CMS Manual System	Department of Health & Human Services (DHHS)
Pub 100-04 Medicare Claims Processing	Centers for Medicare & Medicaid Services (CMS)
Transmittal 11208	Date: January 20, 2022
	Change Request 12573

SUBJECT: Healthcare Common Procedure Coding System (HCPCS) Codes Subject to and Excluded from Clinical Laboratory Improvement Amendments (CLIA) Edits

I. SUMMARY OF CHANGES: This Change Request (CR) informs contractors about the new HCPCS codes for 2022 that are subject to and excluded from CLIA edits. This Recurring Update Notification applies to Chapter 16, section 70.9.

EFFECTIVE DATE: April 1, 2022

*Unless otherwise specified, the effective date is the date of service. IMPLEMENTATION DATE: April 4, 2022

Disclaimer for manual changes only: The revision date and transmittal number apply only to red italicized material. Any other material was previously published and remains unchanged. However, if this revision contains a table of contents, you will receive the new/revised information only, and not the entire table of contents.

II. CHANGES IN MANUAL INSTRUCTIONS: (N/A if manual is not updated) R=REVISED, N=NEW, D=DELETED-*Only One Per Row.*

R/N/D	CHAPTER / SECTION / SUBSECTION / TITLE			
N/A	N/A			

III. FUNDING:

For Medicare Administrative Contractors (MACs):

The Medicare Administrative Contractor is hereby advised that this constitutes technical direction as defined in your contract. CMS does not construe this as a change to the MAC Statement of Work. The contractor is not obligated to incur costs in excess of the amounts allotted in your contract unless and until specifically authorized by the Contracting Officer. If the contractor considers anything provided, as described above, to be outside the current scope of work, the contractor shall withhold performance on the part(s) in question and immediately notify the Contracting Officer, in writing or by e-mail, and request formal directions regarding continued performance requirements.

IV. ATTACHMENTS:

Recurring Update Notification

Attachment - Recurring Update Notification

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I. GENERAL INFORMATION

A. Background: The Clinical Laboratory Improvement Amendments (CLIA) regulations require a facility to be appropriately certified for each test performed. To ensure that Medicare & Medicaid only pay for laboratory tests performed in certified facilities, each claim for a HCPCS code that is considered a CLIA laboratory test is currently edited at the CLIA certificate level.

The HCPCS codes that are considered a laboratory test under CLIA change each year. Contractors need to be informed about the new HCPCS codes that are both subject to CLIA edits and excluded from CLIA edits.

The following HCPCS codes were discontinued on March 31, 2021:

- 0098U Test for detection of respiratory disease-causing organism using amplified probe, 14 target organisms;
- 0099U Test for detection of respiratory disease-causing organism using amplified probe, 20 target organisms (adenovirus, coronavirus 229E, coronavirus HKU1, coronavirus, coronavirus OC43, human metapneumovirus, influenza A, influenza A subtype, influenza A subtype H3, influenza A subtype H1-2009, influenza, parainfluenza virus, parainfluenza virus 2, parainfluenza virus 3, parainfluenza virus 4, human rhinovirus/enterovirus, respiratory syncytial virus, Bordetella pertussis, Chlamydophila pneumonia, Mycoplasma pneumoniae); and
- 0100U Test for detection of respiratory disease-causing organism using amplified probe, 20 target organisms (adenovirus, coronavirus 229E, coronavirus HKU1, coronavirus NL63, coronavirus OC43, human metapneumovirus, human rhinovirus/enterovirus, influenza A, including subtypes H1, H1-2009, and H3, influenza B, parainfluenza virus 1, parainfluenza virus 2, parainfluenza virus 3, parainfluenza virus 4, respiratory syncytial virus, Bordetella parapertussis [IS1001], Bordetella pertussis [ptxP], Chlamydia pneumoniae, Mycoplasma pneumoniae).

The following HCPCS codes were discontinued on September 30, 2021:

- 0139U Measurement of 6 central carbon metabolite biomarkers for autism spectrum disorder in plasma; and
- 0168U DNA analysis for detection of abnormal chromosome number of fetus in maternal plasma specimen.

The following HCPCS codes were discontinued on December 31, 2021:

• 0208U - mRNA gene analysis of 108 genes in fine needle aspiration thyroid specimen, reported as positive or negative for medullary thyroid carcinoma;

- 80500 Clinical pathology consultation; and
- 80502 Comprehensive, clinical pathology consultation.

