



List of Mutations Detectable with the OncoFOCUS™ Panel

| GENE          | NCBI Ref Seq  | HGVS Nomenclature                  |                     | COSMIC ID |
|---------------|---------------|------------------------------------|---------------------|-----------|
|               |               | CDS Mutation                       | AA Mutation         |           |
| EGFR          | NM_005228     | c.323G>A                           | p.R108K             | 21683     |
|               |               | c.787A>C                           | p.T263P             | 21684     |
|               |               | c.866C>T                           | p.A289V             | 21687     |
|               |               | c.866C>A                           | p.A289D             | 21685     |
|               |               | c.1793G>T                          | p.G598V             | 21690     |
|               |               | c.2126A>T                          | p.E709V             | 12371     |
|               |               | c.2125_2127GAA>CAT                 | p.E709H             | 12428     |
|               |               | c.2125G>A                          | p.E709K             | 12988     |
|               |               | c.2126A>G                          | p.E709G             | 13009     |
|               |               | c.2125G>C                          | p.E709Q             | 116882    |
|               |               | c.2126A>C                          | p.E709A             | 13427     |
|               |               | c.2155G>A                          | p.G719S             | 6252      |
|               |               | c.2156G>A                          | p.G719D             | 18425     |
|               |               | c.2155G>T                          | p.G719C             | 6253      |
|               |               | c.2156G>C                          | p.G719A             | 6239      |
|               |               | c.2233_2247del15                   | p.K745_E749delKELRE | 26038     |
|               |               | c.2237_2255>T                      | p.E746_S752>V       | 12384     |
|               |               | c.2237_2250>TCCCT                  | p.E746_A750>VP      | 28623     |
|               |               | c.2235_2246del(12)<br>GGAATTAAGAGA | p.E746_E749del      | NA        |
|               |               | c.2235_2248>AATTC                  | p.E746_A750>IP      | 13550     |
|               |               | c.2237_2252>T                      | p.E746_T751>V       | 12386     |
|               |               | c.2238_2249>TCC                    | p.E746_A750>DP      | 18428     |
|               |               | c.2237_2238AA>TT                   | p.E746V             | 51497     |
|               |               | c.2237_2258>TTCA                   | p.E746_P753>VQ      | 51524     |
| c.2236_2251>T | p.E746_T751>S | 26513                              |                     |           |

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|-------------|------------------|-----------------------------------|-----------------------|------------|
|             |                  | CDS Mutation                      | AA Mutation           |            |
| <b>EGFR</b> | <b>NM_005228</b> | c.2235_2251>AG                    | p.E746_T751>A         | 13549      |
|             |                  | c.2235_2252del18                  | p.E746_T751delELREAT  | 24869      |
|             |                  | c.2236_2248>CAAC                  | p.E746_A750>QP        | 13557      |
|             |                  | c.2237_2257>TCT                   | p.E746_P753>VS        | 18427      |
|             |                  | c.2235_2255>AAT                   | p.E746_S752>I         | 12385      |
|             |                  | c.2237_2251>TGG                   | p.E746_T751>VA        | 53205      |
|             |                  | c.2238_2255del18                  | p.E746_S752>D         | 6220       |
|             |                  | c.2236_2256del21                  | p.E746_S752delELREATS | 133189     |
|             |                  | c.2236G>A                         | p.E746K               | 13184      |
|             |                  | c.2236_2253>CTA                   | p.E746_T751>L         | 133187     |
|             |                  | c.2235_2252>AAT                   | p.E746_T751>I         | 13551      |
|             |                  | c.2235_2246del12                  | p.E746_E749delELRE    | 28517      |
|             |                  | c.2235_2251>AATTC                 | p.E746_T751>IP        | 13552      |
|             |                  | c.2237_2251>TTC                   | p.E746_T751>VP        | 18421      |
|             |                  | c.2236_2253>CAA                   | p.E746_T751>Q         | 22999      |
|             |                  | c.2237_2254del18                  | p.E746_S752>A         | 12367      |
|             |                  | c.2236_2259>ATCTCG                | p.E746_P753>IS        | 133191     |
|             |                  | c.2235_2249del15                  | p.E746_A750delELREA   | 6223       |
|             |                  | c.2236_2248>AGAC                  | p.E746_A750>RP        | 12413      |
|             |                  | c.2236_2257>CTCT                  | p.E746_P753>LS        | 13200      |
|             |                  | c.2239_2258>CA                    | p.L747_P753>Q         | 12387      |
|             |                  | c.2239_2253>GCT                   | p.L747_T751>A         | 23572      |
|             |                  | c.2240_2261>CGAC                  | p.L747_K754>ST        | 20883      |
|             |                  | c.2238_2251>GC                    | p.L747_T751>P         | 22944      |
|             |                  | c.2240_2254del15/c.2239_2253del15 | p.L747_T751delLREAT   | 12369/6254 |
|             |                  | c.2238_2255>GCAACA                | p.L747_S752>QH        | 12421      |
|             |                  | c.2239_2247del(9)TTAAGAGAA        | p.L747_E749del        | 6218       |
|             |                  | c.2240T>C                         | p.L747S               | 26704      |
|             |                  | c.2240_2257del18                  | p.L747_P753>S         | 12370      |

