

## Sequence of the *lktD* gene from *Actinobacillus actinomycetemcomitans*

Janet M.Guthmiller, Ellen Kraig, Marianna P.Cagle and David Kolodrubetz<sup>1\*</sup>

Departments of Cellular and Structural Biology and <sup>1</sup>Microbiology, University of Texas Health Science Center at San Antonio, 7703 Floyd Curl Drive, San Antonio, TX 78284, USA

Submitted July 25, 1990

EMBL accession no. X53956

Members of the RTX family of protein toxins are secreted without the concomitant cleavage of a signal peptide<sup>1, 2</sup>. In *Pasteurella haemolytica* and *E. coli*, the structural gene for the RTX toxin is upstream of two genes, *lktB/hlyB* and *lktD/hlyD*, required for toxin secretion<sup>1, 3</sup>. In the preceding paper, we reported the sequence of the *lktB* gene from the leukotoxin operon of *Actinobacillus actinomycetemcomitans*; it shared a high degree of sequence similarity with the equivalent proteins from *P. haemolytica* and *E. coli*. This was unexpected since the RTX toxins from these bacteria are secreted while the leukotoxin from *A. actinomycetemcomitans* is reportedly retained in the periplasm<sup>4</sup>. Thus, to determine whether a sequence analogous to *lktD/hlyD* was present in *A. actinomycetemcomitans*, a 6 kb Bam HI fragment containing 2.5 kb of DNA 3' to *lktB* was cloned from strain JP-2. The region immediately downstream of the *lktB* gene was sequenced and an open reading frame was identified. The deduced protein, LktD, demonstrates significant homology to the equivalent polypeptides from related bacteria: it shares 61% identity with the LktD protein from *P. haemolytica*<sup>1</sup> and 68% identity with the HlyD protein from *E. coli*<sup>3</sup>. Identities are found in all three proteins at 52% of the residues. Moreover, the *lktD*

gene is expressed in *A. actinomycetemcomitans* as shown by Northern blot analysis using an *lktD*-specific probe (Spitznagel, Kraig, and Kolodrubetz, in preparation). The fact that the LktB and LktD proteins of *A. actinomycetemcomitans* are so similar to the leukotoxin/hemolysin secretion proteins from other organisms suggests that further investigations into the possible secretion of LktA from *A. actinomycetemcomitans* are necessary.

### ACKNOWLEDGEMENTS

We wish to thank Terry Dailey for valuable technical assistance. This work was supported by Public Health Service grants DE08521 and DE00152. J.M.G. is a Dentist-Scientist Awardee.

