

The RNA virome of parasitic nematodes

Shannon Quek^{1*}, Amber Hadermann^{2*}, Yang Wu^{1*}, Lander De Coninck³, Shrilakshmi Hegde¹, Jordan R. Boucher¹, Jessica Cresswell¹, Ella Foreman¹, Andrew Steven¹, E. James LaCourse¹, Stephen A. Ward¹, Samuel Wanji^{4,5}, Grant L. Hughes⁶, Edward I. Patterson⁷, Simon C. Wagstaff¹, Joseph D. Turner¹, Eva Heinz⁸, Jelle Matthijssens³, Robert Colebunders², Mark J. Taylor¹

1. Centre for Neglected Tropical Diseases, Department of Tropical Disease Biology, Liverpool School of Tropical Medicine, Pembroke Place, Liverpool, L3 5QA, UK.
 2. Global Health Institute, University of Antwerp, Doornstraat 331, 2610 Antwerp, Belgium.
 3. KU Leuven, Department of Microbiology, Immunology & Transplantation, Rega Institute, Leuven, Belgium.
 4. Parasite & Vector Biology Research Unit, Department of Microbiology & Parasitology, University of Buea, Buea, Cameroon.

5. Research Foundation for Tropical Diseases and the Environment (REFOTDE), Buea, Cameroon
 6. Centre for Neglected Tropical Diseases, Departments of Tropical Disease Biology & Vector Biology, Liverpool School of Tropical Medicine, Pembroke Place, Liverpool, L3 5QA, UK.
 7. Department of Biological Sciences, Brock University, St. Catharines, ON, L2S 3A1, Canada.
 8. Departments of Vector Biology & Clinical Sciences, Liverpool School of Tropical Medicine



Shannon.Quék@lstm.ac.uk

*Authors contributed equally to this work.

