



CEBPE (NM_001805.4) - cDNA - 2025-04-03

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 CATACTTCG 286 delTGACC
GCCCAGACAG GAAGGCGCTG GGGCCTGGCA TCTACAGCAG CCCAGGGAGC 336
TACGACCCCA GGGCTGTGGC GGTGAAGGAG GAGCCCCGGG GGCCAGAGGG 386
CAGCCGAGCT GCCAGCCGAG GCAGCTACAA TCCCCGTCAG TACCAAGTGG 436
CACACTGTGG GCAGACAGCC ATGCACCTGC CCCCAACTCT GGCAGCACCC 486
GGCCAGCCTC TGC GCGTTCT CAAGGCCCT TTGGCCACTG CCGCACCCCC 536 c.508_509insA
CTGCAATCCC CTCCTGAAGG CGCCCTCCCC GGCTGGCCCC TTACACAAGG 586
GCAAGAAGGC AGTGAACAAA GATAGCCTTG AGTACCGGCT GAGGCGGGAG 636
CGCAACAACA TCGCCGTGCG CAAGAGCCGA GACAAGGCCA AGAGGCGCAT 686 Y218A R219H
TCTGGAGACG CAGCAGAAGG TGCTGGAGTA CATGGCAGAG AACGAGCGCC 736
TCCGCAGCCG CGTGGAGCAG CTCACCCAGG AGCTAGACAC CCTCCGCAAC 786 c.739_744delCGCAGC
CTCTTCCGCC AGATTCTGA GGC GGCCAAC CTCATCAAGG GCGTGGGGGG 836
TTGCAGCTGA GGCTGGCTGG TGGATTGTGG GCACCAGGCT CCCTGGCACG *40
GCCTAACTCT GCGGACCCCC ATCCTGCTGG GGGCCTAGAA CCCTGAGACA *90

TAGACCATGG ATAAATGGCA ACCGGGGTGG CAAAGAGGGC AGGACCCAGC *140

ATAATGATTA TATGGCTGAA TAAAGTTGCA CTGTGACTGG G

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