Hereditary varionendocrine tumor disorders 81449 Hereditary varionendocrine tumor disorders 81440 Nuclear encoded mitochondrial genes, genomic sequence panel, must include sequencing of at least 100 genes, including BCS1L, C10orl2, COQ2, COX10, DGUOK, MPU17, OPA1, PDSS2, PO 81441 Blackfan anemia, Shwachman-Diamond syndrome, GATA2 deficiency syndrome, congenital amegakaryocytic 81442 Noonan spectrum disorders, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF, CBL, HRAS, KRAS, AMPAXI, MAPAX, NARS, ITRAIT, RAF1, 81443 Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 15 genes 81445 Solid organ neoplasm, genomic sequence analysis panel, must include sequencing of at least 15 genes 81446 Hereditary peripheral neuropathites, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathites, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathites, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathites, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathity related genes 81449 ReBB2, KTI, KRAS, MET, NRAS, POGFRA, POGFRB, POR, PIR3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis 81450 Arageted genomic sequence analysis panel, hematolympholid neoplasm or disorder, DNA analysis, & RNA 81451 Arageted genomic sequence analysis panel, hematolympholid neoplasm or disorder, 5-50 genes (eg. BRF, CEBRA, NITAR, SNA), Interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if perform	01433	genes, including APC, BMPR1A, CDH1, MLH1, MSH2, MSH6, MUTYH, PTEN, S		
genes, including MLHI, MSH2, EPCAM, SMAD4, & STK11  81437 Hereditary neuroendocrine tumor disorders  81438 Hereditary cardiomyopathy, genomic sequence analysis panel, must include sequencing of at least 5 acrdiomyopathy-related genes.  81449 Nuclear encoded mitochondrial spens, genomic sequence panel, must include analysis of at least 10 genes, including BCSLI, Clohortz, COZO, COX10, DGUOK, MPV17, OPA1, PDSS2, PO intenses under intensor variative Syntomics (Liphy 21) (Epc. Paral. Deliance, SMAD4) (Clohort, COZO, COX10, DGUOK, MPV17, OPA1, PDSS2, PO intenses under intensor variative Syntomics (Liphy 21) (Epc. Paral. Deliance, SMAD4) (Clohort, COZO, COX10, DGUOK, MPV17, OPA1, PDSS2, PO intenses under intensor variative Syntomics (Clohort, COZO, COX10, DGUOK, MPV17, OPA1, PDSS2, PO intenses under intensor variative Syntomics (Clohort, CoZO, COX10, DGUOK, MPV17, OPA1, PDSS2, PO intenses under intensor variative Syntomics, CoZO, COX10, DGUOK, MPV17, OPA1, PDSS2, PO intenses under variative Syntomics, Shadchan analysis panel, must include analysis of at least 12 genes, including BRAF, CBL, HRAS, KRAS, MAP2K1, MAP2K2, NRAS, PTPN11, RAF1, Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 15 genes.  81445 Solid organ neoplasm, genomic sequence analysis panel, solid organ neoplasm, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  81448 Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  81449 Figure and properties of the properties of	91/36	Hereditary colon cancer disorders; duplication/deletion gene analysis panel, must include analysis of at least 5		
Hereditary neuroendocrine tumor disorders Hereditary cardiomyopathy, genomic sequence analysis panet, must include sequencing of at least 5 cardiomyopathy, related genes  Nuclear encoded mitochondriat genes, genomic sequence panet, must include analysis of at least 100 genes, including BCS1L, C10or12, COQ2, COX10, DGUOK, MPV17, OPA1, PDSS2, PO miterinetry observations relationship of the mitochondriat genes, genomic sequence panet, must include analysis of at least 100 genes, including BCS1L, C10or12, COQ2, COX10, DGUOK, MPV17, OPA1, PDSS2, PO miterinetry observationship of the mitochondriat genes, including BCS1L, C10or12, COQ2, COX10, DGUOK, MPV17, OPA1, PDSS2, PO miterinetry observationship of the mitochondriat genes, including BRAF, CBL, HABS, KRAS, MAP2KL, MAP2K2, NRAS, PTPN11, RAF1,  B1442  Noonan spectrum disorders, genomic sequence analysis panet, must include sequencing of at least 12 genes, including BRAF, CBL, HABS, KRAS, MAP2KL, MAP2K2, NRAS, PTPN11, RAF1,  Genetic testing for severe inherited conditions, genomic sequence analysis panet, must include sequencing of at least 12 genes, number variants/rearrangements; DNA or combined DNA/RNA analysis  B1443  B1445  B1446  B1447  B1448  Hereditary peripheral neuropathies, genomic sequence analysis panet, must include sequencing of at least 5 peripheral neuropathy-related genes  Targeted genomic sequence analysis panet, solid organ neoplasm, 5-50 genes (eg. ALK, BRAF, CDKN2A, EGFR, EBB2, KIT, KRAS, MT, NRAS, PDGFRA, PDGFRB, PGR, PIKSCA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, in performed; RNA analysis analysis sheep performed; RNA analysis panet, analysis panet, bematolympholid neoplasm or disorder, 5-50 genes (eg. BRAF, CBRA, DRAF,	81430	genes, including MLH1, MSH2, EPCAM, SMAD4, & STK11		
Hereditary cardiomyopathy, genomic sequence analysis panel, must include sequencing of at least 5 cardiomyopathy-related genes  Nuclear encoded mitochondrial genes, genomic sequence panel, must include analysis of at least 100 genes, including BCS1L, C100rt2, C002, C0X1D, DGUDK, MPV17, OPA1, PDSS2, PO including BCS1L, C100rt2, C002, C0X1D, DGUDK, MPV17, OPA1, PDSS2, PO including BCS1L, C100rt2, C002, C0X1D, DGUDK, MPV17, OPA1, PDSS2, PO including BCS1L, C100rt2, C002, C0X1D, DGUDK, MPV17, OPA1, PDSS2, PO including BCS1L, C100rt2, C002, C0X1D, DGUDK, MPV17, OPA1, PDSS2, PO including BRAF, CBL, HRAS, KRAS, MAPZK1, MAPZK2, NRAS, PTPN11, RAF1,  Noonan spectrum disorders, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF, CBL, HRAS, KRAS, MAPZK1, MAPZK2, NRAS, PTPN11, RAF1,  Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 15 genes  Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for seq variants and copy number variants/rearrangements; DNA or combined DNA/RNA analysis  Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, POG. PIKSCA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNNT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLI, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression revels, if performed; RNA analysis	81437	Hereditary neuroendocrine tumor disorders		
actiomyopathy-related genes  Nuclear encoded mitochondrial genes, genomic sequence panel, must include analysis of at least 100 genes, including BSGSL, C100rtz, COQ2, COX10, DGUOK, MPV17, OPA1, PDSS2, PO internet under introver rainter syntromes (IDHT-3) (Fig. Fancturi attention, dysseriaus) congenita, pramition including BSGSL, C100rtz, COQ2, COX10, DGUOK, MPV17, OPA1, PDSS2, PO internet under introver rainter syntromes (IDHT-3) (Fig. Fancturi attention, dysseriaus) congenita, pramition including BRAF, CBL, HRAS, KRAS, MAP2K1, MAP2K2, NRAS, PTPN11, RAF1,  B1442  B1443  B1443  B14443  B14445  B1445  B1445  B1446  B1448  B1448  B1448  B1449  B1440  B1450  B1450  B1450  B1450  B1455  B1455  B1455  B1455  B1455  B1465  B1465  B1465  B1465  B1465  B1465  B1465  B1466  B1466  B1467  B146	81438	Hereditary neuroendocrine tumor disorders		
Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for sequencing of at least 5 peripheral neuropathy-related genes  1444 Branch and a copy number variants/rearrangements; DNA or combined DNA/RNA analysis  1445 Targeted genomic sequence analysis panel, performent; RNA analysis  1446 Targeted genomic sequence analysis panel, permatory not recognized analysis, and copy number variants/rearrangements; DNA por FRINGA, POFRB, POFR, PIKSCA, PTEN, RET, interrogation for sequence variants and copy number variants or rearrangements; DNA por sequence variants, and copy number variants or rearrangements, if performed; RNA analysis  1445 Targeted genomic sequence analysis panel, performed; RNA analysis  1450 Targeted genomic sequence analysis panel, performed; RNA analysis  1451 Targeted genomic sequence analysis panel, performed; RNA analysis  1451 Targeted genomic sequence analysis panel, performed; RNA analysis  1451 Targeted genomic sequence analysis panel, performed; RNA analysis  1455 Targeted genomic sequence analysis panel, performed; RNA analysis  1455 Targeted genomic sequence analysis panel, performed; RNA analysis  1455 Targeted genomic sequence analysis panel, permatorymphoid neoplasm or disorder, 5-50 genes (eg. BAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLT, NPAS, MDFRA, PDGFRA,	81/39	Hereditary cardiomyopathy, genomic sequence analysis panel, must include sequencing of at least 5		
181440 including BCSLIL, C10ortz, COV2. COX10, DCUOK, MPV17, OPAL, PDSS2, PO interior or uniterior variative syntromes (torm); (tg.; Particul aniema, viyaseratusis congenital amegakaryocytic blackfan aniema, Shwachman-Diamond syndrome, GATA2 deficiency syndrome, congenital amegakaryocytic consideration of the control of	01400	cardiomyopathy-related genes		
including BCS1L, C10orf2, COQ2, COX10, DGUOK, MPV17, OPA1, PDSS2, PO minerinet or unremitural variants syntromies (prior 1) (gs., reactions are state)  81442 Blackfan anemia, Shwachman-Diamond syndrome, GATA2 deficiency syndrome, congenital amegakaryocytic  81442 including BRAF, CBL, HRAS, KRAS, MAP2K1, MAP2K2, NRAS, PTPN11, RAF1,  Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF, CBL, HRAS, KRAS, MAP2K1, MAP2K2, NRAS, PTPN11, RAF1,  Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 15 genes  81445 Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for seq variants and copy number variants/rearrangements; DNA or combined DNA/RNA analysis  81448 Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  1 argeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKNZA, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  81450 Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA  1455 Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression tevels, it performed; RNA analysis  18455 Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	91//0	Nuclear encoded mitochondrial genes, genomic sequence panel, must include analysis of at least 100 genes,		
81441 Blackfan anemia, Shwachman-Diamond syndrome, GATA2 deficiency syndrome, congenital amegakaryocytic  81442 Noonan spectrum disorders, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF, CBL, HRAS, KRAS, MAP2K1, MAP2K2, NRAS, PTPN11, RAF1,  81443 Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 15 genes  81445 Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for seq variants and copy number variants/rearrangements; DNA or combined DNA/RNA analysis  81448 Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg. ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  81450 Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, & Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg. BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression levels, if performed; RNA analysis	01440	including BCS1L, C10orf2, COQ2, COX10, DGUOK, MPV17, OPA1, PDSS2, PO		
Noonan spectrum disorders, genomic sequence analysis panel, must include sequencing of at least 12 genes, including BRAF, CBL, HRAS, KRAS, MAP2K1, MAP2K2, NRAS, PTPN11, RAF1,  81443 Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 15 genes  81445 Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for seq variants and copy number variants/rearrangements; DNA or combined DNA/RNA analysis  81448 Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  81449 ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  81450 Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression levels, if performed; RNA analysis	Q1 <i>/</i> //1			
including BRAF, CBL, HRAS, KRAS, MAPZK1, MAPZK2, NRAS, PTPN11, RAF1,  Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 15 genes  Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for seq variants and copy number variants/rearrangements; DNA or combined DNA/RNA analysis  Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBSZ, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, hematolymphoid neoplasm, DNA analysis, & RNA	01441	thrombooutopopie) acquence englysic panel, must include acquencing of at least 20 capes, including PDCA2		
including BRAF, CBL, HRAS, KRAS, MAPZK1, MAPZK2, NRAS, PTPN11, RAF1,  Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at least 15 genes  Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for seq variants and copy number variants/rearrangements; DNA or combined DNA/RNA analysis  Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression levels, if performed; RNA analysis	81442	Noonan spectrum disorders, genomic sequence analysis panel, must include sequencing of at least 12 genes,		
least 15 genes  Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for seq variants and copy number variants/rearrangements; DNA or combined DNA/RNA analysis  Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	01442	including BRAF, CBL, HRAS, KRAS, MAP2K1, MAP2K2, NRAS, PTPN11, RAF1,		
least 15 genes  Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for seq variants and copy number variants/rearrangements; DNA or combined DNA/RNA analysis  ### Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  ##################################	81443	Genetic testing for severe inherited conditions, genomic sequence analysis panel, must include sequencing of at		
number variants/rearrangements; DNA or combined DNA/RNA analysis  Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	01440	least 15 genes		
Inumber variants/rearrangements; DNA or combined DNA/RNA analysis  Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5 peripheral neuropathy-related genes  Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	81///5	Solid organ neoplasm, genomic sequence analysis panel, 5-50 genes, interrogation for seq variants and copy		
peripheral neuropathy-related genes  Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	01440	number variants/rearrangements; DNA or combined DNA/RNA analysis		
Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  81450 Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, & Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  81455 Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	81448	Hereditary peripheral neuropathies, genomic sequence analysis panel, must include sequencing of at least 5		
81449 ERBB2, KIT, KRAS, MET, NRAS, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis  81450 Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, & Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression levels, if performed; RNA analysis  81455 Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	313	peripheral neuropathy-related genes		
and copy number variants or rearrangements, if performed; RNA analysis  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA				
Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, DNA analysis, & RNA analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	81449			
analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA		and copy number variants or rearrangements, if performed; RNA analysis		
analysis when performed, 5-50 genes, interrogation for sequence variants, &  Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	81450			
CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA		analysis when performed, 5-50 genes, interrogation for sequence variants, &		
sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA		Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF,		
sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis  Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	81451			
Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, & RNA	32.52			
81455				
analysis when performed, 51 or greater genes, interrogation for sequence	81455			
	31400	analysis when performed, 51 or greater genes, interrogation for sequence		



bcbstmedicare.com  Procedure codes related to the CMS Inpatient Only		t have been removed, however the inpatient admission care requires an authorization.	
81456	Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm or disorder, 51 or greater genes (eg, ALK, BRAF, CDKN2A, CEBPA, DNMT3A, EGFR, ERBB2, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MET, MLL, NOTCH1, NPM1, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis		
81457	Solid organ neoplasm, genomic sequence analysis panel, interrogation for sequence variants; DNA analysis, microsatellite instability		
81458	Solid organ neoplasm, genomic sequence analysis panel, interrogation for sequence variants; DNA analysis, copy number variants and microsatellite instability		
81459	Solid organ neoplasm, genomic sequence analysis panel, interrogation for sequence variants; DNA analysis or combined DNA and RNA analysis, copy number variants, microsatellite instability, tumor mutation burden, and rearrangements		
81462	Solid organ neoplasm, genomic sequence analysis panel, cell-free nucleic acid (eg, plasma), interrogation for sequence variants; DNA analysis or combined DNA and RNA analysis, copy number variants and rearrangements		
81463	Solid organ neoplasm, genomic sequence analysis panel, cell-free nucleic acid (eg, plasma), interrogation for sequence variants; DNA analysis, copy number variants, and microsatellite instability		
81464	Solid organ neoplasm, genomic sequence analysis panel, cell-free nucleic acid (eg, plasma), interrogation for sequence variants; DNA analysis or combined DNA and RNA analysis, copy number variants, microsatellite instability, tumor mutation burden, and rearrangements		
81479	Unlisted molecular pathology procedure		
81490	Autoimmune, analysis of 12 biomarkers using immunoassays, utilizing serum, prognostic algorithm reported as disease activity score		
81504	Oncology (tissue of origin), microarray gene expression profiling > 2000 genes, utilizing formalin-fixed paraffinembedded tissue		
81513	Infectious disease, bacterial vaginosis, quantitative real-time amplification of RNA markers for Atopobium vaginae, Gardnerella vaginalis		
81514	Infectious disease, bacterial vaginosis & vaginitis, quantitative real-time amplification of DNA markers for Gardnerella vaginalis, Atopobium		
81517	Liver disease, analysis of 3 biomarkers, using immunoassays, utilizing serum, prognostic algorithm reported as risk score & risk of liver fibrosis & liver-related clinical eve		
81518	Oncology, mRNA, gene expression profiling by real-time RT-PCR of 11 genes, utilizing formalin-fixed paraffinembedded tissue, algorithms reported as percentage risk for metast		
81519	Oncology , mRNA, gene expression profiling by real-time RT-PCR of 21 genes, utilizing formalin-fixed paraffin embedded tissue, algorithm		
81520	Oncology, mRNA gene expression profiling by hybrid capture of 58 genes, utilizing formalin-fixed paraffinembedded tissue, algorithm reported as recurrence risk score		
81521	Oncology, mRNA, microarray gene expression profiling of 70 content genes & 465 housekeeping genes, utilizing fresh frozen or formalin-fixed paraffin-embed		
81522	Oncology, mRNA, gene expression profiling by RT-PCR of 12 genes, utilizing formalin-fixed paraffin-embedded tissue, algorithm reported		
81523	Oncology, mRNA, next-generation sequencing gene expression profiling of 70 content genes & 31 housekeeping genes, utilizing formalin-fixed paraffin-embedded tissue, algorithm		



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	tmedicare.com  Procedure codes related to the CMS Inpatient Only L	ist have been removed, however the inpatient admission care requires an authorization.	
81525	Oncology, mRNA gene expression profiling real-time RT-PCR 12 genes utilizing formalin-fixed paraffin-embedd	ed	
01020	tissue, algorithm		
81528	Oncology screening quantitative real-time target & signal amplification of 10		
	Oncology (cutaneous melanoma), mRNA, gene expression profiling by real-time RT- PCR of 31 genes (28 conte	nt	
81529	and 3 housekeeping), utilizing formalin-fixed paraffin- embedded tissue, algorithm reported as recurrence risk		
	including likelihood of sentinel lymph node metastasis		
81539	Oncology, biochemical assay 4 proteins, utilizing plasma/serum, prognostic algorithm reported as probability		
01505	score		
81542	Oncology, mRNA, microarray gene expression profiling of 22 content genes, utilizing formalin-fixed paraffin-		
010-12	embedded tissue, algorithm reported		
81546	Oncology (thyroid), mRNA, gene expression analysis of 10,196 genes, utilizing fine needle aspirate, algorithm		
01040	reported as a categorical result (eg, benign or suspicious)		
81551	Oncology, promoter methylation profiling real-time PCR 3 genes, formalin-fixed		
81552	Oncology, mRNA, gene expression profiling by real-time RT-PCR of 15 genes, utilizing fine needle aspirate or		
01552	formalin-fixed paraffin-embedded tissue, algorithm reported as ris		
	Pulmonary disease (idiopathic pulmonary fibrosis [IPF]), mRNA, gene expression analysis of 190 genes, utilizin		
81554	transbronchial biopsies, diagnostic algorithm reported as categorical result (eg, positive or negative for high		
	probability of usual interstitial pneumonia [UIP])		
81558	palymetroes obein reaction (aDCD) of 100 gapes, utilizing whole blood, elegations reported as a binary		
81560	Transplantation medicine, measurement of donor & third-party-induced CD154+T-cytotoxic memory cells,		
01500	utilizing whole peripheral blood		
81595	Cardiology mRNA gene expression profiling real-time quantitative PCR 20		
81596	Infectious disease, chronic hepatitis C virus infection, six biochemical assays utilizing serum, prognostic		
01000	algorithm reported as scores for fibrosis & necroinflammatory activ		
81599	Unlisted multianalyte assay with algorithmic analysis		
83700	Lipoprotein, blood; electrophoretic separation & quantitation		
84999	Unlisted chemistry procedure (BR)		
87154	Culture, typing; identification of blood pathogen & resistance typing, when performed, by nucleic acid probe,		
0,104	multiplexed amplified probe technique including multiplex reverse		
87468	Infectious agent detection by nucleic acid (DNA or RNA); Anaplasma phagocytophilum, amplified probe		
07400	technique		
87469	Infectious agent detection by nucleic acid (DNA or RNA); Babesia microti, amplified probe technique		
87478	Infectious agent detection by nucleic acid (DNA or RNA); Borrelia miyamotoi, amplified probe technique		
87484	Infectious agent detection by nucleic acid (DNA or RNA); Ehrlichia chaffeensis, amplified probe technique		
88245	Chromosome analysis for breakage syndromes baseline SCE 20-25 cells		
88248	Chromosome analysis for breakage syndromes baseline score 50-100 cells		
88249	Chromosome anaylsis for breakage syndromes score 100 cells clastogen stress		
88271	Molecular cytogenetics DNA probe each		
88272	Molecular cytogenetics chromosomal in situ hydridization analyze 3-5 cells		
88273	Molecular cytogenetics chromosomal in situ hydridization analyze 10-30 cells		
88274	Molecular cytogenetics interphase in situ hydridization analyze 25-99 cells		
88275	Molecular cytogenetics interphase in situ hydridization analyze 100-300 cells		
88280	Chromosome analysis additional karyotypes each study		



bcbstmedicare.com  Procedure codes related to the CMS Inpatient Only List have been removed, ho		ist have been removed, however the inpatient admission care requires an authorization.		
88283	Chromosome analysis additional specialized banding technique			
88285	Chromosome analysis additional cells counted each study			
88289	Chromosome analysis additional high resolution study			
88291	Cytogenetics & molecular cytogenetics interpretation and report			
88299	Unlisted cytogenetic study (BR)			
88326	Morphometric analysis, tumor immunohistochemistry, quantitative orsemiquantitative, per specimen, each			
88320	single antibody stain procedure; using computer assisted technology			
88361	Morphometric analysis, tumor immunohistochemistry, quantitative orsemiquantitative, per specimen, each			
	single antibody stain procedure; using computer assisted technology			
0005U	Oncology gene expression profile by real-time RT-PCR of 3 genes, urine, algorithm reported as risk score			
0008U	Helicobacter pylori detection & antibiotic resistance, DNA, 16S & 23S rRNA, gyrA, pbp1, rdxA & rpoB, next			
	generation sequencing, formalin-fixed paraffin-embedded or fresh tiss			
0013M	Oncology, mRNA, gene expression profiling by real-time quantitative PCR of 5 genes, utilizing urine, algorithm			
	reported as risk score for having recurrent urothelial carcinoma			
0016M	Oncology, mRNA, microarray gene expression profiling of 219 genes, utilizing formalin-fixed paraffin-embedde			
	tissue, algorithm reported as molecular subtype			
	Oncology (central nervous system), analysis of 30000 DNA methylation loci by methylation array, utilizing DNA			
0020M	extracted from tumor tissue, diagnostic algorithm reported as probability of matching a reference tumor			
	subclass			
0023U	Oncology, DNA, genotyping of internal tandem duplication, p.D835, p.1836, using mononuclear cells, reported	38		
	detection or non-detection of FLT3 mutation & indication for or			
0026U	Oncology, DNA & mRNA 112 genes, next-generation sequencing, fine needle aspirate thyroid nodule, algorithm			
	analysis reported categorical result			
0037U	Targeted genomic sequence analysis, solid organ neoplasm, DNA analysis of 324genes, interrogation for			
000011	sequence variants, gene copy number amplifications, generearrangements,			
0039U 0040U	Deoxyribonucleic acid antibody, double stranded, high avidity			
00400	BCR/ABL1 translocationanalysis, major breakpoint, quantitative			
0045U	Oncology, mRNA, gene expression profiling by real-time RT-PCR of 12 genes, utilizing formalin-fixed paraffinembedded tissue, algorithm reported as recurrence score			
0046U	FLT3 internal tandem duplication variants, quantitative			
00400	Oncology, mRNA, gene expression profiling by real-time RT-PCR of 17 gene utilizing formalin-fixed paraffin-			
0047U	embedded tissue algorithm			
	Oncology, DNA, targeted sequencing of protein-coding exons of 468 cancer- associated genes, including			
0048U	interrogation for somatic mutations & microsatellite instability, matched			
0049U				
	Targeted genomic sequence analysis panel, acute myelogenous leukemia, DNA analysis, 194 genes,			
0050U	nterrogation for sequence variants, copy number variants or rearrangements			
0055U				
	Oncology, mRNA, gene expression profiling by massively parallel sequencing for analysis of 51 genes, utilizing			
0057U	formalin-fixed paraffin-embedded tissue, algorithm reported as n			
0058U	Oncology, detection of antibodies to Merkel cell polyoma virus oncoprotein, serum, quantitative			
005011	Oncology, detection of antibodies to Merkel cell polyoma virus capside protein, serum, reported as positive or			
0059U	negative			