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|------|--------------|-----------------------------------|------------------------|------------|
|      |              | CDS Mutation                      | AA Mutation            |            |
| EGFR | NM_005228    | c.2238_2248>GC                    | p.L747_A750>P          | 12422      |
|      |              | c.2238_2252>GCA                   | p.L747_T751>Q          | 12419      |
|      |              | c.2240_2251del12                  | p.L747_T751>S          | 6210       |
|      |              | c.2239_2262del24                  | p.L747_K754delLREATSPK | 24970      |
|      |              | c.2239_2240TT>CC                  | p.L747P                | 24267      |
|      |              | c.2238_2256>GCAA                  | p.L747_S752>Q          | 26441      |
|      |              | c.2239_2247delTTAAGAGAA           | p.L747_E749delLRE      | 6218       |
|      |              | c.2239_2253>CCAACG                | p.L747_T751>PT         | 1235296    |
|      |              | c.2239_2256del18                  | p.L747_S752delLREATS   | 6255       |
|      |              | c.2239_2264>GCCAA                 | p.L747_A755>AN         | 85891      |
|      |              | c.2251_2277>TCT                   | p.T751_I759>S          | 22945      |
|      |              | c.2252C>T                         | p.T751I                | 13185      |
|      |              | c.2252_2276>A                     | p.T751_I759>N          | 96856      |
|      |              | c.2252_2275del24                  | p.T751_E758delTSPKANKE | 133207     |
|      |              | c.2252_2277>GAGAAGCG              | p.T751_I759>REA        | 22956      |
|      |              | c.2254T>C                         | p.S752P                | 29274      |
|      |              | c.2253_2276del24/c.2254_2277del24 | p.S752_I759delSPKANKEI | 13556/6256 |
|      |              | c.2255C>A                         | p.S752Y                | 13186      |
|      |              | c.2257_2277del21                  | p.P753_I759del         | 24269      |
|      |              | c.2257C>T                         | p.P753S                | 6268       |
|      |              | c.2258C>A                         | p.P753Q                | NA         |
|      |              | c.2257_2277del21                  | p.P753_I759delIPKANKEI | 24269      |
|      |              | c.2276T>A                         | p.I759N                | 23633      |
|      |              | c.2281G>A                         | p.D761N                | 13188      |
|      |              | c.2281G>T                         | p.D761Y                | 21984      |
|      |              | c.2298_2299insGCCATA              | p.M766_A767insAI       | 13559      |
|      |              | c.2302_2303insCGCTGGCCA           | p.A767_S768insTLA      | 12425      |
|      |              | c.2303G>A                         | p.S768N                | 12989      |
|      |              | c.2303G>T                         | p.S768I                | 6241       |