1. Background

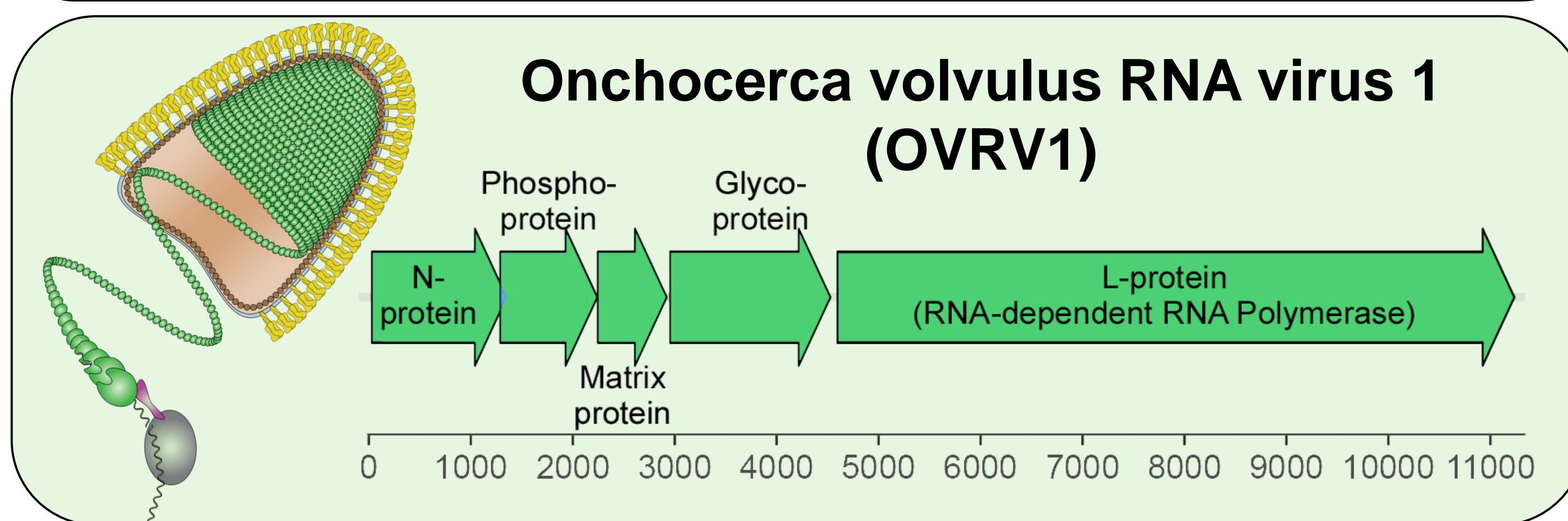
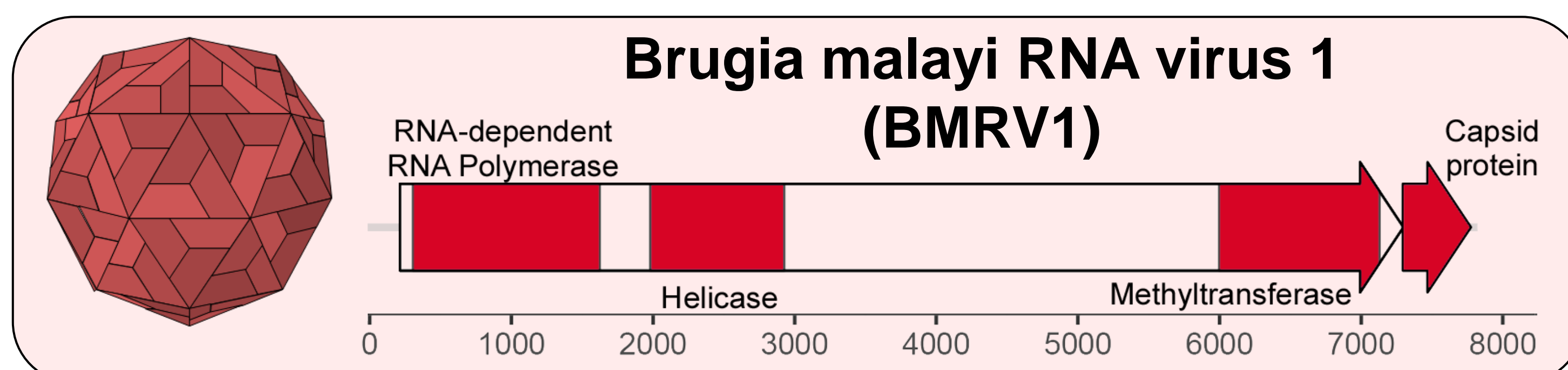
- Parasitic nematodes infect billions of humans and animals worldwide
- They cause significant impacts on mental and physical health, as well as economic productivity
- Basic biology knowledge for several species is lacking, e.g. role of microbiome, and specifically their “RNA virome”

2. Results

Order	Nematode Species	Definitive host	Viruses
Spirurida	<i>Brugia malayi</i>	Humans, monkeys, cats, dogs	1
	<i>Brugia pahangi</i>	Cats, dogs	2
	<i>Onchocerca volvulus</i>	Humans	6
	<i>Onchocerca ochengi</i>	Cattle	11
	<i>Anguillicola crassus</i>	Eels	1
Ascaridida	<i>Ascaris suum/lumbricoides</i>	Pigs, humans	4
	<i>Toxocara canis</i>	Dogs	7
	<i>Ancylostoma ceylanicum</i>	Humans, dogs, cats	4
	<i>Angiostrongylus cantonensis</i>	Rodents	2
	<i>Dictylocalus viviparus</i>	Cattle	2
Strongyloida	<i>Haemonchus contortus</i>	Ruminants	2
	<i>Heligmosomoides polygyrus</i>	Rodents	3
	<i>Nippostrongylus brasiliensis</i>	Rodents	5
	<i>Oesophagostomum dentatum</i>	Swine	2
	<i>Ostertagia ostertagi</i>	Ruminants	3
Trichinellida	<i>Teladorsagia circumcincta</i>	Sheep, goats	1
	<i>Encapsulated Trichinella spp.</i>	Human, swine, ursine	2
	<i>Trichinella pseudospiralis</i>	Human, swine, ursine	7
	<i>Trichuris muris</i>	Mouse	10
	<i>Trichuris suis</i>	Swine	5
	<i>Trichuris trichiura</i>	Humans, primates	3

Bioinformatic analysis of previously published transcriptome data identified 91 different RNA viruses across 28 species of parasitic nematodes of medical and veterinary importance.

