

Table S4: PCR primer sequences

Quantitative PCR primers		
HD86	Dnmt2-RA forward	5'- GTTCGTCGATCATGATACATGCG -3'
HD87	Dnmt2-RC forward	5'- CTATTTAGTGGCATTGGCGGC-3'
HD88	Dnmt2 reverse	5'- GCAGCATGTTGGCCTGAAGC -3'
HD174	rp49 forward	5'- CGGATCGATATGCTAAGCTGT -3'
HD175	rp49 reverse	5'- GCGCTTGTTCGATCCGTA -3'
HD672	Drosocin forward	5'-GTCCACCACTCCAAGCAC- 3'
HD673	Drosocin reverse	5'-CTTTAGGCGGGCAGAATGGG -3'
HD674	Metchnikowin forward	5'-CGCCACCGAGCTAAGATGC -3'
HD675	Metchnikowin reverse	5'-GGTTAGGATTGAAGGGCGAC -3'
HD611N	Vago forward	5'-TGCAACTCTGGGAGGATAGC -3'
HD821N	Vago reverse	5'-AATTGCCCTGCGTCAGTTT -3'
HD822N	Vir-1 forward	5'-GATCCCAATTTTCCCATCAA -3'
HD823N	Vir-1 reverse	5'-GATTACAGCTGGGTGCACAA-3'
HD824N	IM1 forward	5'-TCCACTGTGCCCCGATCC -3'
HD825N	IM1 reverse	5'-CTTGGGTTGAAACTTCCTACTTGC -3'
HD826N	Cecropin A1 forward	5'-GCGTTGGTCAGCACACTC -3'
HD827N	Cecropin A1 reverse	5'-CAGTTGCGGGCAGATTGG -3'
HDT047	DCV forward	5'-TCATCGGTATGCACATTGCT -3'
HDT048	DCV reverse	5'-CGCATAACCATGCTCTTCTG -3'
HDT053	Turandot A forward	5'-GGTTTGCTTCAGCGTTCCAAAAAGTCATAACC -3'
HDT054	Turandot A reverse	5'-ATTAAACAATATTAACCAGTGAATAATTGAG -3'
ZD123	Nora virus forward	5'-AACCTCGTAGCAATCCTCTCAAG -3'
ZD124	Nora virus reverse	5'-TTCTTGTCCGGTGTATCCTGTATC -3'
ZD119	DXV forward	5'- GGAGTTGAAGCCACGGTTTG-3'
ZD120	DXV reverse	5'- GACGATCTTGCCAGTTGGCTCATCG-3'
ZD121	Sigma virus forward	5'-ATGTAACCTCGGGTGTGACAG -3'
ZD122	Sigma virus reverse	5'- CCTTCGTTTCATCCTCCTGAG-3'
Diagnostic qPCR primers		
ZD245	fwd A. pomorum	5'-TGTTTCCGCAAGGGACCTCT-3'
ZD246	rev A. pomorum	5'-AGAGTGCCAGCCCAACCTGA-3'
ZD247	fwd A. tropicalis	5'-TAGCTAACGCGATAAGCACA-3'
ZD248	rev A. tropicalis	5'-ACAGCCTACCCATACAAGCC-3'
ZD249	fwd L. fructivorans	5'-AACCTGCCAGAGAAGGGGA-3'
ZD250	rev L. fructivorans	5'-GCGCCGCGGATCCATCCAAA-3'
ZD251	fwd L. plantarum	5'-TGTTCTCAGTCCCAATGTGGCCG-3'
ZD252	rev L. plantarum	5'-GGCTATCACTTTTGGATGGTCCCGC-3'
Northern probes		
HD174	rp49 forward	5'- CGGATCGATATGCTAAGCTGT -3'
HD175	rp49 reverse	5'- GCGCTTGTTCGATCCGTA -3'
HD672	Drosocin forward	5'-GTCCACCACTCCAAGCAC- 3'
HD673	Drosocin reverse	5'-CTTTAGGCGGGCAGAATGGG -3'
HD674	Metchnikowin forward	5'-CGCCACCGAGCTAAGATGC -3'
HD675	Metchnikowin reverse	5'-GGTTAGGATTGAAGGGCGAC -3'
HDT047	DCV forward	5'-TCATCGGTATGCACATTGCT -3'
HDT048	DCV reverse	5'-CGCATAACCATGCTCTTCTG -3'
stem loop RT primer		
HD649	tRNA-Asp reverse	5'-CTCAACTGGATTGGCTNNNNNGATAAATCCAGT TGAGTGGCTCCC-3'
RIP primers		
HD254	tRNAAsp fwd	5'-TCCTCGATAGTATAGT-3'
HD616	tRNAAsp rev	5'-CTCAACTGGATTGGCT-3'
ZD127	DCV 698-1435 fwd	5'-GTGTATGATCTATACGCAC-3'
ZD128	DCV 698-1435 rev	5'-CAATCTTCATAATATCCTC-3'
ZD137	DCV 4443-5163 fwd	5'-GAAGAATATTTCTCAC-3'
ZD138	DCV 4443-5163 rev	5'-GGCGAAATGCAACTACAAAATG-3'
ZD149	DCV 8841-9264 fwd	5'-CAATTAATAGTGTCTTGCCTG-3'
ZD150	DCV 8841-9264 rev	5'-AGAAAACAACATATTTTAAATTC-3'