# **Proprietary Laboratory Analyses (PLA)**

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Proprietary Laboratory Analyses (PLA) codes represent proprietary laboratory services. The following codes may include a range of laboratory tests including, but not limited to multianalyte assays with algorithmic analyses (MAAA) and genomic sequencing procedures (GSP). MAAAs are procedures that utilize multiple results derived from assays of various types, including molecular pathology assays, fluorescent in situ hybridization assays and non-nucleic acid-based assays (for example., proteins, polypeptides, lipids, carbohydrates). Consistent with CPT<sup>®</sup> coding guidelines, when a PLA code is available, the specific PLA code takes precedence.

# Table of Proprietary Laboratory Analyses (PLA) Codes

Code and Code Description	TAR and/or Billing Requirements	Frequency
0014M	N/A	N/A
Liver disease, analysis of 3 biomarkers (hyaluronic acid [HA], procollagen III amino terminal peptide [PIIINP], tissue inhibitor of metalloproteinase 1 [TIMP-1]), using immunoassays, utilizing serum, prognostic algorithm reported as a risk score and risk of liver fibrosis and liver-related clinical events within 5 years		
<<0017M	One of the following ICD-10-CM diagnosis codes is	Once in a
Oncology (diffuse large b-cell lymphoma [dlbcl]), mRNA, gene expression profiling by fluorescent probe hybridization of 20 genes, formalin-fixed paraffin embedded tissue, algorithm reported as cell of origin	required on the claim: C83.30, C83.31, C83.32, C83.33, C83.34, C83.35, C83.36, C83.37, C83.38, C83.39	lifetime>>

Code and Code Description	TAR and/or Billing Requirements	Frequency
0001U	N/A	N/A
Red blood cell antigen typing, DNA, human erythrocyte antigen gene analysis of 35 antigens from 11 blood groups, utilizing whole blood, common RBC alleles reported		
0003U	The following ICD-10-CM diagnosis code is required on the claim: R19.09	Once in a
Oncology (ovarian) biochemical assays of five		lifetime, with
proteins (apolipoprotein A-1, CA 125 II, follicle stimulating hormone, human epididymis protein	Reimbursable for females who meet the following criteria:	TAR/SAR
4, transferrin), utilizing serum, algorithm reported as a likelihood score	<ul> <li>18 years of age or older</li> </ul>	override
	<ul> <li>Ovarian adnexal mass present for which surgery is planned, and not yet referred to an oncologist</li> </ul>	
0007U	N/A	N/A
Drug test(s), presumptive, with definitive confirmation of positive results, any number of drug classes, urine, includes specimen verification including DNA authentication in comparison to buccal DNA, per date of service		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0016U Oncology (hematolymphoid neoplasia), RNA, NCR/ABL1 major and minor breakpoint fusion transcripts, quantitative PCR amplification, blood or bone marrow, report of fusion not detected or detected with quantitation	One of the following ICD-10-CM diagnosis codes is required on the claim: C92.10 thru C92.12	N/A
0017U Oncology (hematolymphoid neoplasia), JAK2 mutation, DNA, PCR amplification of exons 12-14 and sequence analysis, blood or bone marrow, report pf JAK2 mutation not detected or detected	One of the following ICD-10-CM diagnosis codes is required on the claim: D45, D47.1 or D47.3	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0018U	The service requires a TAR.	Once in a
Oncology (thyroid), microRNA profiling by	A TAR requires documentation of the following criteria:	lifetime, with TAR/SAR
RT-PCR of 10 microRNA sequences, utilizing fine needle aspirate, algorithm reported as a positive or negative result for moderate to high risk of malignancy	The patient is under evaluation for thyroid nodule(s)	
	The cytopathology result from fine needle aspiration is indeterminate, defined as one of the following:	override
	Follicular lesion of undetermined significance (FLUS), Bethesda III, or	
	Atypia of undetermined significance (AUS), Bethesda III, or	
	Follicular neoplasm, Bethesda IV	
	The diagnostic or treatment strategy will be contingent on test results	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0022U	The service requires a TAR.	Once in a
Targeted genomic sequence analysis panel, non-small cell lung neoplasia, DNA and RNA analysis, 23 genes, interrogation for sequence variants and rearrangements, reported as presence/absence of variants and associated therapy(ies) to consider.	<ul> <li>A TAR requires documentation of the following criteria:</li> <li>Patient has a diagnosis of non-small cell lung cancer (NSCLC).</li> <li>Treatment is contingent on test results</li> </ul>	lifetime, with TAR/SAR override

Code and Code Description	TAR and/or Billing Requirements	Frequency
0023U Oncology (acute myelogenous leukemia), DNA, gentotyping of internal tandem duplication, p.D835, p.I836, using mononuclear cells, reported as detection or non-detection of FLT3 mutation and indication for or against the use of midostaurin.	One of the following ICD-10-CM diagnosis codes is required on the claim: C92.00 thru C92.02, C92.60 thru C92.62, C92.A0 thru C92.A2	N/A
0026U	The service requires a TAR.	Once in a
Oncology (thyroid), DNA and mRNA of 112 genes, next-generation sequencing, fine needle aspirate of thyroid nodule, algorithmic analysis reported as a categorical result ("Positive, high probability of malignancy" or "Negative, low probability of malignancy").	A TAR requires documentation of the following criteria:	lifetime, with TAR/SAR override
	<ol> <li>The patient is under evaluation for thyroid nodule(s)</li> </ol>	
	The cytopathology result from fine needle aspiration is indeterminate, defined as one of the following:	
	a. Follicular lesion of undetermined significance (FLUS), Bethesda III, or	
	Atypia of undetermined significance (AUS), Bethesda III, or	
	Follicular neoplasm, Bethesda IV	
	The diagnostic or treatment strategy will be contingent on test results	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0027U JAK2 (Janus kinase 2) (e.g., myeloproliferative disorder) gene analysis, targeted sequence analysis exons 12-15.	One of the following ICD-10-CM diagnosis codes is required on the claim: D45, D47.1 or D47.3	N/A
0034U	The service requires a TAR.	N/A
TPMT (thiopurine S-methyltransferase), NUDT15 (nudix hydroxylase 15)(e.g., thiopurine metabolism), gene analysis, common variants (ie, TPMT *2, *3A, *3B, *3C, *4, *5, *6, *8, *12; NUDT15 *3, *4, *5).	<ul> <li>A TAR requires documentation of the following criteria:</li> <li>That the patient is undergoing thiopurine therapy, and</li> <li>The patient has severe or prolonged myelosuppression</li> </ul>	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0035U	The service requires a TAR.	Once in a
Neurology (prion disease), cerebrospinal fluid,	A TAR requires documentation of the following criteria:	lifetime, with
detection of prion protein by quaking-induced conformational conversion, qualitative	1. Rapidly progressive dementia, and	TAR/SAR
	At least two out of the following four clinical features:	override
	b. Myoclonus	
	Visual or cerebellar signs	
	Pyramid/extrapyramidal signs	
	Akinetic mutism	
	A positive result on at least one of the following tests:	
	c. A typical EEG (periodic sharp wave complexes) during an illness of any duration	
	High signal in caudate/putamen in magnetic resonance imaging (MRI) brain scan or at least two cortical regions (temporal, parietal occipital) either on diffusion-weighted imaging (DWI) or fluid attenuated inversion recovery (FLAIR)	
	No routine investigations indicating an alternative diagnosis	

