

Comprehensive Genomic Profiling and Genetic Testing for Solid Tumors



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Medical Coverage Policy

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Related Medical/Pharmacy Coverage Policies

[Genetic Testing](#)
[Genetic Testing for Hematologic Malignancies and Suspected Myeloid Disorders](#)
[Liquid Biopsy](#)
[Pharmacogenomics and Companion Diagnostics](#)

Description

Comprehensive genomic profiling (CGP) (also referred to as comprehensive molecular profiling) is a type of somatic (tumor) test that involves a combination of laboratory methodologies to detect genetic alterations and the simultaneous evaluation of large numbers (hundreds to thousands) of biomarkers in tumor tissue to aid in the management of advanced solid tumors, including guiding treatment decisions as well as determination of clinical trial eligibility. Techniques can vary from test to test and may include next-generation sequencing (NGS), fluorescence in situ hybridization (FISH) and immunohistochemistry (IHC) and often provides information on tumor mutational burden (TMB), microsatellite instability (MSI) and homologous recombination deficiency (HRD). Examples include **Altera Tumor Genomic Profiling**, **Guardant360**, **NeoGenomics Solid Tumor NGS Fusion Panel** and **TissueNext**.

Some CGP tests analyze both deoxyribonucleic acid (DNA) and ribonucleic acid (RNA). **NeoTYPE DNA & RNA – Lung** is an NGS profiling test that detects single nucleotide variants, insertions/deletions, copy number variants, and/or RNA fusions in a total of 50 genes (44 genes analyzed by DNA and 19 by RNA), plus MSI and TMB. **Tempus xT** is another example of a CGP somatic test performed for the management of advanced cancer. Previously Tempus xT conducted sequencing for both ribonucleic acid (RNA) and deoxyribonucleic acid (DNA) within a single panel. However, a new test, **Tempus xR**, is now available as an independent assay dedicated solely to RNA sequencing. This means that Tempus xT focuses on DNA analysis while Tempus xR specializes in RNA analysis, and they are no longer combined into one test.

Single gene testing can be utilized to diagnose and monitor cancer including, but may not be limited to, cholangiocarcinoma, gallbladder cancer, gastrointestinal stromal tumor, glioblastoma, melanoma and thyroid cancer. This type of testing is indicated for an individual who exhibits disease symptoms and may be necessary to diagnose or rule out suspected cancer.

DNA Specimen Provenance Assignment (DSPA) Testing (eg, know error System) is a molecular diagnostic test intended for the protection and control of tissue samples to purportedly decrease the incidence of diagnostic mistakes due to the misidentification, specimen transposition or cell contamination of samples, also known as specimen provenance complications (SPCs). Breast and prostate tissues are most often tested but other tissue types, such as bone marrow, may also be examined.

LungOI is an artificial-intelligence (AI)-based molecular profiling that uses a digitized biopsy image and is proposed for the diagnosis of lung cancer. **(Refer to Coverage Limitations section)**

Genomic profiling in conjunction with in vitro chemoresistance and chemosensitivity assays has been proposed to guide treatment decisions. **(Refer to Coverage Limitations section)**

Coverage Determination

Any state mandates for comprehensive genomic profiling and genetic testing for solid tumors take precedence over this medical coverage policy.

Genetic testing may be excluded by certificate. Please consult the member's individual certificate regarding Plan coverage.

Comprehensive Genomic Profiling for Solid Tumors

Humana members may be eligible under the Plan for **comprehensive genomic profiling for solid tumors** (81445, 81455, 81456, 81449) (eg, LungHDPK [0478U], OncoReveal Dx Lung & Colon Cancer Assay [0448U], OptiSeq Colorectal Cancer NGS Panel [0498U], OptiSeq Dual Cancer Panel Kit [0499U]) when the following criteria are met:

- Diagnosed with recurrent, relapsed, refractory, metastatic or advanced stage III or IV cancer; **AND**
- Has not previously received comprehensive molecular profiling for the same tumor type; **AND**
- Treatment with anticancer therapy is being considered

IDH1, IDH2 and TERT Mutation Analysis Panel

Humana members may be eligible under the Plan for **IDH1, IDH2 and TERT Mutation Analysis Panel** (0481U) for glioma.

