



## List of Mutations Detectable with the UltraSEEK™ Lung Panel

| GENE                                 | NCBI Ref Seq  | HGVS Nomenclature                    |                | COSMIC ID |
|--------------------------------------|---------------|--------------------------------------|----------------|-----------|
|                                      |               | CDS Mutation                         | AA Mutation    |           |
| <b>BRAF</b>                          | NM_004333     | c.1406G>C                            | p.G469A        | 460       |
|                                      |               | c.1406G>T                            | p.G469V        | 459       |
|                                      |               | c.1781A>G                            | p.D594G        | 467       |
|                                      |               | c.1799T>A                            | p.V600E        | 476       |
| <b>EGFR</b>                          | NM_005228     | c.2125G>A                            | p.E709K        | 12988     |
|                                      |               | c.2126A>C                            | p.E709A        | 13427     |
|                                      |               | c.2126A>G                            | p.E709G        | 13009     |
|                                      |               | c.2126A>T                            | p.E709V        | 12371     |
|                                      |               | c.2155G>T                            | p.G719C        | 6253      |
|                                      |               | c.2155G>A                            | p.G719S        | 6252      |
|                                      |               | c.2156G>C                            | p.G719A        | 6239      |
|                                      |               | c.2233_2247del(15)AAGGAATTAAGAGAA    | p.K745_E749del | 26038     |
|                                      |               | c.2235_2249del(15)GGAATTAAGAGAAGC    | p.E746_A750del | 6223      |
|                                      |               | c.2235_2248>AATTC                    | p.E746_A750>IP | 13550     |
|                                      |               | c.2235_2251>AATTC                    | p.E746_T751>IP | 13552     |
|                                      |               | c.2236_2250del(15)GAATTAAGAGAAGCA    | p.E746_A750del | 6225      |
|                                      |               | c.2236_2253del(18)GAATTAAGAGAAGCAACA | p.E746_T751del | 12728     |
|                                      |               | c.2237_2251del(15)AATTAAGAGAAGCAA    | p.E746_T751>A  | 12678     |
|                                      |               | c.2237_2254del(18)AATTAAGAGAAGCAACAT | p.E746_S752>A  | 12367     |
|                                      |               | c.2237_2252>T                        | p.E746_T751>V  | 12386     |
|                                      |               | c.2237_2253>TTGCT                    | p.E746_T751>VA | 12416     |
|                                      |               | c.2237_2253>TTCCT                    | p.E746_T751>VP | 52935     |
|                                      |               | c.2237_2257>TCT                      | p.E746_P753>VS | 18427     |
|                                      |               | c.2237_2255>T                        | p.E746_S752>V  | 12384     |
| c.2238_2255del(18)ATTAAGAGAAGCAACATC | p.E746_S752>D | 6220                                 |                |           |

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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.2239_2247del(9)TTAAGAGAA           | p.L747_E749del    | 6218      |
|      |              | c.2239_2256del(18)TTAAGAGAAGCAACATCT | p.L747_S752del    | 6255      |
|      |              | c.2239_2256>CAA                      | p.L747_S752>Q     | 12403     |
|      |              | c.2239_2248TTAAGAGAAG>C              | p.L747_A750>P     | 12382     |
|      |              | c.2239_2251TTAAGAGAAGCAA>C           | p.L747_T751>P     | 12383     |
|      |              | c.2239_2258>CA                       | p.L747_P753>Q     | 12387     |
|      |              | c.2240_2251del(12)TAAGAGAAGCAA       | p.L747_T751>S     | 6210      |
|      |              | c.2240_2254del(15)TAAGAGAAGCAACAT    | p.L747_T751del    | 12369     |
|      |              | c.2240_2257del(18)TAAGAGAAGCAACATCTC | p.L747_P753>S     | 12370     |
|      |              | c.2303G>T                            | p.S768I           | 6241      |
|      |              | c.2307_2308ins(9)GCCAGCGTG           | p.V769_D770insASV | 12376     |
|      |              | c.2308_2309ins(9)CCAGCGTGG           | p.V769_D770insASV | 12426     |
|      |              | c.2309_2310AC>CCAGCGTGGAT            | p.V769_D770insASV | 13558     |
|      |              | c.2310_2311insGGT                    | p.D770-N771insG   | 12378     |
|      |              | c.2311_2312ins(9)GCGTGGACA           | p.D770_N771insSVD | 13428     |
|      |              | c.2319_2320ins(9)AACCCCCAC           | p.H773_V774insNPH | 12381     |
|      |              | c.2319_2320insCAC                    | p.H773-V774insH   | 12377     |
|      |              | c.2369C>T                            | p.T790M           | 6240      |
|      |              | c.2389T>A                            | p.C797S           | N/A       |
|      |              | c.2390G>C                            | p.C797S           | 5945664   |
|      |              | c.2573T>G                            | p.L858R           | 6224      |
|      |              | c.2582T>A                            | p.L861Q           | 6213      |
|      |              | c.2582T>G                            | p.L861R           | 12374     |

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|---------------|--------------|--------------------------------|--------------------|-----------|
|               |              | CDS Mutation                   | AA Mutation        |           |
| <i>ERBB2</i>  | NM_004448    | c.2324_2325ins(12)ATACGTGATGGC | p.A775_G776insYVMA | 20959     |
|               |              | c.2325_2326ins(12)TACGTGATGGCT | p.A775_G776insYVMA | 12558     |
|               |              | c.2326_2327ins(3)TGT           | p.G776>VC          | 12553     |
|               |              | c.2326_2327ins(3)TTT           | p.G776>VC          | 12552     |
| <i>KRAS</i>   | NM_004985    | c.34G>A                        | p.G12S             | 517       |
|               |              | c.34G>C                        | p.G12R             | 518       |
|               |              | c.34G>T                        | p.G12C             | 516       |
|               |              | c.35G>A                        | p.G12D             | 521       |
|               |              | c.35G>C                        | p.G12A             | 522       |
|               |              | c.35G>T                        | p.G12V             | 520       |
|               |              | c.37G>T                        | p.G13C             | 527       |
|               |              | c.38G>A                        | p.G13D             | 532       |
|               |              | c.183A>C                       | p.Q61H             | 554       |
|               |              | c.183A>T                       | p.Q61H             | 555       |
|               |              | c.181C>A                       | p.Q61K             | 549       |
|               |              | c.181C>G                       | p.Q61E             | 550       |
|               |              | c.182A>C                       | p.Q61P             | 551       |
|               |              | c.182A>G                       | p.Q61R             | 552       |
| c.182A>T      | p.Q61L       | 553                            |                    |           |
| <i>PIK3CA</i> | NM_006218    | c.1624G>A                      | p.E542K            | 760       |
|               |              | c.1633G>A                      | p.E545K            | 763       |
|               |              | c.3140A>G                      | p.H1047R           | 775       |
|               |              | c.3140A>T                      | p.H1047L           | 776       |

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