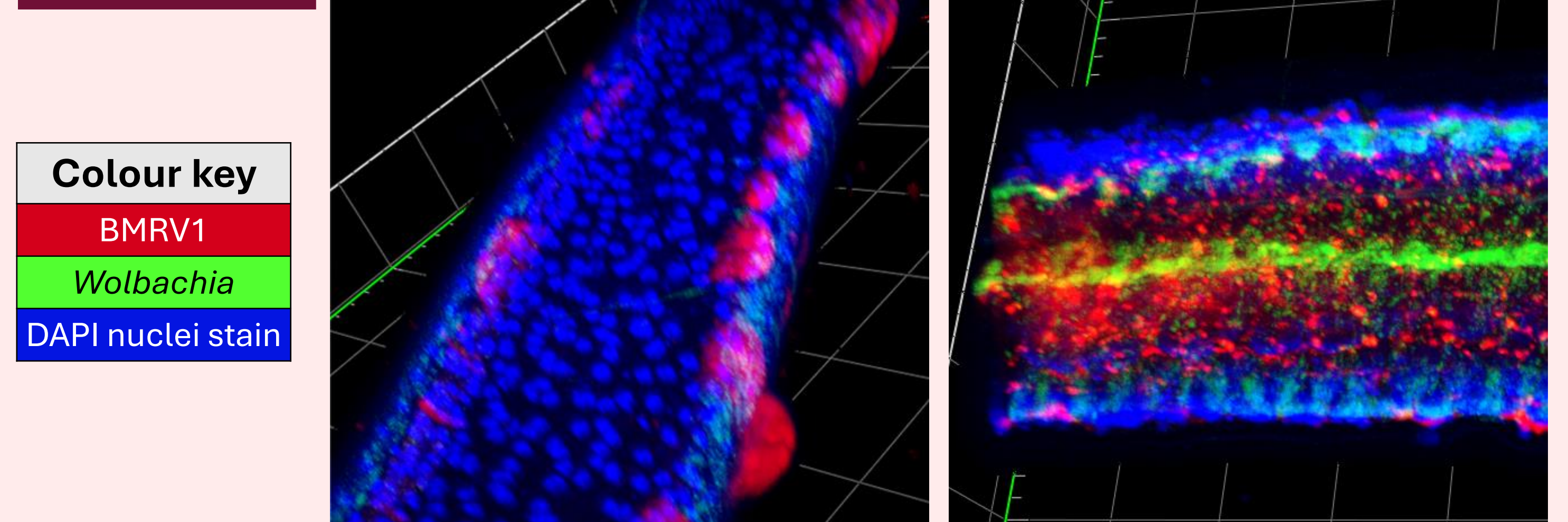
Further characterization of two viruses



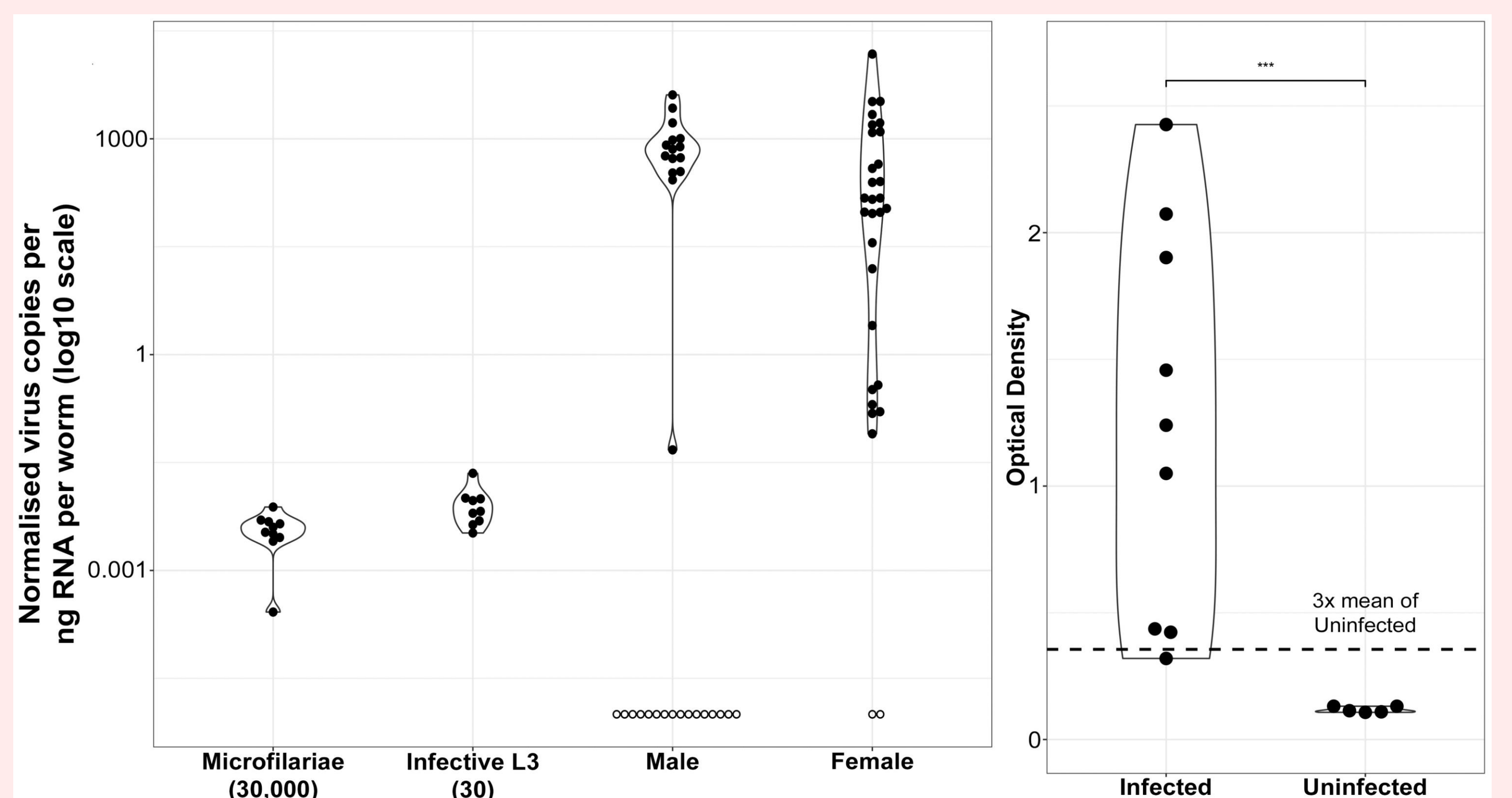
3. Major conclusions

- From published transcriptome data, we have identified 91 RNA viruses from 28 different parasitic nematode species
- We find extensive diversity and conserved global spread of virus-nematode associations across multiple continents, suggesting ancestral acquisition and host-virus co-evolution
- Viruses of *Brugia malayi* (BMRV1) and *Onchocerca volvulus* (OVRV1) are found in the reproductive tract suggesting sexual/vertical transmission
- BMRV1 RNA can be found in epicuticular inflations in older parasites
- BMRV1 and OVRV1 elicit antibody responses from the vertebrate host, demonstrating direct exposure to the immune system
- Interactions between parasite-virus are unknown, with potential effects on parasite biology/disease pathogenesis unknown
- Identified viruses are a novel paradigm for understanding parasitic nematode diseases

