

>EL-010 EGFR (A767_dupASV)

ATGCGACCCTCCGGGACGGCCGGGGCAGCGCTCCTGGCGCTGCTGGCTGCGCTCTGCCCCG
GCGAGTCGGGCTCTGGAGGAAAAGAAAGTTTGCCAAGGCACGAGTAACAAGCTCACGCAGTT
GGGCACTTTTGAAGATCATTTTCTCAGCCTCCAGAGGATGTTCAATAACTGTGAGGTGGTCCTTG
GGAATTTGGAAATTACCTATGTGCAGAGGAATTATGATCTTTCCTTCTTAAAGACCATCCAGGAGGT
GGCTGGTTATGTCCCTCATTGCCCTCAACACAGTGGAGCGAATTCCTTTGAAAAACCTGCAGATC
ATCAGAGGAAATATGTACTACGAAAATTCCTATGCCTTAGCAGTCTTATCTAACTATGATGCAAATAA
AACCGGACTGAAGGAGCTGCCCATGAGAAATTTACAGGAAATCCTGCATGGCGCCGTGCGGTT
CAGCAACAACCCTGCCCTGTGCAACGTGGAGAGCATCCAGTGGCGGGACATAGTCAGCAGTG
ACTTTCTCAGCAACATGTGCATGGACTTCCAGAACCACCTGGGCAGCTGCCAAAAGTGTGATCC
AAGCTGTCCCAATGGGAGCTGCTGGGGTGCAGGAGAGGAGAACTGCCAGAACTGACCAAAA
TCATCTGTGCCCAGCAGTGCTCCGGGCGCTGCCGTGGCAAGTCCCCCAGTGACTGCTGCCAC
AACCAGTGTGCTGCAGGCTGCACAGGCCCCCCGGGAGAGCGACTGCCTGGTCTGCCGCAAT
TCCGAGACGAAGCCACGTGCAAGGACACCTGCCCCCCTCATGCTCTACAACCCACCAC
GTACCAGATGGATGTGAACCCCGAGGGCAAATACAGCTTTGGTGCCACCTGCGTGAAGAAGTG
TCCCCGTAATTATGTGGTGACAGATCACGGCTCGTGCGTCCGAGCCTGTGGGGCCGACAGCTA
TGAGATGGAGGAAGACGGCGTCCGCAAGTGTAAGAAGTGCGAAGGGCCTTGCCGCAAAGTGT
GTAACGGAATAGGTATTGGTGAATTTAAAGACTCACTCTCCATAAATGCTACGAATATTAAACACTT
CAAAAACCTGCACCTCCATCAGTGGCGATCTCCACATCCTGCCGGTGGCATTAGGGGTGACTC
CTTACACATACTCCTCCTCTGGATCCACAGGAACTGGATATTCTGAAAACCGTAAAGGAAATCA
CAGGGTTTTTGCTGATTGAGGCTTGGCCTGAAAACAGGACGGACCTCCATGCCTTTGAGAACCT
AGAAATCATACGCGGCAGGACCAAGCAACATGGTCAGTTTTCTCTTGCAAGTCGTCAGCCTGAAC
ATAACATCCTTGGGATTACGCTCCCTCAAGGAGATAAGTGATGGAGATGTGATAATTCAGGAAAC
AAAAATTTGTGCTATGCAAATACAATAAACTGGAAAAAACTGTTTGGGACCTCCGGTCAGAAAACC
AAAATTATAAGCAACAGAGGTGAAAACAGCTGCAAGGCCACAGGCCAGGTCTGCCATGCCTTG
TGCTCCCCCGAGGGGCTGCTGGGGCCCGGAGCCCAGGGACTGCGTCTCTTGCCGGAATGTCA
GCCGAGGCAGGGAATGCGTGGACAAGTGCAACCTTCTGGAGGGTGAGCCAAGGGAGTTTGTG
GAGAACTCTGAGTGCATACAGTGCCACCCAGAGTGCTGCCTCAGGCCATGAACATCACCTGC
ACAGGACGGGGACCAGACAACTGTATCCAGTGTGCCCACTACATTGACGGCCCCCACTGCGT
CAAGACCTGCCCAGGAGTGCATGGGAGAAAACAACACCCTGGTCTGGAAGTACGCAGACG
CCGGCCATGTGTGCCACCTGTGCCATCCAACTGCACCTACGGATGCACTGGGCCAGGTCTT
GAAGGCTGTCCAACGAATGGGCCTAAGATCCCGTCCATCGCCACTGGGATGGTGGGGGCCCT
CCTCTTGCTGCTGGTGGTGGCCCTGGGGATCGGCCTCTTCATGCGAAGGCGCCACATCGTTC
GGAAGCGCACGCTGCGGAGGCTGCTGCAGGAGAGGGAGCTTGTGGAGCCTCTTACACCCAG
TGGAGAAGCTCCCAACCAAGCTCTCTTGAGGATCTTGAAGGAACTGAATTCAAAAGATCAAA
GTGCTGGGCTCCGGTGCGTTCGGCACGGTGTATAAGGGACTCTGGATCCCAGAAGGTGAGAAA
GTTAAAATTCCCGTCGCTATCAAGGAATTAAGAGAAGCAACATCTCGAAAGCCAACAAGGAAAT

