
Supplementary information

Genomes of six viruses that infect Asgard archaea from deep-sea sediments

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Supplementary Information for
**Genomes of six viruses that infect Asgard archaea from deep-sea
sediments**

Ian M. Rambo, Marguerite V. Langwig, Pedro Leão, Valerie De Anda, and Brett J. Baker

Correspondence to: brett_baker@utexas.edu

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Fig. 2 abbreviations

AdoMet MTase, S-adenosyl-l-methionine-dependent methyltransferase; *cas1*, CRISPR-associated endonuclease Cas1; *cas2*, CRISPR-associated endoribonuclease Cas2; *cas3*, CRISPR-associated endonuclease/helicase Cas3; *cas4*, CRISPR-associated exonuclease Cas4; *cas5*, CRISPR system Cascade subunit Cas5; *cas6*, CRISPR-associated endonuclease Cas6; *cas7*, CRISPR-Cas Type I effector complex subunit Cas7; *cas8*, CRISPR-associated protein Cas8; *csm2*, Type III CSM-effector complex small subunit Csm2; Coil, coiled-coil protein; *csm3*, Type III RAMP superfamily CSM-effector complex Csm3; *csm4*, Type III RAMP superfamily CSM-effector complex Csm4; *csm5*, Type III RAMP superfamily CSM-effector complex Csm5; *csx1*, CRISPR system endoribonuclease Csx1; *csx14*, Subtype III-U associated protein Csx14; *cft2*, Cft2 family RNA processing exonuclease; DUF 362, domain of unknown function 362; EamA, probable amino-acid metabolite efflux pump; HEPN, higher eukaryotes and prokaryotes nucleotide-binding domain; kb, kilobase; NT, nucleotidyl transferase; NTPH, nucleoside triphosphate hydrolase; TA, toxin-antitoxin; vWA, Von Willebrand factor type A protein; WH, winged helix DNA binding domain.