



LYN (NM_002350.4) - cDNA - 2024-05-18

AGTTCCTCTC CCGCCGCGCC GGGCCGCGCT GCCGCTCGCT CCCC GGCCGT -225
GGCGCCTCCG GGCCAGACGC GCTGCAGCCT CCAGCCCGCG GCAAGCGGGG -175
CGGCCGCGCC ACCCCCGGCC CCGCGCCAGC AGCCCCTCGC 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 CAACGTCCAA TAAACAGCAA 126
AGGCCAGTTC CAGAATCTCA GCTTTTACCT GGACAGAGGT TTCAAACATA 176
AGATCCAGAG GAACAAGGAG ACATTGTGGT AGCCTTGTAC CCCTATGATG 226
GCATCCACCC GGACGACTTG TCTTTCAAGA 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 AGGAACTATG TCTGTGCAAG CCTTCCTGGA AGAAGCCAAC 876
 CTCATGAAGA CCCTGCAGCA TGACAAGCTC GTGAGGCTCT ACGCTGTGGT 926
 CACCAGGGAG GAGCCCATT T ACATCATCAC CGAGTACATG GCCAAGGGCA 976
 GTTTGCTGGA TTTCTGAAG AGCGATGAAG GTGGCAAAGT GCTGCTTCCA 1026
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 CCGAGTCACT CATGTGCAAA ATTGCAGATT TTGGCCTTGC TAGAGTAATT 1176
 GAAGATAATG AGTACACAGC AAGGGAAGGT GCTAAGTTCC CTATTAAGTG 1226
 GACGGCTCCA GAAGCAATCA ACTTTGGATG TTTCACTATT AAGTCTGATG 1276
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 CTACAGGATG CCCCGTGTGG AGAACTGCCC AGATGAGCTC TATGACATTA 1426
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 TTACAGAGCG TCCTGGATGA TTTCTACACA GCCACGGAAG GGCAATACCA 1526 Q507* Y508H Y508E Y508*
 GCAGCAGCCT TAGAGCACAG GGAGACCCGT CCATTTGGCA GGGGTGGCTG *37
 CCTCATTTAG AGAGGAAAAG TAACCATCAC TGGTTGCACT TATGATTTCA *87
 TGTGCGGGGA TCATCTGCCG TGCCTGGATC CTGAAATAGA GGCTAAATTA *137
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 ATTCTCCACT CAGTTGCAAC TTGGACTTGT CCTCAGCAGC TGGTAATCTT *237
 GCTCTGCTTG ACAACATCTG AGTGCAGCCG TTTGAGAAGA AAACATCTAT *287
 TCTCTCCAAA AATGCACCCA ACTAGCTCTA TGTTTACAAA TGGACATAGG *337
 ACTCAAAGTT TCAGAGACCA TTGCAATGAA TCCCAATAA TTGCAGAACT *387
 AAACTCATTT ATAAAGCTAA AATAACCGGA TATATACATA GCATGACATT *437
 TCTTTGTGCT TTGGCTTACT TGTTTAAAAA AAAAAAAAAA CTAATCCAAC *487
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 TTTCAATGAT TTTTTTCCCC CTACCTCCCA AAATCTGAGA CTGTTAAAAC *587
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 GGCGTGCACA TCTCTCTCTC TTCCAGCAGG AGGAGCCCGT GAGCACGCAC *687

AGCTGCCCTG TCTGCTCACC CGAAGGCACC GGGCTCACCT GGACCTCCCA *737
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TTGACAATGT AGTTTGGAAG AACTAAGATT CTAATCTCTG AAGAACCTTA *887
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TGTGATCCTT ACCTCCATGT GGGCCCTTCA CCAGCTTGGG CCTCATCTCT *2037
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CAGTCTCCTT ACTTAGCTAT AGGTTTCCAG CCTCCCTGTG ACAGACAGGC *2137
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