### REFERENCES

- Strathdee,C.A. and Lo,R.Y.C. (1989) *J. of Bacteriol.* **171**, 916–928.
  - Mackman,N., Nicaud,J.-M., Gray,L. and Holland,I.B. (1986) *Curr. Topics Microbiol. and Immunol.* **125**, 159–181.
  - Felmlee,T., Pellett,S. and Welch,R.A. (1985) *J. of Bacteriol.* **163**, 94–105.
  - DiRienzo,J.M. (1986) In A.J.L. Macario and E.C. de Macario (eds). *Monoclonal Antibodies against Bacteria* **3**, 250–293.
- ```

-12 GAACGAACGATAATGAAAACCTTGGCTTTGGCATTTGTATGATGTGCTTCCCGTTATAAAACGTTGGAATGAAACTTGGAAAATTCTGTAACAGCTTGGATT
   M K T W L L A L Y D V L S R Y K N V W N E T W K I R K Q L D S P V R

  103 GAAAAAGATGAAAATGATTTCTGCCCGCCTTGGAATTAAATTGAAACACCTGTATAATGCAACGCCCCCTTCAGGAAAATTGCACTAAAGCCTTGGGAGAGCAAAGAAATTAAACCGATTGAAAATTCTATGGTTAACTT
   E K D E N E F L P A H L E L I E T P V S N A P R F V S Y S I M L F L T L A I

  217 ATTGTTCTATCTTAGCAATGTGGAAAATTATCGCACCGGCTTCAGGAAAATTGCACTAAAGCCTTGGGAGAGCAAAGAAATTAAACCGATTGAAAATTCTATGGTTAAACATATT
   I V S I F S N V E I I A T A S G K F A L S G R S K E I K P I E N S L V K H I

  331 TTGTTGAAAGGGCCAATACGTGAAAAGGGTGAACCTATTGTTAAAAGCCTGACCTTGGTGCCTTGGGAGCGGATACGTTAAAACGAAAGACATCGCTTCTCAAGCTAAACTG
   F V K E G E Y V K K G E L L L K L T A L G A E A D T L K T K T S L S Q A K L

  445 GAGGAATTCTGTTATAAACTTCTTGGAAAGCTGTGAAAAGATCAATTACCAATTAGATTGTTCTAAATTGATTTCTAAATTGTTACCTTTATGACGGAAAAGCATCAAAGAGTG
   E E F R Y K S L L E A V E K D Q L P I L D F S K I D L P F M T E N D Q K R V

  559 ACTTTACTGATTGAAGAACATTCTCTACTTGGCAAAAACAACGCCATCAGAAAACCTGAACTTAAATAAAAAGAAGCGGGAAAACCTGAGTTATGGCACGAATAAAAAA
   T L L I E E Q F S T W Q K Q R H Q K T L N L N K K E A E K L S Y L A R I K K

  673 TATGAAGGTCTGATTAATACAGAGCAAGCTGGATTGGATGTTAGGCCATTATAAGGAACATGCTATTGCAAAACATACAGTTAGATGAAGAGAATAATATCAGGAT
   Y E G L I N T E Q V R L D D F R A L Y K E H A I A K H T V L D E E N K Y Q D

  787 GCAATCAATGAGCTTGGGTATAAGGGCGATTAACTGCAAGTTGAAACGAAGTTTATTGGCAAAAGAAGGCCAGGAAATTAGTCACACAATTGTTAGATGAAGAGAATAATATCAGGAT
   A I N E L E V Y K A S L H M Q V E N E V L L A K E E Q E L V T Q L F K N D I L

  901 GACAAGCTAAAACAGGGCAGCGATAATGTGAATTATTGACATTGCAACTGGACAAAACAATCAACGCCAGCAAGTCTCTGAAATTGAGCTCTGTATCAGGTACTGTACAA
   D K L K Q A T D N V N L L T F E L D K N N Q R Q Q V S E I R A P V S G T V Q

  1015 CAATTAAAAGTTACACAATAGATGGCGCTTGTACTACGGCTGAAACATTAATGGTAGTTGTCAGGAAAGATTCCCTGAAAGTCACTGCACTTATTCAAATAAAGATATT
   Q L K V H T I D G V V T T A E T L M V V V P E E D S L E V T A L I Q N K D I

  1129 GGTTTTGTGAAAGAAGGGCAAGAGGTTGATTAAGGTGAAAGCCTTCCCCTATACCCGTTATGGTTATTAACGGGAAAAGTGAAGAAATTACTTGTGCGATTGAAACAT
   G F V K E G O E V V I K V E A F P Y T R Y G Y L T G K V K N I T L D A I E H

  1243 CCAAAGCTTGGACTGTATTTAATACGATTATTGAAAGTAAACACTTACCGAAGAAAAGAAAATCCCACTTCCCGCAGGGATGGAATTACTGCAGAAATTAG
   P K L G L V F N T I E L D K K T L S T E E K E I P L S A G M E I T A E I K

  1357 ACAGGCATGAGAAGCGTTATAAGCTATCTACTTAGCCATTAGAAGAATCTATTGATAAAAGTTAGAGAAGCTTAAATATAAGAGATAATAACATTCT
   T G M R S V I S Y L L S P L E E S I D K S L R E R term

```

\* To whom correspondence should be addressed