# 2013 CPT Codes - PATHOLOGY

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*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, adoponectin, 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