Procedure codes related to the CMS Inpatient		have been removed, however the inpatient admission care requires an authorization.
0060U	Twin zygosity, genomic targeted sequence analysis of chromosome 2, using circulating cell-free fetal DNA in maternal blood	
0061U	Transcutaneous measurement of 5 biomarkers & multi-spectral analysis	
0070U	CYP2D6 gene analysis, common & select rare variants	
0071U	CYP2D6 gene analysis, full gene sequence	
0072U	CYP2D6 gene analysis, targeted sequence analysis	
0073U	CYP2D6 gene analysis, targeted sequence analysis	
0074U	CYP2D6 gene analysis, targeted sequence analysis	
0075U	CYP2D6 gene analysis, targeted sequence analysis	
0076U	CYP2D6 gene analysis, targeted sequence analysis	
0077U	Immunoglobulin paraprotein, qualitative, immunoprecipitation & mass spectrometry, blood or urine, including isotype	
0078U	Pain management genotyping panel, 16 common variants, buccal swab or other germline tissue sample, algorithm reported as positive or negative risk of opioid-use disorder	
0079U	Comparative DNA analysis using multiple selected single-nucleotide polymorphisms, urine & buccal DNA, for specimen identity verification	
0089U	Oncology, gene expression profiling by RTqPCR, PRAME & LINC00518, superficial collection using adhesive patch	
0094U	Genome, rapid sequence analysis	
0101U	Hereditary colon cancer disorders, genomic sequence analysis panel utilizing combination of NGS, Sanger, MLPA, & array CGH, with mRNA analytics to resolve variants of unknown	
0105U	Nephrology, multiplex electrochemiluminescent immunoassay of tumor necrosis factor receptor 1A, receptor superfamily 2, & kidney injury molecule-1 combined with longitudinal c	
0111U	Oncology, targeted KRAS & NRAS gene analysis utilizing formalin-fixed paraffin-embedded tissue	
0113U	Oncology, measurement of PCA3 & TMPRSS2-ERG in urine & PSA in serum following prostatic massage, by RNA amplification & fluorescence-based detection, algorithm reported as ris	
0114U	Gastroenterology, VIM & CCNA1 methylation analysis, esophageal cells, algorithm reported as likelihood for Barrett's esophagus	
0118U	Transplantation medicine, quantification of donor-derived cell-free DNA using whole genome next-generation sequencing, plasma, reported as percentage of donor-derived cell-fre	
0120U	Oncology, mRNA, gene expression profiling by fluorescent probe hybridization of 58 genes, formalin-fixed paraffin-embedded tissue, algorithm reported as likelihood for primary	
0129U	Hereditary breast cancer-related disorders, genomic sequence analysis & deletion/duplication analysis panel	
0152U	U Infectious disease, microbial cell-free DNA, plasma, untargeted next-generation sequencing, report for significant positive pathogens	
0153U	Oncology, mRNA, gene expression profiling by next-generation sequencing of 101 genes, utilizing formalin-fixed paraffin-embedded tissue	
0154U	Oncology, RNA, analysis by real-time RT-PCR of the FGFR3 gene analysis utilizing formalin-fixed paraffinembedded urothelial cancer tumor tissue, reported as FGFR gene alterat	
0155U	Oncology, DNA, PIK3CA gene analysis, utilizing formalin-fixed paraffinembedded breast tumor tissue, reported as PIK3CA gene mutation status	
0156U	Copy number, sequence analysis	



0238U

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. Neurology, RNA, next-generation sequencing, saliva, algorithmic analysis, & results reported as predictive 0170U probability of ASD diagnosis Targeted genomic sequence analysis panel, acute myeloid leukemia, myelodysplastic syndrome, & 0171U myeloproliferative neoplasms, DNA analysis, 23 genes, interrogation for sequence Oncology, somatic mutation analysis of BRCA1, BRCA2 & analysis of homologous recombination deficiency 0172U pathways, DNA, formalin-fixed paraffin-embedded tissue, algorithm quantif 0175U Psychiatry; genomic analysis panel, variant analysis of 15 genes Oncology, cell free DNA, targeted sequence analysis of 23 genes [single nucleotide variations, insertions & 0179U deletions, fusions without prior knowledge of partner/breakpoint, c Autoimmune, mRNA, gene expression profiling by quantitative RT-PCR, 17 genes, whole blood, reported as 0203U continuous risk score & classification of inflammatory bowel disease agg Oncology, mRNA, gene expression analysis of 593 genes for sequence variants & rearrangements, utilizing fine 0204U needle aspirate, reported as detected or not detected AR (androgen receptor) (eg, spinal and bulbar muscular atrophy, Kennedy disease, X chromosome inactivation), full sequence analysis, including small sequence changes in exonic and intronic regions, deletions, duplications, 0230U short tandem repeat (STR) expansions, mobile element insertions, and variants in non-uniquely mappable regions CACNA1A (calcium voltage-gated channel subunit alpha 1A) (eg, spinocerebellar ataxia), full gene analysis, 0231U including small sequence changes in exonic and intronic regions, deletions, duplications, short tandem repeat (STR) gene expansions, mobile element insertions, and variants in non-uniquely mappable regions CSTB (cystatin B) (eg, progressive myoclonic epilepsy type 1A, Unverricht-Lundborg disease), full gene analysis, 0232U including small sequence changes in exonic and intronic regions, deletions, duplications, short tandem repeat (STR) expansions, mobile element insertions, and variants in non-uniquely mappable regions FXN (frataxin) (eg, Friedreich ataxia), gene analysis, including small sequence changes in exonic and intronic 0233U regions, deletions, duplications, short tandem repeat (STR) expansions, mobile element insertions, and variants in non-uniquely mappable regions MECP2 (methyl CpG binding protein 2) (eg, Rett syndrome), full gene analysis, including small sequence 0234U changes in exonic and intronic regions, deletions, duplications, mobile element insertions, and variants in nonuniquely mappable regions PTEN (phosphatase and tensin homolog) (eg, Cowden syndrome, PTEN hamartoma tumor syndrome), full gene 0235U analysis, including small sequence changes in exonic and intronic regions, deletions, duplications, mobile element insertions, and variants in non-uniquely mappable regions SMN1 (survival of motor neuron 1, telomeric) and SMN2 (survival of motor neuron 2, centromeric) (eg, spinal 0236U muscular atrophy) full gene analysis, including small sequence changes in exonic and intronic regions, duplications and deletions, and mobile element insertions Cardiac ion channelopathies (eg, Brugada syndrome, long QT syndrome, short QT syndrome, catecholaminergic polymorphic ventricular tachycardia), genomic sequence analysis panel including ANK2, CASQ2, CAV3, KCNE1, 0237U KCNE2, KCNH2, KCNJ2, KCNQ1, RYR2, and SCN5A, including small sequence changes in exonic and intronic

regions, deletions, duplications, mobile element insertions, and variants in non-uniquely mappable regions

Oncology (Lynch syndrome), genomic DNA sequence analysis of MLH1, MSH2, MSH6, PMS2, and EPCAM,

including small sequence changes in exonic and intronic regions, deletions, duplications, mobile element

insertions, and variants in non- uniquely mappable regions



DCD:	Procedure codes related to the CMS Inpatient Only Li	st have been removed, however the inpatient admission care requires an authorization.
0239U	Targeted genomic sequence analysis panel, solid organ neoplasm, cell-free DNA, analysis of 311 or more genes interrogation for sequence variants, including substitutions, insertions, deletions, select rearrangements, and copy number variations	
0240U	Infectious disease, pathogen-specific RNA, 3 targets (severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), influenza A, influenza B), upper respiratory specimen, each	
0242U	Targeted genomic sequence analysis panel, solid organ neoplasm, cell-free circulating DNA analysis of 55-74 genes, interrogation for sequence variants, gene copy number amplif	
0244U	Oncology, DNA, comprehensive genomic profiling, 257 genes, interrogation for single-nucleotide variants, insertions/deletions, copy number alterations, gene rearrangements, tu	
0245U	Oncology, mutation analysis of 10 genes & 37 RNA fusions & expression of 4 mRNA markers using next- generation sequencing, fine needle aspirate, report includes associated risk	
0246U	Red blood cell antigen typing, DNA, genotyping of at least 16 blood groups with phenotype prediction of at least 51 red blood cell antigens	
0250U	Oncology, targeted genomic sequence DNA analysis of 505 genes, interrogation for somatic alterations, microsatellite instability & tumor-mutation burden	
0252U	Fetal aneuploidy short tandem repeat comparative analysis, fetal DNA from products of conception, reported as normal, monosomy, trisomy, or partial deletion/duplication, mosai	
0258U	Autoimmune (psoriasis), mRNA, next-generation sequencing, gene expression profiling of 50-100 genes, skin- surface collection using adhesive patch, algorithm reported as likelihood of response to psoriasis biologics	
0260U	Rare diseases (constitutional/heritable disorders), identification of copy number variations, inversions, insertions, translocations, and other structural variants by optical genome mapping	
0262U	Oncology (solid tumor), gene expression profiling by real-time RT-PCR of 7 gene pathways (ER, AR, PI3K, MAPK, HH, TGFB, Notch), formalin-fixed paraffin-embedded (FFPE), algorithm reported as gene pathway activity score	
0264U	Rare diseases (constitutional/heritable disorders), identification of copy number variations, inversions, insertions, translocations, and other structural variants by optical genome mapping	
0265U	Rare constitutional and other heritable disorders, whole genome and mitochondrial DNA sequence analysis, blood, frozen and formalin-fixed paraffin-embedded (FFPE) tissue, saliva, buccal swabs or cell lines, identification of single nucleotide and copy number variants	
0266U	Unexplained constitutional or other heritable disorders or syndromes, tissue-specific gene expression by whole transcriptome and next-generation sequencing, blood, formalin-fixed paraffin-embedded (FFPE) tissue or fresh frozen tissue, reported as presence or absence of splicing or expression changes	
0267U	Rare constitutional and other heritable disorders, identification of copy number variations, inversions, insertions translocations, and other structural variants by optical genome mapping and whole genome sequencing	,
0268U	Hematology (atypical hemolytic uremic syndrome [aHUS]), genomic sequence analysis of 15 genes, blood, buccal swab, or amniotic fluid	
0269U	Hematology (autosomal dominant congenital thrombocytopenia), genomic sequence analysis of 14 genes, blood, buccal swab, or amniotic fluid	
0270U	Hematology (congenital coagulation disorders), genomic sequence analysis of 20 genes, blood, buccal swab, o amniotic fluid	



0298U

fixed paraffin-embedded tissue, blood or bone marrow, comparative sequen

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. Hematology (congenital neutropenia), genomic sequence analysis of 23 genes, blood, buccal swab, or amniotic 0271U Hematology (genetic bleeding disorders), genomic sequence analysis of 51 genes, blood, buccal swab, or 0272U amniotic fluid, comprehensive Hematology (genetic hyperfibrinolysis, delayed bleeding), genomic sequence analysis of 8 genes (F13A1, F13B, 0273U FGA, FGB, FGG, SERPINA1, SERPINE1, SERPINF2, PLAU), blood, buccal swab, or amniotic fluid Hematology (genetic platelet disorders), genomic sequence analysis of 43 genes, blood, buccal swab, or 0274U amniotic fluid ernatorogy (inneriteu tiirorinoocytopenia), genomic sequence anatysis or 23 genes, ntoou, nuccat swan, o 0276U Hematology (genetic platelet function disorder), genomic sequence analysis of 31 genes, blood, buccal swab, or 0277U amniotic fluid Hematology (genetic thrombosis), genomic sequence analysis of 12 genes, blood, buccal swab, or amniotic 0278U Red blood cell antigen typing, DNA, genotyping of 12 blood group system genes to predict 44 red blood cell 0282U antigen phenotypes Oncology, response to radiation, cell-free DNA, quantitative branched chain DNA amplification, plasma, reported as radiation toxicity score 0285U 0286U CEP72, NUDT15 & TPMT gene analysis, common variants Oncology, DNA & mRNA, next-generation sequencing analysis of 112 genes, fine needle aspirate or formalin-0287U fixed paraffin-embedded tissue, algorithmic prediction of cancer recur Oncology, mRNA, quantitative PCR analysis of 11 genes & 3 reference genes, formalin-fixed paraffin-embedded 0288U tumor tissue, algorithmic interpretation reported as recurrence ris Neurology, mRNA, gene expression profiling by RNA sequencing of 24 genes, whole blood, algorithm reported as 02890 predictive risk score Pain management, mRNA, gene expression profiling by RNA sequencing of 36 genes, whole blood, algorithm 0290U reported as predictive risk score Psychiatry, mRNA, gene expression profiling by RNA sequencing of 144 genes, whole blood, algorithm reported 0291U as predictive risk score - Deny INV/N10 Psychiatry, mRNA, gene expression profiling by RNA sequencing of 72 genes, whole blood, algorithm reported as 0292U predictive risk score - Deny INV/N10 Psychiatry, mRNA, gene expression profiling by RNA sequencing of 54 genes, whole blood, algorithm reported as 0293U predictive risk score - Deny INV/N10 Longevity & mortality risk, mRNA, gene expression profiling by RNA sequencing of 18 genes, whole blood, 0294U algorithm reported as predictive risk score Oncology, protein expression profiling by immunohistochemistry of 7 proteins, with 4 clinicopathologic factors, 0295U utilizing formalin-fixed paraffin-embedded Oncology, gene expression profiling by RNA sequencing of at least 20 molecular features, saliva, algorithm 0296U reported as positive or negative for signature associated with mali Oncology, whole genome sequencing of paired malignant & normal DNA specimens, fresh or formalin-fixed 0297U paraffin-embedded tissue, blood or bone marrow Oncology, whole transcriptome sequencing of paired malignant & normal RNA specimens, fresh or formalin-



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bcbs	tmedicare.com	Procedure codes related to the CMS Inpatient Only List	t have been removed, however the inpatient admission care requires an authorization.
	Oncology, whole genome optical genome	mapping of paired malignant & normal DNA specimens, fresh frozen	
0299U	tissue, blood, or bone marrow, comparat	ve structural	
	Oncology, whole genome sequencing & o	ptical genome mapping of paired malignant & normal DNA specimens,	
0300U	fresh tissue, blood, or bone marrow, com	parative seq	
0301U	· · · · · · · · · · · · · · · · · · ·	, Bartonella henselae & Bartonella quintana, droplet digital PCR;	
	,	l, Bartonella henselae & Bartonella quintana, droplet digital PCR;	
0302U	following liquid enrichment		
		encing analysis, cell-free DNA, initial assessment to determine a patient	
0306U	specific panel for future comparisons to		
	Oncology, next-generation targeted sequ	encing analysis of a patient-specific panel, cell-free DNA, subsequent	
0307U	assessment with comparison to previous	y analyzed patient specime	
		sequencing analysis of 74 genes & analysis of CEA gene expression,	
0313U	pancreatic cyst fluid, algorithm reported	as a categorical result	
	Oncology, mRNA gene expression profilir	g by RT-PCR of 35 genes, utilizing formalin-fixed paraffin-embedded	
0314U	tissue, algorithm reported as a categorica		
		g by RT-PCR of 40 genes, utilizing formalin-fixed paraffin-embedded	
0315U	tissue, algorithm reported as a categorica		
	Nephrology, RNA expression by select tra	nscriptome sequencing, using pretransplant peripheral blood,	
0319U	algorithm reported as a risk score for earl	•	
		nscriptome sequencing, using posttransplant peripheral blood,	
0320U	algorithm reported as a risk score for acu		
		l, genitourinary pathogens, identification of 20 bacterial & fungal	
0321U	organisms & identification of 16 associate		
0323U		l, central nervous system pathogen, metagenomic next-generation	
	sequencing, cerebrospinal fluid, identific	ation of pathogenic bacteria	
0326U		el, solid organ neoplasm, cell-free circulating DNA analysis of 83 or	
	more genes, interrogation for sequence v	ariants, gene copy number a	
0329U	Oncology, exome & transcriptome seque	nce analysis for sequence variants, gene copy number amplifications &	
	deletions, gene rearrangements, microsa		
0330U	Infectious agent detection by nucleic acid	l, vaginal pathogen panel, identification of 27 organisms, amplified	
	probe technique, vaginal swab		
0331U		py number alterations & gene rearrangements utilizing DNA from blood	
	or bone marrow, report of clinically signif		
0332U		latory markers by quantitative polymerase chain reaction, whole blood,	
	reported as a high or low probability of re-	-	
0333U		carcinoma in high risk patients, analysis of methylation patterns on	
	circulating cell-free DNA plus measurem		
0334U		alysis, formalin-fixed paraffin embedded tumor tissue, DNA analysis,	
	84 or more genes, interrogation for seque		
0335U		analysis, including small sequence changes, copy number variants,	
	deletions, duplications, mobile element i		
0336U	· · · · · · · · · · · · · · · · · · ·	analysis, including small sequence changes, copy number variants,	
	deletions, duplications, mobile element i	nsertions, uniparental disomy,	



Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. Oncology, circulating plasma cell immunologic selection, identification, morphological characterization, & 0337U enumeration of plasma cells based on differential CD138, CD38, CD19, Oncology, circulating tumor cell selection, identification, morphological characterization, detection & 0338U enumeration based on differential EpCAM, cytokeratins -Oncology, mRNA expression profiling of HOXC6 & DLX1, reverse transcription polymerase chain reaction, first-0339U void urine following digital rectal examination, algorithm reported Oncology, analysis of minimal residual disease from plasma, with assays personalized to each patient based on 0340U prior next-generation sequencing of the patients tumor & germlin Fetal aneuploidy DNA sequencing comparative analysis, fetal DNA from products of conception, reported as 0341U normal, monosomy, trisomy, or partial deletion/duplication, mosaicism, Oncology, multiplex immunoassay of C5, C4, cystatin C, factor B, osteoprotegerin, gelsolin, IGFBP3, CA125 & 034211 multiplex electrochemiluminescent immunoassay for CA19-9, serum, di Oncology, exosome-based analysis of 442 small noncoding RNAs by quantitative reverse transcription 0343U polymerase chain reaction, urine, reported as molecular evidence of no-, low Hepatology, semiquantitative evaluation of 28 lipid markers by liquid chromatography with tandem mass 0344U spectrometry, serum, reported as at-risk for nonalcoholic steatohepatitis Psychiatry, genomic analysis panel, variant analysis of 15 genes, including deletion/duplication analysis of 0345U 0346U Beta amyloid, AB40 & AB42 by liquid chromatography with tandem mass spectrometry, ratio, plasma Drug metabolism or processing, whole blood or buccal specimen, DNA analysis, 16 gene report, with variant 0347U analysis & reported phenotypes Drug metabolism or processing, whole blood or buccal specimen, DNA analysis, 25 gene report, with variant 0348U analysis & reported phenotypes Drug metabolism or processing, whole blood or buccal specimen, DNA analysis, 27 gene report, with variant 0349U analysis, including reported phenotypes & impacted gene-drug interact Drug metabolism or processing, whole blood or buccal specimen, DNA analysis, 27 gene report, with variant 0350U analysis & reported phenotypes Infectious disease, biochemical assays, tumor necrosis factor related apoptosis-inducing ligand, interferon 0351U gamma-induced protein-10, & C-reactive protein, serum, algorithm re Infectious disease, multiplex amplified probe technique, for detection of bacterial vaginosis associated bacteria. 0352U algorithm reported nfectious agent detection by nucleic acid, Chlamydia trachomatis & Neisseria gonorrhoeae, multiplex amplified 0353U probe technique, urine, vaginal 0355U APOL1 (apolipoprotein L1) (eg, chronic kidney disease), risk variants (G1, G2) Oncology (oropharyngeal/anal), eval of 17 DNA biomarkers using droplet digital PCR, cell-free DNA, algorithm 0356U reported as a prognostic risk score for cancer recurrence Oncology (papillary thyroid cancer), gene-expression profiling via targeted hybrid capture-enrichment RNA 0362U sequencing of 82 content genes and 10 housekeeping genes, formalin-fixed Oncology, mRNA, geneexpression profiling by real-time quantitative PCR of 5 genes, utilizing urine, algorithm 0363U incorporates age, sex, smoking history, and macrohematuria freque Oncology (hematolymphoid neoplasm), genomic sequence analysis using multiplex (PCR) and next-generation 0364U sequencing with algorithm, quantification of dominant clonal sequence(s), reported as presence or absence of

minimal residual disease (MRD) with quantitation of disease burden, when appropriate



Updated: 07/01/2025

Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization.

