



LYN (NM_002350.4) - cDNA - 2025-04-02

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 CAACGTCCAA 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 TTTCTGAAAG AGCGATGAAG GTGGCAAAGT GCTGCTTCCA 1026
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GCGGAAGAAC TACATTACCC GGGACCTGCG AGCAGCTAAT GTTCTGGTCT 1126
CCGAGTCACT CATGTGAAA ATTGCAGATT TTGGCCTTGC TAGAGTAATT 1176
GAAGATAATG AGTACACAGC AAGGGAAGGT GCTAAGTTCC CTATTAAGTG 1226
GACGGCTCCA GAAGCAATCA ACTTTGGATG TTTCACTATT AAGTCTGATG 1276
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CTACAGGATG CCCCCTGTGG AGAACTGCCC AGATGAGCTC TATGACATTA 1426
TGAAAATGTG CTGGAAAGAA AAGGCAGAAG AGAGACCAAC GTTTGACTAC 1476
TTACAGAGCG TCCTGGATGA TTTCTACACA GCCACGGAAG GGCAATACCA 1526 Q507* Y508H Y508F Y508*
GCAGCAGCCT **TAG**AGCACAG 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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ACTCAAAGTT TCAGAGACCA TTGCAATGAA TCCCAATAA TTGCAGAACT *387
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CTGTTAGATT TTGCAGGTGA AGTCAGCAGC TTAAAAATGT CTTTCCCAGA *537
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 AAGAACCCTA *887
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TGTGATCCTT ACCTCCATGT GGGCCCTTCA CCAGCTTGGG CCTCATCTCT *2037
GGTGTCCAGC ATGTGTGGAA GTCACACGTT CCCTTGATGA ACAGCACACA *2087

CAGTCTCCTT ACTTAGCTAT AGGTTTCCAG CCTCCCTGTG ACAGACAGGC *2137
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