

Sequence of the adenine methylase gene of the *Streptococcus faecalis* plasmid pAM β 1John Brehm¹, George Salmond² and Nigel Minton¹¹Microbial Technology Laboratory, PHLS Centre for Applied Microbiology and Research, Porton Down, Salisbury and ²Department of Biological Sciences, University of Warwick, Coventry, UK

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120
 CAAAGAAAAACGAAATGATACACCAACTAGTCGAAAAAAAAGATAATGGAGATAAGACCGTTCTGTTCCGCTGACTTGACCATATCATAAAAATCGAACACGAAAGAATGGCG
 CATTCTTAAATGGCACAACAGGTACCGTTATTGAGGTATTTCTTATCTATGGGTTAACATGGATTATTCATTTAAATCATGATATTGTCGAGAGTGATTGGCTTCGCGTA
 -35 -10 240
 GAAACGTTAAAGAAGTTATGGAAAAGACTTAGAAGCAAACCTTAAAGAGTGTGTTGATAGTGCAGTTCTTAAATTTGTATAATAGGAATTGAAGTTAAATTAGATGCTAAAAATTGG
 TGGTTAACCTTA ORF1 S.D. M L V F Q M R N V D K T S T V L K Q T K N S D Y A D K Y V R L I
 TAAATTAAGGAGGAGG 600TGTCTATGGTATTCCAATGGTAAATGAGATAAAACATCTACTGTTGAAACAGACTAAACAGTGAATTACGGAGATAAAACGTTAGATTAAAC
 P T S D 275 S.D. M N E N I ORF2 275
 CTACAGTGACTAACTTATGACTTTAACAGATAACTAAACAAACATCGTTAACCTCTGTTATTTATTTATAGATGATCTTCAGGAG 600TATTACATGAAACAAAAATAT
 395 515 635
 K Y S Q N F L T S E K V L N Q I I K Q L N L K E T D T V Y E I G T G K Q H L T T
 AAAAATTCTCAAACCTTTAACCGAGTAAAGACTCAACCCAAAATAAAACAAATTGAAATTAAAGAACCGATACCCGTTACCGAAATTGGACAGGTTAAAGCCATTAAACGAC
 A G I
 K L A E K I S K Q V T S I E L D S H L F N C F L N L S S E K L K L N T R V T L I H Q D I L
 GAAACTGGCTAAATTAAGTAAACAGGTAACCTCTATTGAAATTAGACAGTCATCTTCAACTTACCTCTGAGAAATTAAACTGAAACTCTGTTACCTTAAATCACCAGATATCTGAT
 C T
 755
 Q F Q A P N K Q R Y K I V G S I P Y H L S T Q I K K V V F E S H A S D I Y L I
 ACAGTTCTTCAACACAGGTTAAATTGTTGGAGTATTCTTACCAATTGACACAAACAAATTAAAGTGTITTTGAAAGCCATCGCTGTCATCTTAAATCACCAGATATCTGAT
 A
 V E E G F Y K R T L D I H R T L G L L L H T Q V S I Q Q L L K L P A E C F H P K
 TCTTGAAGAAGGATTCTCAAGCGTACTCTGGTATTACCGAACACTAGGTTCTGCTGCACTCAAGTCCTGCAACTAGCATTGCTTAAGCTGCCAGGGAAATGCTTCTACCTTAA
 tt
 P K V N S V L I K L T R H T T D V P D K Y W K L Y T Y F V S K W V N R E Y R Q L
 ACCAAAAGTAAACAGTGTCTTAAAAACTACCCGGCCATACACAGATGTTCCAGATAATAATGGAAAGCTTACAGTACTTGTCTTAAAGGTCATGAGAATATCTGCAACT
 1095
 F T K N Q F H Q A M K H A K V N N L S T V T Y E Q V L S I F N S Y L L F N G R K
 GTTAACTTCTATGAGTCGCTTTGTAATTTGGAAAGTTACACGTTACTAAAGGGAAATGAGATAAAATTAGGTATACTACTGACAGCTCCAAGGAGCTAAAGGCTCCAGGCTA
 ORF3 M S R F C K F G K L H V T K G N V D K L L G I L L T A S K E L K R S L A P
 ATAAATCTATGAGTCGCTTTGTAATTTGGAAAGTTACACGTTACTAAAGGGAAATGAGATAAAATTAGGTATACTACTGACAGCTCCAAGGAGCTAAAGGCTCCAGGCTA
 CGGGGAATTG 1226

Figure Legend. The complete nucleotide sequence of the adenine methylase gene of the *Streptococcus faecalis* plasmid pAM β 1. The gene was cloned from the plasmid PR29 (Vasseghi, H. and Claverys, J. 1983, Gene 21:285-292) as a c.1200 bp *Hha*I fragment, blunt-ended with T4 DNA polymerase and inserted into the *Sma*I site of M13mp8 and mp9. Nucleotide sequencing was by the dideoxy procedure employing specific 17 mer oligonucleotide primers. The illustrated sequence extends 1226 bp beginning at the last base of the cloned *Hha*I site, with the adenine methylase gene labelled as ORF2. A comparison with the equivalent regions from pAM77 (Horinouchi, S. et al. J.Bacteriol. 154:1252-1262) and Tn917 (Shaw, J.H. and Clewell, D.B. J.Bacteriol. 164:782-796) is also presented. Homology with Tn917 does not begin until nt 133. The previously determined sequence of pAM77 corresponds to nt 150 (*) onwards. Differences at the amino acid and nucleotide sequence level are indicated by upper (Tn917) and lower (pAM77) case letters above and below the sequence, respectively. The major overall difference is the presence of a deletion in the pAM β 1 sequence (nt 254) which removes ORF1, known to be responsible for induction of Em resistance in the case of Tn917 and pAM77. Its absence explains the observed constitutive expression of Em resistance in cells harbouring pAM β 1 and its derivatives.