The following HCPCS codes are excluded from CLIA edits, and do not require a facility to have any CLIA certificate:

- 0254U reimplantation genetic assessment of embryo by gene sequence analysis of 24 chromosomes for abnormal chromosome number Effective July 1, 2021;
- 0294U Longevity and mortality risk, mrna, gene expression profiling by rna sequencing of 18 genes, whole blood, algorithm reported as predictive risk score Effective January 1, 2022;
- 80503 Pathology clinical consultation for clinical problem, 5-20 minutes Effective January 1, 2022;
- 80504 Pathology clinical consultation for moderately complex clinical problem, 21-40 minutes Effective January 1, 2022;
- 80505 Pathology clinical consultation for complex clinical problem, 41-60 minutes Effective January 1, 2022; and
- 80506 Pathology clinical consultation, additional 30 minutes Effective January 1, 2022.

The HCPCS codes that follow are all subject to CLIA edits. These lists do not include new HCPCS codes for waived tests or provider-performed microscopy procedures. All these HCPCS codes require a facility to have either a CLIA certificate of registration (certificate type code 9), a CLIA certificate of compliance (certificate type code 1), or a CLIA certificate of accreditation (certificate type code 3). A facility without a valid, current, CLIA certificate, with a current CLIA certificate of waiver (certificate type code 2) or with a current CLIA certificate for provider-performed microscopy procedures (certificate type code 4) must not be permitted to be paid for these tests, unless a facility with a current CLIA certificate of waiver (certificate type code 4) must not be permitted to be paid for these tests, unless a facility with a current CLIA certificate of waiver (certificate type code 4) bills the appropriate HCPCS service code with a QW modifier.

1. The HCPCS codes listed below were added on April 1, 2021 and are subject to CLIA edits.

- 0242U Gene analysis of 55-74 genes associated with solid organ cancer in cell-free circulating DNA, targeted genomic sequence;
- 0243U Time-resolved fluorescence immunoassay of placental-growth factor in maternal serum to evaluate risk of preeclampsia;
- 0244U Gene analysis of 257 genes associated with solid organ cancer in tumor tissue sample, comprehensive genomic profiling;
- 0245U Gene analysis of 10 genes and 37 RNA fusions and expression of 4 mRNA markers, nextgeneration sequencing, in fine needle aspirate of thyroid to evaluate risk of thyroid cancer;
- 0246U Blood typing for 16 or more blood groups with phenotype prediction of 51 or more red blood cell antigens; and
- 0247U Quantitative measurement of insulin-like growth factor-binding protein 4 and sex hormonebinding globulin (SHBG) in maternal serum by LC-MS/MS to evaluate risk of premature birth.
- 2. The HCPCS codes listed below were added on July 1, 2021 and are subject to CLIA edits.
 - 0248U Culture of brain cancer cells with 12 drug panel testing for tumor response prediction;
 - 0249U Analysis of 32 phosphoproteins and protein analytes associated with breast cancer, with interpretation and report;
 - 0250U Gene analysis of 505 genes associated with solid organ cancer in tumor tissue sample, targeted genomic sequence interrogation for somatic alterations, microsatellite instability and tumor-mutation burden;

