RIPK1 (NM_001354930.2) - cDNA - 2026-11-26

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CCGGAGCGCG GCGACTCCAG GGGACCCACA GCTGGGGCGC CAGAGCGCGG -129

CCATCCGGGC GGGGCCGACG GAGCGCGGCA GGACTTGGCT GGACGGCGCG -79

GCCACGGAGA AGGGCGCGGG TACAGCTCTG CCGGGGGGGG AAAAAGTGGT -29

ACCATTTTGG GCGTTCTTGA GCTTCAGAAT GCAACCAGAC ATGTCCTTGA 22

ATGTCATTAA GATGAAATCC AGTGACTTCC TGGAGAGTGC AGAACTGGAC 72

AGCGGAGGCT TTGGGAAGGT GTCTCTGTGT TTCCACAGAA CCCAGGGACT 122

CATGATCATG AAAACAGTGT ACAAGGGGCC CAACTGCATT GAGCACAACG 172

AGGCCCTCTT GGAGGAGGCG AAGATGATGA ACAGACTGAG ACACAGCCGG 222

GTGGTGAAGC TCCTGGGCGT CATCATAGAG GAAGGGAAGT ACTCCCTGGT 272

GATGGAGTAC ATGGAGAAGG GCAACCTGAT GCACGTGCTG AAAGCCGAGA 322

TGAGTACTCC GCTTTCTGTA AAAGGAAGGA TAATTTTGGA AATCATTGAA 372

GGAATGTGCT ACTTACATGG AAAAGGCGTG ATACACAAGG ACCTGAAGCC 422

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GCCTTGCCTC CTTTAAGATG TGGAGCAAAC TGAATAATGA AGAGCACAAT 522

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AGTCGGATGT GTACAGCTTT GCTGTAGTAC TCTGGGCGAT ATTTGCAAAT 672 Y212*

AAGGAGCCAT ATGAAAATGC TATCTGTGAG CAGCAGTTGA TAATGTGCAT 722 c.688_688+20del

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GAGAAATTAT CAGTCTCATG AAGCTCTGCT GGGAAGCGAA TCCGGAAGCT 822

CGGCCGACAT TTCCTGGCAT TGAAGAAAAA TTTAGGCCTT TTTATTTAAG 872 Y289*

TCAATTAGAA GAAAGTGTAG AAGAGGACGT GAAGAGTTTA AAGAAAGAGT 922

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GGTTTGCTCC TTCCCTGGAG CACCCACAAG AAGAGAATGA GCCCAGCCTG 1122

CAGAGTAAAC TCCAAGACGA AGCCAACTAC CATCTTTATG GCAGCCGCAT 1172 K377E R390G

GGACAGGCAG ACGAAACAGC AGCCCAGACA GAATGTGGCT TACAACAGAG 1222

AGGAGGAAAG GAGACGCAGG GTCTCCCATG ACCCTTTTGC ACAGCAAAGA 1272

CCTTACGAGA ATTTTCAGAA TACAGAGGGA AAAGGCACTG CTTATTCCAG 1322 Y426*

TGCAGCCAGT CATGGTAATG CAGTGCACCA GCCCTCAGGG CTCACCAGCC 1372

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GGAACAAGAC CACTGGATCC AGGAACAGCA GGTCCCAGAG TTTGGTACAG 1472

GCCAATTCCA AGTCATATGC CTAGTCTGCA TAATATCCCA GTGCCTGAGA 1522

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GGATCGACCT TCTGAGCAGC TTGATTTACG TCAGCCAGAA CTAACCCTGG *6

ATGGGCTACG GCAGCTGAAG TGGACGCCTC ACTTAGTGGA TAACCCCAGA *56

AAGTTGGCTG CCTCAGAGCA TTCAGAATTC TGTCCTCACT GATAGGGGTT *106

CTGTGTCTGC AGAAATTTTG TTTCCTGTAC TTCATAGCTG GAGAATGGGG *156

AAAGAAATCT GCAGCAAAGG GGTCTCACTC TGTTGCCAGG CTGGTCTCAA *206

ACTTCTGGAC TCAAGTGATC CTCCCGCCTC GGCCTTCCAA AGTGCTGGGA *256

TATCAGGCAC TGAGCCACTG CGCCCAGCCA ACAATCCGCT CTGAGGAAAG *306

CGTAAGCAGG AAGACCTCTT AATGGCATAG CACCAATAAA AAAATGACTC *356

CTAGTTGTGT TTGGAAAGGG AGAGAAGAGA TGTCTGAGGA AGGTCATGTT *406

CTTTCAGCTT ATGGCATTTC CTAGAGTTTT GTTGAAGCAA GAAGAAAAAC *456

TCAGAGAATA TAAAATCAAC TTTTAAAATT GTGTGCTCTC TTCTTCACGT *506

AGGCTCCTGT TAAAAACAAA GTGCAGTCAG ATTCTAAGCC CTGTTCAGAG *556

ACTTCGTGGA TCACAGCTGC AGCTCACCGC CACATCACAG GATCCGTTAA *606

CGTTAATACC CAATACTCTG TCAGCCACTG TAGGCTCTAA GAACCACGTG *656

CAGTCTTCAG CCCATTAAAT TATCGATTAT TTTTTAATGA ATTGAATTTA *706

TATTGAGTCT TCAAATTAAC TGAATGGATT TAAAGGGGTA CCAAGGAGGG *756

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CGTGACTACA ACAAAAATTC TTGATTGACT TTTAAAGTTA TTTCCTGGCA *856

TTCTGGTACC TTCACCCAGC CTGAGTGCCC TGGAGAGGGA ACAGGAAATG *906

CTGATCTCTA CCCCTGGGTG AGACCAGAAC CTCAGGGCTG ATACTGTTGA *956

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TCTGTGTGCC AGAGTGAAAA AGGACAGCTT CTGAGTGTGG TAATTGTGCC *1056

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nfevers