

Supplementary Table 11. Information regarding *Pegivirus hominis* identified by metagenomic shotgun sequencing.

Sample ID	Total reads*	Reads mapped to virus *	RPM*	Genome coverage (NT)*	Average depth	Identity (NT)*	Genotype	Sex	Age	Date	Location	GenBank_ID
CRN_174	2,446,090	30	13.9	14.1%	0.3	80%	NA	F	23	2022-07-21	Manaus, AM	SAMN52632121
CRN_188	2,028,368	1,716	846.3	97.9%	11.5	96.9%	II	M	26	2022-08-24	Manaus, AM	SAMN52632123

RPM: reads per million; NT: nucleotide; F: female; M: male; AM: Amazonas, Brazil.

*Total reads, reads mapped to virus, and RPM were obtained from CZID (Chan Zuckerberg ID platform).

Genome coverage, average depth, and identity values (NT) were calculated based on mapping to the *Pegivirus hominis* 1 reference genome (ON340918.1). NA: not applicable (not sequenced or not detected).