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Code and Code Description	TAR and/or Billing Requirements	Frequency
0037U	The service requires a TAR.	N/A
Targeted genomic sequence analysis, solid organ neoplasm, DNA analysis of 324 genes, interrogation for sequence variants, gene copy number amplifications, gene rearrangements, microsatellite instability and tumor mutational burden.	<ul> <li><a criteria:<="" documentation="" following="" li="" of="" requires="" tar="" the=""> <li>The patient has either recurrent, relapsed, refractory, metastatic or advanced stages III or IV cancer, and</li> <li>The patient either has not been previously tested using the same Next Generation Sequencing (NGS) test for the same primary diagnosis of cancer or repeat testing using the same NGS test only when a new primary cancer diagnosis is made by the treating physician, and</li> <li>The decision for additional cancer treatment is contingent on the test results&gt;&gt;</li> </a></li></ul>	

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Code and Code Description	TAR and/or Billing Requirements	Frequency
0038U	N/A	N/A
Vitamin D, 25 hydroxy D2 and D3, by LC-MS/MS, serum microsample, quantitative		
0039U	N/A	N/A
Deoxyribonucleic acid (DNA) antibody, double stranded, high avidity		
0040U	The following ICD-10-CM diagnosis code is required on	N/A
BCR/ABL1 (t(9;22)) (e.g., chronic myelogenous leukemia) translocation analysis, major breakpoint, quantitative	the claim: C92.10. Allow TAR/SAR override.	
0041U	N/A	N/A
Borrelia burgdorferi, antibody detection of 5 recombinant protein groups, by immunoblot, IgM		
0042U	N/A	N/A
Borrelia burgdorferi, antibody detection of 12 recombinant protein groups, by immunoblot, IgG		

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Code and Code Description	TAR and/or Billing Requirements	Frequency
0043U	N/A	N/A
Tick-borne relapsing fever Borrelia group, antibody detection to 4 recombinant protein groups, by immunoblot, IgM		
0044U	N/A	N/A
Tick-borne relapsing fever Borrelia group, antibody detection to 4 recombinant protein groups, by immunoblot, IgG		
0046U	One of the following ICD-10-CM diagnosis codes is	N/A
FLT3 (fms-related tyrosine kinase 3) (e.g., acute myeloid leukemia) internal tandem duplication (ITD) variants, quantitative	required on the claim: C92.00 thru C92.02, C92.60 thru C92.62, C92.A0 thru C92.A2	

# Table of Proprietary Laboratory Analyses (PLA) Codes (continued)

Code and Code Description	TAR and/or Billing Requirements	Frequency
0047U Oncology (prostate), mRNA, gene expression profiling by real-time RT-PCR of 17 genes (12 content and 5 housekeeping), utilizing formalin-fixed paraffin-embedded tissue,	The coverage policy for Gene Expression Profile (GEP) for prostate cancer is based on the 2019 American Society of Clinical Oncologist (ASCO) Guideline titled, "Molecular Biomarkers in Localized Prostate Cancer: ASCO Guideline.	Once in a lifetime. Allow TAR/SAR override
algorithm reported as a risk score	The service requires a TAR.	
	A TAR requires documentation of the following criteria:	
	<ol> <li>For identification of patients with prostate cancer who are most likely to benefit from active surveillance or treatment:</li> </ol>	
	d. Coverage is limited to Oncotype Dx Prostate.	
	<ul> <li>i. Oncotype DX Prostate – Use PLA code 0047U</li> </ul>	
	The patient must have one of the following:	
	<ul> <li>i. Higher volume Grade Group 1</li> <li>ii. Favorable intermediate risk (e.g., Grade Group 2, percentage of positive biopsy cores, 50 percent, and no more than one NCCN intermediate-risk factor)</li> <li>iii. Discordant features in their risk stratification (e.g., palpable mass with Grade Group 1)</li> </ul>	

(Code 0047U continued on next page)

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Code and Code Description	TAR and/or Billing Requirements	Frequency
0047U (continued) Oncology (prostate), mRNA, gene expression profiling by real-time RT-PCR of 17 genes (12 content and 5 housekeeping), utilizing formalin-fixed paraffin-embedded tissue, algorithm reported as a risk score	<ul> <li>iv. Other features associated with progression while on active surveillance (e.g., high PSA density and certain germline or somatic mutations).</li> <li>v. Unfavorable intermediate-risk when considering decisions to proceed with treatment (i.e. add androgen deprivation therapy to radiation).</li> </ul>	Once in a lifetime
	Result of the test, when considered as a whole with routine clinical factors, is likely to influence the decision to proceed with surveillance or treatment.	
	For post-prostatectomy patients who seek guidance on adjuvant vs. salvage radiation:	
	<ul> <li>Coverage is limited to Decipher Genomic Classifier</li> </ul>	
	<ul> <li>Result of the test, when considered as a whole without routine clinical factors, is likely to affect treatment.</li> </ul>	
0049U NPM1 (nucleophosmin) (e.g., acute myeloid leukemia) gene analysis, quantitative	One of the following ICD-10-CM diagnosis codes is required on the claim: C92.00 thru C92.02, C92.60 thru C92.62, C92.A0 thru C92.A2	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0050U Targeted genomic sequence analysis panel, acute myelogenous leukemia, DNA analysis, 194 genes, interrogation for sequence variants, copy number variants or rearrangements	One of the following ICD-10-CM diagnosis codes is required on the claim: C92.00 thru C92.02, C92.60 thru C92.62, C92.A0 thru C92.A2	N/A
0051U	N/A	N/A
Prescription drug monitoring, evaluation of drugs present by LC-MS/MS, urine, 31 drug panel, reported as quantitative results, detected or not detected, per date of service		
0052U	N/A	N/A
Lipoprotein, blood, high resolution fractionation and quantitation of lipoproteins, including all five major lipoprotein classes and subclasses of HDL, LDL, and VLDL by vertical auto profile ultracentrifugation		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0054U	N/A	N/A
Prescription drug monitoring, 14 or more classes of drugs and substances, definitive tandem mass spectrometry with chromatography, capillary blood, quantitative report with therapeutic and toxic ranges, including steady-state range for the prescribed dose when detected, per date of service		
0056U Hematology (acute myelogenous leukemia), DNA, whole genome next-generation sequencing to detect gene rearrangement(s), blood or bone marrow, report of specific gene rearrangement(s).	One of the following ICD-10-CM diagnosis codes is required on the claim: C92.00 thru C92.02, C92.60 thru C92.62, C92A0 thru C92.A2	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0058U	One of the following ICD-10-CM diagnosis codes is	Once in a
Oncology (Merkel cell carcinoma), detection of	required on the claim: C4A.0, C4A.10 thru C4A.12,	lifetime,
antibodies to the Merkel cell polyoma virus	C4A.20 thru C4A.22, C4A.30 thru C4A.39, C4A.51 thru	with
oncoprotein (small T antigen), serum,	C4A.59, C4A.60 thru C4A.62, C4A.70 thru C4A.72,	TAR/SAR
quantitative	C4A.8, C4A.9	override
0059U	One of the following ICD-10-CM diagnosis codes is	Once in a
Oncology (Merkel cell carcinoma), detection of	required on the claim: C4A.0, C4A.10 thru C4A.12,	lifetime,
antibodies to the Merkel cell polyoma virus	C4A.20 thru C4A.22, C4A.30 thru C4A.39, C4A.51 thru	with
capsid protein (VP1), serum, reported as	C4A.59, C4A.60 thru C4A.62, C4A.70 thru C4A.72,	TAR/SAR
positive or negative	C4A.8, C4A.9	override

