Please See the Coverage Summary for July 2025 New HCPCS Codes Below:

Not covered experimental and investigational for Commercial and ASO:

 (coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 40 mcg/0.4 mL dosage, for intramuscular use Influenza virus vaccine, H5N1, derived from cell cultures, adjuvanted, for intramuscular use Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 10 mcg/0.2 mL dosage, for intramuscular use Interrogation device evaluation (remote), up to 90 days, cardiac contractility modulation system with integral analysis, review and report(s) by a physician or other qualified health care professional 	'	voi covered experimental and investigational for commercial and A50.
Q4370 Aeroguard, per square centimeter Q4371 Neoguard, per square centimeter Q4372 Amchoplast excel, per square centimeter Q4373 Membrane wrap lite, per square centimeter Q4375 Duograft ac, per square centimeter Q4376 Duograft ap, per square centimeter Q4377 Trigraft ft, per square centimeter Q4378 Renew ft matrix, per square centimeter Q4380 Advograft one, per square centimeter Q4382 Advograft dual, per square centimeter 90382 Respiratory syncytial virus, monoclonal antibody, seasonal dose, 0.7 mL, for intramuscular use 90612 Influenza virus vaccine, trivalent, and severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 31.7 mcg/0.32 mL dosage, for intramuscular use 90613 Influenza virus vaccine, quadrivalent, and severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 40 mcg/0.4 mL dosage, for intramuscular use 90635 Influenza virus vaccine, H5N1, derived from cell cultures, adjuvanted, for intramuscular use 91323 Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 10 mcg/0.2 mL dosage, for intramuscular use 0948T Interrogation device evaluation (remote), up to 90 days, c	Q4368	Amchothick, per square centimeter
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Q4373 Amchoplast excel, per square centimeter Q4375 Duograft ac, per square centimeter Q4376 Duograft ac, per square centimeter Q4377 Trigraft ft, per square centimeter Q4378 Renew ft matrix, per square centimeter Q4379 Amniodefend ft matrix, per square centimeter Q4379 Advograft one, per square centimeter Q4380 Advograft one, per square centimeter Q4381 Respiratory syncytial virus, monoclonal antibody, seasonal dose, 0.7 mL, for intramuscular use Q4382 Respiratory syncytial virus, monoclonal antibody, seasonal dose, 0.7 mL, for intramuscular use Q4381 Influenza virus vaccine, trivalent, and severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 31.7 mcg/0.32 mL dosage, for intramuscular use Q4383 Influenza virus vaccine, quadrivalent, and severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 40 mcg/0.4 mL dosage, for intramuscular use Q4384 Influenza virus vaccine, H5N1, derived from cell cultures, adjuvanted, for intramuscular use Q4485 Influenza virus vaccine, H5N1, derived from cell cultures, adjuvanted, for intramuscular use Q4486 Interrogation device evaluation (remote), up to 90 days, cardiac contractility modulation system with inte analysis, review and report(s) by a physician or other qualified health care professional Q4497 Interrogation device evaluation (remote), up to 90 days, cardiac contractility modulation system, remote acquisition(s), receipt of transmissions, technician review, technical support, and distribution of results	Q4370	Aeroguard, per square centimeter
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analysis, review and report(s) by a physician or other qualified health care professional O949T Interrogation device evaluation (remote), up to 90 days, cardiac contractility modulation system, remote acquisition(s), receipt of transmissions, technician review, technical support, and distribution of results	91323	Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 10 mcg/0.2 mL dosage, for intramuscular use
acquisition(s), receipt of transmissions, technician review, technical support, and distribution of results	0948T	Interrogation device evaluation (remote), up to 90 days, cardiac contractility modulation system with interim analysis, review and report(s) by a physician or other qualified health care professional
0950T Ablation of benign prostate tissue, transrectal, with high intensity-focused ultrasound (HIFU), including	0949T	Interrogation device evaluation (remote), up to 90 days, cardiac contractility modulation system, remote data acquisition(s), receipt of transmissions, technician review, technical support, and distribution of results
ultrasound guidance	0950T	
	0951T	Totally implantable active middle ear hearing implant; initial placement, including mastoidectomy, placement

0952T	Totally implantable active middle ear hearing implant; revision or replacement, with mastoidectomy and replacement of sound processor
0953T	Totally implantable active middle ear hearing implant; revision or replacement, without mastoidectomy and replacement of sound processor
0954T	Totally implantable active middle ear hearing implant; replacement of sound processor only, with attachment to existing transducers
0955T	Totally implantable active middle ear hearing implant; removal, including removal of sound processor and all implant components
0956T	Partial craniectomy, channel creation, and tunneling of electrode for sub-scalp implantation of an electrode array, receiver, and telemetry unit for continuous bilateral electroencephalography monitoring system, including imaging guidance
0957T	Revision of sub-scalp implanted electrode array, receiver, and telemetry unit for electrode, when required, including imaging guidance
0958T	Removal of sub-scalp implanted electrode array, receiver, and telemetry unit for continuous bilateral electroencephalography monitoring system, including imaging guidance
0959T	Removal or replacement of magnet from coil assembly that is connected to continuous bilateral electroencephalography monitoring system, including imaging guidance
0960T	Replacement of sub-scalp implanted electrode array, receiver, and telemetry unit with tunneling of electrode for continuous bilateral electroencephalography monitoring system, including imaging guidance
0961T	Shortwave infrared radiation imaging, surgical pathology specimen, to assist gross examination for lymph node localization in fibroadipose tissue, per specimen (List separately in addition to code for primary procedure)
0962T	Assistive algorithmic analysis of acoustic and electrocardiogram recording for detection of cardiac dysfunction (eg, reduced ejection fraction, cardiac murmurs, atrial fibrillation), with review and interpretation by a physician or other qualified health care professional
0963T	Anoscopy with directed submucosal injection of bulking agent into anal canal
0964T	Impression and custom preparation of jaw expansion oral prosthesis for obstructive sleep apnea, including initial adjustment; single arch, without mandibular advancement mechanism
0965T	Impression and custom preparation of jaw expansion oral prosthesis for obstructive sleep apnea, including initial adjustment; dual arch, with additional mandibular advancement, non-fixed hinge mechanism
0966T	Impression and custom preparation of jaw expansion oral prosthesis for obstructive sleep apnea, including initial adjustment; dual arch, with additional mandibular advancement, fixed hinge mechanism
0967T	Transanal insertion of endoluminal temporary colorectal anastomosis protection device, including vacuum anchoring component and flexible sheath connected to external vacuum source and monitoring system