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|------|--------------|---|---|-------------|
|      |              | CDS Mutation  | AA Mutation                               |             |
| EGFR | NM_005228    | c.2303G>C   | p.S768T                                   | 291998      |
|      |              | c.2303_2305GCG>TCT                                  | p.S768_V769>IL                            | 85750       |
|      |              | c.2308_2309insGCAGCGTGG                             | p.V769_D770insGSV                         | 18429       |
|      |              | c.2307_2308insTGCGTG                                | p.V769_D770insCV                          | 12379       |
|      |              | c.2308_2309insGGGTCGTGG                             | p.V769_D770insGVV                         | 18430       |
|      |              | c.2309_2310AC>CCAGCGTGGAT                           | p.V769_D770insASV                         | 13558       |
|      |              | c.2308_2309insGTT                                   | p.D770>GY                                 | 12427       |
|      |              | c.2311_2312insGCGTGGACA                             | p.D770_N771insSVD                         | 13428       |
|      |              | c.2310_2311insGGC                                   | p.D770_N771insG                           | 13004       |
|      |              | c.2308_2315GACAACCC><br>CCAGCGTGGATAACCG            | p.D770_P772>ASVDNR                        | NA          |
|      |              | c.2310_2311insGGGTTA                                | p.D770_N771insGL                          | 48921       |
|      |              | c.2310_2311insGCACCGTGG                             | p.D770_N771insAPW                         | 20886       |
|      |              | c.2310_2311insGGGTTT                                | p.D770_N771insGF                          | 655155      |
|      |              | c.2308G>A   | p.D770N                                   | 14068       |
|      |              | c.2309_2312ACAA>CTGGTGG                             | p.D770_N771>AGG                           | 12737       |
|      |              | c.2311_2312ins(12)TGCCACCCCA                        | p.D770_N771insMATP<br>(5' Detection Only) | 26719       |
|      |              | c.2310_2311insGGGGAC                                | p.D770_N771insGD                          | 85795       |
|      |              | c.2311_2312AA>GGGTT                                 | p.N771>GF                                 | 18431       |
|      |              | c.2311_2312insACCGGC                                | p.N771_P772insRH                          | 166390      |
|      |              | c.2312_2315ACCC>GCGTGGACAACCG                       | p.N771_P772>SVDNR                         | 13554       |
|      |              | c.2311_2312insGTC                                   | p.N771>SH                                 | 24434       |
|      |              | c.2311_2312insCAC                                   | p.N771>TH                                 | 22946       |
|      |              | c.2311A>GGTT  | p.N771>GY                                 | 53189       |
|      |              | c.2319_2320insCAC                                   | p.H773_V774insH                           | 12377       |
|      |              | c.2319_2320insAACCCCCAC/<br>c.2310_2311insAACCCCCAC | p.H773_V774insNPH                         | 12381/48920 |
|      |              | c.2319_2320insCAG                                   | p.H773_V774insQ                           | 131552      |

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|----------------------|-----------------------------------|--|------------------|-----------------------------|
|                      |                                   | CDS Mutation   | AA Mutation      |                             |
| <b>EGFR</b>          | NM_005228                         | c.2319_2320insCCCCAC   | p.H773_V774insPH | 12380                       |
|                      |                                   | c.2320G>A  | p.V774M          | 13006                       |
|                      |                                   | c.2316_2317insCACGTG   | p.V774_C775insHV | 133201                      |
|                      |                                   | c.2320G>T  | p.V774L          | 25090                       |
|                      |                                   | c.2326C>T  | p.R776C          | 6226                        |
|                      |                                   | c.2327G>A  | p.R776H          | 22940                       |
|                      |                                   | c.2369C>T  | p.T790M          | 6240                        |
|                      |                                   | c.2560A>G  | p.T854A          | 28537                       |
|                      |                                   | c.2572C>A  | p.L858M          | 12366                       |
|                      |                                   | c.2573T>G/c.2573_2574TG>GA/<br>c.2573_2574TG>GT/c.2572_2573CT>AG | p.L858R          | 6224/133630/<br>12429/13553 |
|                      |                                   | c.2572_2573CT>AA   | p.L858K          | 24268                       |
|                      |                                   | c.2582T>A  | p.L861Q          | 6212                        |
|                      |                                   | c.2582T>G  | p.L861R          | 12374                       |
|                      |                                   | c.2125_2129delGAAAC  | p.E709fs*1       | 24435                       |
|                      |                                   | c.2252_2275>G  | p.T751fs*4       | 12410                       |
| c.2317_2317C>AACCCCT | p.H773>NPY<br>(3' Detection Only) | NA   |                  |                             |
| <b>BRAF</b>          | NM_004333                         | c.1405_1406GG>TC   | p.G469S          | 458                         |
|                      |                                   | c.1406G>C  | p.G469A          | 460                         |
|                      |                                   | c.1406G>A  | p.G469E          | 461                         |
|                      |                                   | c.1405G>C  | p.G469R          | 455                         |
|                      |                                   | c.1406G>T  | p.G469V          | 459                         |
|                      |                                   | c.1781A>G  | p.D594G          | 467                         |
|                      |                                   | c.1781A>T  | p.D594V          | 466                         |
|                      |                                   | c.1789_1790CT>TC   | p.L597S          | 1126                        |
|                      |                                   | c.1790T>A  | p.L597Q          | 1125                        |
|                      |                                   | c.1789C>G  | p.L597V          | 470                         |
|                      |                                   | c.1790T>G  | p.L597R          | 471                         |

