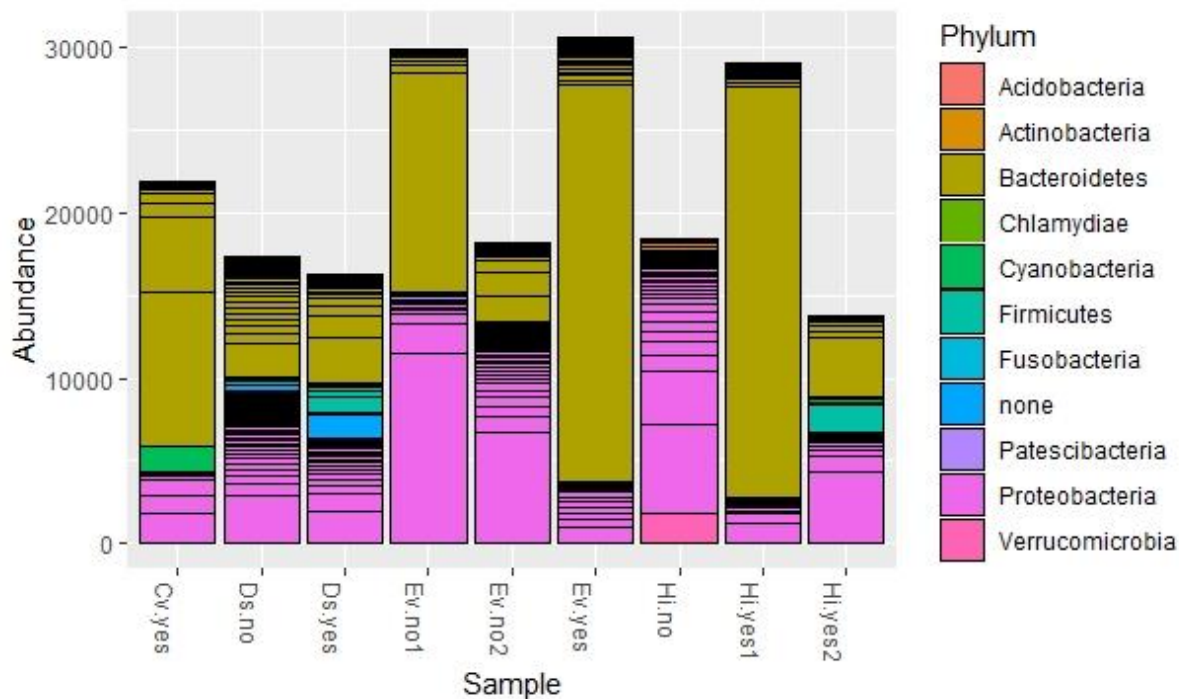
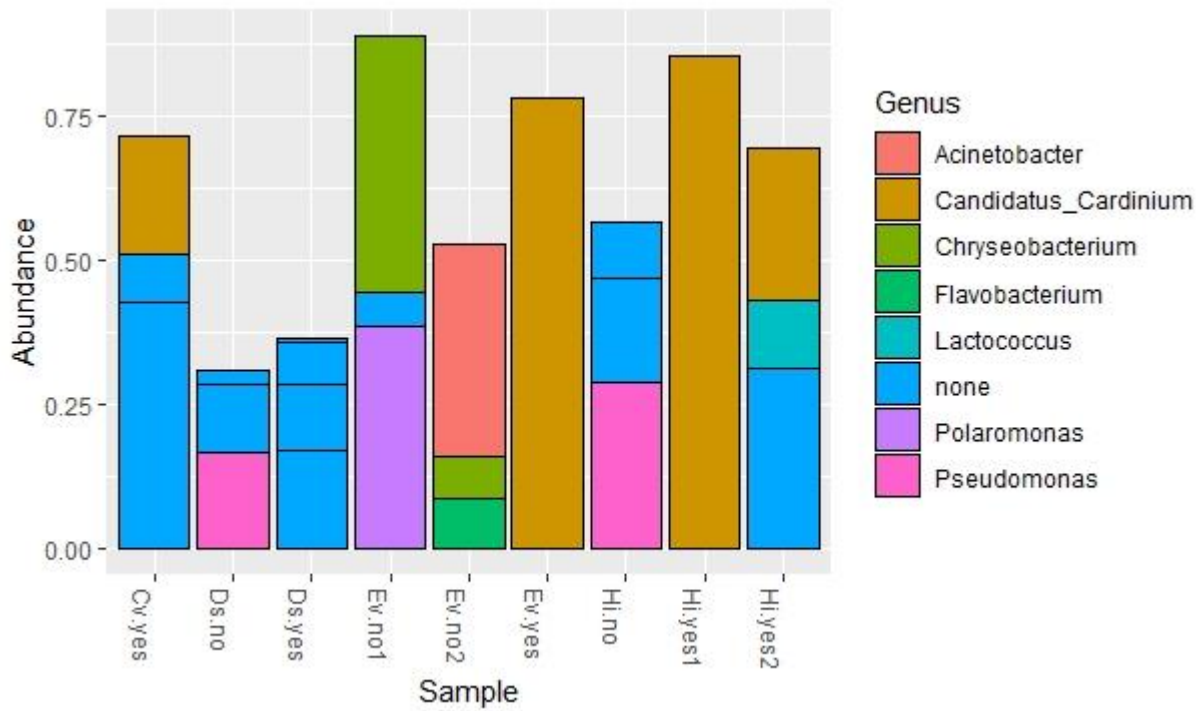