Code and Code Description	TAR and/or Billing Requirements	Frequency
0064U	N/A	N/A
Antibody, Treponema pallidum, total and rapid plasma reagin (RPR), immunoassay, qualitative		
0065U	N/A	N/A
Syphilis test, non-treponemal antibody, immunoassay, qualitative (RPR)		
0068U	N/A	N/A
Candida species panel (C. albicans, C. glabrata, C. parapsilosis, C. kruseii, C. tropicalis, and C. auris), amplified probe technique with qualitative report of the presence or absence of each species		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0077U	N/A	N/A
Immunoglobulin paraprotein (M-protein), qualitative, immunoprecipitation and mass spectrometry, blood or urine, including isotype		
0081U Oncology (uveal melanoma), mRNA, gene-expression profiling by real-time RT-PCR of 15 genes (12 content and 3 housekeeping genes), utilizing fine needle aspirate or formalin- fixed paraffin-embedded tissue, algorithm reported as risk of metastasis	One of the following ICD-10-CM diagnosis codes is required on the claim: C69.30 thru C69.32, C69.40 thru C69.42	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0082U	N/A	N/A
Drug test(s), definitive, 90 or more drugs or substances, definitive chromatography with mass spectrometry, and presumptive, any number of drug classes, by instrument chemistry analyzer (utilizing immunoassay), urine, report of presence or absence of each drug, drug metabolite or substance with description and severity of significant interactions per date of service		
0084U	N/A	N/A
Red blood cell antigen typing, DNA, genotyping of 10 blood groups with phenotype prediction of 37 red blood cell antigens		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0087U Cardiology (heart transplant), mRNA gene expression profiling by microarray of 1283 genes, transplant biopsy tissue, allograft rejection and injury algorithm reported as a probability score	The following ICD-10-CM diagnosis code is required on the claim: Z94.1	N/A
0088U Transplantation medicine (kidney allograft rejection), microarray gene expression profiling of 1494 genes, utilizing transplant biopsy tissue, algorithm reported as a probability score for rejection	The following ICD-10-CM diagnosis code is required on the claim: Z94.0	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0107U	N/A	N/A
Clostridium difficile toxin(s) antigen detection by immunoassay technique, stool, qualitative, multiple-step method.		
0109U	N/A	N/A
Infectious disease (Aspergillus species), real-time PCR for detection of DNA from 4 species (A. fumigatus, A. terreus, A. niger, and A. flavus), blood, lavage fluid, or tissue, qualitative reporting of presence or absence of each species		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0120U Oncology (B-cell lymphoma classification), mRNA, gene expression profiling by fluorescent probe hybridization of 58 genes (45 content and 13 housekeeping genes), formalin-fixed paraffin-embedded tissue, algorithm reported as likelihood for primary mediastinal B-cell lymphoma (PMBCL) and diffuse large B-cell lymphoma (DLBCL) with cell of origin subtyping in the latter	One of the following ICD-10-CM diagnosis codes is required on the claim: C83.30 thru C83.39, C85.20 thru C85.29	N/A
0140U Infectious disease (fungi), fungal pathogen identification, DNA (15 fungal targets), blood culture, amplified probe technique, each target reported as detected or not detected	N/A	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0141U	N/A	N/A
Infectious disease (bacteria and fungi), gram- positive organism identification and drug resistance element detection, DNA (20 gram- positive bacterial targets, 4 resistance genes, 1 pan gram-negative bacterial target, 1 pan Candida target), blood culture, amplified probe technique, each target reported as detected or not detected		
0142U	N/A	N/A
Infectious disease (bacteria and fungi), gram- negative bacterial identification and drug resistance element detection, DNA (21 gram- negative bacterial targets, 6 resistance genes, 1 pan gram-positive bacterial target, 1 pan Candida target), amplified probe technique, each target reported as detected or not detected		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0154U Oncology (urothelial cancer), RNA, analysis by real-time RT-PCR of the FGFR3 (fibroblast growth factor receptor 3) gene analysis (ie, p.R248C [c.742C>T], p.S249C [c.746C>G], p.G370C [c.1108G>T], p.Y373C [c.1118A>G], FGFR3-TACC3v1, and FGFR3-TACC3v3)	One of the following ICD-10-CM diagnosis codes is required on the claim: C67.0 thru C67.9	Once in a lifetime, with TAR/SAR override
0155U Oncology (breast cancer), DNA, PIK3CA (phosphatidylinositol-4,5-bisphosphate 3-kinase, catalytic subunit alpha) (e.g., breast cancer) gene analysis (i.e., p.C420R, p.E542K, p.E545A, p.E545D [g.1635G>T only], p.E545G, p.E545K, p.Q546E, p.Q546R, p.H1047L, p.H1047R, p.H1047Y)	One of the following ICD-10-CM diagnosis codes is required on the claim: C50.011 thru C50.929	Once in a lifetime, with TAR/SAR override

