

2013 CPT Codes - PATHOLOGY

Category	Code Range	New	Deleted	Revised	Total
PATHOLOGY & LABORATORY	80047 - 89398	41	24	19	84

*Underlined words are new for 2013

*Words with a ~~Strikethrough~~ are deleted for 2013

New Codes

81201 - APC (adenomatous polyposis coli) (eg. familial adenomatosis polyposis (FAP), attenuated FAP gene analysis; full gene sequence

81202 - ; know familial variants

81203 - ; duplication/deletion variants

81236 - EGFR (epidermal growth factor receptor) (eg, non-small cell lung cancer) gene analysis, common variants (eg, exon 19 LREA deletion, L858R, T790M, G719A, G719S, L861Q)

81252 - GJB2 (gap junction, beta 2, 26KDa; connexin 26) (eg, nonsyndromic hearing loss) gene analysis; full gene sequence

81253 - ; known familial variants

81254 - GJB6 (gap junction protein, beta6, 30KDa, connexin 30) (eg, nonsyndromic hearing loss) gene analysis, common variants (eg, 309kb [del (GJB6-D13S1830)] and 232kb [del (GJB6-D13S1854)])

81321 - PTEN (phosphatase and tensin homolog) (eg, Cowden syndrome, PTEN hamartoma tumor syndrome) gene analysis; full sequence analysis

81322 - ; known familial variant

81323 - ; duplication/deletion variant

81324 - PMP22 (peripheral myelin protein 22) (eg, Charcot-Marie-Tooth, hereditary neuropathy with liability to pressure palsies) gene analysis; duplication/deletion analysis

81325 - ; full sequence analysis

81326 - ; known familial variant

81479 - Unlisted molecular pathology procedure

81500 - Oncology (ovarian) biochemical assays of two proteins (Ca-125 and HE4), utilizing serum, with menopausal status, algorithm reported as a risk score

81503 - Oncology (ovarian) biochemical assays of five proteins (Ca-125, apolipoprotein A1, beta-2 microglobulin, transferrin, and pre-albumin), utilizing serum, algorithm reported as a risk score

81506 - Endocrinology (type 2 diabetes), biochemical assays of seven analytes (glucose, HbA1c, insulin, hs-CRP, adiponectin, ferritin, interleukin 2-receptor alpha), utilizing serum or plasma, algorithm reporting a risk score

81508 - Fetal congenital abnormalities, biochemical assays of two proteins (PAPP-A, hCG [any form]) utilizing maternal serum, algorithm reported as a risk score

81509 - Fetal congenital abnormalities, biochemical assays of two proteins (PAPP-A, hCG [any form], DIA) utilizing maternal serum, algorithm reported as a risk score

81510 - Fetal congenital abnormalities, biochemical assays of three analytes (AFP, uE3, hCG [any form]) utilizing maternal serum, algorithm reported as a risk score

81511 - Fetal congenital abnormalities, biochemical assays of four analytes (AFP, uE3, hCG [any form], DIA) utilizing maternal serum, algorithm reported as a risk score (may included additional results from previous biochemical testing)

81312 - Fetal congenital abnormalities, biochemical assays of five analytes (AFP, uE3, hCG , hyperglycosylated hCG, DIA) utilizing maternal serum, algorithm reported as a risk score