CCTCGATGAAGCCTACGTGATG**GCCAGCGTGGCCAGCGTG**GACAACCCCCACGTGTGCCGC
CTGCTGGGCATCTGCCTCACCTCCACCGTGCAGCTCATCACGCAGCTCATGCCCTTCGGCTG
CCTCCTGGACTATGTCCGGGAACACAAAGACAATATTGGCTCCCAGTACCTGCTCAACTGGTGT
GTGCAGATCGCAAAGGGCATGAACTACTTGGAGGACCGTCGCTTGGTGCACCGCGACCTGGC
AGCCAGGAACGTACTGGTGAAAACACCGCAGCATGTCAAGATCACAGATTTTGGGCTGGCCAA
ACTGCTGGGTGCGGAAGAGAAAGAATACCATGCAGAAGGAGGCAAAGTGCCTATCAAGTGGAT
GGCATTGGAATCAATTTTACACAGAATCTATACCCACCAGAGTGATGTCTGGAGCTACGGGGTGA
CTGTTTGGGAGTTGATGACCTTTGGATCCAAGCCATATGACGGAATCCCTGCCAGCGAGATCTC
CTCCATCCTGGAGAAAGGAGAACGCCTCCCTCAGCCACCCATATGTACCATCGATGTCTACATG
ATCATGGTCAAGTGCTGGATGATAGACGCAGATAGTCGCCCCAAAGTTCCGTGAGTTGATCATCGA
ATTCTCCAAAATGGCCCGAGACCCCCAGCGCTACCTTGTCATTCAGGGGGATGAAAGAATGCA
TTTGCCAAGTCCTACAGACTCCAATTCTACCGTGCCCTGATGGATGAAGAAGACATGGACGAC
GTGGTGGATGCCGACGAGTACCTCATCCACAGCAGGGCTTCTTCAGCAGCCCCTCCACGTC
ACGGACTCCCCTCCTGAGCTCTCTGAGTGCAACCAGCAACAATTCCACCGTGGCTTGCATTGAT
AGAAATGGGCTGCAAAGCTGTCCCATCAAGGAAGACAGCTTCTTGCAGCGATACAGCTCAGAC
CCCACAGGCGCCTTGACTGAGGACAGCATAGACGACACCTTCCTCCCAGTGCCTGAATACATA
AACCAGTCCGTTCCCAAAGGCCCGCTGGCTCTGTGCAGAATCCTGTCTATCACAAATCAGCCT
CTGAACCCCGCACCCAGCAGAGACCCACACTACCAGGACCCCCACAGCACTGCAGTGGGC
AACCCCGAGTATCTCAAACTGTCCAGCCCACCTGTGTCAACAGCACATTTCGACAGCCCTGCC
CACTGGGCCCCAGAAAGGCAGCCACCAAATTAGCCTGGACAACCCTGACTACCAGCAGGACTT
CTTTCCCAAGGAAGCCAAGCCAAATGGCATCTTTAAGGGCTCCACAGCTGAAAATGCAGAATAC
CTAAGGGTCGCGCCACAAAGCAGTGAATTTATTGGAGCA