	Procedure codes related to the CMS inpatient Only List	have been removed, however the inpatient admission care requires an authorization.
0365U	Oncology (bladder), analysis of 10 protein biomarkers (A1AT, ANG, APOE, CA9, IL8, MMP9, MMP10, PAI1, SDC1 and VEGFA) by immunoassays, urine, algorithm reported as a probability of bladder cancer	
0366U	Oncology (bladder), analysis of 10 protein biomarkers (A1AT, ANG, APOE, CA9, IL8, MMP9, MMP10, PAI1, SDC1	
	and VEGFA) by immunoassays, urine, algorithm reported as a probability of recurrent bladder cancer  Oncology (bladder), analysis of 10 protein biomarkers (A1AT, ANG, APOE, CA9, IL8, MMP9, MMP10, PAI1, SDC1	
0367U	and VEGFA) by immunoassays, urine, diagnostic algorithm reported as a risk score for probability of rapid	
	recurrence of recurrent or persistent cancer following transurethral resection	
	Oncology (colorectal cancer), evaluation for mutations of APC, BRAF, CTNNB1, KRAS, NRAS, PIK3CA, SMAD4,	
0368U	and TP53, and methylation markers (MYO1G, KCNQ5, C9ORF50, FL11, CLIP4, ZNF132 and TWIST1), multiplex quantitative polymerase chain reaction (qPCR), circulating cell-free DNA (cfDNA), plasma, report of risk score	
	for advanced adenoma or colorectal cancer	
	Infectious agent detection by nucleic acid (DNA and RNA), gastrointestinal pathogens, 31 bacterial, viral, and	
0369U	parasitic organisms and identification of 21 associated antibiotic-resistance genes, multiplex amplified probe	
	technique	
0370U	Infectious agent detection by nucleic acid (DNA and RNA), surgical wound pathogens, 34 microorganisms and	
	identification of 21 associated antibiotic resistance genes, multiplex amplified probe technique, wound swab	
	Infectious agent detection by nucleic acid (DNA or RNA), genitourinary pathogen, semiquantitative identification,	
0371U	DNA from 16 bacterial organisms and 1 fungal organism, multiplex amplified probe technique via quantitative	
	polymerase chain reaction (qPCR), urine  Infectious disease (genitourinary pathogens), antibiotic-resistance gene detection, multiplex amplified probe	
0372U	technique, urine, reported as an antimicrobial stewardship risk score	
	Infectious agent detection by nucleic acid (DNA and RNA), respiratory tract infection, 17 bacteria, 8 fungus, 13	
0373U	virus, and 16 antibiotic-resistance genes, multiplex amplified probe technique, upper or lower respiratory	
	specimen  Infectious agent detection by nucleic acid (DNA or RNA), genitourinary pathogens, identification of 21 bacterial	
0374U	and fungal organisms and identification of 21 associated antibiotic-resistance genes, multiplex amplified probe	
	technique, urine	
	Oncology (ovarian), biochemical assays of 7 proteins (follicle stimulating hormone, human epididymis protein 4,	
0375U	apolipoprotein A-1, transferrin, beta-2 macroglobulin, prealbumin [ie, transthyretin], and cancer antigen 125),	
	algorithm reported as ovarian cancer risk score  Oncology (prostate cancer), image analysis of at least 128 histologic features and clinical factors, prognostic	
0376U	algorithm determining the risk of distant metastases, and prostate cancer specific mortality, includes predictive	
	algorithm to androgen deprivation therapy response, if appropriate	
0378U	RFC1 (replication factor C subunit 1), repeat expansion variant analysis by traditional and repeat-primed PCR,	
	blood, saliva, or buccal swab  Targeted genomic sequence analysis panel, solid organ neoplasm, DNA (523 genes) and RNA (55 genes) by	
0379U	next generation sequencing, interrogation for sequence variants, gene copy number amplifications, gene	
	rearrangements, microsatellite instability, and tumor mutational burden	
0380U	Drug metabolism (adverse drug reactions and drug response), targeted sequence analysis, 20 gene variants and	
00000	CYP2D6 deletion or duplication analysis with reported genotype and phenotype	



BCB.	Procedure codes related to the CMS Inpatient Only Lis	t have been removed, however the inpatient admission care requires an authorization.
	Oncology (non-small cell lung cancer), next-generation sequencing with identification of single nucleotide	
0388U	variants, copy number variants, insertions and deletions, and structural variants in 37 cancer-related genes,	
	plasma, with report for alteration detection	
	Pediatric febrile illness (Kawasaki disease [KD]), interferon alpha inducible protein 27 (IFI27) and mast cell-	
0389U	expressed membrane protein 1 (MCEMP1), RNA, using reverse transcription polymerase chain reaction (RT-	
	qPCR), blood, reported as a risk score for KD	
	Oncology (solid tumor), DNA and RNA by next-generation sequencing, utilizing formalin-fixed paraffin-embedded	
0391U	(FFPE) tissue, 437 genes, interpretive report for single nucleotide variants, splice site variants,	
	insertions/deletions, copy number alterations, gene fusions, tumor mutational burden, and microsatellite	
	instability, with algorithm quantifying immunotherapy response score	
0392U	Drug metabolism (depression, anxiety, attention deficit hyperactivity disorder [ADHD]), gene-drug interactions, variant analysis of 16 genes, including deletion/duplication analysis of CYP2D6, reported as impact of gene-drug	
03920	interaction for each drug	
	Oncology (lung), multi-omics (microbial DNA by shotgun next generation sequencing and carcinoembryonic	
0395U	antigen and osteopontin by immunoassay), plasma, algorithm reported as malignancy risk for lung nodules in	
00000	early-stage disease	
	Oncology (non-small cell lung cancer), cell-free DNA from plasma, targeted sequence analysis of at least 109	
0397U	genes, including sequence variants, substitutions, insertions, deletions, select rearrangements, and copy	
	number variations	
	Gastroenterology (Barrett esophagus), P16, RUNX3, HPP1, and FBN1 DNA methylation analysis using PCR,	
0398U	formalin-fixed paraffin-embedded (FFPE) tissue, algorithm reported as risk score for progression to high-grade	
	dysplasia or cancer	
0401U	Cardiology (coronary heart disease [CAD]), 9 genes (12 variants), targeted variant genotyping, blood, saliva, or	
04010	buccal swab, algorithm reported as a genetic risk score for a coronary event	
	Oncology (prostate), mRNA, gene expression profiling of 18 genes, first-catch post-digital rectal examination	
0403U	urine (or processed first-catch urine), algorithm reported as percentage of likelihood of detecting clinically	
	significant prostate cancer	
0405U	Oncology (pancreatic), 59 methylation haplotype block markers, next-generation sequencing, plasma, reported	
	as cancer signal detected or not detected	
0.40011	Oncology (solid tumor), DNA (80 genes) and RNA (36 genes), by next-generation sequencing from plasma,	
0409U	including single nucleotide variants, insertions/deletions, copy number alterations, microsatellite instability, and	
<del></del>	fusions, report showing identified mutations with clinical actionability	
0410U	Oncology (pancreatic), DNA, whole genome sequencing with 5-hydroxymethylcytosine enrichment, whole blood or plasma, algorithm reported as cancer detected or not detected	
	Psychiatry (eg, depression, anxiety, attention deficit hyperactivity disorder [ADHD]), genomic analysis panel,	
0411U	variant analysis of 15 genes, including deletion/duplication analysis of CYP2D6	
	Oncology (hematolymphoid neoplasm), optical genome mapping for copy number alterations, aneuploidy, and	
0413U	balanced/complex structural rearrangements, DNA from blood or bone marrow, report of clinically significant	
	alterations	
	Oncology (lung), augmentative algorithmic analysis of digitized whole slide imaging for 8 genes (ALK, BRAF,	
0414U	EGFR, ERBB2, MET, NTRK1-3, RET, ROS1), and KRAS G12C and PD-L1, if performed, formalin-fixed paraffin-	
	embedded (FFPE) tissue, reported as positive or negative for each biomarker	
•		•



-	Procedure codes related to the CMS Inpatient Only List have b	been removed, however the inpatient admission care requires an authorization.
0416U	Infectious agent detection by nucleic acid (DNA), genitourinary pathogens, identification of 20 bacterial and fungal organisms, including identification of 20 associated antibiotic-resistance genes, if performed, multiplex amplified probe technique, urine	
0417U	Rare diseases (constitutional/heritable disorders), whole mitochondrial genome sequence with heteroplasmy detection and deletion analysis, nuclear-encoded mitochondrial gene analysis of 335 nuclear genes, including sequence changes, deletions, insertions, and copy number variants analysis, blood or saliva, identification and categorization of mitochondrial disorder-associated genetic variants	
0419U	Neuropsychiatry (eg, depression, anxiety), genomic sequence analysis panel, variant analysis of 13 genes, saliva or buccal swab, report of each gene phenotype	
0420U	Oncology (urothelial), mRNA expression profiling by real-time quantitative PCR of MDK, HOXA13, CDC2, IGFBP5, and CXCR2 in combination with droplet digital PCR (ddPCR) analysis of 6 single-nucleotide polymorphisms (SNPs) genes TERT and FGFR3, urine, algorithm reported as a risk score for urothelial carcinoma	
0421U	Oncology (colorectal) screening, quantitative real-time target and signal amplification of 8 RNA markers (GAPDH, SMAD4, ACY1, AREG, CDH1, KRAS, TNFRSF10B, EGLN2) and fecal hemoglobin, algorithm reported as a positive or negative for colorectal cancer risk	
0422U	Oncology (pan-solid tumor), analysis of DNA biomarker response to anti-cancer therapy using cell-free circulating DNA, biomarker comparison to a previous baseline pre-treatment cell-free circulating DNA analysis using next-generation sequencing, algorithm reported as a quantitative change from baseline, including specific alterations, if appropriate	
0423U	Psychiatry (eg, depression, anxiety), genomic analysis panel, including variant analysis of 26 genes, buccal swab, report including metabolizer status and risk of drug toxicity by condition	
0424U	Oncology (prostate), exosome based analysis of 53 small noncoding RNAs (sncRNAs) by quantitative reverse transcription polymerase chain reaction (RT qPCR), urine, reported as no molecular evidence, low-, moderate-or elevated-risk of prostate cancer	
0425U	Genome (eg, unexplained constitutional or heritable disorder or syndrome), rapid sequence analysis, each comparator genome (eg, parents, siblings)	
0426U	Genome (eg, unexplained constitutional or heritable disorder or syndrome), ultra-rapid sequence analysis	
0428U	Oncology (breast), targeted hybrid-capture genomic sequence analysis panel, circulating tumor DNA (ctDNA) analysis of 56 or more genes, interrogation for sequence variants, gene copy number amplifications, gene rearrangements, microsatellite instability, and tumor mutation burden	
0430U	Gastroenterology, malabsorption evaluation of alpha-1-antitrypsin, calprotectin, pancreatic elastase and reducing substances, feces, quantitative	
0431U	Glycine receptor alpha1 IgG, serum or cerebrospinal fluid (CSF), live cell-binding assay (LCBA), qualitative	
0433U	Oncology (prostate), 5 DNA regulatory markers by quantitative PCR, whole blood, algorithm, including prostate-specific antigen, reported as likelihood of cancer	
0434U	Drug metabolism (adverse drug reactions and drug response), genomic analysis panel, variant analysis of 25 genes with reported phenotypes	
0435U	Oncology, chemotherapeutic drug cytotoxicity assay of cancer stem cells (CSCs), from cultured CSCs and primary tumor cells, categorical drug response reported based on cytotoxicity percentage observed, minimum of 14 drugs or drug combinations	
0436U	Oncology (lung), plasma analysis of 388 proteins, using aptamer based proteomics technology, predictive algorithm reported as clinical benefit from immune checkpoint inhibitor therapy	



0452U

specific real-time PCR (LTE-qMSP), urine, reported as likelihood of bladder cancer

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. Psychiatry (anxiety disorders), mRNA, gene expression profiling by RNA sequencing of 15 biomarkers, whole 0437U blood, algorithm reported as predictive risk score Drug metabolism (adverse drug reactions and drug response), buccal specimen, gene-drug interactions, variant 0438U analysis of 33 genes, including deletion/duplication analysis of CYP2D6, including reported phenotypes and impacted gene drug interactions Cardiology, DNA, analysis of 5 single-nucleotide polymorphisms & 3 DNA methylation markers, qPCR & digital 0439U PCR, whole blood, algorithm reported as 4-tiered risk score for 3-year risk of symptomatic CHD Cardiology (coronary heart disease [CHD]), DNA, analysis of 10 single-nucleotide polymorphisms & 6 DNA 0440U methylation markers, qPCR & digital PCR, whole blood, algorithm reported as detected or not detected for CHD Infectious disease (bacterial, fungal, or viral infection), semiquantitative biomechanical assessment (via 0441U deformability cytometry), whole blood, with algorithmic analysis and result reported as an index Infectious disease (respiratory infection), Myxovirus resistance protein A (MxA) and C-reactive protein (CRP), 0442U fingerstick whole blood specimen, each biomarker reported as present or absent 0443U Neurofilament light chain (NfL), ultra-sensitive immunoassay, serum or cerebrospinal fluid Oncology (solid organ neoplasia), targeted genomic sequence analysis panet or 301 genes, interrogation or gene 0444U fusions, translocations, or other rearrangements, using DNA from formalin-fixed paraffin-embedded (FFPE) β-amyloid (Abeta42) and phospho tau (181P) (pTau181), electrochemiluminescent immunoassay (ECLIA), 0445U cerebral spinal fluid, ratio reported as positive or negative for amyloid pathology Autoimmune diseases (systemic lupus erythematosus [SLE]), analysis of 10 cytokine soluble mediator 0446U biomarkers by immunoassay, plasma, individual components reported with an algorithmic risk score for current disease activity Autoimmune diseases (systemic lupus erythematosus [SLE]), analysis of 11 cytokine soluble mediator 0447U biomarkers by immunoassay, plasma, individual components reported with an algorithmic prognostic risk score for developing a clinical flare Oncology (lung and colon cancer), DNA, qualitative, nextgeneration sequencing detection of single-nucleotide 0448U variants and deletions in EGFR and KRAS genes, formalin-fixed paraffinembedded (FFPE) solid tumor samples, reported as presence or absence of targeted mutation(s), with recommended therapeutic options Carrier screening for severe inherited conditions (eg, cystic fibrosis, spinal muscular atrophy, beta 0449U hemoglobinopathies [including sickle cell disease], alpha thalassemia), regardless of race or self-identified ancestry, genomic sequence analysis panel, must include analysis of 5 genes (CFTR, SMN1, HBB, HBA1, HBA2) Oncology (multiple myeloma), liquid chromatography with 0450U tandem mass spectrometry (LCMS/MS), monoclonal paraprotein sequencing analysis, serum, results reported as baseline presence or absence of detectable clonotypic peptides Oncology (multiple myeloma), LCMS/MS, peptide ion 0451U quantification, serum, results compared with baseline to determine monoclonal paraprotein abundance Oncology (bladder), methylated PENK DNA detection by linear target enrichment-quantitative methylation-



Updated: 07/01/2025

	Procedure codes related to the CMS Inpatient Unity List	t have been removed, however the inpatient admission care requires an authorization.
0453U	Oncology (colorectal cancer), cellfree DNA (cfDNA), methylationbased quantitative PCR assay (SEPTIN9, IKZF1, BCAT1, Septin9-2, VAV3, BCAN), plasma, reported as presence or absence of circulating tumor DNA (ctDNA)	
0454U	Rare diseases (constitutional/heritable disorders), identification of copy number variations, inversions, insertions, translocations, and other structural variants by optical genome mapping	
0455U	Infectious agents (sexually transmitted infection), Chlamydia trachomatis, Neisseria gonorrhoeae, and Trichomonas vaginalis, multiplex amplified probe technique, vaginal, endocervical, gynecological specimens, oropharyngeal swabs, rectal swabs, female or male urine, each pathogen reported as detected or not detected	
0456U	Autoimmune (rheumatoid arthritis), next-generation sequencing (NGS), gene expression testing of 19 genes, whole blood, with analysis of anticyclic citrullinated peptides (CCP) levels, combined with sex, patient global assessment, and body mass index (BMI), algorithm reported as a score that predicts nonresponse to tumor necrosis factor inhibitor (TNFi) therapy	
0457U	Perfluoroalkyl substances (PFAS)(eg, perfluorooctanoic acid, perfluorooctane sulfonic acid), 9 PFAS compounds by LC-MS/MS, plasma or serum, quantitative	
0458U	Oncology (breast cancer), S100A8 and S100A9, by enzymelinked immunosorbent assay	
0459U	β-amyloid (Abeta42) and total tau (tTau), electrochemiluminescent immunoassay (ECLIA), cerebral spinal fluid, ratio reported as positive or negative for amyloid pathology	
0460U	Oncology, whole blood or buccal, DNA single-nucleotide polymorphism (SNP) genotyping by real-time PCR of 24 genes, with variant analysis and reported phenotypes	
0461U	Oncology, pharmacogenomic analysis of single-nucleotide polymorphism (SNP) genotyping by real-time PCR of 24 genes, whole blood or buccal swab, with variant analysis, including impacted gene-drug interactions and reported phenotypes	
0462U	Melatonin levels test, sleep study, 7 or 9 sample melatonin profile (cortisol optional), enzyme-linked immunosorbent assay (ELISA), saliva, screening/preliminary	
0464U	Oncology (colorectal) screening, quantitative real-time target and signal amplification, methylated DNA markers, including LASS4, LRRC4 and PPP2R5C, a reference marker ZDHHC1, and a protein marker (fecal hemoglobin), utilizing stool, algorithm reported as a positive or negative result	
0465U	Oncology (urothelial carcinoma), DNA, quantitative methylationspecific PCR of 2 genes (ONECUT2, VIM), algorithmic analysis reported as positive or negative	
0466U	Cardiology (coronary artery disease [CAD]), DNA, genomewide association studies (564856 single-nucleotide polymorphisms [SNPs], targeted variant genotyping), patient lifestyle and clinical data, buccal swab, algorithm reported as polygenic risk to acquired heart disease	
0467U	Oncology (bladder), DNA, nextgeneration sequencing (NGS) of 60 genes and whole genome aneuploidy, urine, algorithms reported as minimal residual disease (MRD) status positive or negative and quantitative disease burden	
0468U	Hepatology (nonalcoholic steatohepatitis [NASH]), miR-34a-5p, alpha 2-macroglobulin, YKL40, HbA1c, serum and whole blood, algorithm reported as a single score for NASH activity and fibrosis	