Code and Code Description	TAR and/or Billing Requirements	Frequency
0157U APC (APC regulator of WNT signaling pathway) (e.g., familial adenomatosis polyposis [FAP]) mRNA sequence analysis (List separately in addition to code for primary procedure)	One of the following ICD-10-CM diagnosis codes is required on the claim: C18.0 thru C18.9, D12.0 thru D12.9, K63.5, Z86.010	Once in a lifetime, with TAR/SAR override
0158U	One of the following ICD-10-CM diagnosis codes is	Once in a
MLH1 (mutL homolog 1) (e.g., hereditary non-	required on the claim: C17.0 thru C20, C24.0 thru	lifetime,
polyposis colorectal cancer, Lynch syndrome)	C25.9, C54.0 thru C54.9, C65.1 thru C66.9, C71.0 thru	with
mRNA sequence analysis (List separately in	C71.9, D23.0 thru D23.9, Z80.0, Z80.49, Z85.030,	TAR/SAR
addition to code for primary procedure)	Z85.038, Z85.040, Z85.048, Z85.42	override
0159U	One of the following ICD-10-CM diagnosis codes is	Once in a
MSH2 (mutS homolog 2) (e.g., hereditary colon	required on the claim: C17.0 thru C20, C24.0 thru	lifetime,
cancer, Lynch syndrome) mRNA sequence	C25.9, C54.0 thru C54.9, C65.1 thru C66.9, C71.0 thru	with
analysis (List separately in addition to code for	C71.9, D23.0 thru D23.9, Z80.0, Z80.49, Z85.030,	TAR/SAR
primary procedure)	Z85.038, Z85.040, Z85.048, Z85.42	override

Code and Code Description	TAR and/or Billing Requirements	Frequency
0160U MSH6 (mutS homolog 6) (e.g., hereditary colon cancer, Lynch syndrome) mRNA sequence analysis (List separately in addition to code for primary procedure)	One of the following ICD-10-CM diagnosis codes is required on the claim: C17.0 thru C20, C24.0 thru C25.9, C54.0 thru C54.9, C65.1 thru C66.9, C71.0 thru C71.9, D23.0 thru D23.9, Z80.0, Z80.49, Z85.030, Z85.038, Z85.040, Z85.048, Z85.42	Once in a lifetime, with TAR/SAR override
0161U PMS2 (PMS1 homolog 2, mismatch repair system component) (e.g., hereditary non-polyposis colorectal cancer, Lynch syndrome) mRNA sequence analysis (List separately in addition to code for primary procedure)	One of the following ICD-10-CM diagnosis codes is required on the claim: C17.0 thru C20, C24.0 thru C25.9, C54.0 thru C54.9, C65.1 thru C66.9, C71.0 thru C71.9, D23.0 thru D23.9, Z80.0, Z80.49, Z85.030, Z85.038, Z85.040, Z85.048, Z85.42	Once in a lifetime, with TAR/SAR override

Code and Code Description	TAR and/or Billing Requirements	Frequency
0162U Hereditary colon cancer (Lynch syndrome), targeted mRNA sequence analysis panel (MLH1, MSH2, MSH6, PMS2) (List separately in addition to code for primary procedure)	One of the following ICD-10-CM diagnosis codes is required on the claim: C17.0 thru C20, C24.0 thru C25.9, C54.0 thru C54.9, C65.1 thru C66.9, C71.0 thru C71.9, D23.0 thru D23.9, Z80.0, Z80.49, Z85.030, Z85.038, Z85.040, Z85.048, Z85.42	Once in a lifetime, with TAR/SAR override
0165U Peanut allergen-specific quantitative assessment of epitopes using enzyme-linked immunosorbent assay (ELISA), blood, individual epitope results and probability of peanut allergy.	One of the following ICD-10-CM diagnosis codes is required on the claim: Z01.82, Z91.010	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0166U	N/A	N/A
Liver disease, 10 biochemical assays (α2-macroglobulin, haptoglobin, apolipoprotein A1, bilirubin, GGT, ALT, AST, triglycerides, cholesterol, fasting glucose) and biometric and demographic data, utilizing serum, algorithm reported as scores for fibrosis, necroinflammatory activity, and steatosis with a summary interpretation		
0167U	N/A	N/A
Gonadotropin, chorionic (hCG), immunoassay with direct optical observation, blood		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0169U	The service requires a TAR.	N/A
NUDT15 (nudix hydrolase 15) and TPMT	A TAR requires documentation of the following criteria:	
(thiopurine S-methyltransferase) (e.g., drug metabolism) gene analysis, common variants	<ul> <li>That the patient is undergoing thiopurine therapy, and</li> </ul>	
	<ul> <li>The patient has severe or prolonged myelosuppression.</li> </ul>	
0171U	One of the following ICD-10-CM diagnosis codes is	N/A
Targeted genomic sequence analysis panel, acute myeloid leukemia, myelodysplastic syndrome, and myeloproliferative neoplasms, DNA analysis, 23 genes, interrogation for sequence variants, rearrangements and minimal residual disease, reported as presence/absence	required on the claim: C92.00, C92.01, C92.02, C92.10 thru C92.22, C95.10, D45, D46.0, D46.1, D46.20 thru D46.22, D46.4, D46.9, D46.A, D46.B, D46.C, D46.Z, D47.1, D47.3.	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0172U Oncology (solid tumor as indicated by the label), somatic mutation analysis of BRCA1 (BRCA1, DNA repair associated), BRCA2 (BRCA2, DNA repair associated) and analysis of homologous recombination deficiency pathways, DNA, formalin-fixed paraffin-embedded tissue, algorithm quantifying tumor genomic instability score	<ul> <li>The service requires a TAR.</li> <li>A TAR requires documentation of the following criteria:</li> <li>1. The patient has advanced ovarian, fallopian tube or primary peritoneal cancer</li> <li>2. Treatment is contingent on the result of the test</li> </ul>	Once in a lifetime, with a TAR/SAR override
0177U Oncology (breast cancer), DNA, PIK3CA (phosphatidylinositol-4,5-bisphosphate 3-kinase catalytic subunit alpha) gene analysis of 11 gene variants utilizing plasma, reported as PIK3CA gene mutation status	<ul> <li>The service requires a TAR.</li> <li>A TAR requires documentation of the following criteria:</li> <li>1. The patient has confirmed diagnosis of breast cancer</li> <li>2. Treatment is contingent the result of the test</li> </ul>	Once in a lifetime, with a TAR/SAR override