- 0251U ELISA assay for hepacidin-25 in serum or plasma;
- 0252U Analysis of fetal DNA, short tandem-repeat comparative analysis, for abnormal chromosome number
- 0253U RNA gene expression profiling of 238 genes by next-generation sequencing specimen from lining of womb to evaluate window of implantation for embryo transfer; and
- G0327 Colorectal cancer screening; blood-based biomarker.
- 3. The HCPCS codes listed below were added on October 1, 2021 and are subject to CLIA edits.
 - 0018M Measurement of kidney donor and third-party-induced CD154+T-cytotoxic memory cells in whole peripheral blood specimen, algorithm reported as kidney transplant rejection risk score;
 - 0255U Evaluation of sperm using fluorescence microscopic evaluation of ganglioside GM1 distribution patterns, reported as percentage of capacitated sperm and probability of generating pregnancy score;
 - 0256U Tandem mass spectroscopy (MS/MS) profile of trimethylamine/trimethylamine N-oxide (TMA/TMAO) profile in urine, with algorithmic analysis and interpretive report;
 - 0257U Evaluation of very long chain acyl-coenzyme A (CoA) dehydrogenase (VLCAD) white blood cell enzyme activity in whole blood;
 - 0258U mRNA gene expression profiling of 50-100 genes in skin surface specimen, algorithm reported as likelihood of response to psoriasis biologics;
 - 0259U Nuclear MR spectroscopy measurement of myo-inositol, valine, and creatinine, algorithmically combined with cystatin C (by immunoassay) and demographic data to evaluate kidney function;
 - 0260U Optical genome mapping for detection of abnormalities associated with rare heritable diseases;
 - 0261U Image analysis with artificial intelligence assessment of 4 cellular and immune features in colorectal cancer tumor tissue specimen, reported as immune response and recurrence-risk score;
 - 0262U mRNA gene expression profiling of 7 gene pathways in solid organ tumor tissue specimen, algorithm reported as gene pathway activity score;
 - 0263U LC-MS/MS spectroscopy of 16 central carbon metabolites associated with autism spectrum disorders (ASD) in plasma specimen, algorithmic analysis with result reported as negative or positive (with metabolic subtypes of ASD);
 - 0264U Detection of abnormalities associated with rare heritable diseases by optical genome mapping;
 - 0265U Whole genome and mDNA sequence analysis for detection of abnormalities associated with rare constitutional/heritable diseases;
 - 0266U Gene expression profiling by whole transcriptone and next-generation sequencing for detection of unexplained heritable disease;
 - 0267U Optical genome mapping and whole genome sequencing for detection of abnormalities associated with rare heritable diseases;
 - 0268U Genomic sequence analysis of 15 genes for detection of abnormalities associated with atypical hemolytic uremic syndrome;
 - 0269U Genomic sequence analysis of 14 genes for detection of abnormalities associated with autosomal dominant congenital thrombocytopenia (low platelet count);
 - 0270U Genomic sequence analysis of 20 genes for detection of abnormalities associated with congenital coagulation disorders (blood clotting disorders);
 - 0271U Genomic sequence analysis of 23 genes for detection of abnormalities associated with congenital neutropenia (low white blood cell count);
 - 0272U Comprehensive genomic sequence analysis of 51 genes for detection of abnormalities associated with congenital bleeding disorders;
 - 0273U Genomic sequence analysis of 8 genes for detection of abnormalities associated with genetic hyperfibrinolysis and delayed bleeding;
 - 0274U Genomic sequence analysis of 43 genes for detection of abnormalities associated with genetic platelet disorders;