	stmedicare.com	Procedure codes related to the CMS Inpatient Only List	have been removed, however the inpatient admission care requires an authorization.
	Rare diseases (const	titutional/heritable disorders), whole genome sequence analysis for chromosomal	
	abnormalities, copy	number variants, duplications/deletions, inversions, unbalanced translocations, regions of	
	homozygosity (ROH)	, inheritance pattern that indicate uniparental disomy (UPD), and aneuploidy, fetal sample	
0469U	(amniotic fluid, chor	ionic villus sample, or products of conception), identification and categorization of genetic	
	variants, diagnostic r	report of fetal results based on phenotype with maternal sample and paternal sample, if	
	performed, as compa	arators and/or	
	maternal cell contan	nination	
	Oncology (oropharyr	ngeal), detection of minimal residual	
0470U	disease by next-gene	eration sequencing (NGS) based	
	quantitative evaluati	on of 8 DNA targets, cell-free HPV 16 and 18 DNA from plasma	
0471U	Oncology (colorecta	l cancer), qualitative real-time PCR of 35 variants of KRAS and NRAS genes (exons 2, 3, 4),	
04/10	formalinfixed paraffii	n-embedded (FFPE), predictive, identification of detected mutations	
	Carbonic anhydrase	VI (CA VI), parotid specific/secretory protein (PSP) and salivary protein (SP1) IgG, IgM, and	
0472U	IgA antibodies, enzyr	me-linked immunosorbent assay (ELISA), semiqualitative, blood, reported as predictive	
	evidence of early Sjö	gren syndrome	
0473U	Oncology (solid tumo	or), nextgeneration	
	sequencing (NGS) of	DNA from formalin-fixed paraffinembedded (FFPE) tissue with comparative sequence	
0474U	analysis from a mato	hed normal specimen (blood or saliva), 648 genes, interrogation for sequence variants,	
04740	insertion and deletio	n alterations, copy number variants, rearrangements, microsatellite instability, and tumor-	
	mutation burden		
	Hereditary prostate of	cancerrelated disorders, genomic sequence analysis panel using next-generation	
0.47511	sequencing (NGS), S	anger sequencing, multiplex ligation-dependent probe amplification (MLPA), and array	
0475U	comparative genomi	c hybridization (CGH), evaluation of 23 genes and duplications/deletions when indicated,	
	pathologic mutations	s reported with a genetic risk score for prostate cancer	
	Drug metabolism, ps	sychiatry (eg, major depressive disorder, general anxiety disorder, attention deficit	
0476U	hyperactivity disorde	er [ADHD], schizophrenia), whole blood, buccal swab, and pharmacogenomic genotyping of	
	14 genes and CYP2D	6 copy number variant analysis and reported phenotypes	
	Drug metabolism, ps	sychiatry (eg, major depressive disorder, general anxiety disorder, attention deficit	
0477U	hyperactivity disorde	er [ADHD], schizophrenia), whole blood, buccal swab, and pharmacogenomic genotyping of	
04770	14 genes and CYP2D	6 copy number variant analysis, including impacted gene-drug interactions and reported	
	phenotypes		
	Oncology (non-small	l cell lung cancer), DNA and RNA, digital PCR analysis of 9 genes (EGFR, KRAS, BRAF, ALK,	
0478U	ROS1, RET, NTRK 1/2	2/3, ERBB2, and MET) in formalin-fixed paraffin-embedded (FFPE) tissue, interrogation for	
04760	single-nucleotide var	riants, insertions/deletions, gene rearrangements, and reported as actionable detected	
	variants for therapy s	selection	
0479U	Tau, phosphorylated, pTau217		
<u>-</u>	Infectious disease (b	acteria, viruses, fungi, and parasites), cerebrospinal fluid (CSF), metagenomic next-	
0480U	generation sequenci	ng (DNA and RNA),	
	bioinformatic analys	is, with positive pathogen identification	
	IDH1 (isocitrate dehy	ydrogenase 1 [NADP+]), IDH2 (isocitrate dehydrogenase 2 [NADP+]), and TERT (telomerase	
0481U	reverse transcriptase	e) promoter (eg, central nervous system [CNS] tumors), next-generation sequencing (single-	
	nucleotide variants [	SNV], deletions, and insertions)	



5655	Procedure codes related to the CMS Inpatient Only Li	st have been removed, however the inpatient admission care requires an authorization.
0482U	Obstetrics (preeclampsia), biochemical assay of soluble fms- like tyrosine kinase 1 (sFlt-1) and placental growt factor (PIGF), serum, ratio reported for sFlt- 1/PIGF, with risk of progression for preeclampsia with severe features within 2 weeks	1
0483U	Infectious disease (Neisseria gonorrhoeae), sensitivity, ciprofloxacin resistance (gyrA S91F point mutation), ora rectal, or vaginal swab, algorithm reported as probability of fluoroquinolone resistance	,
0484U	Infectious disease (Mycoplasma genitalium), macrolide sensitivity (23S rRNA point mutation), oral, rectal, or vaginal swab, algorithm reported as probability of macrolide resistance	
0485U	Oncology (solid tumor), cell-free DNA and RNA by next-generation sequencing, interpretative report for germline mutations, clonal hematopoiesis of indeterminate potential, and tumor-derived single-nucleotide variants, smainsertions/deletions, copy number alterations, fusions, microsatellite instability, and tumor mutational burden	
0486U	Oncology (pan-solid tumor), next-generation sequencing analysis of tumor methylation markers present in cell-free circulating tumor DNA, algorithm reported as quantitative measurement of methylation as a correlate of tumor fraction	
0487U	Oncology (solid tumor), cell-free circulating DNA, targeted genomic sequence analysis panel of 84 genes, interrogation for sequence variants, aneuploidycorrected gene copy number amplifications and losses, gene rearrangements, and microsatellite instability	
0488U	Obstetrics (fetal antigen noninvasive prenatal test), cellfree DNA sequence analysis for detection of fetal presence or absence of 1 or more of the Rh, C, c, D, E, Duffy (Fya), or Kell (K) antigen in alloimmunized pregnancies, reported as selected antigen(s) detected or not detected	
0489U	Obstetrics (single-gene noninvasive prenatal test), cellfree DNA sequence analysis of 1 or more targets (eg, CFTR, SMN1, HBB, HBA1, HBA2) to identify paternally inherited pathogenic variants, and relative mutation-dosage analysis based on molecular counts to determine fetal inheritance of maternal mutation, algorithm reported as a fetal risk score for the condition (eg, cystic fibrosis, spinal muscular atrophy, beta hemoglobinopathies [including sickle cell disease], alpha thalassemia)	
0490U	Oncology (cutaneous or uveal melanoma), circulating tumor cell selection, morphological characterization and enumeration based on differential CD146, high molecular–weight melanomaassociated antigen, CD34 and CD45 protein biomarkers, peripheral blood	
0491U	Oncology (solid tumor), circulating tumor cell selection, morphological characterization and enumeration base on differential epithelial cell adhesion molecule (EpCAM), cytokeratins 8, 18, and 19, CD45 protein biomarkers, and quantification of estrogen receptor (ER) protein biomarker–expressing cells, peripheral blood	
0492U	Oncology (solid tumor), circulating tumor cell selection, morphological characterization and enumeration base on differential epithelial cell adhesion molecule (EpCAM), cytokeratins 8, 18, and 19, CD45 protein biomarkers, and quantification of PD-L1 protein biomarker– expressing cells, peripheral blood	1
0493U	Transplantation medicine, quantification of donor-derived cell-free DNA (cfDNA) using next-generation sequencing, plasma, reported as percentage of donor-derived cell-free DNA	
0494U	Red blood cell antigen (fetal RhD gene analysis), next-generation sequencing of circulating cell-free DNA (cfDNA) of blood in pregnant individuals known to be RhD negative, reported as positive or negative	



Beb	Procedure codes related to the CMS Inpatient Only List have been removed, however the in	patient admission care requires an authorization.
0495U	Oncology (prostate), analysis of circulating plasma proteins (tPSA, fPSA, KLK2, PSP94, and GDF15), germline U polygenic risk score (60 variants), clinical information (age, family history of prostate cancer, prior negative prostate biopsy), algorithm reported as risk of likelihood of detecting clinically significant prostate cancer	
0496U	Oncology (colorectal), cell-free DNA, 8 genes for mutations, 7 genes for methylation by real-time RT-PCR, and 4 U proteins by enzyme-linked immunosorbent assay, blood, reported positive or negative for colorectal cancer or advanced adenoma risk	
0497U	Oncology (prostate), mRNA gene-expression profiling by real-time RT-PCR of 6 genes (FOXM1, MCM3, MTUS1, TTC21B, ALAS1, and PPP2CA), utilizing formalin-fixed paraffin-embedded (FFPE) tissue, algorithm reported as a risk score for prostate cancer	
0498U	Oncology (colorectal), next-generation sequencing for mutation detection in 43 genes and methylation pattern in 45 genes, blood, and formalin-fixed paraffin-embedded (FFPE) tissue, report of variants and methylation pattern with interpretation	
0499U	Oncology (colorectal and lung), DNA from formalin-fixed paraffin-embedded (FFPE) tissue, next-generation sequencing of 8 genes (NRAS, EGFR, CTNNB1, PIK3CA, APC, BRAF, KRAS, and TP53), mutation detection	
0500U	U Autoinflammatory disease (VEXAS syndrome), DNA, UBA1 gene mutations, targeted variant analysis (M41T, M41V, M41L, c.118-2A>C, c.118-1G>C, c.118-9_118-2del, S56F, S621C)	
0501U	U Oncology (colorectal), blood, quantitative measurement of cell-free DNA (cfDNA)	
0503U	Neurology (Alzheimer disease), beta amyloid (Aβ40, Aβ42, Aβ42/40 ratio) and tau-protein (ptau217, np-tau217, ptau217/nptau217 ratio), blood, immunoprecipitation with quantitation by liquid chromatography with tandem mass spectrometry (LC-MS/MS), algorithm score reported as likelihood of positive or negative for amyloid plaques	
0504U	U Infectious disease (urinary tract infection), identification of 17 pathologic organisms, urine, real-time PCR, reported as positive or negative for each organism	
0505U	U Infectious disease (vaginal infection), identification of 32 pathogenic organisms, swab, real-time PCR, reported as positive or negative for each organism	
0506U	Gastroenterology (Barrett's esophagus), esophageal cells, DNA methylation analysis by next-generation  U sequencing of at least 89 differentially methylated genomic regions, algorithm reported as likelihood for Barrett's esophagus	
0507U	U Oncology (ovarian), DNA, whole-genome sequencing with 5- hydroxymethylcytosine (5hmC) enrichment, using whole blood or plasma, algorithm reported as cancer detected or not detected	
0508U	Transplantation medicine, quantification of donor-derived cell-free DNA using 40 singlenucleotide  U polymorphisms (SNPs), plasma, and urine, initial evaluation reported as percentage of donor-derived cell-free  DNA with risk for active rejection	
0509U	Transplantation medicine, quantification of donor-derived cell-free DNA using up to 12 single-nucleotide  U polymorphisms (SNPs) previously identified, plasma, reported as percentage of donor-derived cell-free DNA with risk for active rejection	
0510U	U Oncology (pancreatic cancer), augmentative algorithmic analysis of 16 genes from previously sequenced RNA whole-transcriptome data, reported as probability of predicted molecular subtype	
0511U	U Oncology (solid tumor), tumor cell culture in 3D microenvironment, 36 or more drug panel, reported as tumor-response prediction for each drug	



Updated: 07/01/2025

Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization.

	Frocedure codes related to the GMS inpatient Only List	t nave been removed, nowever the inpatient admission care requires an authorization.
054011	Oncology (prostate), augmentative algorithmic analysis of digitized whole-slide imaging of histologic features for	
0512U	microsatellite instability (MSI) status, formalin-fixed paraffin-embedded (FFPE) tissue, reported as increased or	
	decreased probability of MSI-high (MSI-H)	
	Oncology (prostate), augmentative algorithmic analysis of digitized whole-slide imaging of histologic features for	
0513U	microsatellite instability (MSI) and homologous recombination deficiency (HRD) status, formal-infixed paraffin-	
	embedded (FFPE) tissue, reported as increased or	
	decreased probability of each biomarker	
	Gastroenterology (irritable bowel disease [IBD]), immunoassay for quantitative determination of adalimumab	
0514U	(ADL) levels in venous serum in patients undergoing adalimumab therapy, results reported as a numerical value	
	as micrograms per milliliter (μg/mL)	
	Gastroenterology (irritable bowel disease [IBD]), immunoassay for quantitative determination of infliximab (IFX)	
0515U	levels in venous serum in patients undergoing infliximab therapy, results reported as a numerical value as	
	micrograms per milliliter (μg/mL)	
0516U	Drug metabolism, whole blood, pharmacogenomic genotyping of 40 genes and CYP2D6 copy number variant	
00100	analysis, reported as metabolizer status	
	Therapeutic drug monitoring, 80 or more psychoactive drugs or substances, LC-MS/MS, plasma, qualitative and	
0517U	quantitative therapeutic minimally and maximally effective dose of prescribed and non-prescribed medications	
	,	
	Therapeutic drug monitoring, 90 or more pain and mental health drugs or substances, LC-MS/MS, plasma,	
0518U	qualitative and quantitative therapeutic minimally effective range of prescribed and non-prescribed medications	
0519U	Therapeutic drug monitoring, medications specific to pain, depression, and anxiety, LCMS/ MS, plasma, 110 or	
	more drugs or substances, qualitative and quantitative therapeutic minimally effective range of	
	prescribed, non-prescribed, and illicit medications in circulation	
0520U	Therapeutic drug monitoring, 200 or more drugs or substances, LCMS/MS, plasma, qualitative and quantitative	
	therapeutic minimally effective range of prescribed and non-prescribed medications	
1		

	High Tech Imaging		
Code	Code Description	Comments/Allowable Billed Groupings  **Note:  HTI Codes in this column will also require a Prior Authorization	
C8900	MRA Abdomen w/ or w/o Contrast	74185	
C8901	MRA Abdomen w/ or w/o Contrast	74185	
C8902	MRA Abdomen w/ or w/o Contrast	74185	
C8903	MRI Breast w/ or w/o Contrast (Unilateral)	77048	
C8905	MRI Breast w/ or w/o Contrast (Unilateral)	77048	
C8906	MRI Breast w/ or w/o Contrast (Bilateral)	77049	
C8908	MRI Breast w/ or w/o Contrast (Bilateral)	77049	
C8909	MRA Chest w/ or w/o Contrast	71555	
C8910	MRA Chest w/ or w/o Contrast	71555	
C8911	MRA Chest w/ or w/o Contrast	71555	
C8912	MRA LE w/ or w/o Contrast	73725	
C8913	MRA LE w/ or w/o Contrast	73725	



Chatt	neron Hill Circle Ianooga, Tennessee 37402	
bcbst	medicare.com  Procedure codes related to the CMS Inpatient Only List	have been removed, however the inpatient admission care requires an authorization
C8914	MRA LE w/ or w/o Contrast	73725
C8918	MRA Pelvis w/ or w/o Contrast	72198
C8919	MRA Pelvis w/ or w/o Contrast	72198
C8920	MRA Pelvis w/ or w/o Contrast	72198
C8928	Stress Echocardiography	93350
C8930	Stress Echocardiography	93351
C8931	MRA Spinal Canal w/ or w/o Contrast	72159
C8932	MRA Spinal Canal w/ or w/o Contrast	72159
C8933	MRA Spinal Canal w/ or w/o Contrast	72159
C8934	MRA Upper Extremity w/ or w/o Contrast	73225
C8935	MRA Upper Extremity w/ or w/o Contrast	73225
C8936	MRA Upper Extremity w/ or w/o Contrast	73225
C8937	MRI Breast w/ or w/o Contrast (Unilateral or Bilateral)	77048 or 77049
70336	MRI Temporomandibular Joint	70336
70450	CT Head/Brain	70450, 70460, 70470, 0722T
70480	CT Orbit	70480, 70481, 70482, 0722T
70486	CT Maxillofacial/Sinus	70486, 70487, 70488, 76380,0722T
70490	CT Soft Tissue Neck	70490, 70491, 70492, 0722T
70496	CT Angiography, Head	70496
70498	CT Angiography, Neck	70498
70540	MRI Orbit, Face, and/or Neck	70540, 70542, 70543, 0698T
70544	MRA Head	70544, 70545, 70546
70547	MRA Neck	70547, 70548, 70549
70551	MRI Brain (with or without Internal Auditory Canal views)	70551, 70552, 70553, 0698T
70554	Functional MRI Brain	70554, 70555
71250	CT Chest	71250, 71260, 71270, 0722T
71275	CT Angiography, Chest (non coronary)	71275
71550	MRI Chest	71550, 71551, 71552, 0698T
71555	MRA Chest(excluding myocardium)	71555
72125	CT Cervical Spine	72125, 72126, 72127, 0722T
72128	CT Thoracic Spine	72128, 72129, 72130, 0722T
72131	CT Lumbar Spine	72131, 72132, 72133, 0722T
72141	MRI Cervical Spine	72141, 72142, 72156, 0698T
72146	MRI Thoracic Spine	72146, 72147, 72157, 0698T
72148	MRI Lumbar Spine	72148, 72149, 72158, 0698T
72159	MRA Spinal Canal	72159
72191	CT Angiography, Pelvis	72191
72192	CT Pelvis	72192, 72193, 72194, 0722T
72196	MRI Pelvis	72195, 72196, 72197, 0698T
72198	MRA Pelvis	72198
73200	CT Upper Extremity	73200, 73201, 73202, 0722T
73206	CT Angiography, Upper Extremity	73206
73220	MRI Upper Extremity, other than Joint	73218, 73219, 73220, 0698T



S8037

MR Cholangiopancreatography

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. 73221 MRI Upper Extremity Joint 73221, 73222, 73223, 0698T 73225 MRA Upper Extremity 73225 73700 CT Lower Extremity 73700, 73701, 73702, 0722T 73706 CT Angiography, Lower Extremity 73706 73720 MRI Lower Extremity 73718, 73719, 73720, 73721, 73722, 73723, 0698T 72195, 72196, 72197, 73721, 73722, 73723, 0698T 73721 MRI Hip 73725 MRA Lower Extremity 73725 74150, 74160, 74170, 0722T 74150 CT Abdomen 74174 74174 CT Angiography, Abdomen and Pelvis 74175 CT Angiography, Abdomen 74175 74176 CT Abdomen and Pelvis Combination 74176, 74177, 74178, 0722T 74181 MRI Abdomen 74181, 74182, 74183, S8037, 0698T, 0724T 74185 MRA Abdomen 74185 74261 Diagnostic CT Colonoscopy (Virtual Colonoscopy, CT Colonography) 74261, 74262, 0722T 74263 Screening CT Colonoscopy (Virtual Colonoscopy, CT Colonography) 74263, 0722T 74712 Fetal MRI 74712, 74713 75557 MRI Heart 75557, 75559, 75561, 75563, 75565, 0698T 75571 Coronary Artery Ca Score, Heart Scan, Ultrafast CT Heart, Electron Beam CT 75571, S8092, 0722T 75572 CT Heart 75572, 0722T CT Heart congenital studies, non-coronary arteries 75573 75573, 0722T 75574 CTA coronary arteries (CCTA) 75574 75635 CT Angiography, Abdominal Aorta with Lower Extremity Runoff 75635 76380 Follow Up, Limited or Localized CT 76380, 70486, 70487, 70488 76390 MR Spectroscopy 76390, 0698T 76497 Unlisted Computed Tomography Procedure 76497, 0722T Unlisted Magnetic Resonance Procedure 76498 76498, 0698T 77046 MRI Breast 77046, 77047, 77048, 77049, 06981 77078 77078 CT Bone Density Study 77084 MRI Bone Marrow 77084 78429 Heart PET Scan with CT for Attenuation 78459, 78491, 78492, 78434, 78429, 78430, 78431, 78432, 78433 Myocardial Perfusion Imaging - Nuclear Cardiology Study 78451, 78452, 78453, 78454, 78466, 78468, 78469, 78481, 78483, 78499. 0742T 78451 78459 PET Scan, Heart 78459, 78491, 78492, 78434 78472 MUGA Scan 78472, 78473, 78494, 78496 78608 PET Scan, Brain 78608, 78609 78811, 78812, 78813, 78814, 78815, 78816 78813 PET Scan 78816 PET Scan with concurrently acquired CT for attenuation correction and anatomic, localization. 78811, 78812, 78813, 78814, 78815, 78816 93350, 93351, 93320, 93321, 93352, 93356 (PA only 93350 Stress Echocardiography required for 99356 and 93321 when related to a Stress Echo Test) 0042T Cerebral Perfusion Analysis CT 0042T G0219 PET imaging whole body, melanoma for non-covered indications G0219 PET imaging, any site, not otherwise specified G0235 G0235 G0252 PET imaging, initial diagnosis of breast cancer and/or surgical planning for breast cancer G0252

\$8037, 74181, 74182, 74183



S8042

MRI low field

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization.