Single Gene Testing

IDH1/IDH2 Gene Analysis

Humana members may be eligible under the Plan for **IDH1 (81120) and/or IDH2 (81121) gene analysis** for any of the following indications:

- Cholangiocarcinoma, locally advanced, metastatic or unresectable (**IDH1 only**); **OR**
- Gallbladder cancer, metastatic or unresectable (**IDH1 only**); **OR**
- Glioma

KIT (c-KIT) Gene Analysis

Humana members may be eligible under the Plan for **KIT (c-KIT) gene analysis** (81272) for any of the following indications:

- Gastrointestinal stromal tumor (GIST); **OR**
- Melanoma, metastatic or unresectable

KRAS Gene Analysis

Humana members may be eligible under the Plan for **KRAS gene analysis** (81275, 81276) for any of the following indications:

- Colorectal cancer, metastatic; **OR**
- Non-small cell lung cancer, metastatic

MGMT Promoter Methylation Testing

Humana members may be eligible under the Plan for **MGMT promoter methylation testing** (81287) for glioma.

NRAS Gene Analysis

Humana members may be eligible under the Plan for **NRAS gene analysis** (81311) for any of the following indications:

- Colorectal cancer, metastatic; **OR**
- Melanoma, metastatic

PDGFRA Gene Analysis

Humana members may be eligible under the Plan for **PDGFRA gene analysis** (81314) an individual presents with a mass known or clinically suspected to be GIST.

TERT Gene Analysis

Humana members may be eligible under the Plan for **TERT gene analysis** (81345) for any of the following indications:

- Glioma; **OR**
- Melanoma; **OR**
- Uterine sarcoma

TP53 Gene Analysis

Humana members may be eligible under the Plan for **TP53 gene analysis** (81351, 81352) for any of the following indications:

- Endometrial carcinoma; **OR**
- Medulloblastoma; **OR**
- Neuroendocrine tumors; **OR**
- Occult primary; **OR**
- Pancreatic adenocarcinoma; **OR**
- Uterine sarcoma

Repeat CGP

Humana members may be eligible under the Plan for **repeat CGP** when the following criteria are met:

- Individual diagnosed with advanced or metastatic solid tumor; **AND**
- Individual diagnosed with recurrence, relapse or is nonresponsive to treatment; **AND**
- Examination of a new sample of the primary tumor

Coverage Limitations

Humana members may **NOT** be eligible under the Plan for **comprehensive genomic profiling or genetic testing for solid tumors** for any indications or tests other than those listed above including, but may not be limited to:

- Adjunct to in vitro chemoresistance and chemosensitivity assays
- CureMatch (0794T)
- LungOI (0414U)

These are considered experimental/investigational as they are not identified as widely used and generally accepted for the proposed uses as reported in nationally recognized peer-reviewed medical literature published in the English language.

Coding Information

Any codes listed on this policy are for informational purposes only. Do not rely on the accuracy and inclusion of specific codes. Inclusion of a code does not guarantee coverage and/or reimbursement for a service or procedure.

CPT® Code(s)	Description	Comments
81120	IDH1 (isocitrate dehydrogenase 1 [NADP+], soluble) (eg, glioma), common variants (eg, R132H, R132C)	
81121	IDH2 (isocitrate dehydrogenase 2 [NADP+], mitochondrial) (eg, glioma), common variants (eg, R140W, R172M)	
81272	KIT (v-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog) (eg, gastrointestinal stromal tumor [GIST], acute myeloid leukemia, melanoma), gene analysis, targeted sequence analysis (eg, exons 8, 11, 13, 17, 18)	
81275	KRAS (Kirsten rat sarcoma viral oncogene homolog) (eg, carcinoma) gene analysis; variants in exon 2 (eg, codons 12 and 13)	
81276	KRAS (Kirsten rat sarcoma viral oncogene homolog) (eg, carcinoma) gene analysis; additional variant(s) (eg, codon 61, codon 146)	
81287	MGMT (O-6-methylguanine-DNA methyltransferase) (eg, glioblastoma multiforme) promoter methylation analysis	