- 0275U Flow cytometry detection of platelet antibody reactivity in serum for evaluation of heparininduced thrombocytopenia (low platelet count due to heparin);
- 0276U Genomic sequence analysis of 23 genes for detection of abnormalities associated with inherited thrombocytopenia (low platelet count);
- 0277U Genomic sequence analysis of 31 genes for detection of abnormalities associated with genetic platelet function disorder;
- 0278U Genomic sequence analysis of 12 genes for detection of abnormalities associated with genetic thrombosis (excessive clotting);
- 0279U ELISA detection of von Willebrand factor (VWF) and collagen III binding in plasma specimen, report of collagen III binding;
- 0280U ELISA detection of von Willebrand factor (VWF) and collagen IV binding in plasma specimen, report of collagen IV binding;
- 0281U ELISA measurement of von Willebrand propeptide in plasma specimen, diagnostic report of von Willebrand factor (VWF) propeptide antigen level;
- 0282U Red blood cell antigen genotyping of 12 blood group system genes to predict 44 red blood cell antigen phenotypes;
- 0283U Radioimmunoassay platelet-binding evaluation of von Willebrand factor (VWF), type 2B, in plasma specimen; and
- 0284U ELISA evaluation of von Willebrand factor (VWF), type 2N, factor VIII and VWF binding in plasma specimen.
- 4. The HCPCS codes listed below were added on January 1, 2022 and are subject to CLIA edits.
 - 0285U Oncology, response to radiation, cell-free dna, quantitative branched chain dna amplification, plasma, reported as a radiation toxicity score;
 - 0286U Cep72 (centrosomal protein, 72-kda), nudt15 (nudix hydrolase 15) and tpmt (thiopurine smethyltransferase) (eg, drug metabolism) gene analysis, common variants;
 - 0287U 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);
 - 0288U Oncology (lung), mrna, quantitative pcr analysis of 11 genes (bag1, brca1, cdc6, cdk2ap1, erbb3, fut3, il11, lck, rnd3, sh3bgr, wnt3a) and 3 reference genes (esd, tbp, yap1), formalin-fixed paraffin-embedded (ffpe) tumor tissue, algorithmic interpretation reported as a recurrence risk score;
 - 0289U Neurology (alzheimer disease), mrna, gene expression profiling by rna sequencing of 24 genes, whole blood, algorithm reported as predictive risk score;
 - 0290U Pain management, mrna, gene expression profiling by rna sequencing of 36 genes, whole blood, algorithm reported as predictive risk score;
 - 0291U Psychiatry (mood disorders), mrna, gene expression profiling by rna sequencing 144 genes, whole blood, algorithm reported as predictive risk score;
 - 0292U Psychiatry (stress disorders), mrna, gene expression profiling by rna sequencing of 72 genes, whole blood, algorithm reported as predictive risk score;
 - 0293U Psychiatry (suicidal ideation), mrna, gene expression profiling by rna sequencing of 54 genes, whole blood, algorithm reported as predictive risk score;
 - 0295U Oncology (breast ductal carcinoma in situ), protein expression profiling by immunohistochemistry of 7 proteins (cox2, foxa1, her2, ki-67, p16, pr, siah2), with 4 clinicopathologic factors (size, age, margin status, palpability), utilizing formalin-fixed paraffinembedded (ffpe) tissue, algorithm reported as a recurrence risk score;
 - 0296U Oncology (oral and/or oropharyngeal cancer), gene expression profiling by rna sequencing at least 20 molecular features (eg, human and/or microbial mrna), saliva, algorithm reported as positive or negative for signature associated with malignancy;
 - 0297U Oncology (pan tumor), whole genome sequencing of paired malignant and normal dna specimens, fresh or formalin-fixed paraffin-embedded (ffpe) tissue, blood or bone marrow, comparative sequence analyses and variant identification;

- 0298U Oncology (pan tumor), whole transcriptome sequencing of paired malignant and normal rna specimens, fresh or formalin-fixed paraffin-embedded (ffpe) tissue, blood or bone marrow, comparative sequence analyses and expression level and chimeric transcript identification;
- 0299U Oncology (pan tumor), whole genome optical genome mapping of paired malignant and normal dna specimens, fresh frozen tissue, blood, or bone marrow, comparative structural variant identification;
- 0300U Oncology (pan tumor), whole genome sequencing and optical genome mapping of paired malignant and normal dna specimens, fresh tissue, blood, or bone marrow, comparative sequence analyses and variant identification;
- 0301U Infectious agent detection by nucleic acid (dna or rna), bartonella henselae and bartonella quintana, droplet digital pcr (ddpcr);
- 0302U Infectious agent detection by nucleic acid (dna or rna), bartonella henselae and bartonella quintana, droplet digital pcr (ddpcr); following liquid enrichment;
- 0303U Hematology, red blood cell (rbc) adhesion to endothelial/subendothelial adhesion molecules, functional assessment, whole blood, with algorithmic analysis and result reported as an rbc adhesion index; hypoxic;
- 0304U Hematology, red blood cell (rbc) adhesion to endothelial/subendothelial adhesion molecules, functional assessment, whole blood, with algorithmic analysis and result reported as an rbc adhesion index; normoxic;
- 0305U Hematology, red blood cell (rbc) functionality and deformity as a function of shear stress, whole blood, reported as a maximum elongation index;
- 80220 Measurement of hydroxychloroquine;
- 81349 Genome-wide microarray analysis for copy number and loss-of-heterozygosity variants;
- 81523 Next-generation sequencing of breast cancer profiling 70 content genes and 31 housekeeping genes;
- 81560 Measurement of donor and third-party memory cells for transplantation medicine;
- 82653 Measurement of pancreatic elastase (enzyme) in stool;
- 83521- Measurement of immunoglobulin light chains;
- 83529 Measurement of interleukin-6;
- 86015 Measurement of Actin (smooth muscle) antibody;
- 86036 Screening test for antineutrophil cytoplasmic antibody;
- 86037 Antineutrophil cytoplasmic antibody titer;
- 86051 ELISA detection of aquaporin-4 (neuromyelitis optica [NMO]) antibody;
- 86052 Cell-based immunofluorescence (CBA) detection of aquaporin-4 (neuromyelitis optica [NMO]) antibody;
- 86053 Flow cytometry detection of aquaporin-4 (neuromyelitis optica [NMO]) antibody;
- 86231 Detection of endomysial antibody (EMA);
- 86258 Detection of gliadin (deamidated) (DGP) antibody;
- 86362 Cell-based immunofluorescence (CBA) detection of myelin oligodendrocyte glycoprotein (MOG-IgG1) antibody;
- 86363 Flow cytometry detection of myelin oligodendrocyte glycoprotein (MOG-IgG1) antibody;
- 86364 Measurement of tissue transglutaminase;
- 86381 Measurement of mitochondrial antibody;
- 86596 Measurement of voltage-gated calcium channel antibody; and
- 87154 Amplifed nucleic acid probe typing of disease agent in blood culture specimen.