\$8042

Hyperbaric Oxygen Therapy		
Code	Code Description	Comments
99183	Physician or other qualified health care professional attendance & supervision of hyperbaric oxygen therapy, per	
	session	
G0277	Hyperbaric oxygen under pressure, full body chamber, per 30 min interval	

#### **Inpatient Only Codes (IPO List)**

\*\*All Inpatient Services will require a Prior Authorization

#### **Intensive Outpatient Psych (IOP List)**

\*\*All IOP Services will require a Prior Authorization

Code	Code Description	Comments
20560	General introduction or removal procedures on the musculoskeletal system	
20561	General introduction or removal procedures on the musculoskeletal system	
22206	Osteotomy of spine, posterior or posterolateral approach, 3 columns, 1 vertebral segment (e.g.,	
22206	pedicle/vertebral body subtraction); thoracic	
22207	Osteotomy of spine, posterior approach, 3 columns, 1 vertebral segment (e.g., Pedicle/vertebral body	
22207	subtraction); lumbar	
	Osteotomy of spine, posterior or posterolateral approach, 3 columns, 1 vertebral segment (e.g.,	
22208	pedicle/vertebral body subtraction); each additional vertebral segment (list separately in addition to code for	
	primary procedure)	
22210	Osteotomy of spine, posterior or posterolateral approach, one vertebral segment; cervical	
22212	Osteotomy of spine, posterior or posterolateral approach, 1 vertebral segment; thoracic	
22214	Osteotomy of spine, posterior or posterolateral approach, 1 vertebral segment; lumbar	
22216	each additional vertebral segment (list separately in addition to code for primary procedure)	
22220	Osteotomy of spine, including discectomy anterior approach, single vertebral segment;cervical	
22224	Osteotomy of spine, including discectomy, anterior approach, single vertebral segment; lumbar	
22226	Osteotomy of spine, including discectomy, anterior approach, single vertebral segment; each additional	
22220	vertebral segment (List separately in addition to code for primary procedure	
22510	initiation in alumina of all installed which shall be a six	
22511	Percutaneous vertebroplasty (bone biopsy included when performed), 1 vertebral body, unilateral or bilateral	
22011	injection, inclusive of all imaging guidance; lumbosacral rercutaneous vertebrat body, unitaterat or bitaterat	
22512	initiation in the initiation of all interesting of violence and a stable and the analysis of the stable and the	
	Percutaneous vertebral augmentation, including cavity creation (fracture reduction and bone biopsy included	
22513	when performed) using mechanical device (e.g., kyphoplasty), 1 vertebral body, unilateral or bilateral	
	cannulation, inclusive of all imaging guidance; thoracic	
22514	whon performed) united manhanisal device (e.g. lambanisch) 1 vertebrei bedu uniteterel ex bilaterel	



bcb	Procedure codes related to the CMS Inpatient Only	List have been removed, however the inpatient admission care requires an authorization.
	Percutaneous vertebral augmentation, including cavity creation (fracture reduction and bone biopsy include when performed) using mechanical device (e.g., kyphoplasty), 1 vertebral body, unilateral or bilateral	d
22515	cannulation, inclusive of all imaging guidance; each additional thoracic or lumbar vertebral body (List separa	tely
22532	in addition to code for primary procedure) Arthrodesis, taterat extractavitary technique, including minimat discectority to prepare interspace totale than	101
22332	Arthrodesis, lateral extracavitary technique, including minimal discectomy to prepare interspace (other than	for
22533	decompression); lumbar  Anniouesis, taterat extracavitary technique, including minimat discertionly to prepare interspace (other than decompression); lumbar  Anniouesis, taterat extracavitary technique, including minimat discertionly to prepare interspace (other than decompression); lumbar	
22534	decompression by the region of humber and additional vertabral agreement (List apparently in addition to each of	1VI
22551	Arthrodesis, anterior interbody, including disc space preparation, discectomy, osteophytectomy and decompression of spinal cord and/or nerve roots; cervical below C2	
22552	Arthrodesis, anterior interbody, including disc space preparation, discectomy, osteophytectomy and	
22332	decompression of spinal cord	
22554	Arthrodesis, amenor interbody technique, including minimal discectority to prepare interspace (other than it	
22556	Arthrodesis, anterior interbody technique, including minimal discectomy to prepare interspace (other than for decompression); thoracic	ır
22558	Arthrodesis, anterior interbody technique, including minimal discectomy to prepare interspace (other than for decompression); lumbar	
22585	Arthrodesis, anterior interpody technique, including minima discectority to prepare interspace (other trial in	1
22590	Arthrodesis, posterior technique, craniocervical (occiput-C2)	
22595	Arthrodesis, posterior technique, atlas-axis (C1-C2)	
22600	Arthrodesis, posterior or posterolateral technique, single level; cervical below C2 segment	
22610	Arthrodesis, posterior or posterolateral technique, single level; thoracic (with or without lateral transverse technique)	
22612	Arthrodesis, posterior or posterolateral technique, single level; lumbar (with or without lateral transverse technique)	
22614	Artinouesis, posterior or posterorateral technique, single tevet, each additional vertebral segment (List	
22630	Arthrodesis, posterior interbody technique, including laminectomy and/or discectomy to prepare interspace (other than for decompression), single interspace: lumbar	
22632	(other than for decompression), single interspace; lumbar Antinodesis, posterior interbody technique, including taminectomy and/or discectomy to prepare interspace	
22633	Arthrodesis, componen posieno, or posietorarerar rechnique with postenor nicel botay rechnique incluting to a	
	Arthrodesis, combined posterior or posterolateral technique with posterior interbody technique including	
	laminectomy and/or discectomy sufficient to prepare interspace (other than for decompression), single	
22634	interspace and segment; each additional interspace and segment (List separately in addition to code for prin	nary
	procedure)	
22800	Arthrodesis, posterior, for spinal deformity, with or without cast; up to 6 vertebral segments	
22802	Arthrodesis, posterior, for spinal deformity, with or without cast; 7 to 12 vertebral segments	
22830	Exploration of spinal fusion	
-	Posterior non-segmental instrumentation (e.g., Harrington rod technique, pedicle fixation across 1 interspace	е,
22840	atlantoaxial transarticular screw fixation, sublaminar wiring at C1, facet screw fixation) (List separately in addition to code for primary procedure)	
	Posterior segmental instrumentation (e.g., pedicle fixation, dual rods with multiple hooks and sublaminar win	res);
22842	3 to 6 vertebral segments (List separately in addition to code for primary procedure)	



bcbs	stmedicare.com	Procedure codes related to the CMS Inpatient Only List have b	been removed, however the inpatient admission care requires an authorization.
22843	Posterior segmental i	instrumentation (e.g., pedicle fixation, dual rods with multiple hooks and sublaminar wires);	
22843	7 to 12 vertebral segn	ments (List separately in addition to code for primary procedure)	
20044	Posterior segmental i	instrumentation (e.g., pedicle fixation, dual rods with multiple hooks and sublaminar wires);	
22844	13 or more vertebral s	segments (List separately in addition to code for primary procedure)	
22845	Anterior instrumentat	ation; 2 to 3 vertebral segments (List separately in addition to code for primary procedure)	
22846	Anterior instrumentat	ation; 4 to 7 vertebral segments (List separately in addition to code for primary procedure)	
22847	Anterior instrumentat procedure)	ation; 8 or more vertebral segments (List separately in addition to code for primary	
22849	Reinsertion of spinal 1	fixation device	
22853		y Diometrianical device with integral antenor instrumentation for device anchoring, when	
22854	when performed to	etrarolulineananacia aeviče witinamėtė iartantentu antstronheina nonhibi taevice anchoning,	
	Total disc arthroplast	sty (artificial disc), anterior approach, including discectomy with end plate preparation	
22856	(includes osteophyte	ectomy for nerve root or spinal cord decompression and microdissection); single interspace,	
	cervical		
	Total disc arthroplast	sty (artificial disc), anterior approach, including discectomy with end plate preparation	
22858	(includes osteophyte	ectomy for nerve root or spinal cord decompression and microdissection); second level,	
	cervical (List separate	tely in addition to code for primary procedure)	
22859	Insertion of interverte	ebral biomechanical device to intervertebral disc space or vertebral body defect w/o	
22000	interbody arthrodesis	s, each contiguous defect **ADD ON CODE**	
22869	Insertion of interlamir	inar/interspinous process stabilization/distraction device, w/o open decompression or	
	fusion, including guid	dance when performed, lumbar; single level & 2nd level manninerspinous process stabilization device, who open decompression of	
22870	funion in alumbia or del	den en un ben de en en en el de le en ein els le control de le col	
22899	Under Other Procedu	ures on the Spine (Vertebral Column)	
23040		imeral joint,including exploration, drainage, or removal of foreign body	
23101	Arthrotomy, acromico	oclavicular joint or sternoclavicular joint, including biopsy and/or excision of torn cartilage	
23120	Claviculectomy; parti		
23130	Acromioplasty or acro	rominonectomy, partial with or without coracoacromial ligament release	
23334		nos, motados debridamentana synovestemy when periorinea, namerator general	
23335	Removal of prosthese component	ses, includes debridement and synovectomy when performed; humeral and glenoid	
23410	Repair of ruptured mu	nusculotendinous cuff (e.g., rotator cuff) open; acute	
23412	Repair of ruptured mu	nusculotendinous cuff (e.g., rotator cuff) open; chronic	
23420	Reconstruction of cor	omplete shoulder (rotator) cuff avulsion, chronic (includes acromioplasty)	
23430	Tenodesis of long ten	·	
23470	Arthroplasty, glenohu	umeral joint; hemiarthroplasty umerat joint; totat shoutder (glenoud and proximal numeral reptacement (e.g., total	
23472	A unoplasty, glenonu	amorar joint, total shoulder (Blenoid and proximal numeral replacement (e.g., total	
23473	Revision of total shou	ulder arthroplasty, including allograft when performed; humeral or glenoid component	
23474	Revision of total shou	ulder arthroplasty, including allograft when performed; humeral and glenoid component	
23700	Manipulation under a	anesthesia, shoulder joint, including application of fixation apparatus (dislocation excluded)	
27091	complicated, includir	ing total hip prosthesis, methylmethacrylate with or without insertion of spacer	
27125	Hemiarthroplasty, hip	ip, partial (e.g., femoral stem prosthesis, bipolar arthroplasty)	



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DCD	Procedure codes related to the CMS Inpatient Only List	have been removed, however the inpatient admission care requires an authorization.
27130	Arthroplasty, acetabular and proximal femoral prosthetic replacement (total hip arthroplasty), with or without	
2,100	autograft or allograft)	
27132	Conversion of previous hip surgery to total hip arthroplasty, with or without autograft or allograft	
27134	Revision of total hip arthroplasty; both components, with or without autograft or allograft	
27137	acetabular component only, with or without autograft or allograft	
27138	femoral component only, with or without autograft or allograft	
27275	Manipulation, hip joint, requiring general anesthesia	
27280	Arthrodesis, open, sacroiliac joint including obtaining bone graft, including instrumentation, when performed	
27299	Unlisted procedure, pelvis or hip joint	
27331	including joint exploration, biopsy, or removal of loose or foreign bodies	
27332	Arthrotomy, with excision of semilunar cartilage (meniscectomy) knee; medial or lateral	
27260	Partial excision (craterization, saucerization, or diaphysectomy) bone, femur, proximal tibia and/or fibula (e.g.,	
27360	osteomyelitis or bone abscess)	
27403	Arthrotomy with meniscus repair, knee	
27405	Repair, primary, torn ligament and/or capsule, knee; collateral	
27438	Arthroplasty, patella with prosthesis	
27442	Arthroplasty, femoral condyles or tibial plateau(s), knee;	
27445	Arthroplasty, knee, hinge prosthesis (e.g., Walldius type)	
27446	Arthroplasty, knee, condyle and plateau; medial OR lateral compartment medial AND lateral compartments with	
2/446	or without patella resurfacing (total knee arthroplasty)	
27447	medial AND lateral compartments with or without patella resurfacing (total knee arthroplasty)	
27486	Revision of total knee arthroplasty, with or without allograft; one component	
27487	femoral and entire tibial component	
27570	Manipulation of knee joint under general anesthesia (includes application of traction or other fixation devices)	
27599	Unlisted procedure, femur or knee	
29805	Arthroscopy, shoulder, diagnostic, with or without synovial biopsy	
29806	Arthroscopy, shoulder, surgical; capsulorrhaphy	
29807	Arthroscopy, shoulder, surgical; repair of SLAP lesion	
29819	Arthroscopy, shoulder, surgical; with removal of loose body or foreign body	
29820	Arthroscopy, shoulder, surgical; synovectomy, partial	
29821	Arthroscopy, shoulder, surgical; synovectomy, complete	
29822	Arthroscopy, shoulder, surgical; debridement, limited	
29823	Arthroscopy, shoulder, surgical; debridement, extensive	
29824	Arthroscopy, shoulder, surgical; distal claviculectomy including distal articular surface (Mumford procedure)	
29825	Arthroscopy, shoulder, surgical; with lysis and resection of adhesions, with or without manipulation	
	Arthroscopy, shoulder, surgical; decompression of subacromial space with partial acromioplasty, with	
29826	coracoacromial ligament (ie, arch) release, when performed (list separately in addition to code for primary	
	procedure)	
29827	Arthroscopy, shoulder, surgical; with rotator cuff repair Arthroscopy and treatment or uplantacture, proximal (plateau), unicondytal, includes internat inzation,	
29855	when perfermed (includes orthrospens)	



Lumbar/Sacral)

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. 29861 Arthroscopy, hip, surgical; with removal of loose body or foreign body Arthroscopy, hip, surgical; with debridement/shaving of articular cartilage (chondroplasty), abrasion 29862 arthroplasty, and/or resection of labrum 29863 Arthroscopy, hip, surgical; with synovectomy 29871 Arthroscopy, knee, surgical; for infection, lavage and drainage 29873 Arthroscopy, knee, surgical; with lateral release 29874 29875 Arthroscopy, knee, surgical; synovectomy, limited (e.g., plica or shelf resection) (separate procedure) 29876 Arthroscopy, knee, surgical; synovectomy, major, 2 or more compartments (e.g., medial or lateral) 29877 Arthroscopy, knee, surgical; debridement/shaving of articular cartilage (chondroplasty) 29879 Arthroscopy, knee, surgical; with meniscectomy (medial AND lateral, including any meniscal shaving) including 29880 debridement/shaving of articular cartilage (chondroplasty), same or separate compartment(s), when performed Arthroscopy, knee, surgical; with meniscectomy (medial OR lateral, including any meniscal shaving) including 29881 debridement/shaving of articular cartilage (chondroplasty), same or separate compartment(s), when performed 29882 Arthroscopy, knee, surgical; with meniscus repair (medial OR lateral) 29884 Arthroscopy, knee, surgical; with lysis of adhesions, with or without manipulation (separate procedure) 29888 Arthroscopically aided anterior cruciate ligament repair/augmentation or reconstruction 29914 Arthroscopy, hip, surgical; with femoroplasty (ie, treatment of cam lesion) 29915 Arthroscopy, hip, surgical; with acetabuloplasty (ie, treatment of pincer lesion) 29916 Arthroscopy, hip, surgical; with labral repair 62282 epidural, lumbar, sacral (caudal) Injection, of diagnostic or therapeutic substance, not including neurolytic substances, including needle or 62321 catheter placement, interlaminar epidural or subarachnoid, with imaging (62321-cervical/thoracic, 62323lumbar/sacral) Injection, of diagnostic or therapeutic substance, not including neurolytic substances, including needle or 62322 catheter placement, interlaminar epidural or subarachnoid, w/o imaging (62320-cervical/thoracic, 62322-Lumbar/Sacral) Injection, of diagnostic or therapeutic substance, not including neurolytic substances, including needle or 62323 catheter placement, interlaminar epidural or subarachnoid, with imaging (62321-cervical/thoracic, 62323lumbar/sacral) Injection, including indwelling catheter placement, continuous infusion or intermittent bolus, of diagnostic or 62324 therapeutic substance, not including neurolytic substances, int, w/o imaging (62324-cervical/thoracic, 62326-Lumbar/Sacral) Injection, including indwelling catheter placement, continuous infusion or intermittent bolus, of diagnostic or therapeutic substance, not including neurolytic substances, int, w/o imaging (62324-cervical/thoracic, 62326-62326 Lumbar/Sacral) intermittent bolus, of diagnostic or therapeutic substance, not including neurolytic substances, int, w/o imaging (62324-cervical/thoracic, 62326-Lumbar/Sacral) Injection, including indwelling catheter placement, continuous infusion or intermittent bolus, of diagnostic or 62327 therapeutic substance, not including neurolytic substances, int, with imaging (62325-cervical/thoracic, 62327-



63051

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. 62328 Injection, Drainage, or Aspiration Procedures on the Spine and Spinal Cord 62329 Injection, Drainage, or Aspiration Procedures on the Spine and Spinal Cord Implantation, revision or repositioning of tunneled intrathecal or epidural catheter, for long-term medication 62350 administration via an external pump or implantable reservoir/ infusion pump; without laminectomy 62360 Implantation or replacement of device for intrathecal or epidural drug infusion; subcutaneous reservoir 62361 Implant or replace device for drug infusion; non-programmable pump 62362 Implant or replace device for drug infusion; programmable pump 62365 Removal of subcutaneous reservoir or pump, previously implanted for intrathecal or epidural infusion 62367 62368 Electronic analysis programmable, implanted pump, for infusion, w reprogramming ammettoniv with exploration and/or decomplession or spinar cord and/or cauda equina, without racetectoniv 63001 Laminectomy with exploration and/or decompression of spinal cord and/or cauda equina, without facetectomy, 63003 foraminotomy or discectomy (e.g., spinal stenosis), 1 or 2 vertebral segments; thoracic 63005 Laminectomy with exploration and/or decompression of spinal cord and/or cauda equina, without facetectomy, 63011 foraminotomy or discectomy (e.g., spinal stenosis), 1 or 2 vertebral segments; sacral 63012 Laminectomy with exploration and/or decompression of spinal cord and/or cauda equina, without facetectomy, 63015 foraminotomy or discectomy (e.g., spinal stenosis), more than 2 vertebral segments; cervical anninectorny with exploration and/or decompression or spinal cord and/or cauda equina, without racetectorny, 63016 Laminectomy with exploration and/or decompression of spinal cord and/or cauda equina, without facetectomy, 63017 foraminotomy or discectomy (e.g., spinal stenosis), more than 2 vertebral segments; lumbar Laminotomy (hemilaminectomy), with decompression of nerve root(s), including partial facetectomy, 63020 foraminotomy and/or excision of herniated intervertebral disc. including open and endoscopically-assisted approaches; 1 interspace, cervical ammotorny (nermaninectorny), with decompression of herve root(s), including partial facetectorny, 63030 Laminotomy (hemilaminectomy), with decompression of nerve root(s), including partial facetectomy, foraminotomy and/or excision of herniated intervertebral disc, including open and endoscopically-assisted 63035 approaches; each additional interspace, cervical or lumbar (List separately in addition to code for primary procedure) 63042 Laminotomy (hemilaminectomy), with decompression of nerve root(s), including partial facetectomy, 63044 foraminotomy and/or excision of herniated intervertebral disc,re exploration, single interspace; each additional lumbar interspace (List separately in addition to code for primary procedure) 63045 aminectomy, facetectomy and foraminotomy (unilateral or bilateral with decompression of spinal cord, cauda. 63046 equina and/or nerve root[s], [e.g., spinal or lateral recess stenosis), single vertebral segment; thoracic anninectomy, racetectomy and roraminotomy (unitaterat or bitaterat with decompression or spinat cord, cadda 63047 Laminectomy, facetectomy and foraminotomy (unilateral or bilateral with decompression of spinal cord, cauda 63048 equina and/or nerve root[s], [e.g., spinal or lateral recess stenosis), single vertebral segment; each additional segment, cervical, thoracic, or lumbar (List separately in addition to code for primary procedure)



bcb	Procedure codes related to the CMS Inpatient Only	List have been removed, however the inpatient admission care requires an authorization.
63052	Laminectomy, facetectomy, or foraminotomy, during posterior interbody arthrodesis, lumbar; single vertebra segment	
63053	Learninectorny, racetectorny, or roranimotorny, during posterior interbody artificuesis, tumbar, each additional	
64582	Open implantation of hypoglossal nerve neurostimulator array, pulse generator, & distal respiratory sensor electrode or electrode array - Nurse Review	
64628	Thermat destruction of intraosseous pasiverteprat herve, including attimaging guidance, first 2 verteprat pour	55,
64629	Thermal destruction of intraosseous basivertebral nerve, including all imaging guidance; each additional vertebral body, lumbar or sacral	
63055	Transpedicutal approach with decompression of spinar cord, equina and/or nerve root(s) (e.g., nerniated	
63056	Transpedicular approach with decompression of spinal cord, equina and/or nerve root(s) (e.g., herniated intervertebral disc), single segment; lumbar (including transfacet, or lateral extraforaminal approach) (e.g., falateral herniated intervertebral disc)	r
63057	Transpedicular approach with decompression of spinal cord, equina and/or nerve root(s) (e.g., herniated intervertebral disc), single segment; each additional segment, thoracic or lumbar (List separately in addition toode for primary procedure)	0
63075	code for primary procedure) Discectionity, anticrior, with decompression of spinar cord and/or nerve root(s), including osteophytectomy,	
63077	Discectomy, anterior, with decompression of spinal cord and/or nerve root(s), including osteophytectomy; thoracic, single interspace	
63081	Vertebral corpectomy (vertebral body resection), partial or complete, anterior approach with decompression spinal cord and/or nerve root(s); cervical, single segment	of
63082	Vertebral corpectomy (vertebral body resection), partial or complete, anterior approach with decompression spinal cord and/or nerve root(s); cervical, each additional segment (List separately in addition to code for primary procedure)	of
63087	Vertebral corpectomy (vertebral body resection), partial or complete, combined thoracolumbar approach wit decompression of spinal cord, cauda equina or nerve root(s), lower thoracic or lumbar; single segment	n
63090	Vertebral corpectomy (vertebral body resection), partial or complete, transperitoneal or retroperitoneal approach with decompression of spinal cord, addition to code for primary procedure)	
63091	Vertebral corpectomy (vertebral body resection), partial or complete, transperitoneal or retroperitoneal approach with decompression of spinal cord, cauda equina or nerve root(s), lower thoracic, lumbar, or sacral each additional segment (List separately in addition to code for primary procedure)	;
63101	Vertebral corpectomy (vertebral body resection), partial or complete, lateral extracavitary approach with decompression of spinal cord and/or nerve root(s) (e.g., for tumor or retropulsed bone fragments); thoracic, single segment	
63102	Vertebral corpectomy (vertebral body resection), partial or complete, lateral extracavitary approach with decompression of spinal cord and/or nerve root(s) (e.g., for tumor or retropulsed bone fragments); lumbar, single segment	
63103	vertebrat corpectority (vertebrat body resection), partial or complete, taterat extracavitary approach with	
63172	Laminectomy with drainage of intramedullary cyst/syrinx; to subarachnoid space	
63185	Laminectomy with rhizotomy; 1 or 2 segments	
63251	Laminectomy for excision or occlusion of arteriovenous malformation of spinal cord; thoracic	
63265	Laminectomy for excision or evacuation of intraspinal lesion other than neoplasm, extradural; cervical	
63266	Laminectomy for excision or evacuation of intraspinal lesion other than neoplasm, extradural; thoracic	
63267	Laminectomy for excision or evacuation of intraspinal lesion other than neoplasm, extradural; lumbar	