0968T	Insertion or replacement of epicranial neurostimulator system, including electrode array and pulse generator, with connection to electrode array
0969T	Removal of epicranial neurostimulator system
0970T	Ablation, benign breast tumor (eg, fibroadenoma), percutaneous, laser, including imaging guidance when performed, each tumor
0971T	Ablation, malignant breast tumor(s), percutaneous, laser, including imaging guidance when performed, unilateral
0972T	Assistive algorithmic classification of burn healing (ie, healing or nonhealing) by noninvasive multispectral imaging, including system set-up and acquisition, selection, and transmission of images, with automated generation of report
0973T	Selective enzymatic debridement, partial-thickness and/or full-thickness burn eschar, requiring anesthesia (ie, general anesthesia, moderate sedation), including patient monitoring, trunk, arms, legs; first 100 sq cm
0974T	Selective enzymatic debridement, partial-thickness and/or full-thickness burn eschar, requiring anesthesia (ie, general anesthesia, moderate sedation), including patient monitoring, trunk, arms, legs; each additional 100 sq cm (List separately in addition to code for primary procedure)
0975T	Selective enzymatic debridement, partial-thickness and/or full-thickness burn eschar, requiring anesthesia (ie, general anesthesia, moderate sedation), including patient monitoring, scalp, neck, hands, feet, and/or multiple digits; first 100 sq cm
0976T	Selective enzymatic debridement, partial-thickness and/or full-thickness burn eschar, requiring anesthesia (ie, general anesthesia, moderate sedation), including patient monitoring, scalp, neck, hands, feet, and/or multiple digits; each additional 100 sq cm (List separately in addition to code for primary procedure)
0977T	Upper gastrointestinal blood detection, sensor capsule, with interpretation and report
0978T	Submucosal cryolysis therapy; soft palate, base of tongue, and lingual tonsil
0979T	Submucosal cryolysis therapy; soft palate only
0980T	Submucosal cryolysis therapy; base of tongue and lingual tonsil only
0981T	Transcatheter implantation of wireless inferior vena cava sensor for long-term hemodynamic monitoring, including deployment of the sensor, radiological supervision and interpretation, right heart catheterization, and inferior vena cava venography, when performed
0982T	Remote monitoring of implantable inferior vena cava pressure sensor, physiologic parameter(s) (eg, weight, blood pressure, pulse oximetry, respiratory flow rate), initial set-up and patient education on use of equipment
0983T	Remote monitoring of an implanted inferior vena cava sensor for up to 30 days, including at least weekly downloads of inferior vena cava area recordings, interpretation(s), trend analysis, and report(s) by a physician or other qualified health care professional

0984T	Intravascular imaging of extracranial cerebral vessels using optical coherence tomography (OCT) during diagnostic evaluation and/or therapeutic intervention, including all associated radiological supervision, interpretation, and report; initial vessel (List separately in addition to code for primary procedure)
0985T	Intravascular imaging of extracranial cerebral vessels using optical coherence tomography (OCT) during diagnostic evaluation and/or therapeutic intervention, including all associated radiological supervision, interpretation, and report; each additional vessel (List separately in addition to code for primary procedure)
0986T	Intravascular imaging of intracranial cerebral vessels using optical coherence tomography (OCT) during diagnostic evaluation and/or therapeutic intervention, including all associated radiological supervision, interpretation, and report; initial vessel (List separately in addition to code for primary procedure)
0987T	Intravascular imaging of intracranial cerebral vessels using optical coherence tomography (OCT) during diagnostic evaluation and/or therapeutic intervention, including all associated radiological supervision, interpretation, and report; each additional vessel (List separately in addition to code for primary procedure)
0558U	Oncology (colorectal), quantitative enzyme-linked immunosorbent assay (ELISA) for secreted colorectal cancer protein marker (BF7 antigen), using serum, result reported as indicative of response/no response to therapy or disease progression/regression
0559U	Oncology (breast), quantitative enzyme-linked immunosorbent assay (ELISA) for secreted breast cancer protein marker (BF9 antigen), serum, result reported as indicative of response/no response to therapy or disease progression/regression
0568U	Neurology (dementia), beta amyloid (AB40, AB42, AB42/40 ratio), tau-protein phosphorylated at residue (eg, pTau217), neurofilament light chain (NfL), and glial fibrillary acidic protein (GFAP), by ultra-high sensitivity molecule array detection, plasma, algorithm reported as positive, intermediate, or negative for Alzheimer pathology
0570U	Neurology (traumatic brain injury), analysis of glial fibrillary acidic protein (GFAP) and ubiquitin carboxyl- terminal hydrolase L1 (UCH-L1), immunoassay, whole blood or plasma, individual components reported with the overall result of elevated or non-elevated based on threshold comparison
0573U	Oncology (pancreas), 3 biomarkers (glucose, carcinoembryonic antigen, and gastricsin), pancreatic cyst lesion fluid, algorithm reported as categorical mucinous or non-mucinous
0574U	Mycobacterium tuberculosis, culture filtrate protein-10-kDa (CFP-10), serum or plasma, liquid chromatography mass spectrometry (LC-MS)

Covered when prior authorized via eviCore for Commercial and ASO:

0552U	Reproductive medicine (preimplantation genetic assessment), analysis for known genetic disorders from trophectoderm biopsy, linkage analysis of disease-causing locus, and when possible, targeted mutation analysis for known familial variant, reported as low-risk or high-risk for familial genetic disorder
0553U	Reproductive medicine (preimplantation genetic assessment), analysis of 24 chromosomes using DNA genomic sequence analysis from embryonic trophectoderm for structural rearrangements, aneuploidy, and a mitochondrial DNA score, results reported as normal/balanced (euploidy/balanced), unbalanced structural rearrangement, monosomy, trisomy, segmental aneuploidy, or mosaic, per embryo tested