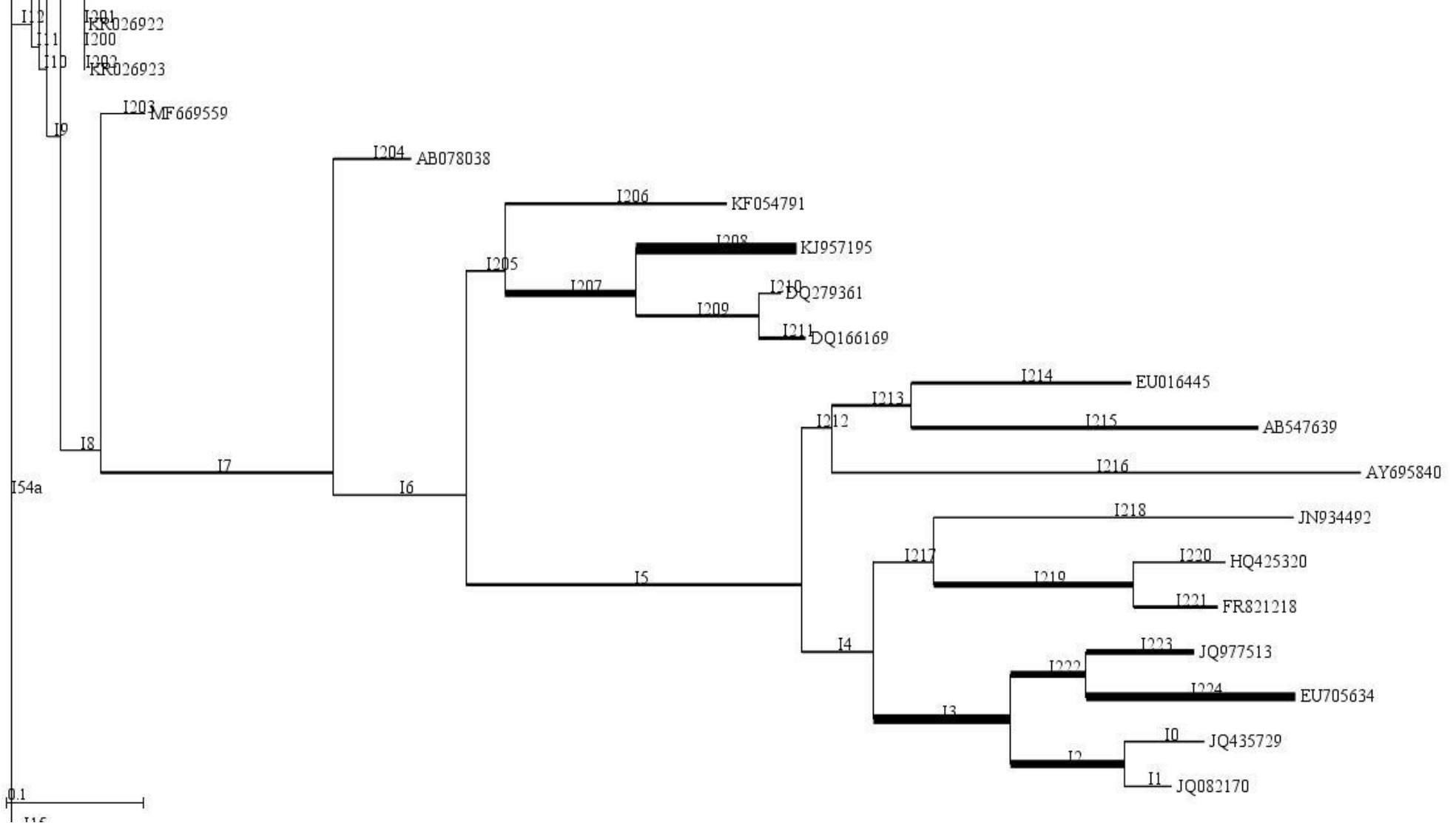
### Supplementary material

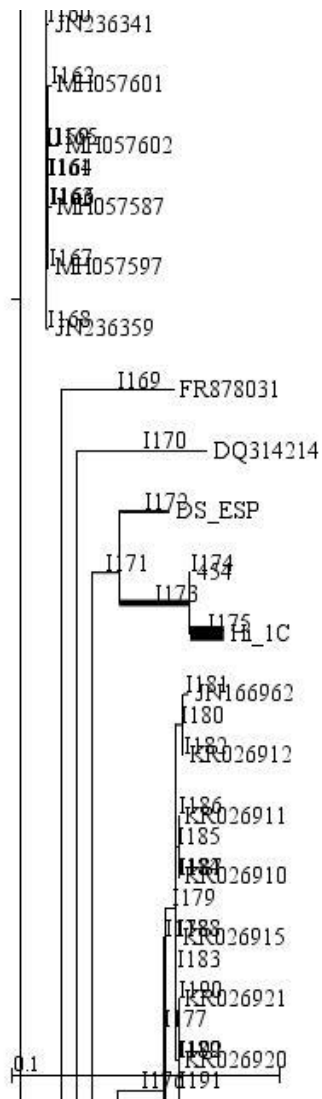


**Figure S1A: Cumulative bar charts of non-marine ostracod microbiomes.** Ds= *Darwinula stevensoni*; Cv= *Cypridopsis vidua*; Ev = *Eucypris virens*; Hi= *Heterocypris incongruens*. See Table S1A for more details on the investigated specimens and Table S4A on details of all classified ASVs. yes = *Cardinium* present according to classic PCR screening. no= *Cardinium* absent according to classic PCR screening. Based on high-throughput 16S amplicon sequencing, analysed with DADA2 (Callahan et al., 2016) and plotted with phyloseq (McMurdie & Holmes, 2013) and ggplot (Wickham, 2016).

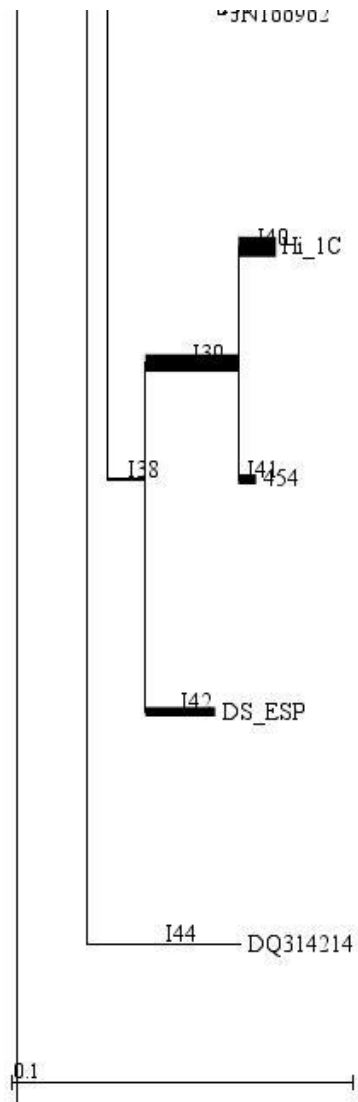


**Figure S1B: Top 20 bacterial ASVs in investigated non-marine ostracods.** Ds= *Darwinula stevensoni*; Cv= *Cypridopsis vidua*; Ev = *Eucypris virens*; Hi= *Heterocypris incongruens*. ASV= 16S Amplicon sequence variant. See Table S1A for more details on the investigated specimens and Table S4A on details of all classified ASVs. yes = *Cardinium* present according to classic PCR screening. no= *Cardinium* absent according to classic PCR screening. Based on high-throughput 16S amplicon sequencing, analysed with DADA2 (Callahan et al., 2016) and plotted with phyloseq (McMurdie & Holmes, 2013) and ggplot (Wickham, 2016).