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bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. 63268 Laminectomy for excision or evacuation of intraspinal lesion other than neoplasm, extradural; sacral 63271 Laminectomy for excision of intraspinal lesion other than neoplasm, intradural; thoracic 63276 Laminectomy for biopsy/excision of intraspinal neoplasm; extradural, thoracic 63277 Laminectomy for biopsy/excision of intraspinal neoplasm; extradural, lumbar 63280 Laminectomy for biopsy/excision of intraspinal neoplasm; intradural, extramedullary, cervical 63281 Laminectomy for biopsy/excision of intraspinal neoplasm; intradural, extramedullary, thoracic 63282 Laminectomy for biopsy/excision of intraspinal neoplasm; intradural, extramedullary, lumbar Laminectomy for biopsy/excision of intraspinal neoplasm; intradural, intramedullary, cervical 63285 Vertebral corpectomy (vertebral body resection), partial or complete, for excision of intraspinal lesion, single 63300 segment; extradural, cervical 63308 63650 Percutaneous implantation of neurostimulator electrode array, epidural 63655 Laminectomy for implantation of neurostimulator electrodes, plate/paddle, epidural Removal of spinal neurostimulator electrode percutaneous array(s), including fluoroscopy, when performed 63661 63662 Revision including replacement, when performed, of spinal neurostimulator electrode percutaneous array(s), 63663 including fluoroscopy, when performed iacement, when performed, or spinal neurostimulator electrode plate/paddiers) placed 63664 63685 Insertion or replacement of spinal neurostimulator pulse generator or receiver, direct or inductive coupling 63688 Revision or removal of implanted spinal neurostimulator pulse generator or receiver 64479 Injection, anesthetic agent and/or steroid, transforaminal epidural; cervical or thoracic, single level 64480 64483 Injection, anesthetic agent and/or steroid, transforaminal epidural; lumbar or sacral, single level 64484 Injection, anesthetic agent and/or steroid, transforaminal epidural; lumbar or sacral, each additional level Injection(s), diagnostic or therapeutic agent, paravertebral facet (zygapophyseal) joint (or nerves innervating that 64490 joint) with image guidance (fluoroscopy or CT), cervical or thoracic; single level njection(s), uragnostic or therapeutic agent, paravertebrat racet (zygapopnyseat) join 64491 Injection(s), diagnostic or therapeutic agent, paravertebral facet (zygapophyseal) joint (or nerves innervating that 64492 joint) with image guidance (fluoroscopy or CT), cervical or thoracic; third and any additional level(s) (List separately in addition to code for primary procedure) (5), ulagnostic or therapeutic agenit, paravertebrat facet (zygapophyseat) joint (or nerves innervating tha 64493 Injection(s), diagnostic or therapeutic agent, paravertebral facet (zygapophyseal) joint (or nerves innervating that 64494 joint) with image guidance (fluoroscopy or CT), lumbar or sacral; second level (List separately in addition to code njection(5), diagnostic of therapeutic agent, paravertebrat facet (zygapopnyseat) joint (or nerves innervating tha 64495 Injection, anesthetic agent; stellate ganglion (cervical sympathetic) 64510 64520 Injection, anesthetic agent; lumbar or thoracic (paravertebral sympathetic) Destruction by neurolytic agent, paravertebral facet joint nerve(s), with imaging guidance (fluoroscopy or CT); 64633 cervical or thoracic, single facet joint struction by neurotytic agent, paravertebrat facet joint nerve(s), with imaging guidance (itdoroscopy or Gr), 64634 Destruction by neurolytic agent, paravertebral facet joint nerve(s), with imaging guidance (fluoroscopy or CT); 64635 lumbar or sacral, single facet joint pestruction by néurotytic agent, paravertebrat racet joint nerve(s), with imaging guidance (ituoroscopy or Gr), 64636



Updated: 07/01/2025 bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. 97810 Acupuncture, one or more needles, without electrical stimulation 97811 Each additional 15 minutes of personal one-on-one contact with the patient 97813 Acupuncture, one or more needles, with electrical stimulation 97814 Each additional 15 minutes of personal one-on-one contact with the patient Posterior vertebral joint(s) arthroplasty (e.g., facet joint[s] replacement), including facetectomy, laminectomy, 0202T foraminotomy, and vertebral column fixation, injection of bone cement, when performed, including fluoroscopy, single level, lumbar spine Intravertebral body fracture augmentation with implant C1062 C1767 Generator, neurostimulator (implantable), nonrechargeable C1778 Lead, neurostimulator (implantable) Arthroscopy, shoulder, surgical; with implantation of subacromial spacer, includes debridement, subacromial C9781 decompression, acromioplasty, & biceps tenodesis when performed Vascular embolization or occlusion procedure with use of a pressure-generating catheter (e.g., one-way valve, C9797 intermittently occluding). **Non-Emergency Ambulance** Code Code Description Comments Modifiers HH, HN, or NH do not require an authorization A0426 Ambulance service, advance life support, non-emergency level 1 Modifiers HH, HN, or NH do not require an authorization A0428 Ambulance service, basic life support, non-emergency (BLS) **Outpatient Substance Abuse Code Description** Code Comments 90846 Family psychotherapy, 50 min 90847 Family psychotherapy, 50 min 90853 Group psychotherapy; other than multiple family Physical Therapy, Occupational Therapy, Speech Therapy and Chiropractic Code **Code Description** Comments 92507 Treatment of speech, language, voice, communication &/or auditory disorder; individual 92508 Treatment of speech, language, hearing disorder; group 92524 Behavioral & qualitative analysis of voice & resonance 92526 Treatment of swallowing dysfunction &/or oral function for feeding 92609 Therapeutic services use of speech-generating device, program & modify 95851 Procedure Canalith repositioning procedure, per day 95992 Assessment of aphasia w interpretation & report 96105 Standard cognitive performance testing per hour of a qualified health care professional time, both face-to-face 96125 with the patient and time interpreting test results 97012 Traction, mechanical 97016 Vasopneumatic devices 97018 Paraffin bath 97022 Whirlpool



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97024	Diathermy	
97032	Application to a modality to 1 or more areas: electrical stimulation (manual) each 15 minutes	
97033	lontophoresis, each 15 minutes	
97034	Contrast baths, each 15 minutes	
97035	Ultrasound, each 15 minutes	
97039	Unlisted modality (specify type and time if constant attendance)	
97110	Therapeutic procedure, 1 or more areas	
97112	Neuromuscular reeducation of movement, balance, coordination, kinesthetic sense, posture, and/	or
37112	proprioception for sitting and/or standing activities	
97113	Aquatic therapy with therapeutic exercises	
97116	Gait training (includes stair climbing)	
97124	Massage, including effleurage, petrissage and/or tapotement (stoking, compression, percussion)	
97129	Therapeutic interventions that focus on cognitive function & compensatory strategies to manage pe	
	activity, direct patient contact; initial 15 minutes	
97130	a skinike, alima skinika a mata skinika shi a sala salaliki a mali 4 Finalime ka	Tormunos or
97139	Unlisted therapeutic procedure (specify)	
97140	Manual therapy techniques, 1 or more regions, each 15 mins	
97150	Therapeutic procedure(s), group ( 2 or more individuals)	
97530	Therapeutic activities, direct (one to one) patient contact by the provider, each 15 minutes	
97533	Sensory integrative techniques to enhance sensory processing and promote adaptive responses to	
	environmental demands, direct (one to one) patient contact by the provider, each 15 minutes	
97535	Self-care/home management training, direct (one to one) patient contact by the provider, each 15 r	
97537	Community/work reintegration training, direct (one to one) patient contact by the provider, each 15	minutes
97542	Wheelchair management, each 15 minutes	
97597	Debridement, when performed and instructions for ongoing care, per session, total wound surface	
07500	square cm or less   Debitu open wound incr topic app(s), wound assess use or w/poot, when pen + inst(s) for ongoing t	Does not require a prior auth when billed with POS 11
97598	And all 1 1 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Does not require a prior auth when bitted with POS 11
97605	Negative pressure wound therapy, including tropical applications, wound assessment, and instruct	ions for
07000	ongoing care, per session, total wound surface area less than or equal to 50 square cm	
97606 97750	Total wound surface area greater than 50 square cm	
97760	Physical performance test or measurement with written report, each 15 minutes	
97761	Orthotic management and training, each 15 minutes  Prosthetic training, each 15 minutes	
97763	Checkout for orthotic/ prosthetic use, established patient, each 15 minutes	
98940	Chiropractic manipulative treatment spinal, 1-2 regions	Prior Authorization for MedAdvantage Only
98941	Chiropractic manipulative treatment spinal, 3-4 regions	Prior Authorization for MedAdvantage Only
98942	Chiropractic manipulative treatment spinal, 5 regions  Chiropractic manipulative treatment spinal, 5 regions	Prior Authorization for MedAdvantage Only  Prior Authorization for MedAdvantage Only
G0151	Services performed by a qualified physical therapist in the home health or hospice setting, each 15	
G0152		
G0153	Services performed by a qualified occupational therapist in the home health or hospice setting, each services performed by a quantied speech-tanguage pathologist in the home health or hospice setting.	g, each 13
G0157	Services performed by a qualified physical therapist in the home health or hospice setting, each 15	minutes
	Services performed by a qualified physical therapist in the home health setting, in the establishmer	
G0158	of a safe and effective physical therapy maintenance program each 15 minutes	
L	1	l



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	Trocodure codes related to the of 16 inputiont only Elec	thave been removed, nowever the inpatient admission care requires an authorization.
G0159	Services performed by a qualified physical therapist, in the home health setting, in the establishment or delivery	
	of a safe and effective physical therapy maintenance program, each 15 minute	
G0160	Services performed by a qualified occupational therapist in the home health setting, in the establishment or	
00100	delivery of a safe and effective physical therapy maintenance program each 15 minutes	
G0162	Skilled services by an RN in the delivery of mgt & eval of the POC; EACH 15 mins (Pt's underlying cond/compl	
00102	requires an RN to ensure that ess non-skilled care achieve	
G0176	Activity therapy, for care & treatment of disabling mental health problem	
	Electrical stimulation (unattended) to one or more areas, for chronic Stage III and Stage IV pressure ulcers,	
G0281	arterial ulcers, diabetic ulcers, and venous stasis ulcers not demonstrating measurable signs of healing after 30	
	days of conventional care, as part of a therapy plan of care	
G0283	Electrical stimulation (unattended), to one or more areas for indication(s) other than wound care, as part of	
00203	therapy plan of care	
G0299	Direct skilled nursing services of RN in HH or hospice setting, each 15 min	
G0300	Direct skilled nursing services of LPN in HH or hospice setting, each 15 min	
G0493	Skilled services of Registered Nurse for observation & assessment of patient's condition, each 15 min	
G0494		
	Skilled services of Licensed Practical Nurse for observation & assessment of patient's condition, each 15 min	
G0495	Skilled services of Registered Nurse, in training and/or education of patient or family member, in HH or hospice	
	setting, each 15 min	
S9123	Nursing care, in the home; RN per hour (use for gen nsg care only (not to be used when 99500-99602 can be	
39123	used)	

	Potential Cosmetic/Investigational/Non-Covered Procedures			
Code	Code Description	Comments		
11980	Sub-Q hormone pellet implantation			
15820	Blepharoplasty, lower eyelid			
15821	Blepharoplasty, lower eyelid; with extensive herniated fat pad			
15822	Blepharoplasty, upper eyelid			
15823	Blepharoplasty, upper eyelid; with excessive skin weighting down lid			
15830	Excision, excessive skin and subcutaneous tissue (includes lipectomy); abdomen, infraumbilical			
15650	panniculectomy			
15832	Excision, excessive skin & subcutaneous tissue; thigh			
15834	Excision, excessive skin & subcutaneous tissue; hip			
15839	Excision, excessive skin & subcutaneous tissue; other area			
15877	Suction assisted lipectomy; trunk			
19300	Mastectomy for gynecomastia			
19316	Mastopexy			
19318	Reduction mammaplasty			
19325	Mammaplasty, augmentation; with prosthetic implant			
19328	Removal of intact mammary implant			
19340	Insert breast prosthesis immediately following surgery			
19342	Delayed insertion of breast prosthesis following mastopexy, mastectomy or in reconstruction			



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19395 Nepticareoristic professor from the Company of the Company o	bebs	Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requ	uires an authorization.
19377 Open periprosthetic capsaudomy, treast 19390 Periprosthetic capsaudomy, treast 19391 Periprosthetic capsaudomy, treast 19391 Periprosthetic capsaudomy, treast 19392 Periprosthetic capsaudomy, treast 1940 Periprosthetic capsaudomy, treast 1940 Periprosthetic capsaudomy, treast 1940 Periprosthetic capsaudomy, treast vestibular stancius by multiple surgical approaches including internal fixation, interdernal fixation, androw wings of dentures or spinits 1940 Periprosthetic capsaudomy, treast vestibular stancius 1940 Periprostatio capsaudomy, treast vestibular stancius 1940 Peri	19350	Nipple/areola reconstruction	
1937. Performsthetic capabilisetic mysuline to design of processing to determine the season of processing the season of processing of the season of processing of design of processing of design of processing of design	19364	Breast reconstruction w free flap	
19380 Revision for constructed preast 21487 Open fraament manadibular fracture; witherdental fixation  21470 Open fraament manadibular fracture by multiple surgical approaches including internal fixation, interdental fixation, under which good produces of splints  30420 Amplitudes, primary including major speat repair  30420 Amplitudes, primary including major speat repair  30520 Septipolisty, or vir occurring sections, contouring or replacement w graft  30520 Septipolisty, or vir occurring sections, contouring or replacement w graft  33880 Heart transplant, with without recipient cardiscorus  33891 Heart transplant, with without recipient cardiscorus  43840 Heart transplant, with without recipient cardiscorus  53840 Heart transplant, with without recipient cardiscorus  538410 Heart transplant, with without recipient cardiscorus  538411 Heart transplant, with without recipient vein, limb or trunk  538417 Heart transplant, with without recipient vein, limb or trunk  538417 Heart transplant, without recipient vein, limb or trunk  538418 Heart transplant, without recipient vein, limb or trunk  538418 Heart transplant transplant vein, limb or trunk  538419 Heart transplant transplant vein, limb or trunk  538419 Heart transplant transplant vein, limb	19370	Open periprosthetic capsulotomy, breast	
21470 Open treatment mandibular fracture; witerdental facilition 21470 Open treatment of complicated mandibular fracture by multiple surgical approaches including internal fixation, interdental fixation, and/or wiring of dentures or splints 30426 Rhitsplasty, primary: including major septat repair 30426 Septoplasty, wo rive cartilage scoring, comouning or replacement wight of spain of maskin visitabular stemonis 30520 Septoplasty, wo rive cartilage scoring, comouning or replacement wight. 32880 International complete scoring, comouning or replacement wight. 32891 International complete scoring, comouning or replacement wight. 32891 International complete scoring, comouning or replacement wight. 33845 International complete scoring, comouning or replacement wight. 33945 International complete scoring, comouning or replacement wight. 34946 International complete scoring, complete sco	19371	Periprosthetic capsulectomy, breast	
Open treatment of complicated manifolish and received by multiple surgical approaches including internal fixation, interested facilitation, and/or writing of entures or splints  30420 Rhinoplasty, primary, including major septal repair  30520 Septipolasty, or vin cartilage scoring, contouring or replacement w graft  Transcatheter implantation writes pulmonary arrey pressure sensor for long-term hemodynamic monitoring, including major replacement w graft  Transcatheter implantation writes pulmonary arrey pressure sensor for long-term hemodynamic monitoring, including major the primary pressure sensor for long-term hemodynamic monitoring, including major the primary pressure sensor for long-term hemodynamic monitoring, including major the primary pressure sensor for long-term hemodynamic monitoring, including major the primary pressure sensor for long-term hemodynamic monitoring, including major the primary pressure sensor for long-term hemodynamic monitoring, including major the primary pressure sensor for long-term hemodynamic monitoring, including major the primary pressure of the primary pressure sensor for long-term hemodynamic monitoring, including major the primary pressure sensor for long-term hemodynamic monitoring, including major the primary pressure sensor for long-term hemodynamic monitoring, general pressure sensor for long-term hemodynamic monitoring, general pressure sensor in the pressure sensor for long-term hemodynamic monitoring, general pressure sensor in the pressure sensor in the pressure sensor in the pressure sensor for long-term hemodynamic monitoring, general sensor in the pressure sensor in the pressure sensor for long-term hemodynamic monitoring, general sensor in the pressure sensor in the	19380	Revision of reconstructed breast	
Intercental fization, and/or writing of dentures or splints	21462	Open treatment mandibular fracture; w interdental fixation	
interdental fastion, and/or wing of dentures or splints 304020 30405 30406 30406 30406 30400 304	21/170	Open treatment of complicated mandibular fracture by multiple surgical approaches including internal fixation,	
Septial of nasal vestibular stenosis		interdental fixation, and/or wiring of dentures or splints	
30520 Septoplasty, wo rwo cartillage scoring, contouring or replacement warft Transcatheter implantation wireless pulmonary aftery pressure sensor for long-term hemodynamic monitoring, include polyments, calibration of sensor, right heart - Deny INV/N10  33845 Heart transplant, with/without recipient cardiectomy injection of non-compounded foam sclerosant with uttrasound compression maneuvers to guide dispersion of the injection, compounded foam sclerosant with uttrasound compression maneuvers to guide dispersion of the injection of officerosant for spider veins, sclerosant with uttrasound compression maneuvers to guide dispersion of the injection of officerosant for spider veins, with uttrasound compression maneuvers to guide dispersion of the injection of sclerosant for spider veins, with uttrasound compression maneuvers to guide uspersion of the injection of sclerosant for spider veins, limb or trunk  36468 Injection of sclerosant for spider veins, limb or trunk  38470 In Sclerosing solution; single vein.  38471 In Sclerosing solution; single vein.  38478 Endowenous ablation this incompretent vein- 1st vein radiofreq trid  38478 Endowenous ablation this incompretent vein; dust using guidance monitoring, percut laser 1st vein and a senting and a senting produce and monitoring,  and a senting produce of the senting guidance monitoring,  and a senting guidance and monitoring,  percutaneous; single vein term the access site, inclusive of all imaging guidance and monitoring,  percutaneous; single vein term of the access site, inclusive of all imaging guidance and monitoring,  percutaneous; subsequent vincing 1 practed of in a single guidance and monitoring,  percutaneous; subsequent vincing 1 practed of in a single guidance and monitoring,  percutaneous; subsequent vincing 1 practed of in a single guidance and monitoring,  percutaneous; subsequent vincing 1 practed of in a single guidance and monitoring,  percutaneous; subsequent vincing 1 practed of in a single guidance and monitoring,  and a sensor of the senso	30420	Rhinoplasty, primary; including major septal repair	
Transcatheter implantation wireless pulmonary artery pressure sensor for long-term hemodynamic monitoring, incl deplayment & calibration of sensor, right heart. Denyi NV/N10  33945 Heart transplant, withwhithout recipient cardiectomy injection of non-compounded foam sclerosant with uttrasound compression maneuvers to guide dispersion of the injectate, incluse of all imaging guidance and monitoring the treatment of a single, incompetent extremity viel with the injectate, incluse of all imaging guidance and monitoring the treatment of a single, incompetent extremity viel miground or non-compounded roam sclerosant with utdrasound compression maneuvers to guide dispersion of the injectate, incluse of all imaging guidance amonitoring the treatment of a single, incompetent extremity viel miground or non-compounded roam sclerosant with utdrasound compression maneuvers to guide dispersion of the injectate, inclusive of all imaging guidance and sclerosant for spider veins, limb or trunk  38466 Miscolarosant for spider veins, limb or trunk  38471 Mis sclerosing solution; single vein.  38475 Endovenous ablation the incompetent vein; 2nd, subset veins  38476 Endovenous ablation the incompetent vein; 2nd, subset veins  38479 Endovenous ablation the reapy of month of a competent vein, extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; site viein interated  Endovenous ablation therapy of incompetent vein, extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; subsequent vein(s) treated in a single active treated in a single active with reated and single active mit of a chemical adhesive (eg. cyanoacrylate) remote	30465	Repair of nasal vestibular stenosis	
337889   Incl. deployment & calibration or sensor, right heart. Deny INV/N10	30520		
incl deployment & calibration of sensor, right heart-Deny INWN30    Assertion   Assertion	33289		
injection of non-compounded foam scierosant with ultrasound compression maneuvers to guide dispersion of the injectate, inclusive of all imaging guidance and monitoring the treatment of a single, incompetent extremity viel injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins, limb or trunk injection of scierosant for spider veins injection of scierosant for s		incl deployment & calibration of sensor, right heart- Deny INV/N10	
the injectate, inclusive of all imaging guidance and monitoring the treatment of a single, incompetent extremity vein  36468 36488 Injection of sclerosant for spider veins, timb or trunk 36470 IN sclerosing solution; single vein. 36471 IN sclerosing solution; single vein. 36472 Endovenous ablation this incompetent vein is their radiofreq tx'd 36475 Endovenous ablation this incompetent vein; 2nd, subs veins 36478 Endovenous ablation this incompetent vein; 2nd, subs veins 36478 Endovenous ablation this incompetent vein; 2nd, subs veins 36479 Endovenous ablation this incompetent vein, extremity, inclusive of all imaging guidance and monitoring, percut laser 1st vein 36479 Endovenous ablation this incompetent vein, extremity, inclusive of all imaging guidance and monitoring, percutal adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; first vein treated 4648 Endovenous ablation therapy of incompetent vein, extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; first vein treated 4648 Endovenous ablation therapy of incompetent vein, extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; subsequent vein(s) treated in a single extremity, each through separate access sites (List separately in addition to code for primary procedure) 47727 Reviscuentzation; envolvascurar, open or percuaneous, tennorar, popurear artery(s), rumaterar, with a substance of the primary procedure) 47728 Ligation, division, a stripping, short saphenous vein 47729 Ligation division, and/or excision of varicose vein cluster(s)-1 leg 47729 Excision of hyperplastic alveolar mucosa, ea quadrant 47820 Excision of hyperplastic alveolar mucosa, ea quadrant	33945		
Vein	_		_
36466 Injection of sclerosant for spider veins, limb or trunk 36471 INJ sclerosing solution; single vein. 36472 Endovenous ablation thx incompetent vein: 1st vein radiofred tx'd 56475 Endovenous ablation thx incompetent vein: 2nd, subs veins 56476 Endovenous ablation thx incompetent vein: 2nd, subs veins 56476 Endovenous ablation thx incompetent vein: 2nd, subs veins 56478 Endovenous ablation thx incompetent vein: 2nd, subs veins 56478 Endovenous ablation thx incompetent vein: 2nd, subs veins 56479 Endovenous ablation thx incompetent vein: 2nd, subs veins 56479 Endovenous ablation thrapy of incompetent vein, extremity, by transcatheter delivery of a chemical adhesive 56482 (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; first vein treated 56483 percutaneous; subsequent vein(s) treated in a single extremity, and through separate access sites (List separately in addition to code for primary procedure) 56483 respectively in addition to code for primary procedure) 57722 Ligation, division, & stripping, short saphenous vein 57755 Stab phlebectomy varicose veins, one ext; 10-20 stab incisions 57765 Stab phlebectomy of Veins one extrem; more than 20 incisions 57765 Ligation, division, and/or excision of varicose vein cluster(s)-1 leg 57765 Excision elsion or tumor dentoalveolar structures; wo repair 57765 Excision elsion or tumor dentoalveolar structures; wo repair 57765 Excision elsion or tumor dentoalveolar structures; wo repair 57765 Excision of hyperplastic alveolar mucosa, ea quadrant 57766 Alteroal division, and/or excision of varicose vein cluster(s)-1 leg 57765 Excision of hyperplastic alveolar mucosa, ea quadrant	36465		
36466 Injection of sclerosant for spider veins, limb or trunk  36470 INJ sclerosing solution; single vein.  36471 INJ sclerosing solution; single vein.  36472 Endovenous ablation this incompetent vein: 1st vein radiofreq tx'd  36476 Endovenous ablation this incompetent vein; 2nd, subs veins  36478 Endovenous ablation this incompetent vein; 2nd, subs veins  36479 Indovenous ablation this incompetent vein; vein, call imaging guidance monitoring, percut laser 1st vein runvenous and anotion trust incomp vein ext, lincl all imaging guidance and monitoring, percut laser 1st vein runvenous and anotion trust incompetent vein, extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; first vein treated  Endovenous ablation therapy of incompetent vein, extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; subsequent vein(s) treated a single extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; subsequent vein(s) treated a single extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; subsequent vein(s) treated a single extremity, each through separate access site (List separately in addition to code for primary procedure)  37222		100 PT 10	
36470 INJ sclerosing solution; single vein. 36471 INJ sclerosing solut, multi-veins, same leg. 36472 Endovenous ablation thx incompetent vein- 1st vein radiofreq bt/d 36476 Endovenous ablation thx incompetent vein; 2nd, subs veins 36478 Endovenous ablation thx incompetent vein; 2nd, subs veins 36479 Endoven ablation thx incompetent vein, extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; first vein treated Endovenous ablation therapy of incompetent vein, extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; first vein treated Endovenous ablation therapy of incompetent vein, extremity, by transcatheter delivery of a chemical adhesive (eg. cyanoacrylate) remote from the access site, inclusive of all imaging guidance and monitoring, percutaneous; subsequent vein(s) treated in a single extremity, each through separate access sites (List separately in addition to code for primary procedure)  37722 revascutarization, entovascutarion, operor of percutameous; emorar, populate artery(s), umaterar, with 37722 Ligation div & stripping long saphen vein saphenodem junct-knee 37765 Stab philebectomy varicose veins, one ext; 10-20 stab incisions 37766 Stab philebectomy of V Veins one extrem; more than 20 incisions 37785 Ligation, division, and/or excision of varicose vein cluster(s)-1 leg 41825 Excision felsion or tumor dentoalveolar structures; wo repair 41826 Excision of hyperplastic alveolar mucosa, a quadrant 41874 Alveoplasty, ea quadrant		At a last state that the first of all line administration and an artificial transmission and training for the first of the	
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41828 Excision of hyperplastic alveolar mucosa, ea quadrant 41874 Alveoplasty, ea quadrant			
41874 Alveoplasty, ea quadrant			
	41874	Alveoplasty, ea quadrant	
43659 Unlisted laparoscopy procedure, stomach Laparoscopy, surgicar, gastric restrictive procedure, removat or adjustable gastric restrictive device component	43659	Unlisted laparoscopy procedure, stomach	
43772 Laparoscopy, surgical, gastric restrictive procedure, removat or adjustable gastric restrictive device component	43772	- Euperiosoppy, surgious, gustine restrictive procedure, removal or aujustable gastric restrictive device component	



