



RIPK1 (NM_003804.6) - cDNA - 2024-11-22

GGGAGTCCGC GCGGAGCGCA GCAGCAGGGC CCGGTCCTGC GCCTCGGGAG -104
TCGGCGTCCA GGCTCGGAGC GCGACACGGA GACTAGGTGG CAGGGTACAG -54
CTCTGCCGGG GGGGGAAAAA GTGGTACCAT TTTGGGCGTT CTTGAGCTTC -4
AGA**ATG**CAAC CAGACATGTC CTTGAATGTC ATTAAGATGA AATCCAGTGA 47
CTTCCTGGAG AGTGCAGAAC TGGACAGCGG AGGCTTTGGG AAGGTGTCTC 97
TGTGTTTCCA CAGAACCCAG GGACTCATGA TCATGAAAAC AGTGTACAAG 147
GGGCCCAACT GCATTGAGCA CAACGAGGCC CTCTTGGAGG AGGCGAAGAT 197
GATGAACAGA CTGAGACACA GCCGGGTGGT GAAGCTCCTG GGCATCATCA 247
TAGAGGAAGG GAAGTACTCC CTGGTGTATGG AGTACATGGA GAAGGGCAAC 297
CTGATGCACG TGCTGAAAGC CGAGATGAGT ACTCCGCTTT CTGTAAAAGG 347
AAGGATAATT TTGGAAATCA TTGAAGGAAT GTGCTACTTA CATGGAAAAG 397
GCGTGATACA CAAGGACCTG AAGCCTGAAA ATATCCTTGT TGATAATGAC 447
TTCCACATTA AGATCGCAGA CCTCGGCCCTT GCCTCCTTTA AGATGTGGAG 497
CAA**A**CTGAAT AATGAAGAGC ACAATGAGCT GAGGGAAGTG GAC**G**GCACCG 547 [G181S](#)
CTAAGAAGAA TGGCGGCACC CTCTACTACA TGGCGCCCGA GCACCTGAAT 597
GACGTCAACG CAAAGCCCAC AGAGAAGTCG GATGTGT**A**CA GCTTTGCTGT 647 [Y212*](#)
AGTACTCTGG GCGATATTTG CAAATAAGGA GCCATATGAA **A**ATGCTATCT 697 [c.688_688+20del](#)
GTGAGCAGCA GTTGATAATG TGCATAAAAT CTGGGAACAG GCCAGATGTG 747
GATGACATCA CTGAGTACTG CCCAAGAGAA ATTATCAGTC TCATGAAGCT 797
CTGCTGGGAA GCGAATCCGG AAGCTCGGCC GACATTTCCCT GGCATTGAAG 847
AAAAATTTAG GCCTTTTT**A**T TTAAGTCAAT TAGAAGAAAG TGTAGAAGAG 897 [Y289*](#)
GACGTGAAGA GTTTAAAGAA AGAGTATTCA AACGAAAATG CAGTTGTGAA 947

