



LYN (NM_002350.4) - cDNA - 2024-11-01

AGTTCCTCTC CCGCCGCGCC GGGCCGCGCT GCCGCTCGCT CCCCggccGT -225
GGCGCCTCCG GGCCAGACGC GCTGCAGCCT CCAGCCCGCG GCAAGCGGGG -175
CGGCCGCGCC ACCCCCGGCC CCGCGCCAGC AGCCCTCGC CGCGCGTCCA -125
GCGTTCCCGG CCAGCAGCCT CCCCATACGC AGGTCCTGCT GGGCCGCCCC -75
GTCGCGCCCC CCACTCTGAA CTCAAGTCAC CGTGGAGCTC CGCCGCCCCG -25
AAACTTTCAC CGCGAGCGGG AAATATGGGA TGTATAAAAT CAAAAGGGAA 26
AGACAGCTTG AGTGACGATG GAGTAGATTT GAAGACTCAA CCAGTACGTA 76
ATACTGAAAG AACTATTTAT GTGAGAGATC CAACGTCCA TAAACAGCAA 126
AGGCCAGTTC CAGAATCTCA GCTTTTACCT GGACAGAGGT TTCAAACATA 176
AGATCCAGAG GAACAAGGAG ACATTGTGGT AGCCTTGTAC CCCTATGATG 226
GCATCCACCC GGACGACTTG TCTTCAAGA AAGGAGAGAA GATGAAAGTC 276
CTGGAGGAGC ATGGAGAATG GTGGAAAGCA AAGTCCCTTT TAACAAAAAA 326
AGAAGGCTTC ATCCCAGCA ACTATGTGGC CAAACTCAAC ACCTTAGAAA 376
CAGAAGAGTG GTTTTTCAAG GATATAACCA GGAAGGACGC AGAAAGGCAG 426
CTTTTGGCAC CAGGAAATAG CGCTGGAGCT TTCCTTATTA GAGAAAGTGA 476
AACATTAAAA GGAAGCTTCT CTCTGTCTGT CAGAGACTTT GACCCTGTGC 526
ATGGTGATGT TATTAAGCAC TACAAAATTA GAAGTCTGGA TAATGGGGGC 576
TATTACATCT CTCCACGAAT CACTTTTCCC TGTATCAGCG ACATGATTAA 626
ACATTACCAA AAGCAGGCAG ATGGCTTGTG CAGAAGATTG GAGAAGGCTT 676
GTATTAGTCC CAAGCCACAG AAGCCATGGG ATAAAGATGC CTGGGAGATC 726
CCCCGGGAGT CCATCAAGTT GGTGAAAAGG CTTGGCGCTG GGCAGTTTGG 776
GGAAGTCTGG ATGGGTTACT ATAACAACAG TACCAAGGTG GCTGTGAAAA 826

CCCTGAAGCC AGGAACATATG TCTGTGCAAG CCTTCCTGGA AGAAGCCAAC 876
CTCATGAAGA CCTGTCAGCA TGACAAGCTC GTGAGGCTCT ACGCTGTGGT 926
CACCAGGGAG GAGCCCATT T ACATCATCAC CGAGTACATG GCCAAGGGCA 976
GTTTGCTGGA TTTCTGGAAG AGCGATGAAG GTGGCAAAGT GCTGCTTCCA 1026
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CCGAGTCACT CATGTGCAA AATTGCAGATT TTGGCCTTGC TAGAGTAATT 1176
GAAGATAATG AGTACACAGC AAGGGAAGGT GCTAAGTTCC CTATTAAGTG 1226
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TGAAAATGTG CTGGAAAGAA AAGGCAGAAG AGAGACCAAC GTTTGACTAC 1476
TTACAGAGCG TCCTGGATGA TTTCTACACA GCCACGGAAG GGCAATACCA 1526 Q507* Y508H Y508F Y508*
GCAGCAGCCT TAGAGCACAG GGAGACCCGT CCATTTGGCA GGGGTGGCTG *37
CCTCATTTAG AGAGGAAAAG TAACCATCAC TGGTTGCACT TATGATTTCA *87
TGTGCGGGGA TCATCTGCCG TGCCCTGGATC CTGAAATAGA GGCTAAATTA *137
CTCAGGAAGA ACACCCTCTA AATGGGAAAG TATTCTGTAC TCTTAGATGG *187
ATTCTCCACT CAGTTGCAAC TTGGACTTGT CCTCAGCAGC TGGTAATCTT *237
GCTCTGCTTG ACAACATCTG AGTGCAGCCG TTTGAGAAGA AAACATCTAT *287
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GGCGTGCACA TCTCTCTCTC TTCCAGCAGG AGGAGCCCGT GAGCACGCAC *687

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