Code and Code Description	TAR and/or Billing Requirements	Frequency
0178U Peanut allergen-specific quantitative assessment of multiple epitopes using enzyme-linked immunosorbent assay (ELISA), blood, report of minimum eliciting exposure for a clinical reaction.	N/A	50/day, with a TAR/SAR override
0180U	N/A	N/A
Red cell antigen (ABO blood group) genotyping (ABO), gene analysis Sanger/chain termination/conventional sequencing, ABO (ABO, alpha 1-3-N- acetylgalactosaminyltransferase and alpha 1-3-galactosyltransferase) gene, including subtyping, 7 exons		
0181U	N/A	N/A
Red cell antigen (Colton blood group) genotyping (CO), gene analysis, AQP1 (aquaporin 1 [Colton blood group]) exon 1		
0182U	N/A	N/A
Red cell antigen (Cromer blood group) genotyping (CROM), gene analysis, CD55 (CD55 molecule [Cromer blood group]) exons 1-10		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0183U	N/A	N/A
Red cell antigen (Diego blood group) genotyping (DI), gene analysis, SLC4A1 (solute carrier family 4 member 1 [Diego blood group]) exon 19		
0184U	N/A	N/A
Red cell antigen (Dombrock blood group) genotyping (DO), gene analysis, ART4 (ADP-ribosyltransferase 4 [Dombrock blood group]) exon 2		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0185U	N/A	N/A
Red cell antigen (H blood group) genotyping (FUT1), gene analysis, FUT1 (fucosyltransferase 1 [H blood group]) exon 4		
0186U	N/A	N/A
Red cell antigen (H blood group) genotyping (FUT2), gene analysis, FUT2 (fucosyltransferase 2) exon 2		
0187U	N/A	N/A
Red cell antigen (Duffy blood group) genotyping (FY), gene analysis, ACKR1 (atypical chemokine receptor 1 [Duffy blood group]) exons 1-2		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0188U	N/A	N/A
Red cell antigen (Gerbich blood group) genotyping (GE), gene analysis, GYPC (glycophorin C [Gerbich blood group]) exons 1-4		
0189U	N/A	N/A
Red cell antigen (MNS blood group) genotyping (GYPA), gene analysis, GYPA (glycophorin A [MNS blood group]) introns 1, 5, exon 2		
0190U	N/A	N/A
Red cell antigen (MNS blood group) genotyping (GYPB), gene analysis, GYPB (glycophorin B [MNS blood group]) introns 1, 5, pseudoexon 3		
0191U	N/A	N/A
Red cell antigen (Indian blood group) genotyping (IN), gene analysis, CD44 (CD44 molecule [Indian blood group]) exons 2, 3, 6		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0192U	N/A	N/A
Red cell antigen (Kidd blood group) genotyping (JK), gene analysis, SLC14A1 (solute carrier family 14 member 1 [Kidd blood group]) gene promoter, exon 9		
0193U	N/A	N/A
Red cell antigen (JR blood group) genotyping (JR), gene analysis, ABCG2 (ATP binding cassette subfamily G member 2 [Junior blood group]) exons 2-26		
0194U	N/A	N/A
Red cell antigen (Kell blood group) genotyping (KEL), gene analysis, KEL (Kell metallo-endopeptidase [Kell blood group]) exon 8		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0195U	N/A	N/A
KLF1 (Kruppel-like factor 1), targeted sequencing (ie, exon 13)		
0196U	N/A	N/A
Red cell antigen (Lutheran blood group) genotyping (LU), gene analysis, BCAM (basal cell adhesion molecule [Lutheran blood group]) exon 3		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0197U	N/A	N/A
Red cell antigen (Landsteiner-Wiener blood group) genotyping (LW), gene analysis, ICAM4 (intercellular adhesion molecule 4 [Landsteiner-Wiener blood group]) exon 1		
0198U	N/A	N/A
Red cell antigen (RH blood group) genotyping (RHD and RHCE), gene analysis Sanger/chain termination/conventional sequencing, RHD (Rh blood group D antigen) exons 1-10 and RHCE (Rh blood group CcEe antigens) exon 5		
0199U	N/A	N/A
Red cell antigen (Scianna blood group) genotyping (SC), gene analysis, ERMAP (erythroblast membrane associated protein [Scianna blood group]) exons 4, 12		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0200U	N/A	N/A
Red cell antigen (Kx blood group) genotyping (XK), gene analysis, XK (X-linked Kx blood group) exons 1-3		
0202U	N/A	N/A
Infectious disease (bacterial or viral respiratory tract infection), pathogen-specific nucleic acid (DNA or RNA), 22 targets including severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), qualitative RT-PCR, nasopharyngeal swab, each pathogen reported as detected or not detected		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0204U	The service requires a TAR.	Once in a
Oncology (thyroid), mRNA, gene expression	A TAR requires documentation of the following criteria:	lifetime
analysis of 593 genes (including BRAF, RAS, RET, PAX8, and NTRK) for sequence variants and rearrangements, utilizing fine needle aspirate, reported as detected or not detected	<ol> <li>The patient is under evaluation for thyroid nodule(s)</li> </ol>	
	<ol> <li>The cytopathology result from fine needle aspiration is indeterminate, defined as one of the following:</li> </ol>	
	a. Follicular lesion of undetermined significance (FLUS), Bethesda III, or	
	<ul> <li>Atypia of undetermined significance (AUS), Bethesda III, or</li> </ul>	
	c. Follicular neoplasm, Bethesda IV.	
	<ol> <li>The diagnostic or treatment strategy will be contingent on test results</li> </ol>	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0210U	N/A	N/A
Syphilis test, non-treponemal antibody, immunoassay, quantitative (RPR)		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0216U Neurology (inherited ataxias), genomic DNA sequence analysis of 12 common genes including small sequence changes, deletions, duplications, short tandem repeat gene expansions, and variants in non-uniquely mappable regions, blood or saliva, identification and categorization of genetic variants	One of the following ICD-10-CM diagnosis codes is required on the claim: G11.0, G11.1, G11.3, G11.9, R26.0, R27.0. Allow TAR/SAR override.	N/A
0217U Neurology (inherited ataxias), genomic DNA sequence analysis of 51 genes including small sequence changes, deletions, duplications, short tandem repeat gene expansions, and variants in non-uniquely mappable regions, blood or saliva, identification and categorization of genetic variants	One of the following ICD-10-CM diagnosis codes is required on the claim: G11.0, G11.1, G11.3, G11.9, R26.0, R27.0. Allow TAR/SAR override.	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0218U	The service requires a TAR.	N/A
Neurology (muscular dystrophy), DMD gene sequence analysis, including small sequence changes, deletions, duplications, and variants in non-uniquely mappable regions, blood or saliva, identification and characterization of genetic variants	<ul> <li>A TAR requires documentation of the following criteria:</li> <li>Patient has a clinical diagnosis of dystrophinopathy based on the history, physical examination and elevated creatine kinase (CK) level</li> <li>Result of the DMD (dystrophin) deletion or duplication is negative</li> </ul>	
0219U Infectious agent (human immunodeficiency virus), targeted viral next-generation sequence analysis (ie, protease [PR], reverse transcriptase [RT], integrase [INT]), algorithm reported as prediction of antiviral drug susceptibility	One of the following ICD-10-CM diagnosis codes is required on the claim: B20, Z21. Allow TAR/SAR override.	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0221U	N/A	N/A
Red cell antigen (ABO blood group) genotyping (ABO), gene analysis, next-generation sequencing, ABO (ABO, alpha 1-3-N-acetylgalactosaminyltransferase and alpha 1-3-galactosyltransferase) gene		
0222U Red cell antigen (RH blood group) genotyping (RHD and RHCE), gene analysis, next-generation sequencing, RH proximal promoter, exons 1-10, portions of introns 2-3	N/A	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0223U	N/A	N/A
Infectious disease (bacterial or viral respiratory tract infection), pathogen-specific nucleic acid (DNA or RNA), 22 targets including severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), qualitative RT-PCR, nasopharyngeal swab, each pathogen reported as detected or not detected		
0224U Antibody, severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (Coronavirus disease [COVID-19]), includes titer(s), when performed	Do Not Report with CPT code 86769.	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0225U	N/A	N/A
Infectious disease (bacterial or viral respiratory tract infection) pathogen-specific DNA and RNA, 21 targets, including severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), amplified probe technique, including multiplex reverse transcription for RNA targets, each analyte reported as detected or not detected		
0226U Surrogate viral neutralization test (sVNT), severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (Coronavirus disease [COVID-19]), ELISA, plasma, serum	N/A	N/A