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|-------------|--------------|----------------------------|--|--------------|
|             |              | CDS Mutation               | AA Mutation                                |              |
| <b>BRAF</b> | NM_004333    | c.1797_1798insACA          | p.T599_V600insT                            | 144982       |
|             |              | c.1797_1797A>TACTACG       | p.T599_V600insTT                           | 1128         |
|             |              | c.1799T>G                  | p.V600G                                    | 6137         |
|             |              | c.1798_1799GT>AA           | p.V600K                                    | 473          |
|             |              | c.1798_1799GT>AG           | p.V600R                                    | 474          |
|             |              | c.1798G>C/c.1798G>T        | p.V600L                                    | 219798/33808 |
|             |              | c.1798G>A                  | p.V600M                                    | 1130         |
|             |              | c.1799T>A/c.1799_1800TG>AA | p.V600E<br>(raremutation=c.1799_1800TG>AA) | 476/475      |
|             |              | c.1798_1798G>TACA          | p.V600>YM                                  | 1159850      |
|             |              | c.1799_1800TG>AT           | p.V600D                                    | 477          |
|             |              | c.1801A>G                  | p.K601E                                    | 478          |
|             |              | c.1801_1803delAAA          | p.K601del                                  | 30594        |
|             |              | c.1803A>T/c.1803A>C        | p.K601N                                    | 6265/1132    |
| <b>KRAS</b> | NM_004985    | c.34_36GGT>TGG             | p.G12W                                     | 36281        |
|             |              | c.34_36GGT>TGC             | p.G12C                                     | 513          |
|             |              | c.35_36GT>AA               | p.G12E                                     | 519          |
|             |              | c.34_35GG>TA               | p.G12Y                                     | 25081        |
|             |              | c.35G>A                    | p.G12D                                     | 521          |
|             |              | c.34_35GG>TT               | p.G12F                                     | 512          |
|             |              | c34-36GGT>AGA/c.34G>C      | p.G12R                                     | 249888/518   |
|             |              | c.34_35GG>AA               | p.G12N                                     | 13643        |
|             |              | c.36T>A                    | p.G12G                                     | 524          |
|             |              | c.34G>A                    | p.G12S                                     | 517          |
|             |              | c.35G>C                    | p.G12A                                     | 522          |
|             |              | c34-35GG>AC                | p.G12T                                     | NA           |
|             |              | c.35G>T                    | p.G12V                                     | 520          |