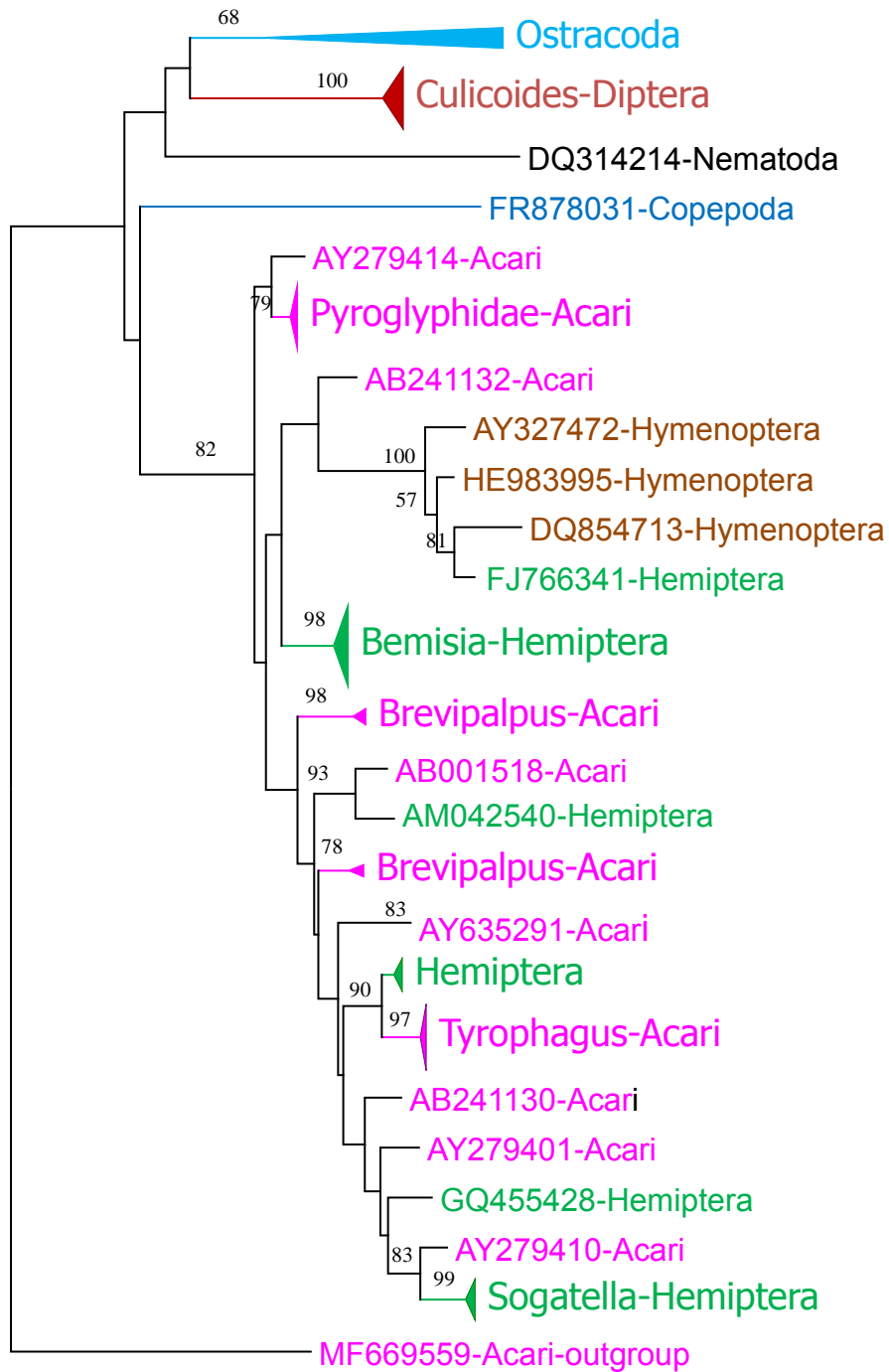




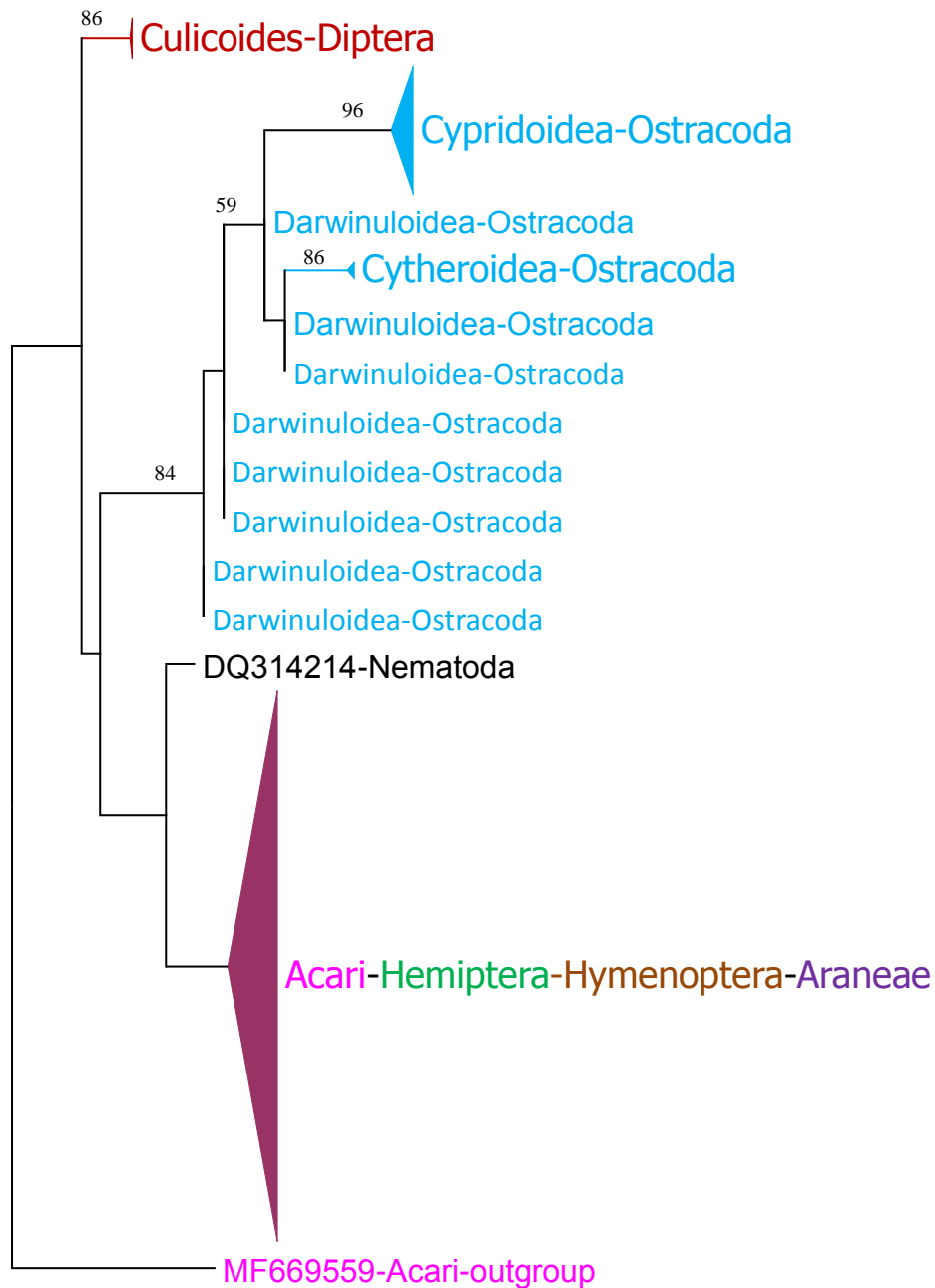
**Figure S2A: Details of the RAxML tree constructed with 16S sequence from non-Cardinium (top) and *Cardinium* (bottom) bacteria with mapped high throughput 16S sequences.** RAxML =Randomized Axelerated Maximum Likelihood (Stamatakis, 2006). Branch width is proportional to the number of mapped sequenced with the Evolutionary Placement Algorithm (EPA; Berger et al., 2011). For more details on the results, see Table S4A. The identity of reference sequences of the tree is provided in Table S1A & B.



**Figure S2B: Details of the RAxML tree constructed with 16S sequences of *Cardinium* with mapped shorter 5' and 3' 16S sequences of non-marine ostracod hosts.