Accepted Manuscript

Recombinant expression and predicted structure of parborlysin, a cytolytic protein from the Antarctic heteronemertine *Parborlasia corrugatus*

Matej Butala, Daniel Šega, Blaž Tomc, Zdravko Podlesek, William R. Kem, Frithjof C. Küpper, Tom Turk

PII: S0041-0101(15)30101-X

DOI: 10.1016/j.toxicon.2015.09.044

Reference: TOXCON 5214

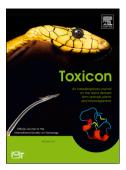
To appear in: Toxicon

Received Date: 15 July 2015

Revised Date: 25 September 2015 Accepted Date: 29 September 2015

Please cite this article as: Butala, M., Šega, D., Tomc, B., Podlesek, Z., Kem, W.R., Küpper, F.C., Turk, T., Recombinant expression and predicted structure of parborlysin, a cytolytic protein from the Antarctic heteronemertine *Parborlasia corrugatus*, *Toxicon* (2015), doi: 10.1016/j.toxicon.2015.09.044.

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.



ACCEPTED MANUSCRIPT

- 1 Recombinant expression and predicted structure of parborlysin, a cytolytic protein from
- 2 the Antarctic heteronemertine Parborlasia corrugatus

3

- ¹Matej Butala, ¹Daniel Šega, ¹Blaž Tomc, ¹Zdravko Podlesek, ²William R. Kem, ³Frithjof C.
- 5 Küpper and ^{1,*}Tom Turk

6

- ¹Department of Biology, Biotechnical Faculty, University of Ljubljana, Ljubljana, Slovenia
- 8 ²Department of Pharmacology and Therapeutics, Collge of Medicine, University of Florida,
- 9 Gainesville, FL 32610, USA
- ³ Oceanlab, University of Aberdeen, Main Street, Newburgh, AB41 6AA, Scotland, UK

11 12

*Corresponding author, email address: tom.turk@bf.uni-lj.si.

13

14

- Abstract
- 15 The heteronemertine *Parborlasia corrugatus* contains a cytolytic protein, parborlysin, which
- after extensive purification was found by Edman sequencing to be a mixture of several
- 17 homologs. To investigate this microheterogeneity and enable the analysis of single toxins, we
- have obtained seven parborlysin isoform genes from *P. corrugatus* collected in Antarctica.
- 19 Total RNA was isolated from the homogenized head region and parborlysin genes were
- 20 identified from a cDNA library using degenerate primers. The translated sequences reveal that
- 21 the isoforms are ~10 kDa basic (pI~10) proteins of which all but one harbour six cysteine
- residues. We generated a model of the three dimensional structure of parborlysins, which
- suggests that they are composed of five alpha-helical segments that include large, exposed
- 24 hydrophobic surfaces. Finally, we constructed plasmids and inserted them into E. coli to
- obtain overexpressed amino- or carboxy-terminal polyhistidine-tagged parborlysin isoforms
- fused to the third domain of the *E. coli* periplasmic-protein TolA to facilitate toxin isolation.
- One of the isoforms adversely affected growth in the E. coli expressing it. Although we
- succeeded in isolating one of the recombinant parborlysin constructs, it lacked haemolytic
- 29 activity.

30

- 31 **Key words:** Cytolysin; Nemertean; Nemertine, Nucleotide sequence; *Parborlasia corrugatus*;
- 32 Parborlysin; Toxin

33

34

Download English Version:

https://daneshyari.com/en/article/8395127

Download Persian Version:

https://daneshyari.com/article/8395127

<u>Daneshyari.com</u>