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Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization.

43774	Laparoscopy, surgical, gastric restrictive procedure; removal of adjustable gastric restrictive device and	·
43774	subcutaneous port components	
43775	Laparoscopy, surgical, gastric restrictive procedure; longitudinal gastrectomy (ie, sleeve gastrectomy)	
47135	Liver allotransplantation, orthotopic, partial or whole, from cadaver or living donor, any age	
55250	Vasectomy, unilateral or bilateral (separate proc), including postop semen examination(s)	
58301	Removal of intrauterine device (IUD)	
58720	Salpingo-oophorectomy, complete or partial, unilateral or bilateral	
64612	Chemodenervation of muscle(s); muscle(s) innervated by facial nerve, unilateral	
04015	ONCE PER DOS Chemodenervation of muscle(s); muscle(s) innervated by facial, trigeminal, cervical spinal &	
64615	accessory nerves, bilateral	
67900	Repair of brow ptosis (supraciliary, mid-forehead or coronal approach)	
67901	Repair blepharoptosis; frontalis muscle technique w suture or other material	
67903	Repair blepharoptosis; levator resection or advancement, internal approach	
67904	Repair blepharoptosis; levator resection or advancement, external approach	
67908	Repair blepharoptosis; conjunctivo- tarso-Muller's muscle levator resection	
67909	Reduction of overcorrection of ptosis	
67911	Correction of lid retraction	

Prosthetics and Orthotics

Code	Description	Comments
A4562	Pessary, reusable, non rubber, any type	
A7040	Vacuum drainage collection unit & tubing kit, including all supplies needed for collection unit change, for use	
A7048	w/implanted catheter, each	
L0120	Cervical, flexible, non-adjustable, prefabricated, off-the-shelf	
L0172	Cervical, collar, semi-rigid thermoplastic foam, two-piece, prefabricated, off-the-shelf	
L0174	Cervical, collar, semi-rigid, thermoplastic foam, two piece w/thoracic extension, prefabricated, off-the-shelf	
L0180	Cervical, multiple post collar, occipital &/or mandibular supports, adjustable	
L0220	Thoracic, rib belt, custom fabricated	
10450	TLSO, flexible, provides trunk support, thoracic region, rigid posterior panel & soft anterior apron, extends	
L0456	fromsacrococcygeal junction & terminates just inferior toscapular	
10400		
L0482	TLSO triplanar control 1 piece rigid plastic shell w interface liner; incl carved plaster/cad-cam model custom fab	
1,0004	SO, flexible, provides pelvic-sacral support, reduces motion aboutsacroiliac joint, includes straps, closures,	
L0621	may include pendulous abdomen design, prefabricated, off-the-she	
1,0007	LSO, sagittal-coronal control, w/rigid anterior & posterior frame/panels, posterior extends from sacrococcygeal	
L0637	junction to T-9 vertebra, lateral strength provided by rigid la	
1,0050	LSO, sagittal-coronal control, w/rigid anterior & posterior frame/panel(s), posterior extends from sacrococcygeal	
L0650	junction to T-9 vertebra, lateral strength provided by rigid	
	Cervical-thoracic-lumbar-sacral-orthoses (ctlso), anterior-posterior-lateral control, prefabricated item that has	
L0720	been trimmed, bent, molded, assembled, or otherwise customized to fit a specific patient by an individual with	
	expertise	
L1660	HO, abduction control of hip joints, static, plastic, prefabricated	
L1686	HO, abduction control of hip joint, post-operative, prefabricated	



Orthopedic shoe addition, rubber insole

L3510

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. L1812 KO, elastic w/joints, prefabricated, off-the-shelf KO, single upright, thigh & calf, w/adjustable flexion & extension joint, medial-lateral & rotation control, w/or w/o L1843 varus/valgus adjustment, prefabricated item that has bee L1846 L1850 KO, Swedish type, prefabricated, off-the-shelf control, with or without varus/valgus adjustment, prefabricat L1851 Knee orthosis, double upright, thigh & calf, with adjustable flexion & extension joint, medial-lateral & rotation L1852 control, with or without varus/valgus adjustment, prefabricat AO, ankle gauntlet or similar, with or without joints, prefabricated, off-the-shelf L1902 L1906 AFO, multiligamentous ankle support, prefabricated, off-the-shelf AFO, plastic or other material, prefabricated, includes fitting & adjustment L1930 AFO, rigid anterior tibial section, total carbon fiber or equal material, prefabricated, off-the-shelf L1933 AFO, molded to patient model, plastic, rigid anterior tibial section, custom L1945 AFO, spiral, (institute of rehabilitative medicine type), plastic or other material, prefabricated, off-the-shelf L1952 AFO, posterior solid ankle, plastic, custom L1960 activation, includes ankle joint, any type, custom fabricated L2005 L2220 Addition to lower extremity, dorsiflexion & plantar assist/resist, ea joint Addition to lower extremity orthosis, suspension sleeve L2397 L2415 Addition to knee lock w integrated release mechanism, any material, ea joint Addition to knee joint, disc or dial lock, adjustable knee flexion, ea joint L2425 L2780 Addition to lower extremity orthosis, noncorrosive finish, per bar L2795 Addition to lower extremity orthosis, knee control, full kneecap L2810 Addition to lower extremity orthosis, knee control, condylar pad Add to lower extremity orthosis, soft interface for molded plastic, above knee L2830 Addition to lower extremity orthosis, tibial length sock, fracture or equal L2840 Addition to lower extremity orthosis, femoral length sock, fracture or equal L2850 L2999 Lower extremity orthosis, not otherwise specified OLH: Foot, insert, removable, molded to model, UCB type, Berkeley shell, ea L3000 OLH: Foot, insert, removable, molded, longitudinal/ metatarsal support L3020 Foot, insert/plate, removable, add to LE orthosis High str light wt material, all hybrid lam L3031 L3040 OLH: Foot, arch support, removable, premolded, longitudinal, ea OLH: Foot, arch support, removable, premolded, longitudinal/metatarsal, ea L3060 OLH: Foot, arch support, nonremovable attached to shoe, longitudinal, ea L3070 OLH: Foot, arch support, nonremovable, longitudinal/metatatarsal, ea L3090 L3100 Hallus-valgus night dynamic splint, prefabricated, off-the-shelf Foot, plastic, silicone or equal, heel stabilizer, prafabricated, off-the-shelf, each L3170 Lift, elevation, heel & sole, neoprene, per inch L3310 L3332 Lift, elevation, inside shoe, tapered, up to one-half inch Lift, elevation, heel, per inch L3334 L3360 Sole wedge, outside sole Sole wedge, between sole L3370 Heel, SACH cushion type L3450



bcbs	estmedicare.com  Procedure codes related to the CMS Inpatient Only Lis	t have been removed, however the inpatient admission care requires an authorization.
L3620	Transfer of an orthosis from one shoe to another, solid stirrup, existing	and the second s
L3702	adjustment	
1.0700	Elbow orthosis, with adjustable position locking joint, prefabricated, item that has been trimmed, bent, molded,	
L3760	assembled, or otherwise customized to fit specific patient by	
L3761	Elbow orthosis, w adjustable position locking joint, prefabricated, off-the-shelf	
L3807	WHFO, w/o joint(s), prefabricated item that has been trimmed, bent, molded, assembled, or otherwise	
L30U/	customized to fitspecific patient byindividual w/expertise	
L3908	WHO, wrist extension control cock-up, non molded, prefabricated, off-the-shelf	
L3915	WHO, includes one or more nontorsion joint(s), elastic bands, turnbuckles, may include soft interface, straps,	
25515	prefabricated item that has been trimmed, bent, molded, assemble	
L3916	WHO, includes 1 or more nontorsion joint(s), elastic bands, turnbuckles, may include soft interface, straps,	
20010	prefabricated, off-the-shelf	
L3917	HO, metacarpal fracture orthosis, prefabricated item that has been trimmed, bent, molded, assembled, or	
	otherwise customized to fitspecific patient byindividual w/expertise	
L3923	HFO, w/o joints, may include soft interface, straps, prefabricated item that has been trimmed, bent, molded,	
	assembled, or otherwise customized to fitspecific patient byindivi	
L3924	HFO, without joints, may include soft interface, straps, prefabricated, off-the-shelf	
L3929	HFO, includes one or more nontorsion joint(s), turnbuckles, elastic bands/springs, may include soft interface	
	material, straps, prefabricated item that has been trimmed, bent,	
L3933	Finder orthogic without ininte movingly described	
	Finger orthosis, without joints, may include soft interface, custom fabricated, includes fitting and adjustment	
L3960	SEWHO, abduction positioning, airplane design, prefabricated	
L3980	Upper extremity fracture orthosis, humeral, prefabricated	<u> </u>
L3982	Upper extremity fracture orthosis, radius or ulnar, prefabricated	
L3995	Addition to upper extremity orthosis, sock, fracture or equal, ea	
L3999	Upper limb orthosis, not otherwise specified  Replace proximal & distal upright for KAFO	
L4070	Replace proximat & distat upright for KAFO  Replace metal bands KAFO, proximal thigh	
L4080	Ankle control orthosis, stirrup style, rigid, includes any type interface, prefabricated, off-the-shelf	
L4350	Walking boot, pneumatic &/or vacuum, w/or w/o joints, w/or w/o interface material, prefabricated item that has	<del> </del>
L4360	been trimmed, bent, molded, assembled, or otherwise customized t	
	Walking boot, pneumatic &/or vacuum, w/or without joints, w/or without interface material, prefabricated, off-	
L4361	the-shelf	
L4370	Pneumatic full leg splint, prefabricated, off-the-shelf	
	<u> </u>	
L4387	Walking boot, non-pneumatic, w/or without joints, w/or without interface material, prefabricated, off-the-shelf	
1.4000	Static or dynamic AFO, including soft interface material, adjustable for fit, for positioning, may be used for	
L4396	minimal ambulation, prefabricated item that has been trimmed, be	
1.4007	Static or dynamic AFO, including soft interface material, adjustable for fit, for positioning, may be used for	
L4397	minimal ambulation, prefabricated, off-the-shelf	
L4398	Foot drop splint, recumbent positioning device, prefabricated, off-the-shelf	
L5540	Prep, below knee PTB socket, nonalignable, laminated socket molded	
L5590	Prep above knee, disarticulation ischial level socket, laminated, molded	



equal)

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. L5622 Addition to lower extremity, test socket, knee disarticulation Lower extremity addition hip disarticulation flexible inner socket external frame L5643 L5677 Addition to lower extremity, below knee, knee joints, polycentric, pair Addition to LE BK/AK custom fab socket insert for congenital/traumatic amputee, INITIAL L5681 L5682 Addition to lower extremity, below knee, thigh lacer, gluteal/ischial molded L5810 Addition, endoskeletal knee-shin system, single axis, manual lock L5814 Endoskeletal knee-shin system, polycentric hydraulic swing phase control L5816 Endoskeletal knee-shin system, polycentric, mechanical stance Endoskeletal knee-shin system, single axis, electromechanical swing and stance phase control, with or without L5827 shock absorption and stance extension damping Endoskeletal ankle foot system, microprocessor controlled feature, dorsiflexion &/or plantar flexion control, incl L5973 power source- Deny INV/N10>=11/13/2010 All lower extremity prostheses, multiaxial ankle, dynamic response foot L5979 L5980 All lower extremity prostheses, flex foot system L5986 All lower extremity prostheses, multi-axial rotation unit, MCP or equal L5988 Addition to lower limb prosthesis, vertical shock reducing pylon feature Transcarpal/metacarpal/partial hand disarticulation prosthesis, extnl power, self-suspended, inner socket L6026 w/removable forearm, electrodes & cables-Deny INV/N10=1/1/15-5/29/18 Below elbow, molded socket, flexible elbow hinge, triceps pad L6100 L6300 Shoulder disarticulation, molded socket, bulkhead, humeral section, forearm Upper extremity prosthesis addition, flexion/extension wrist with or without friction, for use w ext powered L6621 terminal L6629 Upper extremity addition, quick disconnect lamination collar, Otto Bock Upper extremity add shoulder joint multipositional locking use w powered system L6646 Upper extremity addition, harness, chest or shoulder, saddle type L6672 Upper extremity addition, test socket, wrist disarticulation or below elbow L6680 L6684 Upper extremity, test socket, shoulder disarticulation or interscapular thoracic Upper extremity addition, suction socket L6686 Upper extremity, frame type socket, below elbow or wrist disarticulation L6687 Upper extremity addition, frame type socket, shoulder disarticulation L6689 Addition to UE prosthesis, BE/AE, custom fabricated from existing mold or prefabricated, L6694 L6697 congenit Upper extremity addition, external powered feature, myoelectronic control module, additional emg inputs, L6700 pattern-recognition decoding intent movement Terminal device, hand, machanical voluntary opening any material any size L6708 INV/N10=04/11/2013-05/29/2018 L6715 patterns, incl motor-Deny INV/N10 DOS= 12/11/2014 - 5/29/2018 L6880 power L6884 L6890 adjustment Addition to UE prosthesis, glove for terminal device, any material, custom fabricated L6895 L6935 Below elbow, external power, myoelectronic control of terminal device Addition to upper extremity prosthesis, below elbow/wrist disarticulation, u/light mat (titanium, carbon fiber or L7400



session: linear accelerator based

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Updated: 07/01/2025

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	Troccatio coacorotatea to the or is inpution.
L7402	Addition to upper extremity prosthesis, shoulder disarticulation/interscapular thoracic, ultralight material -
	titanium
L7403	Addition to upper extremity prosthesis, below elbow/wrist disarticulation, acrylic material
L7405	Addition to upper extremity prosthesis, shoulder disarticulation/interscapular thoracic, acrylic material
L7406	Addition to upper extremity, user adjustable, mechanical, residual limb volume management system
L8002	Breast prosthesis, mastectomy bra, with integrated breast prosthesis form, bilateral, any size, any type
L8015	External breast prosthesis garment, w mastectomy form, post mastectomy
L8020	Breast prosthesis, mastectomy form
L8040	Nasal prosthesis provided by non-physician
L8300	Truss, single with standard pad
L8501	Tracheostomy speaking valve
L8507	Tracheo-esophageal voice prosthesis, patient inserted, any type, ea
1,0000	Injectable bulking agent, synthetic implant, urinary tract, 1 ml syringe, includes shipping - HCFA RN
L8606	Review=10/25/10-01/15/13
1,0040	Cochlear implant, external speech processor & controller, integrated system, replacement- RN
L8619	Review>=10/1/2004
L8621	each
	Patient programmer for use with implantable programmable neurostimulator pulse generator, replacement only -
L8681	RN Review
L8701	Powered upper extremity range of motion assist device, elbow, wrist, hand with single or double upright,
	includes microprocessor, sensors
L8702	Powered upper extremity range of motion assist device, elbow, wrist, hand, finger, single or double upright,
L0702	includes microprocessor, sensors

