CEBPE (NM_001805.4) - cDNA - 2025-12-07

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ACACAGGAGT GGGTGACAGA GGAGACTGCA GAGGGCAGGT CTAGGGCAGA -115
AGATCGAGAG AGGGCAGGCC CAGGTCAGGA GGAGGTAGAG AGAGGGCAGC -65
CGGAGCACCC CAAGGGGTGC CTCAAGAGCA GGTGGGGGCG GGGAGCCGAG -15
GGGGCGGGCC GGCCATGTCC CACGGGACCT ACTACGAGTG TGAGCCCCGG 36
GGTGGCCAGC AGCCACTCGA GTTCTCAGGG GGCCGAGCTG GGCCCGGGGA 86
GCTAGGGGAC ATGTGTGAGC ATGAGGCCTC CATTGACCTC TCCGCCTACA 136
TCGAGTCTGG GGAAGAGCAG CTTCTCTCCG ATCTCTTTGC CGTGAAGCCA 186
GCGCCTGAGG CCAGAGGCCT CAAGGGCCCC GGAACCCCTG CCTTCCCCCA 236
CTACTTGCCG CCTGACCCTC GGCCCTTTGC CTACCCTCCA CATACCTTCG 286   delTGACC
GCCCAGACAG GAAGGCGCTG GGGCCTGGCA TCTACAGCAG CCCAGGGAGC 336
TACGACCCCA GGGCTGTGGC GGTGAAGGAG GAGCCCCGGG GGCCAGAGGG 386
CAGCCGAGCT GCCAGCCGAG GCAGCTACAA TCCCCTGCAG TACCAAGTGG 436
CACACTGTGG GCAGACAGCC ATGCACCTGC CCCCAACTCT GGCAGCACCC 486
GGCCAGCCTC TGCGCGTTCT CAAGGCCCCT TTGGCCACTG CCGCACCCCC 536   c.508_509insA
CTGCAGTCCC CTCCTGAAGG CGCCCTCCCC GGCTGGCCCC TTACACAAGG 586
GCAAGAAGGC AGTGAACAAA GATAGCCTTG AGTACCGGCT GAGGCGGGAG 636
CGCAACAACA TCGCCGTGCG CAAGAGCCGA GACAAGGCCA AGAGGCGCAT 686   V218A R219H
TCTGGAGACG CAGCAGAAGG TGCTGGAGTA CATGGCAGAG AACGAGCGCC 736
TCCGCAGCCG CGTGGAGCAG CTCACCCAGG AGCTAGACAC CCTCCGCAAC 786   c.739_744delCGCAGC
CTCTTCCGCC AGATTCCTGA GGCGGCCAAC CTCATCAAGG GCGTGGGGGG 836
TTGCAGCTGA GGCTGGCTGG TGGATTGTGG GCACCAGGCT CCCTGGCACG *40
GCCTAACTCT GCGGACCCCC ATCCTGCTGG GGGCCTAGAA CCCTGAGACA *90
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TAGACCATGG ATAAATGGCA ACCGGGGTGG CAAAGAGGGC AGGACCCAGC *140

ATAATGATTA TATGGCTGAA TAAAGTTGCA CTGTGACTGG G

CEBPE (NM_001805.4) - cDNA - 2025-12-07

nfevers