81311	NRAS (neuroblastoma RAS viral [v-ras] oncogene homolog) (eg, colorectal carcinoma), gene analysis, variants in exon 2 (eg, codons 12 and 13) and exon 3 (eg, codon 61)	
81314	PDGFRA (platelet-derived growth factor receptor, alpha polypeptide) (eg, gastrointestinal stromal tumor [GIST]), gene analysis, targeted sequence analysis (eg, exons 12, 18)	
81345	TERT (telomerase reverse transcriptase) (eg, thyroid carcinoma, glioblastoma multiforme) gene analysis, targeted sequence analysis (eg, promoter region)	
81351	TP53 (tumor protein 53) (eg, Li-Fraumeni syndrome) gene analysis; full gene sequence	
81352	TP53 (tumor protein 53) (eg, Li-Fraumeni syndrome) gene analysis; targeted sequence analysis (eg, 4 oncology)	
81449	Targeted genomic sequence analysis panel, solid organ neoplasm, 5-50 genes (eg, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; RNA analysis	
81451	Targeted genomic sequence analysis panel, hematolymphoid neoplasm or disorder, 5-50 genes (eg, BRAF, CEBPA, DNMT3A, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NOTCH1, NPM1, NRAS), interrogation for sequence variants, and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis	
81455	Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm, DNA analysis, and RNA analysis when performed, 51 or greater genes (eg, ALK, BRAF, CDKN2A, CEBPA, DNMT3A, EGFR, ERBB2, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MLL, NPM1, NRAS, MET, NOTCH1, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed	

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81456	Targeted genomic sequence analysis panel, solid organ or hematolymphoid neoplasm or disorder, 51 or greater genes (eg, ALK, BRAF, CDKN2A, CEBPA, DNMT3A, EGFR, ERBB2, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MET, MLL, NOTCH1, NPM1, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; RNA analysis	
81457	Solid organ neoplasm, genomic sequence analysis panel, interrogation for sequence variants; DNA analysis, microsatellite instability	New Code Effective 01/01/2024
81458	Solid organ neoplasm, genomic sequence analysis panel, interrogation for sequence variants; DNA analysis, copy number variants and microsatellite instability	New Code Effective 01/01/2024
81459	Solid organ neoplasm, genomic sequence analysis panel, interrogation for sequence variants; DNA analysis or combined DNA and RNA analysis, copy number variants, microsatellite instability, tumor mutation burden, and rearrangements	New Code Effective 01/01/2024
81479	Unlisted molecular pathology procedure	Not Covered if used to report any test outlined in Coverage Limitations section
0022U	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	
0048U	Oncology (solid organ neoplasia), DNA, targeted sequencing of protein-coding exons of 468 cancer-associated genes, including interrogation for somatic mutations and microsatellite instability, matched with normal specimens, utilizing formalin-fixed paraffin-embedded tumor tissue, report of clinically significant mutation(s)	
0211U	Oncology (pan-tumor), DNA and RNA by next-generation sequencing, utilizing formalin-fixed paraffin-embedded tissue, interpretative report for single nucleotide variants, copy number alterations, tumor mutational burden, and microsatellite instability, with therapy association	Not Covered Test Obsolete/No Longer Available