** Branch width is proportional to the number of mapped sequenced with EPA. For more details on the results, refer to Table S3B. For sequence identities, refer to Table S1A.

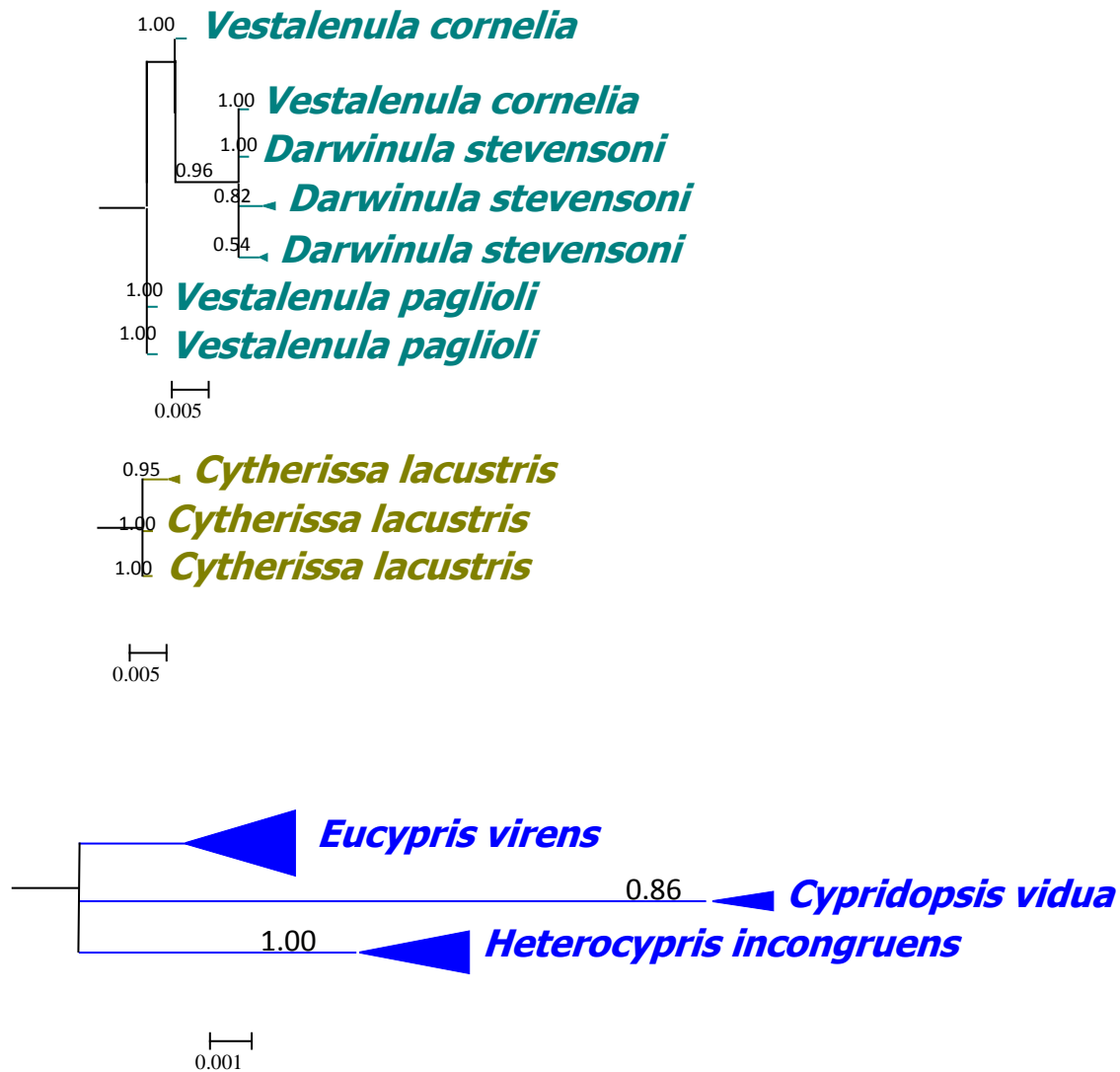


**Figure S3:** Maximum Likelihood (ML) tree constructed from 16S rRNA sequences of *Cardinium* (see Table S1A & 1B for more details on 16S sequence data) with phym (Guindon & Gascuel, 2003). Numbers above branches are bootstrap of 1000 replicates in %. Only bootstraps above 50% are shown. **FigureS3A:** ML tree constructed from 1276bp of the 16S rRNA region of *Cardinium* from various hosts, with the TVM+G model and a gamma parameter of 1.67.