Reproductive medicine (preimplantation genetic assessment), analysis of 24 chromosomes using DNA genomic sequence analysis from trophectoderm biopsy for aneuploidy, ploidy, a mitochondrial DNA score, and embryo quality control, results reported as normal (euploidy), monosomy, trisomy, segmental aneuploidy, triploid, haploid, or mosaic, with quality control results reported as contamination detected or inconsistent cohort when applicable, per embryo tested Reproductive medicine (preimplantation genetic assessment), analysis of 24 chromosomes using DNA genomic sequence analysis from embryonic trophectoderm for structural rearrangements, aneuploidy, ploidy, a mitochondrial DNA score, and embryo quality control, results reported as normal/balanced (euploidy/balanced) unbalanced structural rearrangement, monosomy, trisomy, segmental aneuploidy, triploid, haploid, or mosaic, with quality control results reported as contamination detected or inconsistent cohort when applicable, per embryo tested Oncology (minimal residual disease (MRD)), genomic sequence analysis, cell-free DNA, whole blood and tumor tissue, baseline assessment for design and construction of a personalized variant panel to evaluate current MRD and for comparison to subsequent MRD assessments Oncology (solid tumor), targeted genomic sequence analysis, 33 genes, detection of single-nucleotide variants (SNVs), insertions and deletions, copy-number amplifications, and translocations in human genomic circulating cell-free DNA, plasma, reported as presence of actionable variants Oncology (fung), qPCR-based analysis of 13 differentially methylated regions (CCDC181, HOXA7, LRRC8A, MARCHE11, MIR129-2, NCOR2, PANTR1, PRKCB, SLC9A3, TBR1_2, TRAP1, VWC2, ZNF781), pleural fluid, algorithm reported as a qualitative result insertions, runs of homozygosity, aneuploidy, and inversions, mitochondrial DNA sequence and deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identif		
sequence analysis from embryonic trophectoderm for structural rearrangements, aneuploidy, ploidy, a mitochondrial DNA score, and embryo quality control, results reported as normal/balanced (euploidy/balanced), unbalanced structural rearrangement, monosomy, trisomy, segmental aneuploidy, triploid, haploid, or mosaic, with quality control results reported as contamination detected or inconsistent cohort when applicable, per embryo tested Oncology (minimal residual disease [MRD]), genomic sequence analysis, cell-free DNA, whole blood and tumor tissue, baseline assessment for design and construction of a personalized variant panel to evaluate current MRD and for comparison to subsequent MRD assessments Oncology (minimal residual disease [MRD]), genomic sequence analysis, cell-free DNA, whole blood, subsequent assessment with comparison to initial assessment to evaluate for MRD Oncology (solid tumor), targeted genomic sequence analysis, 33 genes, detection of single-nucleotide variants (SNVs), insertions and deletions, copy-number amplifications, and translocations in human genomic circulating cell-free DNA, plasma, reported as presence of actionable variants Oncology (hepatocellular carcinoma), next-generation sequencing methylation pattern assay to detect 6626 epigenetic alterations, cell-free DNA, plasma, algorithm reported as cancer signal detected or not detected Oncology (lung), qPCR-based analysis of 13 differentially methylated regions (CCDC181, HOXA7, LRRC8A, MARCHF11, MIR129-2, NCOR2, PANTR1, PRKCB, SLC9A3, TBR1_2, TRAP1, VWC2, ZNF781), pleural fluid, algorithm reported as a qualitative result S67U Rare diseases (constitutional/heritable disorders), whole-genome sequence analysis combination of short and long reads, for single-nucleotide variants, insertions/deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and categorization of genetic variants Oncology (solid tumor), next-generation sequenc	0554U	sequence analysis from trophectoderm biopsy for aneuploidy, ploidy, a mitochondrial DNA score, and embryo quality control, results reported as normal (euploidy), monosomy, trisomy, segmental aneuploidy, triploid, haploid, or mosaic, with quality control results reported as contamination detected or inconsistent cohort
tissue, baseline assessment for design and construction of a personalized variant panel to evaluate current MRD and for comparison to subsequent MRD assessments Oncology (minimal residual disease [MRD]), genomic sequence analysis, cell-free DNA, whole blood, subsequent assessment with comparison to initial assessment to evaluate for MRD Oncology (solid tumor), targeted genomic sequence analysis, 33 genes, detection of single-nucleotide variants (SNVs), insertions and deletions, copy-number amplifications, and translocations in human genomic circulating cell-free DNA, plasma, reported as presence of actionable variants Oncology (hepatocellular carcinoma), next-generation sequencing methylation pattern assay to detect 6626 epigenetic alterations, cell-free DNA, plasma, algorithm reported as cancer signal detected or not detected Oncology (lung), qPCR-based analysis of 13 differentially methylated regions (CCDC181, HOXA7, LRRC8A, MARCHF11, MIR129-2, NCOR2, PANTR1, PRKCB, SLC9A3, TBR1_2, TRAP1, VWC2, ZNF781), pleural fluid, algorithm reported as a qualitative result Os67U Rare diseases (constitutional/heritable disorders), whole-genome sequence analysis combination of short and long reads, for single-nucleotide variants, insertions/deletions and characterized intronic variants, copynumber variants, duplications/deletions, mobile element insertions, runs of homozygosity, aneuploidy, and inversions, mitochondrial DNA sequence and deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and categorization of genetic variants Oncology (solid tumor), next-generation sequencing analysis of tumor methylation markers (>20000 differentially methylated regions) present in cell-free circulating tumor DNA (ctDNA), whole blood, algorithm reported as presence or absence of ctDNA with tumor fraction, if appropriate Oncology (solid tumor), DNA (80 genes) and RNA (10 genes), by next-generation sequencing, plasma, including si	0555U	sequence analysis from embryonic trophectoderm for structural rearrangements, aneuploidy, ploidy, a mitochondrial DNA score, and embryo quality control, results reported as normal/balanced (euploidy/balanced), unbalanced structural rearrangement, monosomy, trisomy, segmental aneuploidy, triploid, haploid, or mosaic, with quality control results reported as contamination detected or inconsistent
subsequent assessment with comparison to initial assessment to evaluate for MRD Oncology (solid tumor), targeted genomic sequence analysis, 33 genes, detection of single-nucleotide variants (SNVs), insertions and deletions, copy-number amplifications, and translocations in human genomic circulating cell-free DNA, plasma, reported as presence of actionable variants Oncology (hepatocellular carcinoma), next-generation sequencing methylation pattern assay to detect 6626 epigenetic alterations, cell-free DNA, plasma, algorithm reported as cancer signal detected or not detected Oncology (lung), qPCR-based analysis of 13 differentially methylated regions (CCDC181, HOXA7, LRRC8A, MARCHF11, MIR129-2, NCOR2, PANTR1, PRKCB, SLC9A3, TBR1_2, TRAP1, VWC2, ZNF781), pleural fluid, algorithm reported as a qualitative result Rare diseases (constitutional/heritable disorders), whole-genome sequence analysis combination of short and long reads, for single-nucleotide variants, insertions/deletions and characterized intronic variants, copynumber variants, duplications/deletions, mobile element insertions, runs of homozygosity, aneuploidy, and inversions, mitochondrial DNA sequence and deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and categorization of genetic variants Oncology (solid tumor), next-generation sequencing analysis of tumor methylation markers (>20000 differentially methylated regions) present in cell-free circulating tumor DNA (ctDNA), whole blood, algorithm reported as presence or absence of ctDNA with tumor fraction, if appropriate Oncology (solid tumor), DNA (80 genes) and RNA (10 genes), by next-generation sequencing, plasma, including single-nucleotide variants, insertions/deletions, copy-number alterations, microsatellite instability, and fusions, reported as clinically actionable variants	0560U	tissue, baseline assessment for design and construction of a personalized variant panel to evaluate current
