

Genomic sequence of rat β -globin minor gene

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Submitted May 16, 1989

EMBL accession no. X15160

In order to understand the complexity of rat β -like globin locus a number of clones covering the region was isolated from heterogenous genomic library prepared in EMBL 3 vector from Wistar strain animals carrying Belgrade anemia (1). The following DNA sequence which contains one entire rat β -globin gene ($\beta_B^{mi/hy}$) from clone $\lambda 11$ was determined using Sanger's dideoxy sequencing method. The detailed sequence analysis and comparison with available globin mRNA, globin gene and protein sequences indicate that this is an active β^{min} globin gene which shares extensive homology with the mouse β^{mur} globin gene (2, 3, 4). The coding regions are underlined.

1	<u>CCCTGGTAGT</u>	TATGGCTATC	ATCTCTGAAG	CCTCACCCCTG	CAGAGGCACA
51	CCCTCACATT	GCCCAATCTG	CTCACACAGG	ACAGAGTGAT	CAGGGGCCAG
101	AATTTGGCAT	ATAAAGCAGA	ACAGAACCAG	TTGCTTCTTA	TATTTGCTTC
151	TGATACTGTT	GTGTTGACTC	GCAACCTCAG	GAACAGACAC	CATGGTGCAC
201	<u>CTAACTGATG</u>	<u>CTGAGAAGGC</u>	<u>TACTGTTAGT</u>	<u>GGCCTGTGGG</u>	<u>GAAAGGTGAA</u>
251	<u>TGCTGATAAI</u>	<u>GTTGGCGCTG</u>	<u>AGGCCCTGGG</u>	<u>CAGGTTGGTA</u>	<u>TCCAGGTTAC</u>
301	AAGGTAGCTC	CTAAGTAGAA	GTTTGGTGCT	TGGAGACAGA	GGTCTGCTTT
351	CCAGCAGGCA	CTAACTTTTT	TGCTTCTGG	CTATGTTTCC	CCTTTGTAGG
401	<u>CTGCTGGTTG</u>	<u>TCTACCCTTG</u>	<u>GACCCAGAGG</u>	<u>TACTTTTCTA</u>	<u>AAATTTGGGGG</u>
451	<u>CGCTGCTCTCT</u>	<u>GCCTCTGCTA</u>	<u>TCATGGGTAA</u>	<u>CCCCAGGCTG</u>	<u>AAAGCCCATG</u>
501	<u>GCAAGAAGGT</u>	<u>GATAAATGCC</u>	<u>TTCAATGATG</u>	<u>GCCTGAAACA</u>	<u>CTTGGACAAC</u>
551	<u>CTCAAGGCCA</u>	<u>CCTTTGCTCA</u>	<u>TCTGAGTGAA</u>	<u>CTCCACTGTG</u>	<u>ACAAGCTGCA</u>
601	<u>TGTGGATCCG</u>	<u>GAGAACTTCA</u>	<u>GGTGAATCT</u>	<u>AATGGGCTCC</u>	<u>CCACTGGGTT</u>
651	TCCTTCTGTT	GGCTTTCCTG	CTCAAATFCC	TATCAGAAGG	AAAGAGGAAG
701	CAATTCTAGG	GACGACTTTT	GATGATGATG	TGTTGGATG	CCCTGTGGAG
751	TGTTGACAGG	AGTCCAGTTA	TTTTATCCTC	TATTCACAAT	CACCTTCCCC
801	TCTCACTCTG	TTCTTCTATG	TTGTCAATTC	CTCTTTCTTT	GGTAAACTTT
851	TAAATTTTCT	GTTGCAAGTT	TAAAGTACAT	TTTTTATGTA	CTTTCTCTCT
901	TTTTTTTTAT	TCAGCCATGA	GGGTACCCTC	TAGACTTTAA	AAAACGTAGT
951	ACTTTCCTCT	TTGTTCAAG	TGTTTCCTGC	TACTTTACTC	TGAGGACGTA
1001	AAGATCAATG	ATTCACTCAT	TCCACACCTG	TAAGGAATAG	TAGAACAATA
1051	ATTGGCTTTT	AGGCTAAGAT	GATAGGGAAA	TATATATTTT	GCATATAAAT
1101	TTTGTCTGCT	AGAAGAATTC	TTATCAAAAT	TGACCAGGAG	AACTCAGTAG
1151	TCATTCTGCC	TGTCTTTTAA	GATTATAACT	GCAAACCTCCA	TTTGAATGG
1201	GCCTGCACTG	TCTGATATG	TTGTTCTACT	TCATGTTGAA	ACATCTTCCC
1251	TCTTCCACCA	<u>GCTCCTGGGC</u>	<u>AATATGATG</u>	<u>TGATTGTGTT</u>	<u>GGGCCAGCCAC</u>
1301	<u>CTGGGCAAGG</u>	<u>AATTCACCCC</u>	<u>CTGTGCACAG</u>	<u>GCTGCCTTCC</u>	<u>AGAAGGCGST</u>
1351	<u>GGCTGGAGTG</u>	<u>GCCAGTGCCC</u>	<u>TGGCTCACAA</u>	<u>GTACCACATC</u>	<u>ACCTCTTTTC</u>
1401	CTGCTCTTGT	CTTTGTGCAA	TGGTCAATTTG	TTCCCAAGAG	AGCATCTGTC
1451	AGTTGTGTGC	AAAATGACAA	AGACCTTTGA	AAATCTGTCC	TACTAATTTAA
1501	AGGCATTTAC	TTTCACTGCA	ATGGTGTGTT	AAATTAATTTG	TATCTCATAG
1551	AAGGTTTCAT	GCTTAGGTTT	AAGATACAAA	GCAGTGAGGG	

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