**Figure S3:** Maximum Likelihood (ML) tree constructed from 16S rRNA sequences of *Cardinium* (see Table S1A & 1B for more details on 16S sequence data) with phymI (Guindon & Gascuel, 2003). Numbers above branches are bootstrap of 1000 replicates in %. Only bootstraps above 50% are shown.

**Figure S3B:** ML tree constructed from partial 16S rRNA sequences of *Cardinium* of the 3' end (403bp) from various hosts with the K80 model, kappa=9.10 and ti/tv=4.55.



**Figure S4: Subtree of *Cardinium* from different non-marine ostracod species.**

Details of the collapsed clades of Figure 1B are provided here.



**Table S1: Details of the screened non-marine ostracods and of additional 16S rRNA sequences of *Cardinium* from GenBank that were used for the phylogenetic reconstructions and Bayes Factor tests.**

**A. Information on screened ostracods.**

For each ostracod, the morphospecies, DNA code (ID), geographic origin, infection status, cryptic species identity, latitude and Genbank accession number of the COI sequence is provided if the latter was used for phylogenetic constructions of the hosts. Coordinates are merely indicative for the aquatic habitat and do not include the different sampling points within each habitat. "1" in the column "Cardinium" indicates infection. If no information is provided in this column, the specimen in question was not screened for *Cardinium* but only used to obtain COI DNA sequence data for constructing host phylogenies. For ostracod specimens testing positive with *Cardinium*, also the part of the 16S rRNA that was sequenced and, where applicable, the Genbank accession number is provided. A complete 16S alignment including all 16S sequncens of *Cardinium* from ostracods is available from the first author by request. Ostracod specimens that have been screened for four different symbionts (*Wolbachia*, *Rickettsia*, *Cardinium* & *Spiroplasma*) are indicated in bold. Ostracods that were also screened with high-throughput metagenomics sequencing techniques are underlined. # = number. Genetic species numbers are according to Bode et al. (2010), Schön et al. (2012), Schön et al. in prep. and Cours et al. in prep. Sequences of the 5' end of 16S rRNA of *Cardinium* were used for initial phylogenetic reconstructions and for EPA mapping.

Morpho-species	DNA ID	Population	Habitat	Cardinium	Latitude	Longitude	Genetic species #	GenBank # 16S Cardinium	16S region Cardinium	bp 16S data	GenBank# host COI
<i>Eucypris virens</i>	121	Drongen, Belgium (DR2)	meadow temporary pool	0	51.04936	3.6652778	16				GQ914299
<i>Eucypris virens</i>	119	Drongen, Belgium DR2)	meadow temporary pool	0	51.04936	3.6652778	16				
<i>Eucypris virens</i>	221	Berchem, Belgium (BRC)	temporary pool	1	50.79472	3.5175	16	KC167899	3' V6-V9	403	
<i>Eucypris virens</i>	529	Berchem, Belgium (BRC)	temporary pool	1	50.79472	3.5175	16	KC167902	3' V6-V9	403	
<i>Eucypris virens</i>	<b>FR_FIO 137</b>	Fitou, France (FIO)	temporary pool	1	42.92417	2.99083	31	KC167888	3' V6-V9	403	GQ914305
<i>Eucypris virens</i>	<b>230</b>	Fitou, France (FIT)	temporary pool	0	42.88333	2.96667	1				
<i>Eucypris virens</i>	<b>FR_DU R143</b>	Durfort, France (DUR)	temporary pool	1	43.99083	3.98667	36	KC167896 MH908927	3' V6-V9 5' V1 & V2	403 376	GQ914311
<i>Eucypris virens</i>	198	Urbino, Corsica,	temporary pool	1	42.06094	9.44797	31		3' V6-V9	403	GQ914325

		France (COU)									
<i>Eucypris virens</i>	<b>IT_PUP 207</b>	Punta Grossa, Puglia, Italy (PUP)	temporary pool	1	40.29106	17.79542	26	KC167889 MH908928	3' V6-V9 5' V1-V2	403 376	GQ914333
<i>Eucypris virens</i>	<b>IT_SIM 89</b>	Macarese, Sicily, Italy (SIM)	temporary pool	0	38.1014	12.66080	8				GQ914288
<i>Eucypris virens</i>	<b>499</b>	Piana di Gesturi, Sardinia, Italy (SA5)	temporary pool	0	39.73692	8.99775	1				
<i>Eucypris virens</i>	IT_PUS 122_M	Lecce, Puglia, Italy (PUS)	temporary pool	0	39.9489	18.29920	10				GQ914300
<i>Eucypris virens</i>	IT_PUS 123_M	Lecce, Puglia, Italy (PUS)	temporary pool	0	39.9489	18.29920	10				GQ914301
<i>Eucypris virens</i>	IT_PUS 124_M	Lecce, Puglia, Italy (PUS)	temporary pool	0	39.9489	18.29920	10				GQ914302
<i>Eucypris virens</i>	237	Lecce, Puglia, Italy (PUS)	temporary pool	0	39.9489	18.29920	10				
<i>Eucypris virens</i>	238	Lecce, Puglia, Italy (PUS)	temporary pool	0	39.9489	18.29920	10				
<i>Eucypris virens</i>	IT_SIO4 71_M	S. Fratello, Sicily, Italy (SIO)	temporary pool	0	37.92497	14.67524	33				GQ914491
<i>Eucypris virens</i>	<b>IT_SIP1 63</b>	Sicily, Italy	temporary pool	0	37.92497	13.36417	na				KC167997
<i>Eucypris virens</i>	HR_OM I361	Split, Croatia (OMI)	temporary pool	1	43.45083	16.69583	31		3' V6-V9	403	GQ914434
<i>Eucypris virens</i>	HR_OM I362	Split, Croatia (OMI)	temporary pool	1	43.45083	16.69583	31		3' V6-V9	403	GQ914435
<i>Eucypris virens</i>	HR_OM I363	Split, Croatia (OMI)	temporary pool	1	43.45083	16.69583	31		3' V6-V9	403	GQ914436
<i>Eucypris virens</i>	HR_KR K364	Krk, Croatia (Krk)	temporary pool	1	45.03056	14.59444	31		3' V6-V9	403	GQ914437
<i>Eucypris virens</i>	HR_KR K365	Krk, Croatia (Krk)	temporary pool	1	45.03056	14.59444	31		3' V6-V9	403	GQ914438
<i>Eucypris virens</i>	<b>ES_VA1</b>	El Saler,	temporary pool	0	39.32083	-0.30833	31				KC168003