Code and Code Description	TAR and/or Billing Requirements	Frequency
0227U	N/A	N/A
Drug assay, presumptive, 30 or more drugs or metabolites, urine, liquid chromatography with tandem mass spectrometry (LC-MS/MS) using multiple reaction monitoring (MRM), with drug or metabolite description, includes sample validation		
0230U	The service requires a TAR.	Once in a
AR (androgen receptor) (e.g., spinal and bulbar	A TAR requires documentation of the following criteria:	lifetime
muscular atrophy, Kennedy disease, X chromosome inactivation), full sequence analysis, including small sequence changes in	<ul> <li>The patient has clinical signs or symptoms suspicious for bulbar muscular atrophy, and</li> </ul>	
exonic and intronic regions, deletions, duplications, short tandem repeat (STR) expansions, mobile element insertions, and variants in non-uniquely mappable regions	<ul> <li>The patient requires the service as a confirmatory test for spinal and bulbar muscular atrophy</li> </ul>	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0231U	The service requires a TAR.	N/A
CACNA1A (calcium voltage-gated channel subunit alpha 1A) (e.g., spinocerebellar ataxia), full gene analysis, 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	<ul> <li>A TAR requires documentation of the following criteria:</li> <li>The patient has clinical signs or symptoms suspicious for Episodic ataxia type 2 (EA2), and</li> <li>The patient requires the service as a confirmatory test for EA2</li> </ul>	
0232U	The service requires a TAR.	Once in a
CSTB (cystatin B) (e.g., progressive myoclonic epilepsy type 1A, Unverricht-Lundborg disease), full gene analysis, 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	<ul> <li>A TAR requires documentation of the following criteria:</li> <li>The patient has clinical signs or symptoms suspicious for myoclonic epilepsy type 1 and requires the service as a confirmatory test for myoclonic epilepsy type 1, and</li> <li>Treatment will be contingent on test results</li> </ul>	lifetime

Code and Code Description	TAR and/or Billing Requirements	Frequency
0233U	The service requires a TAR.	Once in a
FXN (frataxin) (e.g., Friedreich ataxia), gene analysis, 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	<ul> <li>A TAR requires documentation of the following criteria:</li> <li>The patient has clinical signs or symptoms suspicious for Friedreich ataxia (FRDA), and</li> <li>The patient requires the service as a confirmatory test for FRDA</li> </ul>	lifetime
0234U MECP2 (methyl CpG binding protein 2) (e.g., Rett syndrome), full gene analysis, including small sequence changes in exonic and intronic regions, deletions, duplications, mobile element insertions, and variants in non-uniquely mappable regions	<ul> <li>The service requires a TAR.</li> <li>A TAR requires documentation of the following criteria:</li> <li>The patient has clinical signs or symptoms suspicious for Rett syndrome, and</li> <li>The patient requires the service as a confirmatory test for Rett syndrome</li> </ul>	Once in a lifetime

Code and Code Description	TAR and/or Billing Requirements	Frequency
0235U	The service requires a TAR.	N/A
PTEN (phosphatase and tensin homolog) (e.g.,	A TAR requires documentation of the following criteria:	
Cowden syndrome, PTEN hamartoma tumor syndrome), full gene analysis, including small	1. Individual with a personal history of:	
sequence changes in exonic and intronic		
regions, deletions, duplications, mobile element insertions, and variants in non-uniquely	<ul> <li>Adult Lhermitte-Duclos disease, or</li> </ul>	
mappable regions	<ul> <li>Autism spectrum disorder AND macrocephaly, or</li> </ul>	
	<ul> <li>Two or more biopsy-proven trichilemmomas, or</li> </ul>	
	<ul> <li>Two or more major criteria (one macrocephaly), or</li> </ul>	
	<ul> <li>Three major criteria without macrocephaly, or</li> </ul>	
	<ul> <li>One major and three or more minor criteria, or</li> </ul>	
	<ul> <li>Four or more minor criteria (please see list below)</li> </ul>	
	2. At-risk individual	
	<ul> <li>With a relative who has a clinical diagnosis of Cowden syndrome or Bannayan-Riley- Ruvalcaba syndrome for whom testing has not been performed AND who has any one major criterion or two minor criteria</li> </ul>	

#### Table of Proprietary Laboratory Analyses (PLA) Codes (continued)

Code and Code Description	TAR and/or Billing Requirements	Frequency
0235U (continued)	Clinical Criteria:	N/A
PTEN (phosphatase and tensin homolog) (e.g.,	Major Criteria	
Cowden syndrome, PTEN hamartoma tumor syndrome), full gene analysis, including small	Breast Cancer	
sequence changes in exonic and intronic	<ul> <li>Mucocutaneous lesions</li> </ul>	
regions, deletions, duplications, mobile element insertions, and variants in non-uniquely	<ul> <li>One biopsy-proven trichilemmoma</li> </ul>	
mappable regions	<ul> <li>Multiple palmoplantar keratosis</li> </ul>	
	Multifocal or extensive oral mucosal papillomatosis	
	Multiple cutaneous facial papules (often verrucous)	
	<ul> <li>Macular pigmentation of glans penis</li> </ul>	
	<ul> <li>Macroencephaly (megalocephaly, ie, ≥97th percentile)</li> </ul>	
	Endometrial cancer	
	<ul> <li>Non-medullary thyroid cancer</li> </ul>	
	<ul> <li>Multiple GI tract hamartomas or ganglioneuromas</li> </ul>	

(Code 0235U continued on next page)