(SNVs), insertions and deletions, copy-number amplifications, and translocations in human genomic circulating cell-free DNA, plasma, reported as presence of actionable variants Oncology (hepatocellular carcinoma), next-generation sequencing methylation pattern assay to detect 6626 epigenetic alterations, cell-free DNA, plasma, algorithm reported as cancer signal detected or not detected Oncology (lung), qPCR-based analysis of 13 differentially methylated regions (CCDC181, HOXA7, LRRC8A, MARCHF11, MIR129-2, NCOR2, PANTR1, PRKCB, SLC9A3, TBR1_2, TRAP1, VWC2, ZNF781), pleural fluid, algorithm reported as a qualitative result Sare diseases (constitutional/heritable disorders), whole-genome sequence analysis combination of short and long reads, for single-nucleotide variants, insertions/deletions and characterized intronic variants, copynumber variants, duplications/deletions, mobile element insertions, runs of homozygosity, aneuploidy, and inversions, mitochondrial DNA sequence and deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and categorization of genetic variants Oncology (solid tumor), next-generation sequencing analysis of tumor methylation markers (>20000 differentially methylated regions) present in cell-free circulating tumor DNA (ctDNA), whole blood, algorithm reported as presence or absence of ctDNA with tumor fraction, if appropriate Oncology (solid tumor), DNA (80 genes) and RNA (10 genes), by next-generation sequencing, plasma, including single-nucleotide variants, insertions/deletions, copy-number alterations, microsatellite instability, and fusions, reported as clinically actionable variants Oncology (prostate), high-throughput telomere length quantification by FISH, whole blood, diagnostic	0561U	
epigenetic alterations, cell-free DNA, plasma, algorithm reported as cancer signal detected or not detected Osciology (lung), qPCR-based analysis of 13 differentially methylated regions (CCDC181, HOXA7, LRRC8A, MARCHF11, MIR129-2, NCOR2, PANTR1, PRKCB, SLC9A3, TBR1_2, TRAP1, VWC2, ZNF781), pleural fluid, algorithm reported as a qualitative result Osciology (selid tunor), and long reads, for single-nucleotide variants, insertions/deletions and characterized intronic variants, copynumber variants, duplications/deletions, mobile element insertions, runs of homozygosity, aneuploidy, and inversions, mitochondrial DNA sequence and deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and categorization of genetic variants Osciology (solid tumor), next-generation sequencing analysis of tumor methylation markers (>20000 differentially methylated regions) present in cell-free circulating tumor DNA (ctDNA), whole blood, algorithm reported as presence or absence of ctDNA with tumor fraction, if appropriate Osciology (solid tumor), DNA (80 genes) and RNA (10 genes), by next-generation sequencing, plasma, including single-nucleotide variants, insertions/deletions, copy-number alterations, microsatellite instability, and fusions, reported as clinically actionable variants Osciology (prostate), high-throughput telomere length quantification by FISH, whole blood, diagnostic	0562U	(SNVs), insertions and deletions, copy-number amplifications, and translocations in human genomic circulating
MARCHF11, MIR129-2, NCOR2, PANTR1, PRKCB, SLC9A3, TBR1_2, TRAP1, VWC2, ZNF781), pleural fluid, algorithm reported as a qualitative result Rare diseases (constitutional/heritable disorders), whole-genome sequence analysis combination of short and long reads, for single-nucleotide variants, insertions/deletions and characterized intronic variants, copynumber variants, duplications/deletions, mobile element insertions, runs of homozygosity, aneuploidy, and inversions, mitochondrial DNA sequence and deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and categorization of genetic variants Oncology (solid tumor), next-generation sequencing analysis of tumor methylation markers (>20000 differentially methylated regions) present in cell-free circulating tumor DNA (ctDNA), whole blood, algorithm reported as presence or absence of ctDNA with tumor fraction, if appropriate Oncology (solid tumor), DNA (80 genes) and RNA (10 genes), by next-generation sequencing, plasma, including single-nucleotide variants, insertions/deletions, copy-number alterations, microsatellite instability, and fusions, reported as clinically actionable variants Oncology (prostate), high-throughput telomere length quantification by FISH, whole blood, diagnostic	0565U	
long reads, for single-nucleotide variants, insertions/deletions and characterized intronic variants, copynumber variants, duplications/deletions, mobile element insertions, runs of homozygosity, aneuploidy, and inversions, mitochondrial DNA sequence and deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and categorization of genetic variants Oncology (solid tumor), next-generation sequencing analysis of tumor methylation markers (>20000 differentially methylated regions) present in cell-free circulating tumor DNA (ctDNA), whole blood, algorithm reported as presence or absence of ctDNA with tumor fraction, if appropriate Oncology (solid tumor), DNA (80 genes) and RNA (10 genes), by next-generation sequencing, plasma, including single-nucleotide variants, insertions/deletions, copy-number alterations, microsatellite instability, and fusions, reported as clinically actionable variants Oncology (prostate), high-throughput telomere length quantification by FISH, whole blood, diagnostic	0566U	MARCHF11, MIR129-2, NCOR2, PANTR1, PRKCB, SLC9A3, TBR1_2, TRAP1, VWC2, ZNF781), pleural fluid,
differentially methylated regions) present in cell-free circulating tumor DNA (ctDNA), whole blood, algorithm reported as presence or absence of ctDNA with tumor fraction, if appropriate Oncology (solid tumor), DNA (80 genes) and RNA (10 genes), by next-generation sequencing, plasma, including single-nucleotide variants, insertions/deletions, copy-number alterations, microsatellite instability, and fusions, reported as clinically actionable variants Oncology (prostate), high-throughput telomere length quantification by FISH, whole blood, diagnostic	0567U	long reads, for single-nucleotide variants, insertions/deletions and characterized intronic variants, copy- number variants, duplications/deletions, mobile element insertions, runs of homozygosity, aneuploidy, and inversions, mitochondrial DNA sequence and deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and
single-nucleotide variants, insertions/deletions, copy-number alterations, microsatellite instability, and fusions, reported as clinically actionable variants Oncology (prostate), high-throughput telomere length quantification by FISH, whole blood, diagnostic	0569U	differentially methylated regions) present in cell-free circulating tumor DNA (ctDNA), whole blood, algorithm
	0571U	single-nucleotide variants, insertions/deletions, copy-number alterations, microsatellite instability, and fusions,
	0572U	

