



Abstract

An Endogenous Retrovirus from Human Hookworm Encodes an Ancient Phlebovirus-Like Class II Envelope Fusion Protein [†]

Monique Merchant 1,2,*, Carlos P. Mata 3 and Yorgo Modis 1,2

- Molecular Immunity Unit, Department of Medicine, University of Cambridge, MRC Laboratory of Molecular Biology, Cambridge Biomedical Campus, Cambridge CB2 0QH, UK; ymodis@mrc-lmb.cam.ac.uk
- ² Cambridge Institute of Therapeutic Immunology & Infectious Disease (CITIID), Department of Medicine, University of Cambridge, Cambridge CB2 0AW, UK
- ³ Astbury Centre for Structural Molecular Biology, School of Molecular and Cellular Biology, Faculty of Biological Sciences, University of Leeds, Leeds LS2 9JT, UK; c.perezmata@leeds.ac.uk
- * Correspondence: monique.merchant.81@gmail.com
- † Presented at Viruses 2020 Novel Concepts in Virology, Barcelona, Spain, 5–7 February 2020.

Published: 10 June 2020

Abstract: Within the parasitic nematode *Ancylostoma ceylanicum*, a ~20 million-year-old Bel/Pao LTR retrotransposon encodes an ancient viral class II envelope fusion protein termed Atlas Gc. Typically, retroviruses and related degenerate retrotransposons encode a hemagglutinin-like class I envelope fusion protein. A subset of Bel/Pao LTR retrotransposons within the phylum *Nematoda* have acquired a phlebovirus-like envelope gene and utilized the encoded fusion machinery to escape the genome as intact exogenous retroviruses. This includes *C. elegans* retroelement 7 virus which was recently reclassified as a member of the genus *Semotivirus*. A 3.76 Å cryoEM reconstruction confirms Atlas Gc as a closely related phleboviral homologue and class II fusion protein in a novel case of gene exaptation. Preliminary biophysical and biochemical characterization indicate Atlas Gc functions under specific physiological conditions targeting late-endosomal membranes, much like modern viral class II envelope fusion proteins. Phylogenetic analyses support the reclassification of the Atlas endogenous retrovirus and five other *A. ceylanicum* ERVs as novel semotiviruses of *Belpaoviridae* of the new viral order of reverse-transcribing viruses *Ortervirales*.

Keywords: transposable element; retrotransposon; endogenous retrovirus (ERV); envelope protein; membrane fusion; horizontal gene transfer (HGT); virus evolution; gene exaptation; nematode



© 2020 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (http://creativecommons.org/licenses/by/4.0/).