Code and Code Description	TAR and/or Billing Requirements	Frequency
0235U (continued)	Minor Criteria	N/A
PTEN (phosphatase and tensin homolog) (e.g.,	<ul> <li>Other thyroid lesions (adenoma, nodule, goiter)</li> </ul>	
Cowden syndrome, PTEN hamartoma tumor syndrome), full gene analysis, including small	<ul> <li>Mental retardation (IQ ≤75)</li> </ul>	
sequence changes in exonic and intronic	Autism spectrum disorder	
regions, deletions, duplications, mobile element insertions, and variants in non-uniquely	<ul> <li>Single GI tract hamartoma or ganglioneuroma</li> </ul>	
mappable regions	<ul> <li>Fibrocystic disease of the breast</li> </ul>	
	• Lipomas	
	• Fibromas	
	Renal cell carcinoma	
	Uterine fibroids	
0236U	The service requires a TAR.	Once in a
SMN1 (survival of motor neuron 1, telomeric) and SMN2 (survival of motor neuron 2, centromeric) (e.g., spinal muscular atrophy) full gene analysis, including small sequence changes in exonic and intronic regions, duplications and deletions, and mobile element insertions	One of the following ICD-10-CM diagnosis codes is required on the claim: 009.00 thru 009.93, Z31.430, Z31.440, Z34.00 thru Z34.03, Z34.80 thru Z34.83, Z34.90 thru Z34.93. Allow TAR/SAR override	lifetime

Code and Code Description	TAR and/or Billing Requirements	Frequency
0237U	The service requires a TAR.	N/A
Cardiac ion channelopathies (e.g., Brugada syndrome, long QT syndrome, short QT syndrome, catecholaminergic polymorphic ventricular tachycardia), genomic sequence analysis panel including ANK2, CASQ2, CAV3,	The TAR must document a copy of the report of the physician interpreted 12-lead electrocardiogram (ECG) with pattern consistent with or suspicious for prolonged QT interval. The TAR must also have clinical documentation of one or more of the following:	
KCNE1, KCNE2, KCNH2, KCNJ2, KCNQ1, RYR2, and SCN5A, including small sequence changes in exonic and intronic regions,	<ol> <li>Torsade de pointes in the absence of drugs known to prolong QT interval</li> </ol>	
deletions, duplications, mobile element	2. T-wave alternans	
insertions, and variants in non-uniquely mappable regions	3. Notched T-wave in three leads	
	4. Syncope	
	5. Family members with long QT syndrome	
	<ol> <li>Sudden death in family members less than 30 years of age without defined cause</li> </ol>	
0238U	One of the following ICD-10-CM diagnosis codes is	Once in a
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	required on the claim: C17.0 thru C20, C24.0 thru C25.9, C54.0 thru C54.9, C65.1 thru C66.9, C71.0 thru C71.9, D23.0 thru D23.9, Z80.0, Z80.49, Z85.030, Z85.038, Z85.040, Z85.048, Z85.42	lifetime

Code and Code Description	TAR and/or Billing Requirements	Frequency
0239U	The service requires a TAR.	N/A
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	<ul> <li>A TAR requires documentation of the following criteria:</li> <li>1. The patient has a diagnosis of either: <ul> <li>Non-small cell lung cancer (plasma), or</li> <li>Metastatic castrate resistant prostate cancer</li> </ul> </li> <li>2. Treatment is contingent on the test result.</li> </ul>	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0240U	N/A	N/A
Infectious disease (viral respiratory tract infection), pathogen-specific RNA, 3 targets (severe acute respiratory syndrome coronavirus 2 [SARS-CoV-2], influenza A, influenza B), upper respiratory specimen, each pathogen reported as detected or not detected		
0241U	N/A	N/A
Infectious disease (viral respiratory tract infection), pathogen-specific RNA, 4 targets (severe acute respiratory syndrome coronavirus 2 [SARS-CoV-2], influenza A, influenza B, respiratory syncytial virus [RSV]), upper respiratory specimen, each pathogen reported as detected or not detected		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0242U	The service requires a TAR	Once in a
Targeted genomic sequence analysis panel,	A TAR requires documentation of the following criteria:	lifetime
solid organ neoplasm, cell-free circulating DNA analysis of 55-74 genes, interrogation for	<ul> <li>Patient has Non-small cell lung cancer</li> </ul>	
sequence variants, gene copy number amplifications, and gene rearrangements	<ul> <li>Treatment is contingent on test result</li> </ul>	
0244U	The service requires a TAR	N/A
Oncology (solid organ), DNA, comprehensive	A TAR requires documentation of the following criteria:	
genomic profiling, 257 genes, interrogation for single-nucleotide variants, insertions/deletions,	For Somatic Testing	
copy number alterations, gene rearrangements, tumor-mutational burden and microsatellite instability, utilizing formalin-fixed paraffin- embedded tumor tissue	<ul> <li>The patient has either recurrent, relapsed, refractory, metastatic or advanced stages III or IV cancer, and</li> </ul>	
	• The patient either has not been previously tested using the same Next Generation Sequencing (NGS) test for the same primary diagnosis of cancer or repeat testing using the same NGS test only when a new primary cancer diagnosis is made by the treating physician, and	
	<ul> <li>The decision for additional cancer treatment is contingent on the test results.</li> </ul>	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0244U (continued)	For Germline Testing	N/A
	<ul> <li>Ovarian or breast cancer; and</li> </ul>	
	<ul> <li>Clinical indication for germline (inherited) testing for hereditary breast or ovarian cancer (ie, American College of Obstetrician and Gynecologists' criteria for further genetic evaluation for hereditary (germline) breast and ovarian cancer) and</li> </ul>	
	<ul> <li>A risk factor for germline (inherited) breast or ovarian cancer; and (BRCAPRO, Myriad, Claus, Boadicea, or Tyrer Cuzick)</li> </ul>	
	<ul> <li>Has not been previously tested with the same germline test using NGS for the same germline genetic content.</li> </ul>	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0245U Oncology (thyroid), mutation analysis of 10 genes and 37 RNA fusions and expression of 4 mRNA markers using next-generation sequencing, fine needle aspirate, report includes associated risk of malignancy expressed as a percentage	<ul> <li>The service requires a TAR</li> <li>A TAR requires documentation of the following criteria: <ol> <li>The patient is under evaluation for thyroid nodule(s)</li> <li>The cytopathology result from fine needle aspiration is indeterminate, defined as one of the following: <ol> <li>Follicular lesion of undetermined significance (FLUS), Bethesda III, or</li> <li>Atypia of undetermined significance (AUS), Bethesda III, or</li> <li>Follicular neoplasm, Bethesda IV.</li> </ol> </li> <li>The diagnostic or treatment strategy will be contingent on test results</li> </ol></li></ul>	Once in a lifetime
0246U	N/A	N/A
Red blood cell antigen typing, DNA, genotyping of at least 16 blood groups with phenotype prediction of at least 51 red blood cell antigens		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0268U Hematology (atypical hemolytic uremic syndrome [aHUS]), genomic sequence analysis of 15 genes, blood, buccal swab, or amniotic	<ul> <li>The service requires a TAR</li> <li>A TAR requires documentation of the following criteria:</li> <li>1. The patient has clinical signs of symptoms for atypical hemolytic uremic syndrome (aHUS), and</li> </ul>	Once in a lifetime
fluid	<ol> <li>The patient requires the service as a diagnostic test for aHUS</li> </ol>	
0269U	The service requires a TAR	Once in a
Hematology (autosomal dominant congenital thrombocytopenia), genomic sequence analysis of 14 genes, blood, buccal swab, or amniotic fluid	<ul> <li>A TAR requires documentation of the following criteria:</li> <li>1. The patient has clinical signs of symptoms suspicious for autosomal dominant congenita thrombocytopenia, and</li> <li>2. The patient requires the service as a diagnostic test for autosomal dominant congenital thrombocytopenia</li> </ul>	lifetime
0271U Hematology (congenital neutropenia), genomic sequence analysis of 23 genes, blood, buccal swab, or amniotic fluid	One of the following ICD-10-CM diagnosis codes is required on the claim: D70.0, D70.1, D70.2, D70.3, D70.4, D70.8, and D70.9. TAR over-ride allowed for ICD-10 codes	Once in a lifetime