Covered with no prior authorization for Commercial and ASO:

0556U	Infectious disease (bacterial or viral respiratory tract infection), pathogen-specific DNA and RNA by real-time PCR, 12 targets, nasopharyngeal or oropharyngeal swab, including multiplex reverse transcription for RNA targets, each analyte reported as detected or not detected
0557U	Infectious disease (bacterial vaginosis and vaginitis), real-time amplification of DNA markers for Atopobium vaginae, Gardnerella vaginalis, Megasphaera types 1 and 2, bacterial vaginosis associated bacteria-2 and -3 (BVAB-2, BVAB-3), Mobiluncus species, Trichomonas vaginalis, Neisseria gonorrhoeae, Candida species (C. albicans, C. tropicalis, C. parapsilosis, C. glabrata, C. krusei), Herpes simplex viruses 1 and 2, vaginal fluid, reported as detected or not detected for each organism
0563U	Infectious disease (bacterial and/or viral respiratory tract infection), pathogen-specific nucleic acid (DNA or RNA), 11 viral targets and 4 bacterial targets, qualitative RT-PCR, upper respiratory specimen, each pathogen reported as positive or negative
0564U	Infectious disease (bacterial and/or viral respiratory tract infection), pathogen-specific nucleic acid (DNA or RNA), 10 viral targets and 4 bacterial targets, qualitative RT-PCR, upper respiratory specimen, each pathogen reported as positive or negative

Not covered experimental and investigational for Medicare Advantage:

90382	Respiratory syncytial virus, monoclonal antibody, seasonal dose, 0.7 mL, for intramuscular use
90362	
	Influenza virus vaccine, trivalent, and severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)
90612	(coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 31.7 mcg/0.32 mL dosage, for intramuscular use
90613	Influenza virus vaccine, quadrivalent, and severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)
	(coronavirus disease [COVID-19]) vaccine, mRNA-LNP, 40 mcg/0.4 mL dosage, for intramuscular use
0558U	Oncology (colorectal), quantitative enzyme-linked immunosorbent assay (ELISA) for secreted colorectal cancer
	protein marker (BF7 antigen), using serum, result reported as indicative of response/no response to therapy or disease progression/regression
0559U	Oncology (breast), quantitative enzyme-linked immunosorbent assay (ELISA) for secreted breast cancer protein
	marker (BF9 antigen), serum, result reported as indicative of response/no response to therapy or disease progression/regression
	Neurology (dementia), beta amyloid (AB40, AB42, AB42/40 ratio), tau-protein phosphorylated at residue (eg,
0568U	pTau217), neurofilament light chain (NfL), and glial fibrillary acidic protein (GFAP), by ultra-high sensitivity
	molecule array detection, plasma, algorithm reported as positive, intermediate, or negative for Alzheimer pathology
	Neurology (traumatic brain injury), analysis of glial fibrillary acidic protein (GFAP) and ubiquitin carboxyl-
0570U	terminal hydrolase L1 (UCH-L1), immunoassay, whole blood or plasma, individual components reported with
	the overall result of elevated or non-elevated based on threshold comparison
057211	Oncology (pancreas), 3 biomarkers (glucose, carcinoembryonic antigen, and gastricsin), pancreatic cyst lesion
0573U	fluid, algorithm reported as categorical mucinous or non-mucinous
	Mycobacterium tuberculosis, culture filtrate protein-10-kDa (CFP-10), serum or plasma, liquid chromatography
0574U	mass spectrometry (LC-MS)
0948T	Interrogation device evaluation (remote), up to 90 days, cardiac contractility modulation system with interim
	analysis, review and report(s) by a physician or other qualified health care professional

0949T	Interrogation device evaluation (remote), up to 90 days, cardiac contractility modulation system, remote data acquisition(s), receipt of transmissions, technician review, technical support, and distribution of results
0950T	Ablation of benign prostate tissue, transrectal, with high intensity-focused ultrasound (HIFU), including ultrasound guidance
0951T	Totally implantable active middle ear hearing implant; initial placement, including mastoidectomy, placement of and attachment to sound processor
0952T	Totally implantable active middle ear hearing implant; revision or replacement, with mastoidectomy and replacement of sound processor
0953T	Totally implantable active middle ear hearing implant; revision or replacement, without mastoidectomy and replacement of sound processor
0954T	Totally implantable active middle ear hearing implant; replacement of sound processor only, with attachment to existing transducers
0955T	Totally implantable active middle ear hearing implant; removal, including removal of sound processor and all implant components
0956T	Partial craniectomy, channel creation, and tunneling of electrode for sub-scalp implantation of an electrode array, receiver, and telemetry unit for continuous bilateral electroencephalography monitoring system, including imaging guidance
0957T	Revision of sub-scalp implanted electrode array, receiver, and telemetry unit for electrode, when required, including imaging guidance
0958T	Removal of sub-scalp implanted electrode array, receiver, and telemetry unit for continuous bilateral electroencephalography monitoring system, including imaging guidance
0959T	Removal or replacement of magnet from coil assembly that is connected to continuous bilateral electroencephalography monitoring system, including imaging guidance
0960T	Replacement of sub-scalp implanted electrode array, receiver, and telemetry unit with tunneling of electrode for continuous bilateral electroencephalography monitoring system, including imaging guidance
0961T	Shortwave infrared radiation imaging, surgical pathology specimen, to assist gross examination for lymph node localization in fibroadipose tissue, per specimen (List separately in addition to code for primary procedure)
0962T	Assistive algorithmic analysis of acoustic and electrocardiogram recording for detection of cardiac dysfunction (eg, reduced ejection fraction, cardiac murmurs, atrial fibrillation), with review and interpretation by a physician or other qualified health care professional
0963T	Anoscopy with directed submucosal injection of bulking agent into anal canal
0964T	Impression and custom preparation of jaw expansion oral prosthesis for obstructive sleep apnea, including initial adjustment; single arch, without mandibular advancement mechanism
0965T	Impression and custom preparation of jaw expansion oral prosthesis for obstructive sleep apnea, including initial adjustment; dual arch, with additional mandibular advancement, fixed hinge mechanism