	<b>353</b>	Albufera, Spain (VA1)									
<i>Eucypris virens</i>	<b>ES_VA4 86</b>	El Saler, Albufera, Spain (VA4)	temporary pool	1	39.32083	-0.31090	31	KC167894	3' V6-V9	403	GQ914285
<i>Eucypris virens</i>	<b>157</b>	El Saler, Albufera, Spain (VA4)	temporary pool	0	39.34328	-0.31090	31				
<i>Eucypris virens</i>	ES_ME 3151	Favaritx, Menorca, Spain (ME3)	temporary pool	1	39.99306	4.25306	31		3' V6-V9	403	GQ914315
<i>Eucypris virens</i>	<b>150</b>	<b>Carniola/Ciutadella, Menorca, Spain (ME1)</b>	temporary pool	0	40.04667	3.90081	16				
<i>Eucypris virens</i>	<b>ES_MF 4436_M</b>	<b>Villarreal de San Carlos, Extremadura, Spain (MF4)</b>	temporary pool	0	39.91167	-6.06167	13				GQ914468
<i>Eucypris virens</i>	<b>MT_M OS200</b>	Mosta, Malta (MOS)	temporary pool	0	35.91289	14.42506	31				GQ914327
<i>Eucypris virens</i>	202	Santa Margherita, Malta (MAR)	temporary pool	0	35.91678	14.43208	31				
<i>Eucypris virens</i>	203	Santa Margherita, Malta (MAR)	temporary pool	0	35.91678	14.43208	31				
<i>Eucypris virens</i>	204	Santa Margherita, Malta (MAR)	temporary pool	0	35.91678	14.43208	31				
<i>Eucypris virens</i>	201	Ghiadira S-safra, Malta (GHI)	temporary pool	0	35.95136	14.44364	31				
<i>Eucypris virens</i>	<b>GR_CO E217</b>	Melisa, Corfu, Greece (COE)	temporary pool	1	39.475	19.91000	31	KC167890	3' V6-V9	403	GQ914338
<i>Eucypris virens</i>	GR_CO E218	Melisa, Corfu, Greece (COE)	temporary pool	1	39.475	19.91000	31		3' V6-V9	403	GQ914339

<i>Eucypris virens</i>	<b>219</b>	Melisa, Corfu, Greece (COE)	temporary pool	1	39.475	19.91000	31	KC167891	3' V6-V9	403	
<i>Eucypris virens</i>	<b>131</b>	Skripera, Corfu, Greece (COA)	temporary pool	0	39.6975	19.78500	34				
<i>Eucypris virens</i>	GR_CO A496_M	Skripera, Corfu, Greece (COA)	temporary pool	0	39.6975	19.78500	34				GQ914592
<i>Eucypris virens</i>	497	Skripera, Corfu, Greece (COA)	temporary pool	0	39.6975	19.78500	34				
<i>Eucypris virens</i>	GR_CO A681_M	Skripera, Corfu, Greece (COA)	temporary pool		39.6975	19.78500	34				GQ914618
<i>Eucypris virens</i>	GR_CO B658_M	Skripera, Corfu, Greece (COB)	temporary pool		39.6975	19.78556	34				GQ914615
<i>Eucypris virens</i>	GR_CO B668_M	Skripera, Corfu, Greece (COB)	temporary pool		39.6975	19.78556	34				
<i>Eucypris virens</i>	132	Skripera, Corfu, Greece (COB)	temporary pool	0	39.69778	19.78556	42				
<i>Eucypris virens</i>	493	Skripera, Corfu, Greece (COB)	temporary pool	0	39.69778	19.78556	34				
<i>Eucypris virens</i>	494	Skripera, Corfu, Greece (COB)	temporary pool	0	39.69778	19.78556	34				
<i>Eucypris virens</i>	495	Skripera, Corfu, Greece (COB)	temporary pool	0	39.69778	19.78556	34				
<i>Eucypris virens</i>	488	Kefalovryso, Corfu, Greece (COD)	temporary pool	0	39.6256	19.79860	34				
<i>Eucypris virens</i>	<b>489</b>	Kefalovryso, Corfu, Greece (COD)	temporary pool	0	39.6256	19.79860	34				

<i>Eucypris virens</i>	<b>212</b>	Kefalovryso, Corfu, Greece (COD)	temporary pool	0	39.6256	19.79860	42				
<i>Eucypris virens</i>	GR_CO D490_M	Kefalovryso, Corfu, Greece (COD)	temporary pool	0	39.6256	19.79860	34				GQ914590
<i>Eucypris virens</i>	491	Kefalovryso, Corfu, Greece (COD)	temporary pool	0	39.6256	19.79860	34				
<i>Eucypris virens</i>	GR_CO D526_M	Kefalovryso, Corfu, Greece (COD)	temporary pool		39.6256	19.79860	34				GQ914594
<i>Eucypris virens</i>	492	Kefalovryso, Corfu, Greece (COD)	temporary pool	0	39.6256	19.79860	34				
<i>Eucypris virens</i>	<b>215</b>	Kefalovryso, Corfu, Greece (COD)	temporary pool	0	39.6256	19.79860	42				
<i>Eucypris virens</i>	<b>225</b>	Gdańsk, Poland (JAB)	temporary pool	1	54.33139	18.56417	37	KC167892 MH908926	3' V6-V9 5' V1 & V2	403 376	
<i>Eucypris virens</i>	223	Gdańsk, Poland (JAB)	temporary pool	1	54.33139	18.56417	37	KC167900	3' V6-V9	403	
<i>Eucypris virens</i>	224	Gdańsk, Poland (JAB)	temporary pool	0	54.33139	18.56417	37				
<i>Eucypris virens</i>	226	Chojnowo, Poland (CHO)	temporary pool	1	52.65583	21.22806	37				
<i>Eucypris virens</i>	227	Chojnowo, Poland (CHO)	temporary pool	0	52.65583	21.22806	37				
<i>Eucypris virens</i>	246	Starkowa Huta, Poland (STA)	temporary pool	0	54.22722	18.17722	37				
<i>Eucypris virens</i>	247	Starkowa Huta, Poland (STA)	temporary pool	0	54.22722	18.17722	37				
<i>Eucypris virens</i>	248	Starkowa Huta, Poland (STA)	temporary pool	0	54.22722	18.17722	37				