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- 81599** - Unlisted multianalyte assay with algorithmic analysis
82777 - Galectin-3
86152 - Cell enumeration using immunologic selection and identification in fluid specimen (eg, circulating tumor cells in blood)
86153 - ; physician interpretation and report, when required
86711 - Antibody, JC (John Cunningham) virus
86828 - Antibody to human leukocyte antigens (HLA), solid phase assays (eg, microspheres or beads, ELISA, Flow cytometry); qualitative assessment of the presence or absence of antibody (ies) to HLA Class I and Class II HLA antigens
86829 - ; qualitative assessment of the presence or absence of antibody (ies) to HLA Class I or Class II HLA antigens
86830 - ; antibody identification by qualitative panel using complete HLA phenotypes, HLA class I
86831 - ; antibody identification by qualitative panel using complete HLA phenotypes, HLA class II
86832 - ; high definition qualitative panel for identification of antibody specificities (eg, individual antigen per bead methodology), HLA Class I
86833 - ; high definition qualitative panel for identification of antibody specificities (eg, individual antigen per bead methodology), HLA Class II
86834 - ; semi-quantitative panel (eg, titer), HLA Class I
86835 - ; semi-quantitative panel (eg, titer), HLA Class II
87631 - Infectious agent detection by nucleic acid (DNA or RNA); respiratory virus (eg, adenovirus, influenza virus, coronavirus, metapneumovirus, parainfluenza virus, respiratory syncytial virus, rhinovirus), multiplex reverse transcription and amplified probe technique, multiple types or subtypes; 3-5 targets
87632 - ; 6 - 11 targets
87633 - ; 12 - 25 targets
87910 - Infectious agent genotype analysis by nucleic acid (DNA or RNA); cytomegalovirus
87912 - Hepatitis B virus
88375 - Optical endomicroscopic image(s), interpretation and report, real-time or referred, each endoscopic session

Deleted Codes

- 83890** - Molecular diagnostics; molecular isolation or extraction, each nucleic acid type (ie, DNA or RNA)
83891 - ; isolation or extraction of highly purified nucleic acid, each nucleic acid type (ie, DNA or RNA)
83892 - ; enzymatic digestion, each enzyme treatment
83893 - ; dot/sot blot production, each nucleic acid preparation
83894 - ; separation by gel electrophoresis (eg, agarose, polyacrylamide), each nucleic acid preparation
83896 - ; nucleic acid probe, each
83897 - ; nucleic acid transfer (eg, Southern, Northern), each nucleic acid preparation
83898 - ; amplification, target, each nucleic acid sequence
83900 - ; amplification, target, multiplex, first 2 nucleic acid sequences
83901 - ; amplification, target, multiplex, each additional nucleic acid sequence beyond 2 (Add-on code)
83902 - ; reverse transcription
83903 - ; mutation scanning by physician properties (eg, single strand conformational polymorphisms (SSCP), heteroduplex denaturing gradient gel electrophoresis (DGGE), RNAase A, single segment, each
83904 - ; mutation identification by sequencing, single segment, each segment
83905 - ; mutation identification by allele specific transcription, single segment, each segment
83906 - ; mutation identification by allele specific translation, single segment, each segment

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- 83907** - ; lysis of cells prior to nucleic acid extraction (eg, stool specimens, paraffin embedded tissue, each specimen
- 83908** - ; amplification, signal, each nucleic acid sequence
- 83909** - ; separation and identification by high resolution technique (eg, capillary electrophoresis, each nucleic acid preparation
- 83912** - ; interpretation and report
- 83913** - ; RNA stabilization
- 83914** - ; Mutation identification by enzymatic ligation or primer extension; single segment, each segment
- 88383** - Array-based evaluation of multiple molecular probes; 11 through 50 probes
- 88385** - ; 51 – 250 probes
- 88386** - ; 251 – 500 probes

Revised Codes

81400 – 81409 - Molecular pathology procedures, Levels 1-8 - Extensive revisions have been made to these molecular pathology procedures. For the complete detailed revisions, see the 2013 AMA CPT Manual.

82009 – ~~Acetone or other ketone bodies~~ Ketone body(s), acetone acetoacetic acid, serum beta hydroxybutyrate; qualitative

82010 - ; quantitative

87498 - enterovirus, reverse transcription and amplified probe technique

87521 - hepatitis C, reverse transcription and amplified probe technique

87522 - hepatitis C, reverse transcription and quantification

87535 - HIV-1, reverse transcription and amplified probe technique

87536 - HIV-1, reverse transcription and quantification

87538 - HIV-2, reverse transcription and amplified probe technique

87539 - HIV-2, reverse transcription and quantification

87901 - HIV-1, reverse transcriptase and protease regions