



STING1 (NM_198282.4) - cDNA - 2024-05-18

GTTCAATTTT CACTCCTCCC TCCTAGGTCA CACTTTTCAG AAAAGAATC -253
TGCATCCTGG AAACCAGAAG AAAAATATGA GACGGGGAAT CATCGTGTGA -203
TGTGTGTGCT GCCTTTGGCT GAGTGTGTGG AGTCCTGCTC AGGTGTTAGG -153
TACAGTGTGT TTGATCGTGG TGGCTTGAGG GGAACCCGCT GTTCAGAGCT -103
GTGACTGCGG CTGCACTCAG AGAAGCTGCC CTTGGCTGCT CGTAGCGCCG -53
GGCCTTCTCT CCTCGTCATC ATCCAGAGCA GCCAGTGTCC GGGAGGCAGA -3
AGATGCCCCA CTCCAGCCTG CATCCATCCA TCCCGTGTCC CAGGGGTCAC 48
GGGGCCCAGA AGGCAGCCTT GGTTCGTGCTG AGTGCCCTGCC TGGTGACCCT 98
TTGGGGGCTA GGAGAGCCAC CAGAGCACAC TCTCCGGTAC CTGGTGCTCC 148
ACCTAGCCTC CCTGCAGCTG GGAAGCTGTG TAAACGGGGT CTGCAGCCTG 198 L189V
GCTGAGGAGC TGCGCACAT CCACTCCAGG TACCGGGGCA GCTACTGGAG 248 H72N
GACTGTGCGG GCCTGCCTGG GCTGCCCCCT CCGCCGTGGG GCCCTGTTGC 298 R94H
TGCTGTCCAT CTATTTCTAC TACTCCCTCC CAAATGCGGT CGGCCCCGCC 348
TTCAC TTGGA TGCTTGCCCT CCTGGGCCTC TCGCAGGCAC TGAACATCCT 398
CCTGGGCCTC AAGGGCCTGG CCCAGCTGA GATCTCTGCA GTGTGTGAAA 448 V147L V147M
AAGGGAATT CAACGTGGCC CATGGCTGG CATGGTCATA TTACATCGGA 498 F153V F153I N154S V155M G158A G166E
TATCTGCGGC TGATCCTGCC AGAGCTCCAG GCCCGGATTG GAACTTACAA 548
TCAGCATTAC AACAACCTGC TACGGGGTGC AGTGAGCCAG CGGCTGTATA 598
TTCTCCTCCC ATTGGACTGT GGGGTGCTGT ATAACCTGAG TATGGCTGAC 648 C206G C206Y G207E
CCCAACATTC GCTTCCTGGA TAAACTGCCC CAGCAGACCG GTGACCATGC 698
TGGCATCAAG GATCGGGTTT ACAGCAACAG CATCTATGAG CTTCTGGAGA 748
ACGGGCAGCG GCGGGGCACC TGTGTCTCTGG AGTACGCCAC CCCCTTGACG 798

ACTTTGTTTG CCATGTCACA ATACAGTCAA GCTGGC**TTA** **GC****CGGGAGGA** 848 **F279L S280R R281W R281Q**
TAGGCTTGAG CAGGCCAAAC TCTTCTGCCG GACACTTGAG GACATCCTGG 898 **R284G R284S**
CAGATGCCCC TGAGTCTCAG AACAACTGCC GCCTCATTGC CTACCAGG**AA** 948
CCTGCAGATG ACAGCAGCTT CTCGCTGTCC CAGGAGGTTT TCCGGGCACCT 998
GCGGCAGGAG GAAA**A**GAAG AGGTTACTGT GGGCAGCTTG AAGACCTCAG 1048 **K338Rfs*9**
CGGTGCCCAG TACCTCCACG ATGTCCCAAG AGCCTGAGCT CCTCATCAGT 1098
GGAATGGAAA AGCCCCTCCC TCTCCGCACG GATTTCTCT**T** **GAG**ACCCAGG *8
GTCACCAGGC CAGAGCCTCC AGTGGTCTCC AAGCCTCTGG ACTGGGGGGCT *58
CTCTTCAGTG GCTGAATGTC CAGCAGAGCT ATTTCCCTTCC ACAGGGGGCC *108
TTGCAGGGAA GGGTCCAGGA CTTGACATCT TAAGATGCGT CTTGTCCCCT *158
TGGGCCAGTC ATTTCCCCTC TCTGAGCCTC GGTGTCTTCA ACCTGTGAAA *208
TGGGATCATA ATCACTGCCT TACCTCCCTC ACGGTTGTTG TGAGGACTGA *258
GTGTGTGGAA GTTTTTCATA AACTTTGGAT GCTAGTGTAC TTAGGGGGTG *308
TGCCAGGTGT CTTTCATGGG GCCTTCCAGA CCCACTCCCC ACCCTTCTCC *358
CCTTCCTTTG CCCGGGGACG CCGAACTCTC TCAATGGTAT CAACAGGCTC *408
CTTCGCCCTC TGGCTCCTGG TCATGTTCCA TTATTGGGGA GCCCCAGCAG *458
AAGAATGGAG AGGAGGAGGA GGCTGAGTTT GGGGTATTGA ATCCCCCGGC *508
TCCCACCCTG CAGCATCAAG GTTGCTATGG ACTCTCCTGC CGGGCAACTC *558
TTGCGTAATC ATGACTATCT CTAGGATTCT GGCACCACTT CCTTCCCTGG *608
CCCCTTAAGC CTAGCTGTGT ATCGGCACCC CCACCCCACT AGAGTACTCC *658
CTCTCACTTG CGGTTTCCTT ATACTCCACC CCTTTCTCAA CGGTCCTTTT *708
TTAAAGCACA TCTCAGATTA