0966T	Transanal insertion of endoluminal temporary colorectal anastomosis protection device, including vacuum anchoring component and flexible sheath connected to external vacuum source and monitoring system
0967T	Insertion or replacement of epicranial neurostimulator system, including electrode array and pulse generator, with connection to electrode array
0968T	Removal of epicranial neurostimulator system
0969T	Ablation, benign breast tumor (eg, fibroadenoma), percutaneous, laser, including imaging guidance when performed, each tumor
0970T	Ablation, malignant breast tumor(s), percutaneous, laser, including imaging guidance when performed, unilateral
0971T	Assistive algorithmic classification of burn healing (ie, healing or nonhealing) by noninvasive multispectral imaging, including system set-up and acquisition, selection, and transmission of images, with automated generation of report
0972T	Selective enzymatic debridement, partial-thickness and/or full-thickness burn eschar, requiring anesthesia (ie, general anesthesia, moderate sedation), including patient monitoring, trunk, arms, legs; first 100 sq cm
0973T	Selective enzymatic debridement, partial-thickness and/or full-thickness burn eschar, requiring anesthesia (ie, general anesthesia, moderate sedation), including patient monitoring, trunk, arms, legs; each additional 100 sq cm (List separately in addition to code for primary procedure)
0974T	Selective enzymatic debridement, partial-thickness and/or full-thickness burn eschar, requiring anesthesia (ie, general anesthesia, moderate sedation), including patient monitoring, scalp, neck, hands, feet, and/or multiple digits; first 100 sq cm
0975T	Selective enzymatic debridement, partial-thickness and/or full-thickness burn eschar, requiring anesthesia (ie, general anesthesia, moderate sedation), including patient monitoring, scalp, neck, hands, feet, and/or multiple digits; each additional 100 sq cm (List separately in addition to code for primary procedure)
0976T	Upper gastrointestinal blood detection, sensor capsule, with interpretation and report
0977T	Submucosal cryolysis therapy; soft palate, base of tongue, and lingual tonsil
0978T	Submucosal cryolysis therapy; soft palate only
0979T	Submucosal cryolysis therapy; base of tongue and lingual tonsil only
0980T	Transcatheter implantation of wireless inferior vena cava sensor for long-term hemodynamic monitoring, including deployment of the sensor, radiological supervision and interpretation, right heart catheterization, and inferior vena cava venography, when performed
0981T	Remote monitoring of implantable inferior vena cava pressure sensor, physiologic parameter(s) (eg, weight, blood pressure, pulse oximetry, respiratory flow rate), initial set-up and patient education on use of equipment
0982T	Remote monitoring of an implanted inferior vena cava sensor for up to 30 days, including at least weekly downloads of inferior vena cava area recordings, interpretation(s), trend analysis, and report(s) by a physician or other qualified health care professional

0983T	Remote monitoring of an implanted inferior vena cava sensor for up to 30 days, including at least weekly
	downloads of inferior vena cava area recordings, interpretation(s), trend analysis, and report(s) by a physician
	or other qualified health care professional
0984T	Intravascular imaging of extracranial cerebral vessels using optical coherence tomography (OCT) during
	diagnostic evaluation and/or therapeutic intervention, including all associated radiological supervision,
	interpretation, and report; initial vessel (List separately in addition to code for primary procedure)
0985T	Intravascular imaging of extracranial cerebral vessels using optical coherence tomography (OCT) during
	diagnostic evaluation and/or therapeutic intervention, including all associated radiological supervision,
	interpretation, and report; each additional vessel (List separately in addition to code for primary procedure)
0986T	Intravascular imaging of intracranial cerebral vessels using optical coherence tomography (OCT) during
	diagnostic evaluation and/or therapeutic intervention, including all associated radiological supervision,
	interpretation, and report; initial vessel (List separately in addition to code for primary procedure)
0987T	Intravascular imaging of intracranial cerebral vessels using optical coherence tomography (OCT) during
	diagnostic evaluation and/or therapeutic intervention, including all associated radiological supervision,
	interpretation, and report; each additional vessel (List separately in addition to code for primary procedure)

Covered with prior authorization for Medicare Advantage:

0552U	Reproductive medicine (preimplantation genetic assessment), analysis for known genetic disorders from trophectoderm biopsy, linkage analysis of disease-causing locus, and when possible, targeted mutation analysis for known familial variant, reported as low-risk or high-risk for familial genetic disorder
0553U	Reproductive medicine (preimplantation genetic assessment), analysis of 24 chromosomes using DNA genomic sequence analysis from embryonic trophectoderm for structural rearrangements, aneuploidy, and a mitochondrial DNA score, results reported as normal/balanced (euploidy/balanced), unbalanced structural rearrangement, monosomy, trisomy, segmental aneuploidy, or mosaic, per embryo tested
0554U	Reproductive medicine (preimplantation genetic assessment), analysis of 24 chromosomes using DNA genomic sequence analysis from trophectoderm biopsy for aneuploidy, ploidy, a mitochondrial DNA score, and embryo quality control, results reported as normal (euploidy), monosomy, trisomy, segmental aneuploidy, triploid, haploid, or mosaic, with quality control results reported as contamination detected or inconsistent cohort when applicable, per embryo tested
0555U	Reproductive medicine (preimplantation genetic assessment), analysis of 24 chromosomes using DNA genomic sequence analysis from embryonic trophectoderm for structural rearrangements, aneuploidy, ploidy, a mitochondrial DNA score, and embryo quality control, results reported as normal/balanced (euploidy/balanced), unbalanced structural rearrangement, monosomy, trisomy, segmental aneuploidy, triploid, haploid, or mosaic, with quality control results reported as contamination detected or inconsistent cohort when applicable, per embryo tested
0560U	Oncology (minimal residual disease [MRD]), genomic sequence analysis, cell-free DNA, whole blood and tumor tissue, baseline assessment for design and construction of a personalized variant panel to evaluate current MRD and for comparison to subsequent MRD assessments

0561U	Oncology (minimal residual disease [MRD]), genomic sequence analysis, cell-free DNA, whole blood, subsequent assessment with comparison to initial assessment to evaluate for MRD
0562U	Oncology (solid tumor), targeted genomic sequence analysis, 33 genes, detection of single-nucleotide variants (SNVs), insertions and deletions, copy-number amplifications, and translocations in human genomic circulating cell-free DNA, plasma, reported as presence of actionable variants
0565U	Oncology (hepatocellular carcinoma), next-generation sequencing methylation pattern assay to detect 6626 epigenetic alterations, cell-free DNA, plasma, algorithm reported as cancer signal detected or not detected
0566U	Oncology (lung), qPCR-based analysis of 13 differentially methylated regions (CCDC181, HOXA7, LRRC8A, MARCHF11, MIR129-2, NCOR2, PANTR1, PRKCB, SLC9A3, TBR1_2, TRAP1, VWC2, ZNF781), pleural fluid, algorithm reported as a qualitative result
0567U	Rare diseases (constitutional/heritable disorders), whole-genome sequence analysis combination of short and long reads, for single-nucleotide variants, insertions/deletions and characterized intronic variants, copy-number variants, duplications/deletions, mobile element insertions, runs of homozygosity, aneuploidy, and inversions, mitochondrial DNA sequence and deletions, short tandem repeat genes, methylation status of selected regions, blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and categorization of genetic variants
0569U	Oncology (solid tumor), next-generation sequencing analysis of tumor methylation markers (>20000 differentially methylated regions) present in cell-free circulating tumor DNA (ctDNA), whole blood, algorithm reported as presence or absence of ctDNA with tumor fraction, if appropriate
0571U	Oncology (solid tumor), DNA (80 genes) and RNA (10 genes), by next-generation sequencing, plasma, including single-nucleotide variants, insertions/deletions, copy-number alterations, microsatellite instability, and fusions, reported as clinically actionable variants
0572U	Oncology (prostate), high-throughput telomere length quantification by FISH, whole blood, diagnostic algorithm reported as risk of prostate cancer

