

Nucleotide sequence of the penicillin-binding protein 2B gene of *Streptococcus pneumoniae* strain R6

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Submitted August 14, 1989

EMBL accession no. X16022

Previously only part of the penicillin-binding protein 2B gene (*penA*) of *Streptococcus pneumoniae* had been cloned and sequenced (1). The whole of *penA* was isolated on a 3.2-kb *EcoRV* fragment and cloned into pBGS9 (2). *E. coli* TG1 containing the resulting plasmid (pCGD3) expressed large amounts of a novel penicillin-binding protein of 74kD. A fragment of the insert, containing the previously unsequenced amino-terminal coding region of *penA*, was subcloned into bacteriophage M13 and sequenced on both strands using a series of oligonucleotide primers.

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GATATCTGGTTGTCCATAGGGAAAGCTCTTACATGAGTCATACTGGAAAGACTAGATCAGGGAAATGTCACACTTCAATTAAACACATAATAAGGATAGATAATAACAGCATC 120
1
M R K
TCTGTTTAAAAACGAAAAAACTGAAAAGCTCTCTCTTCCATAATTCTACTCAAATTGTTGACAATTAAAGAGTAAGATTAAAGTAAATGAGACTGATTGTATGAGAAA 240
20
F N S H S I P I R L N L L F L S I V I L L F M T I I G R L L Y M Q V F N K D F Y E
ATTCACAGCGATTGCGATTGCTGTTACTCTTACGTTGACATTATTGTTGTTGATGAGTTGAGTTTCAACAAGGATTGTTAC 360
60
K K L A S A S Q T K I T S S S A R G E I Y D A S G K P L V E N T L K Q V V S F T
AAAAAAAGTAGGCCCTGAGCTAGCAGACCAAGATTCAACAGAGTCAGCTGCGCTGGAAAATTATGAGCTAGTGAAACCTTGTGAGAAATACGTTAAAGCAGGTTGTTCTTAC 480
100
R S N K M T A T D L K E T A K K L V T Y V S I S S P N L T E R A G D Y Y L A D P
GCGAGACATAAAATGAGCGCTACAGCTTAAAGAACAGCTTAAAGGACTACTGACTCTATGAGCTACAGTTCTCAAAATTGAGCACAGGAGCTGGGGATTACTATTGGCTGATCC 600
120
E I Y K K I V E A L P S E C R L D S G N R L L S E S P L Y L N N A V D S V U O T S Q
TGAATCTATAAAAATGAGCTCCAGCGAGAAACCGCTGGATTCTAGATGGCAATTCTATACGAACTGATGTTAAACATGGCTGATGTCACAAACAGACT 720
140
L N Y T E D E K K E I Y L F S Q L N A V G N F A T G T I A T D P L R N D S Q V A V
ACTAAACATACAGGAGATGAAAAGAAAATCTATCTTCTTCTAGCTTAAATGGCTGTTGAAACCTTGGCGACAGGGACATTGTCACAGATCTCTAAATGATCTCAAGGTTGCTGT 840
160
I A S I S K E M P G I S I S T S D R K V L E T S L S I V G S V S S E A A G L
TATTCGCTTATTCAGGAGATGCGCTGGATTAGTTCTCTGGAGATAAGAAAGTTTGGAAACCTTCTTCTATAGTTGGAGTATCCAGTGAAGAACAGCTGGT 960
180
P A E E A A G L E L K K G Y S L N D R V G T S Y L E K Q Y E E T L L Q G K R S V K E