**Radiation Therapy** Code **Code Description** Comments 77014 CT guidance for placement of radiation therapy fields Respiratory motion management simulation brachyunerapy 1504 Sources of remote aftertoaunig brachytherapy, 1 Channery, 77293 77316 Brachytherapy isodose plan; intermediate made from 5 to 10 sources or remote afterloading brachytherapy, 2-12 77317 channels), includes basic dosimetry calculation brachytherapy isouose plant, complex made over 10 sources of Terriote aftertoading brachytherapy, over 12 77318 77321 Special teletherapy port plan, particles, hemibody, total body 77336 Continuing medical physics consultation support of oncologist per week Multi-leaf collimator (MLC) device(s) for intensity modulated radiation therapy (IMRT), design and construction 77338 77370 Special medical radiation physics consultation Radiation treatment delivery, stereotactic radiosurgery, complete course of treatment of cranial lesion(s) 1 77371 session; multi-source Cobalt 60 based Radiation treatment delivery, stereotactic radiosurgery, complete course of treatment of cranial lesion(s) 1 77372



G6003

#### **Medicare Advantage Master Prior Auth List**

Updated: 07/01/2025

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Updated: 07/01/2025 bcbstmedicare.com Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization. Radiation treatment delivery, 2 separate treatment areas, 3 or more ports on single treatment area, use of G6007 multiple blocks: up to 5 mev iauiation treatment detivery, 5 or more separate treatment areas, custom blocking, tangential ports, wedges, G6012 Radiation treatment delivery, 3 or more separate treatment areas, custom blocking, tangential ports, wedges, G6013 rotational beam, compensators, electron beam; 11-19 mev G6015 Intra-fraction localization & tracking of target or patient motion during delivery of radiation therapy, each fraction G6017 of treatment Radiology Code Code Description Comments 78811 Positron emission tomography (PET) imaging; limited area, eg, chest, head/neck **Sleep Studies Code Description** Code Comments 95805 Multiple sleep latency testing, recording, analysis & interpret ONCE/DOS 95807 Sleep study, recording ventil, ECG & 02 satur technol ONCE/DOS 95808 Polysomnography; any age, sleep staging with 1-3 additional parameters of sleep, attended by a technologist 95810 Polysomnography; age 6 years or older, sleep staging w/4 or more add parameters 95811 Polysomnography; age 6 years or older, sleep staging w/4 or more add parameters **Specialty Pharmacy** PA-Requires a Prior Authorizatio **Code Description** Code Comments Q2055 Abecma PA J9264 Abraxane PA J3262 Actemra IV PA Acthar ST/PA J0801 Adakveo PA J0791 J9042 Adcetris PA J9029 Adstiladrin PA J0172 Aduhelm PA J7192 Advate PA J7207 Adynovate PA J7171 Adzynma PA J7210 Afstyla PA Q5150 Ahzantive st J1931 Aldurazyme PA J9305 Alimta PA J7186 Alphanate PA J7193 Alphanine SD PA



bcbs	tmedicare.com  Procedure codes related to the CMS Inpatient Only Lis	t have been removed, however the inpatient admission care requires an authorization.
J7201	Alprolix PA	
J7214	Altuviiio PA	
J1552	Alyglo <sup>PA</sup>	
Q5126	Alymsys stipa	
J1426	Amondys 45 PA	
J9999	Amtagvi <sup>PA</sup>	
J0225	Amvuttra PA	
J9028	Anktiva PA	
J2277	Aphexda <sup>PA</sup>	
J0256	Aralast PA	
	Aranesp PA	
J0882	(ESRD on dialysis)	
10004	Aranesp PA	
J0881	(non-ESRD)	
J9261	Arranon PA	
J1554	Asceniv PA	
J9118	Asparlas PA	
IOOOF	Asparlas PA Avastin ST/PA	
J9035	(oncology only)	
Q5121	Avsolastipa	
J9025	azacitidine <sup>PA</sup>	
J9023	Bavencio PA	
J9032	Beleodaq PA	
J9036	Belrapzo PA	
J9033	bendamustine PA	
J9036		
J9034	Bendeka PA	
J7195	BeneFIX PA	
J0490	Benlysta IV PA	
J0179	Beovu ST/PA	
J1414	Beqvez PA	
J0597	Berinert PA	
J9229	Besponsa PA	
J1556	Bivigam <sup>PA</sup>	
Q5152	Bkemv	
J9039	Blincyto PA	
J9041		
J9046		
J9048	bortezomib <sup>PA</sup>	
J9049		
J9051		
J9054	Boruzu	
J0585	Botox PA	



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Q2054	Breyanzi <sup>PA</sup>	
J2329	Briumvi <sup>PA</sup>	
Q5124	Byooviz ST/PA	
J9064	cabazitaxel <sup>PA</sup>	
J1952	Camcevi PA	
Q2056	Carvykti PA	
J3392	Casgevy <sup>PA</sup>	
J1786	Cerezyme PA	
Q5128	Cerezyme PA Cimerli ST/PA	
J0717	Cimzia vials PA	
J2786	Cinqair PA	
J0598	Cinryze PA	
J7175	Coagadex PA	
J9286	Columvi <sup>PA</sup>	
J7180	Corifact PA	
J0802	Cortrophin gel ST/PA	
J1448	Cosela PA	
J3247	Cosentyx IV PA	
J0584	Crysvita PA	
J1551	Cutaquig PA	
J1555	Cuvitru PA	
J9308	Cyramza <sup>PA</sup>	
J9348	Danyelza <sup>PA</sup>	
J9145	Darzalex PA	
J9144	Darzalex Faspro PA	
J0589	Daxxify PA	
J7318	Durolane ST/PA	
J0586	Dysport PA	
J1301	edaravone <sup>PA</sup>	
J9063	Elahere PA	
J1743	Elaprase PA	
J3060	Elelyso PA	
J1413	Elevidys PA	
J2508	Elfabrio PA	
J9217	Eligard PA	
J7205	Eloctate PA	
J1323	Elrexfio PA	
J9269	Elzonris PA	
J9176	Empliciti PA	
J9358	Enhertu PA	
J1302	Enjaymo PA	
J3380	Entyvio IV PA	



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Q5149	Enzeevu <sup>st</sup>	
J9321	Epkinly <sup>PA</sup>	
0.4004	Epogen PA	
Q4081	(ESRD)	
IOOOE	Epogen PA	
J0885	(non-ESRD)	
J1325	epoprostenol PA	
Q5151	Epysqli	
J9055	Erbitux PA	
J9179	eribulin mesylate <sup>PA</sup>	
J7204	Esperoct PA	
J7323	Euflexxa PA	
J3111	Evenity ST/PA	
J1305	Evkeeza PA	
J1428	Exondys 51 PA	
J0178	Eylea <sup>ST/PA</sup>	
J0177	Eylea HD ST/PA	
J0180	Fabrazyme <sup>PA</sup>	
J0517	Fasenra PA	
J7198	Feiba NF PA	
J1951	Fensolvi PA	
J1744	Firazyr <sup>PA</sup>	
J9155	Firmagon PA	
J1572	Flebogamma PA	
J1325	Flolan <sup>PA</sup>	
J9307	Folotyn <sup>PA</sup>	
Q5108	Fulphila <sup>PA</sup>	
J0641	Fusilev <sup>PA</sup>	
J9331	Fyarro <sup>PA</sup>	
Q5130	Fylnetra st/PA	
J9210	Gamifant PA	
J1569	Gammagard Liquid PA	
J1566	Gammagard S/D <sup>PA</sup>	
	(powder)	
J1561	Gammaked PA	
J1557	Gammaplex <sup>PA</sup>	
J1561	Gamunex-C <sup>PA</sup> Gazyva <sup>PA</sup>	
J9301	Gazyva <sup>PA</sup>	
J7326	Gel-One ST/PA	
J7328	Gel-Syn <sup>ST/PA</sup>	
J7320	Genvisc ST/PA	
J0223	Givlaari <sup>PA</sup>	



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J0257	Glassia <sup>PA</sup>	
J9179	Halaven PA	
J1411	Hemgenix <sup>PA</sup>	
J7170	Hemlibra <sup>ST/PA</sup>	
J7190	Hemofil M PA	
	Herceptin	
J9356	Hylecta ST/PA	
J9355	Herceptin STIPA	
Q5146	Hercessi <sup>PA</sup>	
Q5113	Herzuma <sup>ST/PA</sup>	
J1559	Hizentra PA	
J7187	Humate-P PA	
J7321	Hyalgan <sup>ST/PA</sup>	
J7322	Hymovis <sup>ST/PA</sup>	
J1575	Hyqvia PA	
J1744	icatibant PA	
J7202	Idelvion PA	
J7355	iDose TR <sup>PA</sup>	
J0638	llaris <sup>PA</sup>	
J3245	Ilumya <sup>PA</sup>	
J7313	Iluvien PA	
J9026	Imdelltra PA	
J9173	Imfinzi <sup>PA</sup>	
J9347	Imjudo <sup>PA</sup>	
J9325	Imlygic PA	
Q5103	Inflectra PA	
J1745	infliximab <sup>PA</sup>	
J9319	Istodax PA	
J9207	Ixempra PA	
J7195	Ixinity PA	
J2782	Izervay PA	
J9281	Jelmyto PA	
J9272	Jemperli <sup>PA</sup>	
J9043	Jevtana <sup>PA</sup>	
J7208	Jivi <sup>PA</sup>	
J9354	Kadcyla <sup>PA</sup>	
J1290	Kalbitor <sup>PA</sup>	
Q5117	Kanjinti <sup>PA</sup>	
J2840	Kanuma <sup>PA</sup>	
J9271	Keytruda <sup>PA</sup>	
J0642	Khapzory PA	
J9274	Kimmtrak <sup>PA</sup>	



bcbst	Procedure codes related to the CMS Inpatient Only Lis	t have been removed, however the inpatient admission care requires an authorization.
J0175	Kisunla <sup>PA</sup>	
J7190	Koate-DVI PA	
J7192	Kogenate FS PA	
J7211	Kovaltry PA	
J2507	Krystexxa PA	
Q2042	Kymriah PA	
J9047	Kyprolis <sup>PA</sup>	
J0217	Lamzede <sup>PA</sup>	
J1932	lanreotide acetate <sup>ST/PA</sup>	
J3490		
J0202	Lemtrada <sup>PA</sup>	
J3590	Lenmeldy PA	
J0174	Leqembi PA	
J1306	Leqvio <sup>PA</sup>	
J2820	Leukine PA	
J1954	leuprolide depot <sup>PA</sup>	
J0641	levoleucovorin calcium <sup>PA</sup>	
J9119	Libtayo <sup>PA</sup>	
J3263	Loqtorzi <sup>PA</sup>	
J2778	Lucentis ST/PA	
J0221	Lumizyme <sup>PA</sup>	
J9350	Lunsumio <sup>PA</sup>	
J9217	Lupron Depot PA	
,021,	(antineoplastic use)	
J1950	Lupron Depot <sup>PA</sup>	
	(endocrine use)	
J3398	Luxturna PA	
J3394	Lyfgenia PA	
J9353	Margenza PA	
J3397	Mepsevii PA	
J0887	Mircera PA	
	(ESRD on dialysis)	
J0888	Mircera PA	
100.40	(non-ESRD)	
J9349	Monjuvi PA	
J7327	Monovisc ST/PA	
Q5107	Mvasi PA	
J9203	Mylotarg PA	
J0587	Myobloc PA	
J1458	Naglazyme PA	
J9261	nelarabine PA	
J2506	Neulasta <sup>PA</sup>	



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J2506	Neulasta Onpro PA	
J0219	Nexviazyme <sup>PA</sup>	
J9038	Niktimvo	
J7182	Novoeight <sup>PA</sup>	
J7189	Novoseven RT PA	
J2802	Nplate PA	
J2182	Nucala <sup>PA</sup>	
J3490	Nulibry <sup>PA</sup>	
J7209	Nuwiq <sup>PA</sup>	
Q5122	Nyvepria <sup>ST/PA</sup>	
J7188	Obizur <sup>PA</sup>	
J2350	Ocrevus <sup>PA</sup>	
J2351	Ocrevus Zunovo	
J1568	Octagam <sup>PA</sup>	
J2353	octreotide acetate inj. susp. PA	
Q5114	Ogivri <sup>PA</sup>	
J2267	Omvoh IV PA	
J9266	Oncaspar PA	
J9205	Onivyde PA	
J0222	Onpattro <sup>PA</sup>	
Q5112	Ontruzant PA	
J9299	Opdivo PA	
J9298	Opdualag PA	
Q5153	Opuviz <sup>ST</sup>	
J0129	Orencia IV PA	
J7324	Orthovisc PA	
Q9999	Otulfi	
J0224	Oxlumo PA	
J7312	Ozurdex <sup>PA</sup>	
J9259	paclitaxel <sup>PA</sup>	
J9264		+
J9177	Padcev PA	+
J1576	Panzyga <sup>PA</sup>	+
Q5147	Pavblu PA	+
J0208	Pedmark <sup>PA</sup>	+
J9294		
J9296		
J9297 J9305		
J9305 J9314	pemetrexed <sup>PA</sup>	
J9314 J9322	решенехей	
J9322 J9353		
J9323		



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J9292	,	
J9304	Pemfexy PA	
J9324	Pemrydi RTU <sup>PA</sup>	
J9306	Perjeta <sup>PA</sup>	
J9316	Phesgo <sup>PA</sup>	
J1307	Piasky <sup>PA</sup>	
J9309	Polivy PA	
J1203	Pombiliti <sup>PA</sup>	
J9295	Portrazza <sup>PA</sup>	
J9204	Poteligeo <sup>PA</sup>	
J9307	pralatrexate <sup>PA</sup>	
J1459	Privigen PA	
0.4004	Procrit <sup>PA</sup>	
Q4081	(ESRD)	
J0885	Procrit PA	
10885	(non-ESRD)	
J7194	Profilnine SD PA	
J0256	Prolastin C PA	
J9015	Proleukin PA	
J0897	Prolia <sup>PA</sup>	
Q2043	Provenge PA	
Q9997	Pyzchiva	
J1304	Qalsody PA	
J1301	Radicava PA	
J7203	Rebinyn PA	
J0896	Reblozyl <sup>PA</sup>	
J3489	Reclast PA	
J7192	Recombinate PA	
J1745	Remicade PA	
J3285	Remodulin PA	
Q5104	Renflexis ST/PA	
Q5105	Retacrit PA	
Q3103	(ESRD on dialysis)	
Q5106	Retacrit PA	
	(non-ESRD)	
J3590	Rethymic PA	
J7311	Retisert PA	
J3590	Revcovi PA	
Q5123	Riabni <sup>ST/PA</sup>	
J9311	Rituxan Hycela <sup>ST/PA</sup>	
J9312	Rituxan <sup>ST/PA</sup>	
J3490	Rivfloza PA	



	dicare.com  Procedure codes related to the CMS Inpatient Only	List have been removed, however the inpatient admission care requires an authorization.
J7200	Rixubis <sup>PA</sup>	
J1412	Roctavian <sup>PA</sup>	
J1449	Rolvedon ST/PA	
J9318	romidepsin <sup>PA</sup>	
J0596	Ruconest PA	
Q5119	Ruxience PA	
J9061	Rybrevant PA	
J9021	Rylaze <sup>PA</sup>	
J2998	Ryplazim <sup>PA</sup>	
J9333	Rystiggo PA	
J0870	Rytelo PA	
J1744	Sajazir <sup>PA</sup>	
J2353	Sandostatin LAR Depot PA	
J0491	Saphnelo PA	
J9227	Sarclisa <sup>PA</sup>	
Q9998	Selarsdi	
J7212	Sevenfact PA	
J2502	Signifor LAR ST/PA	
J1602	Simponi Aria PA	
J2327	Skyrizi IV <sup>PA</sup>	
J3590	Skysona <sup>PA</sup>	
J3490	sodium hyaluronate 1% PA	
J1299	Soliris ST/PA	
J1930	Somatuline Depot PA	
J1747	Spevigo PA	
J2326	Spinraza PA	
J3358	Stelara IV PA	
Q5127	Stimufend ST/PA	
J7321	Supartz FX ST/PA	
J9226	Supprelin LA PA	
J2779	Susvimo PA	
J2781	Syfovre PA	
J2860	Sylvant PA	
J7331	Synojoynt PA	
J7325	Synvisc PA	
J7325	Synvisc-One PA	
J0593	Takhzyro PA	
J3055	Talvey PA	
Q2053	Tecartus PA	
Q2057	Tecelra PA	
J9022	Tecentriq PA	
J9024	Tecentriq	



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J9380	Tecvayli <sup>PA</sup>	
J3490	Tegsedi <sup>PA</sup>	
J9328	Temodar IV PA	
J9330	temsirolimus <sup>PA</sup>	
J3241	Tepezza <sup>PA</sup>	
J3490	Testopel PA	
J9329	Tevimbra PA	
J2356	Tezspire PA	
J3240	Thyrogen PA	
J9273	Tivdak <sup>PA</sup>	
Q5133	Tofidence PA	
J9330	Torisel PA	
Q5116	Trazimera <sup>PA</sup>	
J9033	Treanda PA	
J3315	Trelstar Depot PA	
J1628	Tremfya IV <sup>PA</sup>	
J3285	treprostinil <sup>PA</sup>	
J7181	Tretten PA	
J7332	Triluron PA	
J3316	Triptodur <sup>PA</sup>	
J7329	TriVisc PA	
J9317	Trodelvy PA	
J1746	Trogarzo PA	
Q5115	Truxima PA	
Q5135	Tyenne IV PA	
Q5134	Tyruko PA	
J2323	Tysabri PA	
J9381	Tzield PA	
Q5111	Udenyca ST/PA	
Q5111	Udenyca Onbody ST/PA	
J1303	Ultomiris PA	
J1823	Uplizna PA	
J2777	Vabysmo ST/PA	
J9303	Vectibix PA	
Q5129	Vegzelma <sup>ST/PA</sup>	
J9041	Velcade <sup>PA</sup> Veletri <sup>PA</sup>	
J1325 J9376	Veletri '^ Veopoz <sup>PA</sup>	
J9376 J9025	Veopoz ··· Vidaza <sup>PA</sup>	
J9025 J1427		
J1427 J1322	Viltepso <sup>PA</sup> Vimizim <sup>PA</sup>	
J7322 J7321	Vimizim *** Visco-3 ST/PA	
J/32I	Visco-3	



Code Q4164

### Medicare Advantage Master Prior Auth List Updated: 07/01/2025

Comments

bcbstm	nedicare.com  Procedure codes related to the CMS Inpatie	nt Only List have been removed, however the inpatient admission care requires an authorization.
J3396	Visudyne ST/PA	
J9056	Viivimusta ST/PA	
J7179	Vonvendi <sup>PA</sup>	
J3385	VPRIV ST/PA	
J3032	Vyepti <sup>PA</sup>	
J3401	Vyjuvek <sup>PA</sup>	
J1429	Vyondys 53 PA	
J9332	Vyvgart <sup>PA</sup>	
J9334	Vyvgart Hytrulo PA	
J9153	Vyxeos <sup>PA</sup>	
Q5138	Wezlana IV	
J7183	Wilate PA	
J3590	Winrevair PA	
J1558	Xembify PA	
J0218	Xenpozyme PA	
J0588	Xeomin PA	
J0897	Xgeva <sup>ST/PA</sup>	
J0775	Xiaflex <sup>PA</sup>	
J3299	Xipere PA	
J2357	Xolair <sup>PA</sup>	
J7185	Xyntha <sup>PA</sup>	
J9228	Yervoy PA	
Q2041	Yescarta PA	
J9352	Yondelis <sup>PA</sup>	
J7314	Yutiq <sup>PA</sup>	
J9400	Zaltrap <sup>PA</sup>	
J0256	Zemaira <sup>PA</sup>	
J9223	Zepzelca PA	
Q5120	Ziextenzo ST/PA	
Q5118	Zirabev PA	
J9202	Zoladex <sup>PA</sup>	
J3489	zoledronic acid <sup>PA</sup>	
J3399	Zolgensma <sup>PA</sup>	
J9359	Zynlonta PA	
J3393	Zynteglo <sup>PA</sup>	
J9345	Zynyz <sup>PA</sup>	

**Skin Substitues** 

Code Description

Helicoll PA



Updated: 07/01/2025

Procedure codes related to the CMS Inpatient Only List have been removed, however the inpatient admission care requires an authorization.

#### Surgery

Code	Code Description	Comments	
27702	Arthroplasty, ankle; with implant		
43497	Lower esophageal myotomy, transoral		
69705	Nasopharyngoscopy, surgical, with dilation of eustachian tube; unilateral		
91111	Gastrointestinal tract imaging, intraluminal, esophagus w/interp		