<i>Eucypris virens</i>	249	Trzepowo, Poland (TRZ)	temporary pool	1	54.18	18.29278	37				
<i>Eucypris virens</i>	250	Trzepowo, Poland (TRZ)	temporary pool	1	54.18	18.29278	37				
<i>Eucypris virens</i>	253	Tohela, Estonia (TOH)	temporary pool	1	58.40167	24.00667	37				
<i>Eucypris virens</i>	271	Vara, Estonia (VAR)	temporary pool	0	58.52694	26.95000	38				
<i>Eucypris virens</i>	272	Vara, Estonia (VAR)	temporary pool	0	58.52694	26.95000	38				
<i>Eucypris virens</i>	273	Vara, Estonia (VAR)	temporary pool	0	58.52694	26.95000	38				
<i>Eucypris virens</i>	252	Puikule, Latvia, (PUI)	temporary pool	1	57.65639	24.89389	37				
<i>Eucypris virens</i>	256	Galgauka, Latvia (GAL)	temporary pool	0	57.16667	26.59944	37				
<i>Eucypris virens</i>	257	Gulbene, Latvia (GUL)	temporary pool	0	57.19056	26.89500	38				
<i>Eucypris virens</i>	258	Gulbene, Latvia (GUL)	temporary pool	0	57.19056	26.89500	38				
<i>Eucypris virens</i>	259	Gulbene, Latvia (GUL)	temporary pool	0	57.19056	26.89500	38				
<i>Eucypris virens</i>	260	Gulbene, Latvia (GUL)	temporary pool	0	57.19056	26.89500	38				
<b><i>Eucypris virens</i></b>	<b>TN_TU N384</b>	Ariana, Tunisia (TUN)	temporary pool	1	37.005	10.23100	8	KC167893	3' V6-V9	403	KC168005
<b><i>Eucypris virens</i></b>	<b>TN_TU N386</b>	Ariana, Tunisia (TUN)	temporary pool	1	37.005	10.23100	8	KC167897	3' V6-V9	403	KC168006
<b><i>Eucypris virens</i></b>	<b>392</b>	El Hisiane, Tunisia (HIS)	temporary pool	0	36.995	10.15083	8				
<b><i>Eucypris virens</i></b>	<b>TN_HIS 395_F M</b>	El Hisiane, Tunisia (HIS)	temporary pool	0	36.995	10.15083	8				GQ914443
<b><i>Eucypris virens</i></b>	<b>TN_OU L_376</b>	Ouled Amer, Tunisia (OUL)	temporary pool	0	36.01306	10.31833	30				KC168004
<i>Eucypris virens</i>	377	Ouled Amer, Tunisia (OUL)	temporary pool	0	36.01306	10.31833	30				

<i>Eucypris virens</i>	378	Ouled Amer, Tunisia (OUL)	temporary pool	0	36.01306	10.31833	30				
<i>Eucypris virens</i>	379	Ouled Amer, Tunisia (OUL)	temporary pool	0	36.01306	10.31833	30				
<i>Eucypris virens</i>	383	Ouled Amer, Tunisia (OUL)	temporary pool	0	36.01306	10.31833	30				
<i>Eucypris virens</i>	TN_RA O391_F M	Raoued, Tunisia (RAO)	temporary pool	1	36.95639	10.22083	8		3' V6-V9	403	GQ914440
<b><i>Eucypris virens</i></b>	<b>MA_M B3145_ FM</b>	Youssoufia, Morocco (MB3)	temporary pool	1	32.2875	-8.33222	7				GQ914313
<i>Eucypris virens</i>	429	Chemaia, Morocco (ML1)	temporary pool	0	31.97944	-8.37639	9				
<i>Eucypris virens</i>	MA_ML 1426_ M	Chemaia, Morocco (ML1)	temporary pool		31.97944	-8.37639	9				GQ914463
<i>Eucypris virens</i>	MA_ML 1430_ M	Chemaia, Morocco (ML1)	temporary pool		31.97944	-8.37639	9				GQ914465
<b><i>Eucypris virens</i></b>	<b>146</b>	Chemaia, Morocco (ML1)	temporary pool	0	31.97944	-8.37639	9				
<i>Eucypris virens</i>	431	Chemaia, Morocco (ML1)	temporary pool	0	31.97944	-8.37639	9				
<i>Eucypris virens</i>	432	Chemaia, Morocco (ML1)	temporary pool	0	31.97944	-8.37639	9				
<i>Eucypris virens</i>	433	Chemaia, Morocco (ML1)	temporary pool	0	31.97944	-8.37639	9				
<b><i>Eucypris virens</i></b>	<b>MA_M L1434_ M</b>	Chemaia, Morocco (ML1)	temporary pool	0	31.97944	-8.37639	9				GQ914466
<i>Eucypris virens</i>	278	Chemaia, Morocco	temporary pool	0	31.94111	-8.39889	7				

