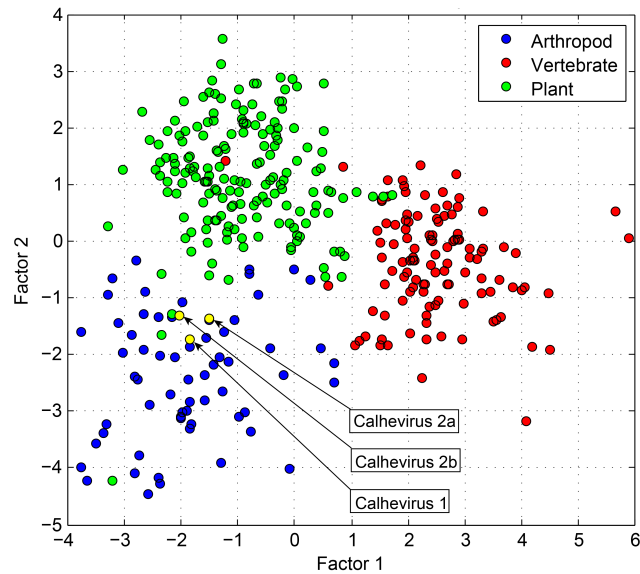




HOKKAIDO UNIVERSITY

Title	Metagenomic analysis of the shrew enteric virome reveals novel viruses related to human stool-associated viruses
Author(s)	Sasaki, Michihito; Orba, Yasuko; Ueno, Keisuke et al.
Citation	Journal of General Virology, 96(2), 440-452 https://doi.org/10.1099/vir.0.071209-0
Issue Date	2015-02
Doc URL	https://hdl.handle.net/2115/60630
Type	journal article
File Information	Supplementary Figure.pdf





Supplementary Figure S1. Canonical discriminat analysis of Calhevirus 2a and Calhevirus 2b.

The graph shows the separation of groups using the two most influential factors based on mononucleotide and dinucleotide frequencies. The genome sequences of 112 vertebrate-derived viruses, 64 arthropod-derived viruses, 171 plant-derived viruses were used as reference sequences for the analysis and shown as red circles, blue circles and green circles, respectively. Calhevirus 1, Calhevirus 2a and Calhevirus 2b are shown as yellow circles.