Code and Code Description	TAR and/or Billing Requirements	Frequency
0275U	N/A	N/A
Hematology (heparin-induced thrombocytopenia), platelet antibody reactivity by flow cytometry, serum		
0276U	The service requires a TAR	Once in a
Hematology (inherited thrombocytopenia), genomic sequence analysis of 23 genes, blood, buccal swab, or amniotic fluid	<ul> <li>A TAR requires documentation of the following criteria:</li> <li>1. The patient has clinical signs or symptoms suspicious for inherited thrombocytopenia, and</li> <li>2. The patient requires the service as a diagnostic test for inherited thrombocytopenia</li> </ul>	lifetime
0279U	N/A	N/A
Hematology (von willebrand disease [vwd]), von willebrand factor (vwf) and collagen iii binding by enzyme-linked immunosorbent assays (elisa), plasma, report of collagen iii binding		
0280U	N/A	N/A>>
Hematology (von willebrand disease [vwd]), von willebrand factor (vwf) and collagen iv binding by enzyme-linked immunosorbent assays (elisa), plasma, report of collagen iv binding		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0281U	N/A	N/A
Hematology (von willebrand disease [vwd]), von willebrand propeptide, enzyme-linked immunosorbent assays (elisa), plasma, diagnostic report of von willebrand factor (vwf) propeptide antigen level		
0282U	N/A	N/A
Red blood cell antigen typing, dna, genotyping of 12 blood group system genes to predict 44 red blood cell antigen phenotypes		
0283U	N/A	N/A
Von willebrand factor (vwf), type 2b, platelet-binding evaluation, radioimmunoassay, plasma		
0284U	N/A	N/A
Von willebrand factor (vwf), type 2n, factor viii and vwf binding evaluation, enzyme-linked immunosorbent assays (elisa), plasma		

Code and Code Description	TAR and/or Billing Requirements	Frequency
0286U	The service requires a TAR	N/A
CEP72 (centrosomal protein, 72-KDa), NUDT15 (nudix hydrolase 15) and TPMT (thiopurine S-methyltransferase) (eg, drug metabolism) gene analysis, common variants	A TAR requires documentation of the following criteria:	
	<ul> <li>That the patient is undergoing thiopurine therapy, and</li> </ul>	
	<ul> <li>The patient has severe or prolonged myelosuppression</li> </ul>	
0287U	The service requires a TAR.	N/A
Oncology (thyroid), DNA and mRNA, next-generation sequencing analysis of 112 genes, fine needle aspirate or formalin- fixed paraffin-embedded (FFPE) tissue, algorithmic prediction of cancer recurrence, reported as a categorical risk result (low, intermediate, high)	A TAR requires documentation of the following criteria:	
	<ol> <li>The patient is under evaluation for thyroid nodule(s), and</li> </ol>	
	2. The cytopathology result from fine needle aspiration is indeterminate, defined as one of the following:	
	a. Follicular lesion of undetermined significance (FLUS), Bethesda III, or	
	<ul> <li>Atypia of undetermined significance (AUS), Bethesda III, or</li> </ul>	
	c. Follicular neoplasm, Bethesda IV.	
	<ol> <li>The diagnostic or treatment strategy will be contingent on test results</li> </ol>	

Code and Code Description	TAR and/or Billing Requirements	Frequency
0301U	N/A	N/A
Infectious agent detection by nucleic acid (DNA or RNA), Bartonella henselae and Bartonella quintana, droplet digital PCR (ddPCR)		
0302U	N/A	N/A
Infectious agent detection by nucleic acid (DNA or RNA), Bartonella henselae and Bartonella quintana, droplet digital PCR (ddPCR); following liquid enrichment		
‹‹0311U	N/A	One unit per
Infectious disease (bacterial), quantitative antimicrobial susceptibility reported as phenotypic minimum inhibitory concentration (MIC)?based antimicrobial susceptibility for each organisms identified		day Allow TAR/SAR override>>

	nued)››
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Code and Code Description	TAR and/or Billing Requirements	Frequency
‹‹0312U	N/A	One unit per day
Autoimmune diseases (eg, systemic lupus erythematosus [SLE]), analysis of 8 IgG autoantibodies and 2 cell-bound complement activation products using enzyme-linked immunosorbent immunoassay (ELISA), flow cytometry and indirect immunofluorescence, serum, or plasma and whole blood, individual components reported along with an algorithmic SLE-likelihood assessment		Allow TAR/SAR override>>
<<0314U Oncology (cutaneous melanoma), mRNA gene expression profiling by RT-PCR of 35 genes (32 content and 3 housekeeping), utilizing formalin-fixed paraffin-embedded (FFPE) tissue, algorithm reported as a categorical result (ie, benign, intermediate, malignant	N/A	One unit per day Allow TAR/SAR override>>

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Code and Code Description	TAR and/or Billing Requirements	Frequency
<<0321U Infectious agent detection by nucleic acid (DNA or RNA), genitourinary pathogens, identification of 20 bacterial and fungal organisms and identification of 16 associated antibiotic-	N/A	One unit per day Allow TAR/SAR override>>
resistance genes, multiplex amplified probe technique		overnue//

# <u>Legend</u>

Symbols used in the document above are explained in the following table.

Symbol	Description
<<	This is a change mark symbol. It is used to indicate where on the page the most recent change begins.
>>	This is a change mark symbol. It is used to indicate where on the page the most recent change ends.