



STING1 (NM_198282.4) - cDNA - 2025-07-19

GTTCATTTT CACTCCTCCC TCCTAGGTCA CACTTTCA GAAAAGAAC -253
TGCATCCTGG AAACCAGAAG AAAATATGA GACGGGAAAT CATCGTGTGA -203
TGTGTGTGCT GCCTTGCT GAGTGTGTGG AGTCCTGCTC AGGTGTTAGG -153
TACAGTGTGT TTGATCGTGG TGGCTTGAGG GGAACCCGCT GTTCAGAGCT -103
GTGACTGCGG CTGCACTCAG AGAAGCTGCC CTTGGCTGCT CGTAGCGCCG -53
GGCCTTCTCT CCTCGTCATC ATCCAGAGCA GCCAGTGTCC GGGAGGCAGA -3
AGATGCCCA CTCCAGCCTG CATCCATCCA TCCCGTGTCC CAGGGGTAC 48
GGGGCCCAGA AGGCAGCCTT GGTTCTGCTG AGTGCCTGCC TGGTGACCT 98
TTGGGGCTA GGAGAGCCAC CAGAGCACAC TCTCCGGTAC CTGGTGTCTCC 148
ACCTAGCCTC CCTGCAGCTG GGACTGCTGT TAAACGGGGT CTGCAGCCTG 198 L189V
GCTGAGGAGC TGCGCACAT CCACTCCAGG TACCGGGCA GCTACTGGAG 248 H72N
GACTGTGCGG GCCTGCCCTGG GCTGCCCTCT CCGCCGTGGG GCCCTGTTGC 298 R94H
TGCTGTCCAT CTATTCTAC TACTCCCTCC CAAATGCGGT CGGCCCGCCC 348
TTCACTTGGA TGCTTGCCCT CCTGGGCCTC TCGCAGGCAC TGAACATCCT 398
CCTGGGCCTC AAGGGCCTGG CCCCAGCTGA GATCTCTGCA GTGTGTGAAA 448 V147L V147M
AAGGAATT CAACGTGGCC CATGGGCTGG CATGGTCATA TTACATCGGA 498 F153V F153I N154S V155M G158A G166E
TATCTGCGGC TGATCCTGCC AGAGCTCCAG GCCCGGATTC GAACTTACAA 548
TCAGCATTAC AACAAACCTGC TACGGGGTGC AGTGAGCCAG CGGCTGTATA 598
TTCTCCTCCC ATTGGACTTGT GGGGTGCCTG ATAACCTGAG TATGGCTGAC 648 C206G C206Y G207E
CCCAACATTC GCTTCCTGGA TAAACTGCC CAGCAGACCG GTGACCATGC 698
TGGCATCAAG GATCGGGTTT ACAGCAACAG CATCTATGAG CTTCTGGAGA 748
ACGGGCAGCG GGCGGGCACC TGTGTCCCTGG AGTACGCCAC CCCCTTGAG 798

ACTTTGTGTTG CCATGTCACA ATACAGTCAA GCTGGC~~TTTA~~CCGGGAGGA 848 F279L S280R R281W R281Q
~~TAGG~~CCTTGAG CAGGCCAAC TCTTCTGCCG GACACTTGAG GACATCCTGG 898 R284G R284S
CAGATGCCCC TGAGTCTCAG AACAACTGCC GCCTCATTGC CTACCAGGAA 948
CCTGCAGATG ACAGCAGCTT CTCGCTGTCC CAGGAGGTTC TCCGGCACCT 998
GCGGCAGGAG GAAA~~AGGAAG~~ AGGTTACTGT GGGCAGCTG AAGACCTCAG 1048 K338Rfs*9
CGGTGCCAG TACCTCCACG ATGTCCAAG AGCCTGAGCT CCTCATCAGT 1098
GGAATGGAAA AGCCCCCTCCC TCTCCGCACG GATTCTCTT ~~GA~~GACCCAGG *8
GTCACCAGGC CAGAGCCTCC AGTGGTCTCC AAGCCTCTGG ACTGGGGGCT *58
CTCTTCAGTG GCTGAATGTC CAGCAGAGCT ATTTCCCTCC ACAGGGGGCC *108
TTGCAGGGAA GGGTCCAGGA CTTGACATCT TAAGATGCGT CTTGTCCCT *158
TGGGCCAGTC ATTTCCCTC TCTGAGCCTC GGTGTCTCA ACCTGTGAAA *208
TGGGATCATA ATCACTGCCT TACCTCCCTC ACGGTTGTTG TGAGGACTGA *258
GTGTGTGGAA GTTTTCATA AACTTTGGAT GCTAGTGTAC TTAGGGGGTG *308
TGCCAGGTGT CTTTCATGGG GCCTTCCAGA CCCACTCCCC ACCCTTCTCC *358
CCTTCCTTG CCCGGGACG CGAACTCTC TCAATGGTAT CAACAGGCTC *408
CTTCGCCCTC TGGCTCCTGG TCATGTTCCA TTATTGGGA GCCCCAGCAG *458
AAGAATGGAG AGGAGGAGGA GGCTGAGTT GGGGTATTGA ATCCCCGGC *508
TCCCACCCCTG CAGCATCAAG GTTGTATGG ACTCTCCTGC CGGGCAACTC *558
TTGCGTAATC ATGACTATCT CTAGGATTCT GGCACCACTT CCTTCCCTGG *608
CCCCCTTAAGC CTAGCTGTGT ATCGGCACCC CCACCCCCACT AGAGTACTCC *658
CTCTCACTTG CGGTTCCCTT ATACTCCACC CCTTTCTCAA CGGTCCCTTT *708
TTAAAGCACA TCTCAGATTA

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