



RIPK1 (NM_003804.6) - cDNA - 2025-04-02

GGGAGTCCGC GCGGAGCGCA GCAGCAGGGC CCGGTCCTGC GCCTCGGGAG -104
TCGGCGTCCA GGCTCGGAGC GCGACACGGA GACTAGGTGG CAGGGTACAG -54
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AGAATGCAAC 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
CAAAC TGAAT AATGAAGAGC ACAATGAGCT GAGGGAAGTG GACGGCACC G 547 G181S
CTAAGAAGAA TGGCGGCACC CTCTACTACA TGGCGCCCGA GCACCTGAAT 597
GACGTCAACG CAAAGCCCAC AGAGAAGTCG GATGTGTACA GCTTTGCTGT 647 Y212*
AGTACTCTGG GCGATATTTG CAAATAAGGA GCCATATGAA AATGCTATCT 697 c.688_688+20del
GTGAGCAGCA GTTGATAATG TGCATAAAAT CTGGGAACAG GCCAGATGTG 747
GATGACATCA CTGAGTACTG CCCAAGAGAA ATTATCAGTC TCATGAAGCT 797
CTGCTGGGAA GCGAATCCGG AAGCTCGGCC GACATTTCCCT GGCATTGAAG 847
AAAAATTTAG GCCTTTTTAT 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
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ACTACCATCT TTATGGCAGC CGCATGGACA GGCAGACGAA ACAGCAGCCC 1197 R390G
AGACAGAATG TGGCTTACAA CAGAGAGGAG GAAAGGAGAC GCAGGGTCTC 1247
CCATGACCCT TTTGCACAGC AAAGACCTTA CGAGAATTTT CAGAATACAG 1297 Y426*
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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
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GGTGGGACGA GTTCATCACT ACTAGACAGC ACAAATACGA ACTTCAAAGA 1697
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TTACGTCAGC CAGAACTAAC CCTGGATGGG CTACGGCAGC TGAAGTGGAC *31
GCCTCACTTA GTGGATAACC CCAGAAAGTT GGCTGCCTCA GAGCATTAG *81
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TGTACTTCAT AGCTGGAGAA TGGGGAAAGA AATCTGCAGC AAAGGGGTCT *181
CACTCTGTTG CCAGGCTGGT CTCAAACCTC TGGACTCAAG TGATCCTCCC *231
GCCTCGGCCT TCCAAAGTGC TGGGATATCA GGCAC TGAGC CACTGCGCCC *281
AGCCAACAAT CCGCTCTGAG GAAAGCGTAA GCAGGAAGAC CTCTTAATGG *331

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GTCCAAGGAA AGGGGAAAAT TGCCCCGTT GAGAAGAGCA CTGCTGTAAA *1581
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CCAGGCTTTG CTGGAGGGC CTGGGTGAGT TCTGTTTGCT CTTGTACCA *1731

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