

Supplementary table 3 Accession numbers and titles of the viral sequences deposited by Gil P. et al. in the NCBI GenBank database along with their corresponding SRS data accession numbers in the NCBI SRA database.

GenBank virus accession number	Virus title	SRA sample accession number(s)
OL870603.1	Culex Iflavi-like virus 4 isolate Tunisia2015, complete genome	SRR18056563, SRR18056564
OL870604.1	Culex Iflavi-like virus 4 isolate Italy2015, complete genome	SRR18056567, SRR18056568
OL870605.1	Culex Iflavi-like virus 4 isolate France2015, complete genome	SRR18056569, SRR18056570
OL870606.1	Culex Iflavi-like virus 4 isolate Spain2015, complete genome	SRR18056565, SRR18056566
OL870607.1	Culex Iflavi-like virus 4 isolate Egypt2015, complete genome	SRR18056571, SRR18056572
OL913368.1	Alphamesonivirus 1 isolate France2015 genomic sequence	SRR18056569, SRR18056570
OL913369.1	Alphamesonivirus 1 isolate Spain2015 genomic sequence	SRR18056565, SRR18056566
OL913370.1	Alphamesonivirus 1 isolate Italy2015 genomic sequence	SRR18056567, SRR18056568
OL913371.1	Alphamesonivirus 1 isolate Tunisia2015 genomic sequence	SRR18056563, SRR18056564
OL913372.1	Alphamesonivirus 1 isolate Egypt2015 genomic sequence	SRR18056571, SRR18056572
OP057135.1	Alilaet virus isolate Egypt_2015 hypothetical protein and RNA dependent RNA polymerase genes, complete cds; and hypothetical protein gene, partial cds	SRR18056571, SRR18056572
OP264878.1	Serbia narna-like virus 1 isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264879.1	Negev-like virus #174 isolate France_2015 genomic sequence	SRR18056569, SRR18056570
OP264880.1	Guadeloupe Culex rhabdovirus isolate Egypt_2015 genomic sequence	SRR18056571, SRR18056572
OP264881.1	Serbia reo-like virus 1 isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264882.1	Hubei virga-like virus 2 isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264883.1	Asum virus isolate France_2015 genomic sequence	SRR18056569, SRR18056570
OP264884.1	Eskilstorp virus isolate France_2015 genomic sequence	SRR18056569, SRR18056570
OP264886.1	Hubei reo-like virus 7 isolate Egypt_2015 genomic sequence	SRR18056571, SRR18056572
OP264887.1	Aedes vexans iflavirus isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264888.1	Vinslov virus isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264889.1	Culex orthophasmavirus isolate Egypt_2015 genomic sequence	SRR18056571, SRR18056572
OP264890.1	Culicidavirus quitotaense isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264891.1	Nackenback virus isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566
OP264892.1	Vivastbo virus isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566
OP264893.1	Culex pipiens-associated Tunisia virus isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264894.1	Rinkaby virus isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264895.1	Hubei mosquito virus 4 isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264896.1	Dezidougou virus isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264897.1	Broome reo-like virus 1 isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264898.1	Culex mononega-like virus 2 isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566
OP264899.1	Thassos sobemo-like virus isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566
OP264900.1	Umatilla virus isolate Italy_2015 genomic sequence	SRR18056567, SRR18056568
OP264901.1	Culex Virga-like virus isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566
OP264902.1	Culex rhabdovirus isolate France_2015 genomic sequence	SRR18056569, SRR18056570
OP264903.1	Aphis gossypii virus isolate Tunisia_2015 genomic sequence	SRR18056563, SRR18056564
OP264904.1	Serbia picorna-like virus 1 isolate France_2015 genomic sequence	SRR18056569, SRR18056570
OP264905.1	Biggievirus Mos11 isolate France_2015 genomic sequence	SRR18056569, SRR18056570
OP264906.1	Shuangao Insect Virus 3 isolate Egypt_2015 genomic sequence	SRR18056571, SRR18056572
OP264907.1	Culex modestus totivirus isolate Tunisia_2015 genomic sequence	SRR18056563, SRR18056564
OP264908.1	Valmbacken virus isolate France_2015 genomic sequence	SRR18056569, SRR18056570
OP264909.1	Xanthi chryso-like virus isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264910.1	Negev virus isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264911.1	Yongsan picorna-like virus 2 isolate Egypt_2015 genomic sequence	SRR18056571, SRR18056572
OP264912.1	Alphachrysovirus shuangaoense isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566
OP264913.1	Big Sioux River virus isolate Tunisia_2015 genomic sequence	SRR18056563, SRR18056564
OP264914.1	Daeseongdong virus 2 isolate France_2015 genomic sequence	SRR18056569, SRR18056570
OP264915.1	Yongsan iflavirus 1 isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264916.1	Daeseongdong virus 1 isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264917.1	Hubei picorna-like virus 23 isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566
OP264918.1	Culex pipiens orthomyxo-like virus isolate Tunisia_2015 genomic sequence	SRR18056563, SRR18056564
OP264919.1	Fako virus isolate Mediterranean_2015 genomic sequence	SRR18056563, SRR18056564, SRR18056565, SRR18056566, SRR18056567, SRR18056568, SRR18056569, SRR18056570, SRR18056571, SRR18056572
OP264920.1	Wuhan Mosquito Virus 4 isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566
OP264921.1	Sonnbo virus isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566
OP264922.1	Culicine-associated Z virus isolate Tunisia_2015 genomic sequence	SRR18056563, SRR18056564
OP264923.1	Almendravirus menghai isolate Tunisia_2015 genomic sequence	SRR18056563, SRR18056564
OP264924.1	Culex modestus orthomyxo-like virus isolate France_2015 genomic sequence	SRR18056569, SRR18056570
OP264925.1	Culex mononega-like virus 1 isolate Spain_2015 genomic sequence	SRR18056565, SRR18056566