

**Nucleotide sequence of a gene encoding the *Borrelia burgdorferi* flagellin**G.S.Gassmann, M.Kramer<sup>2</sup>, U.B.Göbel\* and R.Wallich<sup>1</sup>

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The genes encoding a major flagellar protein (1) from *B. burgdorferi* strains B31 (type strain) and GeHo (isolated from skin biopsy by Dr. Pelz, Freiburg) were identified from a pUEX1 and Agt11 expression library, respectively, by immunoscreening with a flagellin-specific monoclonal antibody. Each sequence has been obtained on both strands by a modified Sanger protocol (2). Fig.1 shows the primary nucleotide sequence of the *B. burgdorferi* B31 flagellin gene. Base changes found within the sequence of strain GeHo are indicated. The structural gene, 1008 bp in length, codes for a protein of 336 amino acids. The correct open reading frame was confirmed by N-terminal amino acid analysis of the purified flagellin (3).

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1 ATGATTATCA ATCATAATAC ATCAGCTTAA AATGCCCTCAA GAAATAATGG CATTAACGCT
61 GCTAACTCTTA GTAAACTCA AGAAAAGCTT TCTAGTGGGT ACAGAATTAA TCGAGCTCT
121 GATGATGCCTG CTGGCATGGG ATGTTCTGGT AACATTAATG CTCAAATAAG AGGTTTGTC
181 CAAGCTTCTA GAAATATTTC AAAGGCTTAA ATTTTTATTC AGACAACAGA AGGGAATTAA
241 AATGAAGTAG AAAAAGCTTT AGTAAGAATG AAGGAATTGG CAGTTCAATC AGGTAACGGC
301 ACATATTCAG ATGCAGACAG AGGTTCTATA CAATTGAAA TAGAGCAACT TACAGACGAA
361 ATTAATAGAA TTGCTGATCA ACCTCAATAT AACCAATGCC ACATGTTATC AAACAAATCT
421 GCTTCTCAA ATGTAAGAAC ACCTGAAGAG CTGGAAATGC ACCCTGCCAA AATTAACACA
          G
          A
481 CCAGCATCAC TTTCAGGGTC TCAAGCGCTT TGGACTTTAA GAGTTCTATG TGGAGCAACC
541 CAAGATGAAG CTATTCCTGT AAATTTTAT GCAGCTTAATG TTGCAAAATCT TTTCCTGGT
601 GAGGGAGCTC AAACTGCTCA GGCTGCACCC GTTCAAGAGG GTGTTCAACA GGAAGGGAGCT
661 CAACAGCCAG CACCTGCTAC ACCACCTCTT CAAGGCCGGAG TTAATTCTCC TGTTAATGTT
721 ACAACTACAG TTGATGCTAA TACATCACTT GCTAAATTC AAAATGCTAT TAGAATGATA
          A
          G
781 AGTGTACAAA GGCCAAATTG AGGTGCTTTC CAAATAGAC TTGAATCTAT AAAGAATAGT
841 ACTGAGTATG CAATTCGAAA TCTAAAGCA TCCTATGCTC AAATAAAAGA TGCCTACAATG
901 ACAGATGAGG TTGTAGGAGC AACAACTAAT AGTATTTTAA CACAATCTGC AATGGCAATG
961 ATTGCCAGG CTAATCAAGT TCCCCATAT GTTTTGTCATG TGCTTAGATA A

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**References:** (1) Barbour, A.G. et al. (1986) Infect. Immun. 52, 549-554. (2) Tabor, S. and Richardson, C.C. (1987) Proc. Natl. Acad. Sci. USA 84, 4767-4771 (3) Gaßmann, G.S. et al., FEMS Lett., in press.