GAGAATGCAG TCTCTTCAAC TTGATTTGTGT GGCAGTACCT TCAAGCCGGT 997 M318fs L321R D324H D324N D324Y D324V D324G C325R
CAAATTCAGC CACAGAACAG CCTGGTTTCC TGCACAGTTC CCAGGGACTT 1047 S333*
GGGATGGGTC CTGTGGAGGA GTCCCTGGTTT GCTCCTTCCC TGGAGCACCC 1097
ACAAGAAGAG AATGAGCCCA GCCTGCAGAG TAAACTCCAA GACGAAGCCA 1147 K377E
ACTACCATCT TTATGGCAGC CGCATGGACA GGCAGACGAA ACAGCAGCCC 1197 R390G
AGACAGAATG TGGCTTACAA CAGAGAGGAG GAAAGGAGAC GCAGGGTCTC 1247
CCATGACCCT TTTGCACAGC AAAGACCTTA CGAGAATTTT CAGAATACAG 1297 Y426*
AGGGAAAAGG CACTGCTTAT TCCAGTGCAG CCAGTCATGG TAATGCAGTG 1347
CACCAGCCCT CAGGGCTCAC CAGCCAACCT CAAGTACTGT ATCAGAACAA 1397
TGGATTATAT AGCTCACATG GCTTTGGAAC AAGACCACTG GATCCAGGAA 1447
CAGCAGGTCC CAGAGTTTGG TACAGGCCAA TTCCAAGTCA TATGCCTAGT 1497
CTGCATAATA TCCCAGTGCC TGAGACCAAC TATCTAGGAA ATACACCCAC 1547
CATGCCATTC AGCTCCTTGC CACCAACAGA TGAATCTATA AAATATACCA 1597
TATACAATAG TACTGGCATT CAGATTGGAG CCTACAATTA TATGGAGATT 1647
GGTGGGACGA GTTCATCACT ACTAGACAGC ACAAATACGA ACTTCAAAGA 1697
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CCATGACTAT GAGCGAGATG GACTGAAAGA AAAGGTTTAC CAGATGCTCC 1897
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GCCCAGGCGC TCCACCAGTG TTCCAGGATC GACCTTCTGA GCAGCTTGAT 1997
TTACGTCAGC CAGAACTAAC CCTGGATGGG CTACGGCAGC TGAAGTGGAC *31
GCCTCACTTA GTGGATAACC CCAGAAAGTT GGCTGCCTCA GAGCATTCAG *81
AATTCTGTCC TCACTGATAG GGGTTCTGTG TCTGCAGAAA TTTTGTTC *131
TGTACTTCAT AGCTGGAGAA TGGGGAAAGA AATCTGCAGC AAAGGGGTCT *181
CACTCTGTTG CCAGGCTGGT CTCAAACCTC TGGACTCAAG TGATCCTCCC *231
GCCTCGGCC TCCAAAGTGC TGGGATATCA GGCAC TGAGC CACTGCGCCC *281
AGCCAACAAT CCGCTCTGAG GAAAGCGTAA GCAGGAAGAC CTCTTAATGG *331

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AGAGATGTCT GAGGAAGGTC ATGTTCTTTC AGCTTATGGC ATTTCCCTAGA *431
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CACTGTAGGC TCTAAGAACC ACGTGCAGTC TTCAGCCCAT TAAATTATCG *681
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GGATTTAAAG GGGTACCAAG GAGGGGGAA ACATCAGAAT TTCCCAGGCA *781
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TGACTTTTAA AGTTATTTCC TGGCATTCTG GTACCTTCAC CCAGCCTGAG *881
TGCCCTGGAG AGGGAACAGG AAATGCTGAT CTCTACCCCT GGGTGAGACC *931
AGAACCCTCAG GGCTGATACT GTTGAGTGGC TTCCTCGGTT TACTCTGTGT *981
ACTGTGAAAG TATTTTCATA TTTTTTCTGT GTGCCAGAGT GAAAAAGGAC *1031
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CAATCTGGCT GTCGGAACAG ATTCTGGTGT CTTGGGCTGA TAACAGTGT *1281
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TTTAGTAGCA TCCCTGCCCC CACCTACTAG ATGCCAGGGG CACTGTTCTC *1481
CCCAGCCCC CGCCCCAGTT GTGACAATAG TCTCTAAACA TTGTCAAATG *1531
GTCCAAGGAA AGGGGAAAAT TGCCCCGTT GAGAAGAGCA CTGCTGTAAA *1581
GTAATGAGCC TCGGCTCTCC TGTCTGCACC TGTCCGTTA CTTACTTGGCC *1631
ACCACGCAGC CTTGGCTCCT ACAGCCCAA AGGGAGAATG GAGGGAGGCT *1681
CCAGGCTTTG CTGGAGGGC CTGGGTGAGT TCTGTTTGCT CTTGTACCA *1731

CCATCCAAAT GGTGTTATCA AATCTCTTAG ATTCCAAAGA GGTGAATAA *1781

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AAAAATTTAT ACCACAA

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