Covered with no prior authorization for Medicare Advantage:

`	Covered with no phot authorization for inedicate Advantage.	
Q4368	Amchothick, per square centimeter	
Q4369	Amnioplast 3, per square centimeter	
Q4370	Aeroguard, per square centimeter	
Q4371	Neoguard, per square centimeter	
Q4372	Amchoplast excel, per square centimeter	
Q4373	Membrane wrap lite, per square centimeter	

Q4375	Duograft ac, per square centimeter
Q4376	Duograft aa, per square centimeter
Q4377	Trigraft ft, per square centimeter
Q4378	Renew ft matrix, per square centimeter
Q4379	Amniodefend ft matrix, per square centimeter
Q4380	Advograft one, per square centimeter
Q4382	Advograft dual, per square centimeter
0556U	Infectious disease (bacterial or viral respiratory tract infection), pathogen-specific DNA and RNA by real-time PCR, 12 targets, nasopharyngeal or oropharyngeal swab, including multiplex reverse transcription for RNA targets, each analyte reported as detected or not detected
0557U	Infectious disease (bacterial vaginosis and vaginitis), real-time amplification of DNA markers for Atopobium vaginae, Gardnerella vaginalis, Megasphaera types 1 and 2, bacterial vaginosis associated bacteria-2 and -3 (BVAB-2, BVAB-3), Mobiluncus species, Trichomonas vaginalis, Neisseria gonorrhoeae, Candida species (C. albicans, C. tropicalis, C. parapsilosis, C. glabrata, C. krusei), Herpes simplex viruses 1 and 2, vaginal fluid, reported as detected or not detected for each organism
0563U	Infectious disease (bacterial and/or viral respiratory tract infection), pathogen-specific nucleic acid (DNA or RNA), 11 viral targets and 4 bacterial targets, qualitative RT-PCR, upper respiratory specimen, each pathogen reported as positive or negative
0564U	Infectious disease (bacterial and/or viral respiratory tract infection), pathogen-specific nucleic acid (DNA or RNA), 10 viral targets and 4 bacterial targets, qualitative RT-PCR, upper respiratory specimen, each pathogen reported as positive or negative

As a reminder, APAD and APEC carve-out drugs are not covered under ACO and will redirect to MassHealth effective 4/1/2025 for MGB ACO Plans:

J3391	Injection, atidarsagene autotemcel, per treatment
Q2058	Obecabtagene autoleucel, 10 up to 400 million cd19 car-positive viable t cells, including leukapheresis and
	dose preparation procedures, per infusion

Not covered under medical benefit, coverage redirected under pharmacy benefit for MGB ACO:

not covered under medical benefit, coverage real edical under pharmacy benefit for med neo.	
J7172	Injection, marstacimab-hncq, 0.5 mg

J7356	Injection, foscarbidopa 0.25 mg/foslevodopa 5 mg
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Not covered under medical benefit, coverage redirected under pharmacy benefit for Commercial:

J7356	Injection, foscarbidopa 0.25 mg/foslevodopa 5 mg

New to Market not covered unless prior authorized for MGB ACO:

J9275	Injection, cosibelimab-ipdl, 2 mg
Q5098	Injection, ustekinumab-srlf (imuldosa), biosimilar, 1 mg
Q5153	Injection, aflibercept-yszy (opuviz), biosimilar, 1 mg

New to Market not covered unless prior authorized for Commercial:

J9174	Injection, docetaxel (beizray), 1 mg
J9275	Injection, cosibelimab-ipdl, 2 mg
Q5098	Injection, ustekinumab-srlf (imuldosa), biosimilar, 1 mg
Q5099	Injection, ustekinumab-stba (steqeyma), biosimilar, 1 mg
Q5100	Injection, ustekinumab-kfce (yesintek), biosimilar, 1 mg
Q5153	Injection, aflibercept-yszy (opuviz), biosimilar, 1 mg

New to Market not covered unless prior authorized for Medicare Advantage:

Q5098	Injection, ustekinumab-srlf (imuldosa), biosimilar, 1 mg
Q5153	Injection, aflibercept-yszy (opuviz), biosimilar, 1 mg

Covered with prior authorization for Commercial:

C9174	Injection, datopotamab deruxtecan-dlnk, 1 mg
J1326	Injection, zolbetuximab-clzb, 2 mg
J3391	Injection, atidarsagene autotemcel, per treatment
J7172	Injection, marstacimab-hncq, 0.5 mg
J9276	Injection, zanidatamab-hrii, 2 mg
J9289	Injection, nivolumab, 2 mg and hyaluronidase-nvhy
J9382	Injection, zenocutuzumab-zbco, 1 mg
Q2058	Obecabtagene autoleucel, 10 up to 400 million cd19 car-positive viable t cells, including leukapheresis and
	dose preparation procedures, per infusion

Covered with prior authorization for MGB ACO:

C9174	Injection, datopotamab deruxtecan-dlnk, 1 mg
J1326	Injection, zolbetuximab-clzb, 2 mg
J9276	Injection, zanidatamab-hrii, 2 mg
J9289	Injection, nivolumab, 2 mg and hyaluronidase-nvhy
J9382	Injection, zenocutuzumab-zbco, 1 mg
Q5099	Injection, ustekinumab-stba (steqeyma), biosimilar, 1 mg
Q5100	Injection, ustekinumab-kfce (yesintek), biosimilar, 1 mg

Covered prior authorization for Medicare Advantage:

	covered prior authorization for Medicare Advantage:
C9174	Injection, datopotamab deruxtecan-dlnk, 1 mg
J1326	Injection, zolbetuximab-clzb, 2 mg
J3391	Injection, atidarsagene autotemcel, per treatment
J7172	Injection, marstacimab-hncq, 0.5 mg
J9275	Injection, cosibelimab-ipdl, 2 mg
J9276	Injection, zanidatamab-hrii, 2 mg
J9289	Injection, nivolumab, 2 mg and hyaluronidase-nvhy
J9341	Injection, thiotepa (tepylute), 1 mg
J9382	Injection, zenocutuzumab-zbco, 1 mg
Q2058	Obecabtagene autoleucel, 10 up to 400 million cd19 car-positive viable t cells, including leukapheresis and dose preparation procedures, per infusion
Q5099	Injection, ustekinumab-stba (steqeyma), biosimilar, 1 mg
Q5100	Injection, ustekinumab-kfce (yesintek), biosimilar, 1 mg
C9174	Injection, datopotamab deruxtecan-dlnk, 1 mg
J1326	Injection, zolbetuximab-clzb, 2 mg
J3391	Injection, atidarsagene autotemcel, per treatment
J7172	Injection, marstacimab-hncq, 0.5 mg
J9275	Injection, cosibelimab-ipdl, 2 mg