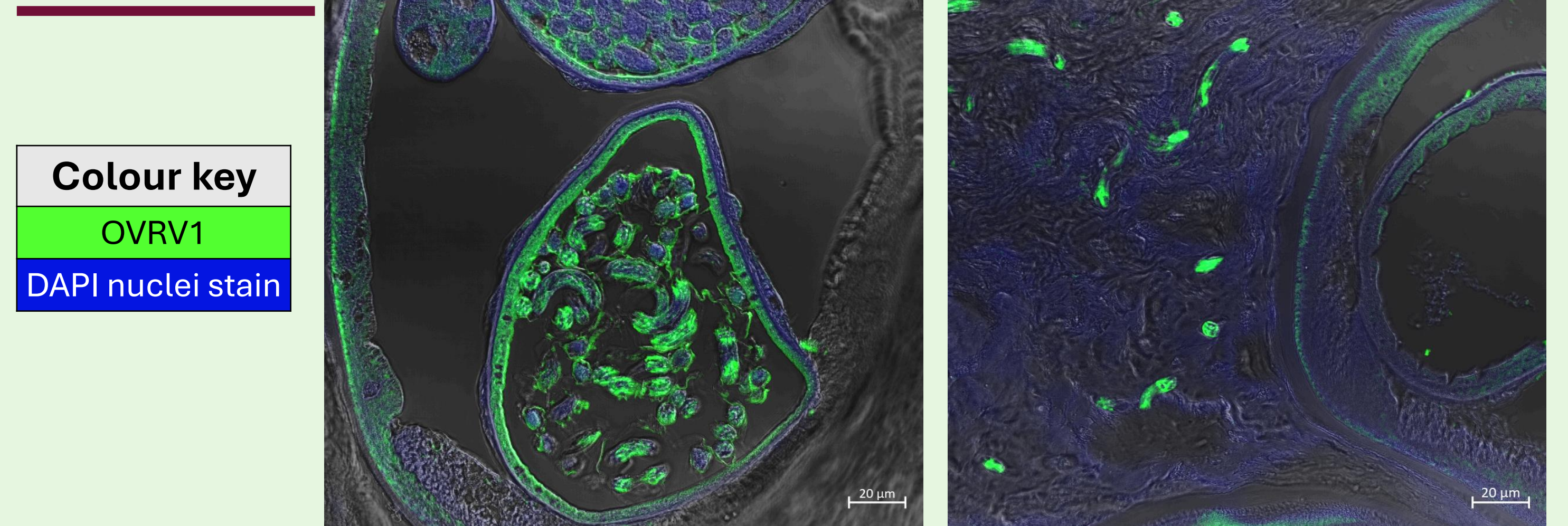
2a. BMRV1



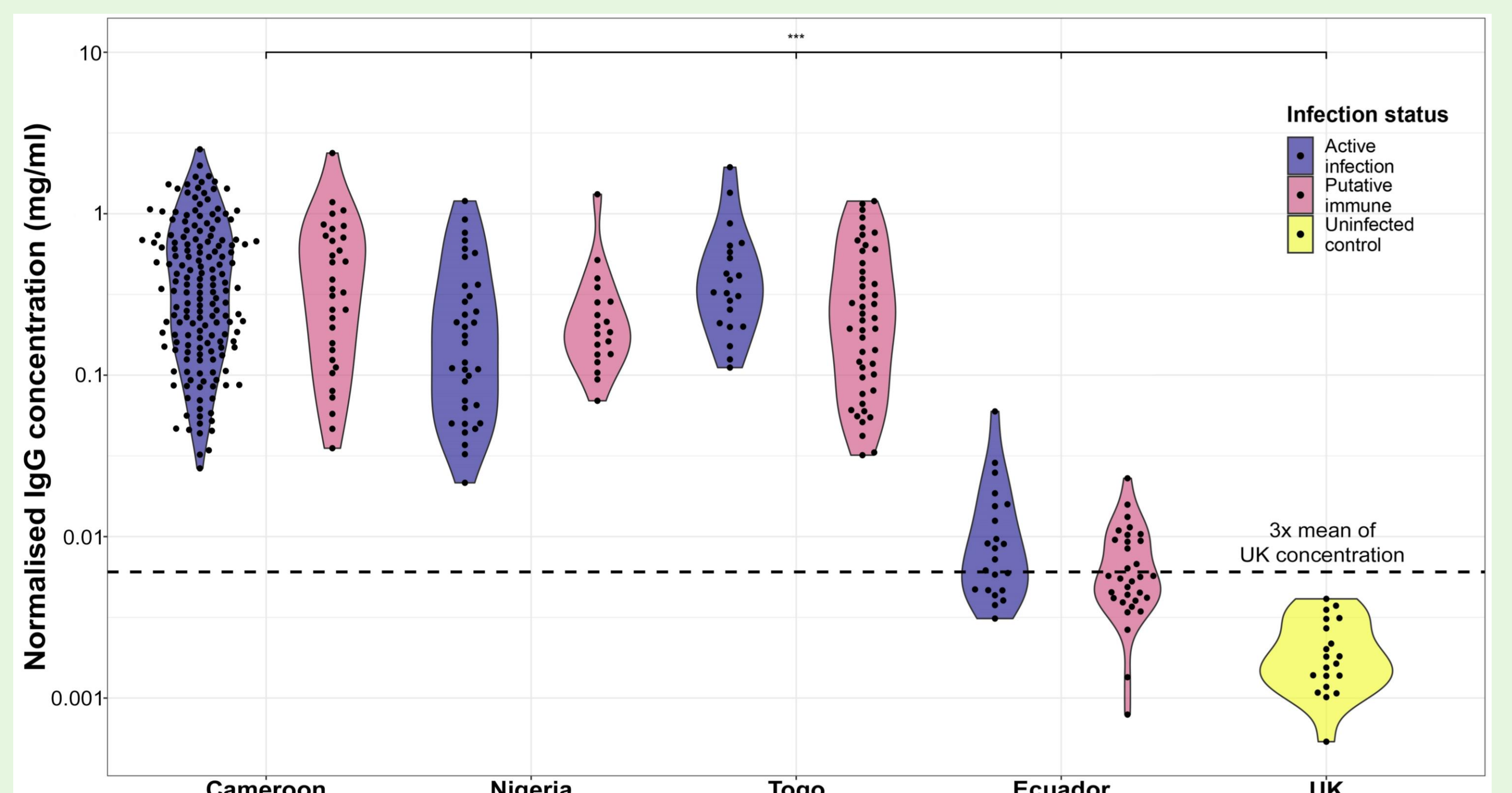
BMRV1 tissue tropism in *Brugia malayi* – confocal microscopy localizes viral signal to ‘worm warts’ within epicuticular inflations (left panel) and within reproductive tissues (right panel).



2b. OVRV1



OVRV1 tissue tropism in *Onchocerca volvulus* – OVRV1 IFA localizes viral signal to reproductive tissues and developing embryos (left) and in microfilariae within nodular tissues (right).



References

- Shi, M. et al. *Redefining the invertebrate RNA virosphere*. Nature 540, 539–543 (2016).
- Dheilly, N. M., Lucas, P., Blanchard, Y. & Rosario, K. A *World of Viruses Nested within Parasites: Unravelling Viral Diversity within Parasitic Flatworms (Platyhelminthes)*. Microbiol. Spectr. 10, (2022).
- Félix, M. A. & Wang, D. *Natural Viruses of Caenorhabditis Nematodes*. Annu. Rev. Genet. 53, 313–326 (2019).

Research was funded by the Liverpool School of Tropical Medicine’s Director’s Catalyst Fund.

Scan for preprint