NOTE This instruction is NOT intended to rescind/replace any previous instructions indicating that a laboratory with a valid CLIA certificate of waiver or CLIA certificate for provider-performed microscopy procedures be allowed to bill the above codes with a QW modifier.

This Recurring Update Notification applies to Chapter 16, Section 70.9.

B. Policy: The CLIA regulations require a facility to be appropriately certified for each test performed. To ensure that Medicare and Medicaid only pay for laboratory tests in a facility with a valid, current CLIA certificate, laboratory claims are currently edited at the CLIA certificate level.

II. BUSINESS REQUIREMENTS TABLE

Number	Requirement	Re	spoi	nsibility						
		A/B MAC DMI			DME	Shared-System Maintainers				Other
		A	В	HHH	MAC	FISS	MCS	VMS	CWF	
12573.1	Contractors shall apply CLIA edits to the HCPCS codes mentioned above as subject to CLIA edits.		Х						X	
12573.2	Contractors shall deny payment for a claim submitted with the HCPCS codes mentioned above as subject to CLIA edits to a provider without valid current CLIA certificate, with a CLIA certificate of waiver (certificate type code 2) (when billed <u>without</u> the 'QW' modifier), or with a CLIA certificate for provider-performed microscopy procedures (certificate type code 4)(when billed <u>without</u> the 'QW' modifier).		X							
12573.3	Contractors shall return a claim as unprocessable if a CLIA number is not submitted on claims by providers for the HCPCS mentioned above as subject to CLIA edits.		X							
12573.4	Contractors need not search their files to either retract payment for claims already paid or to retroactively pay claims. However, contractors shall adjust claims brought to their attention.		X							

III. PROVIDER EDUCATION TABLE

Number	Requirement	Re	spoi	nsibility	,	
			A/ M/		DME MAC	CEDI
		A	В	ННН		
12573.5	Medicare Learning Network® (MLN): CMS will market provider education content through the MLN Connects® newsletter shortly after CMS releases the CR. MACs shall follow IOM Pub. No. 100-09 Chapter 6, Section 50.2.4.1 instructions for distributing the MLN Connects newsletter information to providers and link to relevant information on your website. You may supplement MLN content with your local information after we release the MLN Connects newsletter. Subscribe to the "MLN Connects" listserv to get MLN content notifications. You don't need to separately track and report MLN content releases when you distribute MLN Connects newsletter content per the manual section referenced above.		Х			

IV. SUPPORTING INFORMATION

Section A: Recommendations and supporting information associated with listed requirements:

"Should" denotes a recommendation.

X-Ref Requirement Number	Recommendations or other supporting information:
	N/A

Section B: All other recommendations and supporting information: N/A

V. CONTACTS

Pre-Implementation Contact(s): Kathleen Todd, kathleen.todd@cms.hhs.gov

Post-Implementation Contact(s): Contact your Contracting Officer's Representative (COR).

VI. FUNDING

Section A: For Medicare Administrative Contractors (MACs):

The Medicare Administrative Contractor is hereby advised that this constitutes technical direction as defined in your contract. CMS does not construe this as a change to the MAC Statement of Work. The contractor is not obligated to incur costs in excess of the amounts allotted in your contract unless and until specifically authorized by the Contracting Officer. If the contractor considers anything provided, as described above, to be outside the current scope of work, the contractor shall withhold performance on the part(s) in question and immediately notify the Contracting Officer, in writing or by e-mail, and request formal directions regarding continued performance requirements.

ATTACHMENTS: 0