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0244U	Oncology (solid organ), DNA, comprehensive genomic profiling, 257 genes, interrogation for single-nucleotide variants, insertions/deletions, copy number alterations, gene rearrangements, tumor-mutational burden and microsatellite instability, utilizing formalin-fixed paraffin-embedded tumor tissue	
0250U	Oncology (solid organ neoplasm), targeted genomic sequence DNA analysis of 505 genes, interrogation for somatic alterations (SNVs [single nucleotide variant], small insertions and deletions, one amplification, and four translocations), microsatellite instability and tumor-mutation burden	
0329U	Oncology (neoplasia), exome and transcriptome sequence analysis for sequence variants, gene copy number amplifications and deletions, gene rearrangements, microsatellite instability and tumor mutational burden utilizing DNA and RNA from tumor with DNA from normal blood or saliva for subtraction, report of clinically significant mutation(s) with therapy associations	
0334U	Oncology (solid organ), targeted genomic sequence analysis, formalin-fixed paraffin-embedded (FFPE) tumor tissue, DNA analysis, 84 or more genes, interrogation for sequence variants, gene copy number amplifications, gene rearrangements, microsatellite instability and tumor mutational burden	
0379U	Targeted genomic sequence analysis panel, solid organ neoplasm, DNA (523 genes) and RNA (55 genes) by next-generation sequencing, interrogation for sequence variants, gene copy number amplifications, gene rearrangements, microsatellite instability, and tumor mutational burden	
0391U	Oncology (solid tumor), DNA and RNA by next-generation sequencing, utilizing formalin-fixed paraffin-embedded (FFPE) tissue, 437 genes, interpretive report for single nucleotide variants, splice-site variants, insertions/deletions, copy number alterations, gene fusions, tumor mutational burden, and microsatellite instability, with algorithm quantifying immunotherapy response score	
0414U	Oncology (lung), augmentative algorithmic analysis of digitized whole slide imaging for 8 genes (ALK, BRAF, EGFR, ERBB2, MET, NTRK1-3, RET, ROS1), and KRAS G12C and PD-L1, if performed, formalin-fixed paraffin-embedded (FFPE) tissue, reported as positive or negative for each biomarker	<p style="text-align: center;">Not Covered</p> <p style="text-align: center;">New Code Effective 10/01/2023</p>

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0444U	Oncology (solid organ neoplasia), targeted genomic sequence analysis panel of 361 genes, interrogation for gene fusions, translocations, or other rearrangements, using DNA from formalin-fixed paraffin-embedded (FFPE) tumor tissue, report of clinically significant variant(s)	
0448U	Oncology (lung and colon cancer), DNA, qualitative, next-generation sequencing detection of single-nucleotide variants and deletions in EGFR and KRAS genes, formalin-fixed paraffin-embedded (FFPE) solid tumor samples, reported as presence or absence of targeted mutation(s), with recommended therapeutic options	New Code Effective 04/01/2024
0478U	Oncology (non-small cell lung cancer), DNA and RNA, digital PCR analysis of 9 genes (EGFR, KRAS, BRAF, ALK, ROS1, RET, NTRK 1/2/3, ERBB2, and MET) in formalin-fixed paraffin-embedded (FFPE) tissue, interrogation for single-nucleotide variants, insertions/deletions, gene rearrangements, and reported as actionable detected variants for therapy selection	New code Effective Date 10/01/2024
0481U	IDH1 (isocitrate dehydrogenase 1 [NADP+]), IDH2 (isocitrate dehydrogenase 2 [NADP+]), and TERT (telomerase reverse transcriptase) promoter (eg, central nervous system [CNS] tumors), next-generation sequencing (single-nucleotide variants [SNV], deletions, and insertions)	New code Effective Date 10/01/2024
0498U	Oncology (colorectal), next-generation sequencing for mutation detection in 43 genes and methylation pattern in 45 genes, blood, and formalin-fixed paraffin-embedded (FFPE) tissue, report of variants and methylation pattern with interpretation	New code Effective Date 10/01/2024
0499U	Oncology (colorectal and lung), DNA from formalin-fixed paraffin-embedded (FFPE) tissue, next-generation sequencing of 8 genes (NRAS, EGFR, CTNNB1, PIK3CA, APC, BRAF, KRAS, and TP53), mutation detection	New code Effective Date 10/01/2024
CPT® Category III Code(s)	Description	Comments

0794T	Patient-specific, assistive, rules-based algorithm for ranking pharmaco-oncologic treatment options based on the patient's tumor-specific cancer marker information obtained from prior molecular pathology, immunohistochemical, or other pathology results which have been previously interpreted and reported separately	Not Covered New Code Effective 07/01/2023
HCPSC Code(s)	Description	Comments
No code(s) identified		

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Change Summary

07/25/2024 Annual Review, Coverage Change. Provider Claims Codes Update.

08/29/2024 Update, Coverage Change. Provider Claims Codes Update.