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|-------------|------------------|----------------------|---------------|-----------|
|             |                  | CDS Mutation         | AA Mutation   |           |
| <b>KRAS</b> | <b>NM_004985</b> | c.34_35GG>AT         | p.G12I        | 34144     |
|             |                  | c34-35GG>CC          | p.G12P        | NA        |
|             |                  | c.36_37insGCG        | p.G12_G13insA | 303833    |
|             |                  | c.34_35GG>CT         | p.G12L        | 514       |
|             |                  | c.38G>C              | p.G13A        | 533       |
|             |                  | c.38G>A              | p.G13D        | 532       |
|             |                  | c.37G>C              | p.G13R        | 529       |
|             |                  | c.37_38GG>AA         | p.G13N        | 53283     |
|             |                  | c.37G>A              | p.G13S        | 528       |
|             |                  | c.38G>T              | p.G13V        | 534       |
|             |                  | c.37_38GG>AT         | p.G13I        | 525       |
|             |                  | c.39_40insGGC        | p.G13_V14insG | 219781    |
|             |                  | c.36_37TG>AT/c.37G>T | p.G13C        | 87281/527 |
|             |                  | c.175G>A             | p.A59T        | 546       |
|             |                  | c.181C>A             | p.Q61K        | 549       |
|             |                  | c.181C>G             | p.Q61E        | 550       |
|             |                  | c.182A>G             | p.Q61R        | 552       |
|             |                  | c.183A>C             | p.Q61H        | 554       |
|             |                  | c.182A>C             | p.Q61P        | 551       |
|             |                  | c.182A>T             | p.Q61L        | 553       |
|             |                  | c.437C>G             | p.A146G       | N/A       |
|             |                  | c.436G>A             | p.A146T       | 19404     |
|             |                  | c.437C>T             | p.A146V       | 19900     |
|             |                  | c.436G>C             | p.A146P       | 19905     |
|             |                  | c.176C>G             | p.A59G        | 28518     |
|             |                  | c.176C>A             | p.A59E        | 547       |
|             |                  | c.349A>G             | p.K117E       | N/A       |
|             |                  | c.351A>T             | p.K117N       | 28519     |
|             |                  | c.350A>G             | p.K117R       | 4696722   |

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|             |              | CDS Mutation         | AA Mutation |           |
| <b>NRAS</b> | NM_002524    | c.34G>T              | p.G12C      | 562       |
|             |              | c.34_35GG>TA         | p.G12Y      | 560       |
|             |              | c.34_35GG>AA         | p.G12N      | 12723     |
|             |              | c.35G>C              | p.G12A      | 565       |
|             |              | c.35_36GT>AG         | p.G12E      | 144577    |
|             |              | c35G>T               | p.G12V      | 566       |
|             |              | c.34_35GG>CC         | p.G12P      | 559       |
|             |              | c.38G>C              | p.G13A      | 575       |
|             |              | c.38G>A              | p.G13D      | 573       |
|             |              | c.37G>C              | p.G13R      | 569       |
|             |              | c.38G>T/c.38_39GT>TC | p.G13V      | 574/572   |
|             |              | c.37_38GG>AA         | p.G13N      | 24668     |
|             |              | c.37_38GG>TA         | p.G13Y      | 568       |
|             |              | c.181C>A             | p.Q61K      | 580       |
|             |              | c.182_183AA>GG       | p.Q61R      | 33693     |
|             |              | c.183A>C             | p.Q61H      | 586       |
|             |              | c.182A>C             | p.Q61P      | 582       |
|             |              | c.183A>G             | p.Q61Q      | 587       |
|             |              | c.181C>G             | p.Q61E      | 581       |
|             |              | c.182_183AA>TG       | p.Q61L      | 30646     |
|             |              | c.176C>G             | p.A59G      | 5878737   |
|             |              | c.175G>A             | p.A59T      | 578       |
|             |              | c.349A>G             | p.K117E     | N/A       |
|             |              | c.351G>T/C           | p.K117N     | N/A       |
|             |              | c.350A>G             | p.K117R     | N/A       |
|             |              | c.436G>A             | p.A146T     | 27174     |
|             |              | c.436G>C             | p.A146P     | N/A       |
|             |              | c.436G>T             | p.A146S     | N/A       |
|             |              | c.437C>T             | p.A146V     | 4170228   |
|             |              | c.437C>G             | p.A146G     | N/A       |

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|------------|--------------|-------------------|-------------|-----------|
|            |              | CDS Mutation      | AA Mutation |           |
| <i>KIT</i> | NM_000222    | c.1669T>C         | p.W557R     | 1219      |
|            |              | c.1669T>A         | p.V559A     | 1216      |
|            |              | c.1676T>C         | p.V559D     | 1255      |
|            |              | c.1676T>A         | p.L576P     | 1252      |
|            |              | c.1727T>C         | p.K642E     | 1290      |
|            |              | c.1924A>G         | p.D816H     | 1304      |
|            |              | c.2446G>C         | p.W557G     | 1311      |
|            |              | c.1676T>G         | p.V559G     | 1253      |
|            |              | c.1924A>C         | p.K642Q     | 96871     |
|            |              | c.2446G>T         | p.D816Y     | 1310      |

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