J9276	Injection, zanidatamab-hrii, 2 mg
J9289	Injection, nivolumab, 2 mg and hyaluronidase-nvhy
J9341	Injection, thiotepa (tepylute), 1 mg
J9382	Injection, zenocutuzumab-zbco, 1 mg
Q2058	Obecabtagene autoleucel, 10 up to 400 million cd19 car-positive viable t cells, including leukapheresis and dose preparation procedures, per infusion
Q5099	Injection, ustekinumab-stba (steqeyma), biosimilar, 1 mg
Q5100	Injection, ustekinumab-kfce (yesintek), biosimilar, 1 mg

Covered no prior authorization required for Commercial:

J0165	Injection, epinephrine, not otherwise specified, 0.1 mg
J0166	Injection, epinephrine (bpi), not therapeutically equivalent to j0165, 0.1 mg
J0167	Injection, epinephrine (hospira), not therapeutically equivalent to j0165, 0.1 mg
J0168	Injection, epinephrine (international medication systems), not therapeutically equivalent to j0165, 0.1 mg
J0169	Injection, epinephrine (adrenalin), not therapeutically equivalent to j0165, 0.1 mg
J0616	Injection, metoprolol tartrate, 1 mg
J0618	Injection, calcium chloride, 2 mg
J1163	Injection, diltiazem hydrochloride, 0.5 mg
J2312	Injection, naloxone hydrochloride, not otherwise specified, 0.01 mg
J2313	Injection, naloxone hydrochloride (zimhi), 0.01 mg
J3373	Injection, vancomycin hydrochloride, 10 mg
J3374	Injection, vancomycin hydrochloride (mylan) not therapeutically equivalent to j3373, 10 mg
J3375	Injection, vancomycin hydrochloride (xellia), not therapeutically equivalent to j3373, 10 mg
J9220	Injection, indigotindisulfonate sodium, 1 mg
J9341	Injection, thiotepa (tepylute), 1 mg
J9342	Injection, thiotepa, not otherwise specified, 1 mg
J0165	Injection, epinephrine, not otherwise specified, 0.1 mg
J0166	Injection, epinephrine (bpi), not therapeutically equivalent to j0165, 0.1 mg
J0167	Injection, epinephrine (hospira), not therapeutically equivalent to j0165, 0.1 mg
J0168	Injection, epinephrine (international medication systems), not therapeutically equivalent to j0165, 0.1 mg
J0169	Injection, epinephrine (adrenalin), not therapeutically equivalent to j0165, 0.1 mg
J0616	Injection, metoprolol tartrate, 1 mg
J0618	Injection, calcium chloride, 2 mg

J1163	Injection, diltiazem hydrochloride, 0.5 mg
J2312	Injection, naloxone hydrochloride, not otherwise specified, 0.01 mg
J2313	Injection, naloxone hydrochloride (zimhi), 0.01 mg
J3373	Injection, vancomycin hydrochloride, 10 mg
J3374	Injection, vancomycin hydrochloride (mylan) not therapeutically equivalent to j3373, 10 mg
J3375	Injection, vancomycin hydrochloride (xellia), not therapeutically equivalent to j3373, 10 mg
J9220	Injection, indigotindisulfonate sodium, 1 mg
J9341	Injection, thiotepa (tepylute), 1 mg
J9342	Injection, thiotepa, not otherwise specified, 1 mg

Covered no prior authorization required for MGB ACO:

J0165	Injection, epinephrine, not otherwise specified, 0.1 mg
J0166	Injection, epinephrine (bpi), not therapeutically equivalent to j0165, 0.1 mg
J0167	Injection, epinephrine (hospira), not therapeutically equivalent to j0165, 0.1 mg
J0168	Injection, epinephrine (international medication systems), not therapeutically equivalent to j0165, 0.1 mg
J0169	Injection, epinephrine (adrenalin), not therapeutically equivalent to j0165, 0.1 mg
J0616	Injection, metoprolol tartrate, 1 mg
J0618	Injection, calcium chloride, 2 mg
J1163	Injection, diltiazem hydrochloride, 0.5 mg
J2312	Injection, naloxone hydrochloride, not otherwise specified, 0.01 mg
J2313	Injection, naloxone hydrochloride (zimhi), 0.01 mg
J3373	Injection, vancomycin hydrochloride, 10 mg
J3374	Injection, vancomycin hydrochloride (mylan) not therapeutically equivalent to j3373, 10 mg
J3375	Injection, vancomycin hydrochloride (xellia), not therapeutically equivalent to j3373, 10 mg
J9174	Injection, docetaxel (beizray), 1 mg
J9341	Injection, thiotepa (tepylute), 1 mg
J9342	Injection, thiotepa, not otherwise specified, 1 mg

Covered no prior authorization required for Medicare Advantage:

Covered no prior dutiforization required for intedistrict Advantage.		
Injection, epinephrine, not otherwise specified, 0.1 mg		
Injection, epinephrine (bpi), not therapeutically equivalent to j0165, 0.1 mg		
Injection, epinephrine (hospira), not therapeutically equivalent to j0165, 0.1 mg		
Injection, epinephrine (international medication systems), not therapeutically equivalent to j0165, 0.1 mg		
Injection, epinephrine (adrenalin), not therapeutically equivalent to j0165, 0.1 mg		
Injection, metoprolol tartrate, 1 mg		
Injection, calcium chloride, 2 mg		
Injection, diltiazem hydrochloride, 0.5 mg		
Injection, naloxone hydrochloride, not otherwise specified, 0.01 mg		
Injection, naloxone hydrochloride (zimhi), 0.01 mg		
Injection, vancomycin hydrochloride, 10 mg		
Injection, vancomycin hydrochloride (mylan) not therapeutically equivalent to j3373, 10 mg		
Injection, vancomycin hydrochloride (xellia), not therapeutically equivalent to j3373, 10 mg		
Injection, foscarbidopa 0.25 mg/foslevodopa 5 mg		

J9174	Injection, docetaxel (beizray), 1 mg
J9220	Injection, indigotindisulfonate sodium, 1 mg
J9342	Injection, thiotepa, not otherwise specified, 1 mg