CCCCAGGGAAAGCAGAACGCTCTTAAAGGGCTATTCTTAAATGCTTAAATGAGCTGTTAGGAAACCTTCTATTGGAAAGCATGAGAGACCTTACAAGGAAACGGCTGGTAAAGA 1080
200
I H L D K Y G N E S V D T I E E G S K G N N I K L T I D L A F Q D S V D A L L
AATCAGCTGGATAATGGCAATATGGAAAGGGTGGGATAATTGGAGAACGGAGAGTAGTACAGGAAACAACTAACAGTACAGGCTTGGTTCCAAGATGAGCTGGATGCTTACT 1200
220
K S Y F N S E L E N G G A K Y S E G V Y A V A L N P K T G A V L L S M S G I K H D
GAAAAGTTATTCGAGCTAGAAATGGTGAGGCAACAGGTTCTGGAGGTGTTCTATGAGGTTGCTATGAGGTTGGCTTAACCCAAAACAGGTGGCTTGTATGTCAGGTTAAACATGCA 1320
240
L K T G E L T P D S L G T V N F V P G S V V K U A T I T S S G W E N G V L S G
CTTGAAAGGGAGGTTAAGCTGATTCTGGGAAACGGTAAACCTGTTCTGGCTTCCAGGTGTTGTCAGGGCGGACCATCAGCTTGGGAAATGGAGCTTGTCTAGG 1440
260
N Q T L T D O S I L V F Q G S A P N S W Y T Q A Y G S F F I T A V Q A L E Y S S
AAACACAGACCTGGAGACAGCTGGATTCTGGCTTCAAGGTTCTGGCCCTAACTTCTGGTATCTACAGGCTTACGGCTTCTATCACAGGGTCAAGCTGGAGTATCTAC 1560
280
N T Y M V Q T A L G L M G Q T Y O P N M F V G T S N L E S A M E K L R S T F G E
AAATACCTATGGTCCANACAGCTTAACTGGGAAACCTTCAACCCAAATATGGTTGGCACCAGCAATCTAGATCTGCTATGGAGAAACTGGCTTCAACCTTGGCGA 1680
300
Y G L G T A T G I D L P D E S T G F D V P K E Y S F A N Y I T N A F G Q F D N Y T
ATATGGCTTGGGACTGGCAACGGTACCTGGGATAGTAATCTGGCTTGGCTTCCAAAGGAGTATAGCTTGGCTAACTTACATTAATGCTTGGGAGCTTGTGATAC 1800
320
P M Q L A Q Y V V A T I A N N G V A R V A P R I V E G E I Y G N N D K G G L C O L I Q
GCCGATGCGACTGGCTTGGCTAGTGAACATATTGCAAAATGGTCTGGCTTCTCTGGTATGGTGAAGGCTTGGGATATAATGATAAGGGAGGACTGGTACTGGT 1920
340
Q L Q P T E M N K V N I S D S M S I L H Q G F Y Q V W A H G T S S G L T T G R A F
GCAACTGCAACCGACAGAGATGAAAGTCAATATCCGACTGGCATGAGCTTGGCACCAAGGTTTATCAGGGTGGCATGACTAGTGGATGACACTGGACACTGGCTT 2040
360
S N G A L V S I S G K T T G T A E S Y V A D G Q Q O A T N T N A V A Y A P S D N P Q
TTCAAAATGGCTTGGCTTGGTATCTATTAGCGAAAAACAGGTACAGCGCAAAGCTATGGCGCATGGTCAAGCAGACCAACCAATTGCGGCTTATGCCCATCTGATAATCCC 2160
380
I A A V A V V F P H N T N L T N G V G P S I A R D I I N L Y Q K Y H P M N *
AAATGGCTGTCGAGTGGCTTCTCTCAATAACCACTAACAAATGGTAGGACCTTCCATTGGCGTGACATTCTACATCTATCAAATACCTCCAACTGAACTGAGAAA 2280

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Fig. 1. The sequence shown is that of the *penA* gene of *S. pneumoniae* strain R6. The active site serine is marked with an asterisk.

- REFERENCES** (1) Dowson C.G. et al. (1989) Mol. Microbiol. 3, 95-102;
(2) Spratt B.G. et al. (1986) Gene 41, 337-342.