		(ML2)								
<i>Eucypris virens</i>	MA_ML 2276_F M	Chemaia, Morocco (ML2)	temporary pool		31.94111	-8.39889	7			GQ914389
<i>Eucypris virens</i>	MA_M CH515_ M	Chemaia, Morocco (MCH)	temporary pool	0	31.81167	-8.54528	7			GQ914508
<b><i>Eucypris virens</i></b>	<b><u>MA_M</u></b> <b><u>L2275</u></b>	Chemaia, Morocco (ML2)	temporary pool	0	31.94111	-8.39889	7			KC168002
<i>Eucypris virens</i>	MA_M B1476_ M	Bouchane, Morocco (MB1)	temporary pool	0	32.285	-8.32278	7			GQ914496
<i>Eucypris virens</i>	MA_M B1478_ M	Bouchane, Morocco (MB1)	temporary pool	0	32.285	-8.32278	5			GQ914498
<i>Eucypris virens</i>	MA_M RI479_ M	Berrechid, Morocco (MRI)	temporary pool	0	33.16444	-7.39194	27			GQ914499
<i>Eucypris virens</i>	MA_M RI480_ M	Berrechid, Morocco (MRI)	temporary pool	0	33.16444	-7.39194	21			GQ914500
<i>Eucypris virens</i>	MA_M RI481_ M	Berrechid, Morocco (MRI)	temporary pool	0	33.16444	-7.39194	21			GQ914501
<i>Eucypris virens</i>	MA_M RI482_ M	Berrechid, Morocco (MRI)	temporary pool	0	33.16444	-7.39194	27			GQ914502
<i>Eucypris virens</i>	MA_M RI483_ M	Berrechid, Morocco (MRI)	temporary pool	0	33.16444	-7.39194	27			GQ914503
<i>Eucypris virens</i>	MA_M B2484_ M	Bouchane, Morocco (MB2)	temporary pool	0	32.28833	-8.33444	5			GQ914504
<i>Eucypris virens</i>	MA_M B2485_ M	Bouchane, Morocco (MB2)	temporary pool	0	32.28833	-8.33444	7			GQ914505



<i>Eucypris virens</i>	<b>MA_M B2486_ M</b>	Bouchane, Morocco (MB2)	temporary pool	0	32.28833	-8.33444	7				GQ914506
<i>Eucypris virens</i>	<b>UK_NE W188</b>	Newfield Hall pond, Yorkshire, UK (NEW)	temporary pool	0	54.015	-2.13444	31				GQ914322
<i>Eucypris virens</i>	<b>UK_NE W189</b>	Newfield Hall pond, Yorkshire, UK (NEW)	temporary pool	0	54.015	-2.13444	31				GQ914323
<i>Eucypris virens</i>	<b>UK_NE W190</b>	Newfield Hall pond, Yorkshire, UK (NEW)	temporary pool	0	54.015	-2.13444	31				GQ914324
<i>Eucypris virens</i>	<b>UK_NE W191</b>	Newfield Hall pond, Yorkshire, UK (NEW)	temporary pool	0	54.015	-2.13444	31				KC167999
<i>Eucypris virens</i>	501	Newfield Hall pond, Yorkshire, UK (NEW)	temporary pool	0	54.015	-2.13444	31				
<i>Eucypris virens</i>	502	Newfield Hall pond, Yorkshire, UK (NEW)	temporary pool	0	54.015	-2.13444	31				
<i>Eucypris virens</i>	503	Newfield Hall pond, Yorkshire, UK (NEW)	temporary pool	0	54.015	-2.13444	31				
<i>Eucypris virens</i>	507	Newfield Hall pond, Yorkshire, UK (NEW)	temporary pool	1	54.015	-2.13444	31				
<i>Eucypris virens</i>	508	Newfield Hall pond, Yorkshire, UK	temporary pool	1	54.015	-2.13444	31	KC167901	3' V6-V9	403	

		(NEW)									
<i>Eucypris virens</i>	<b>UK_UP D186</b>	Upper Denton, Cumbria, UK (UPD)	temporary pool	0	54.89472	-2.59333	28				GQ914320
<i>Eucypris virens</i>	UK_UP D187	Upper Denton, Cumbria, UK (UPD)	temporary pool	0	54.89472	-2.59333	28				GQ914321
<i>Eucypris virens</i>	185	Upper Denton, Cumbria, UK (UPD)	temporary pool	0	54.89472	-2.59333	28				
<i>Eucypris virens</i>	<b>PT_CO R232_FM</b>	Monte dos corvos, Portugal (COR)	temporary pool	0	37.43222	-7.95528	15				GQ914350
<i>Eucypris virens</i>	PT_CO R234_M	Monte dos corvos, Portugal (COR)	temporary pool	0	37.43222	-7.95528	15				GQ914352
<i>Eucypris virens</i>	PT_CO R235_M	Monte dos corvos, Portugal (COR)	temporary pool	0	37.43222	-7.95528	15				GQ914353
<i>Eucypris virens</i>	PT_CO R236_M	Monte dos corvos, Portugal (COR)	temporary pool	0	37.43222	-7.95528	15				GQ914354
<i>Eucypris virens</i>	<b>BG_KO B442</b>	Kobleshkovo, Bulgaria (KOB)	temporary pool	0	42.60917	27.53500	31				GQ914472
<i>Eucypris virens</i>	BG_RA D450	Radanovo, Bulgaria (RAD)	temporary pool	1	43.37722	25.64278	16	MH908925	5', 3', V1 & V2, V6-V9	391, 403	GQ914478
<i>Eucypris virens</i>	<b>RO_TA M438</b>	Tamburesti, Romania (TAM)	temporary pool	0	44.04778	23.93167	28				GQ914470

<i>Eucypris virens</i>	RO_LES 451	Leş, Romania (LES)	temporary pool	1	46.96056	21.82944	16		3' V6-V9	403	GQ914479
<b><i>Eucypris virens</i></b>	<b>454</b>	Hrkovce, Slovakia (HRK)	temporary pool	1	48.09528	18.89250	37	KC167898, MH908935	5'-3' V1-V6	1276	
<b><i>Eucypris virens</i></b>	<b>SK_HR K455</b>	Hrkovce, Slovakia (HRK)	temporary pool		48.09528	18.89250	37				GQ914482
<i>Eucypris virens</i>	TR_KAB 349	Kavak, Turkey (KAB)	temporary pool	1	41.00139	35.82028	16		3' V6-V9	403	GQ914428
<i>Eucypris virens</i>	TR_KAB 351	Kavak, Turkey (KAB)	temporary pool	1	41.00139	35.82028	16		3' V6-V9	403	GQ914430
<i>Eucypris virens</i>	TR_LAD 341	Ladik, Turkey (LAD)	temporary pool	0	40.91028	35.93194	31				GQ914424
<i>Eucypris virens</i>	TR_LAD 342	Ladik, Turkey (LAD)	temporary pool	0	40.91028	35.93194	31				
<i>Eucypris virens</i>	TR_YUK 324	Yukiari, Turkey (YUK)	temporary pool	1	41.22667	36.63750	31		3' V6-V9	403	GQ914419
<i>Eucypris virens</i>	262	Alcaria Ruiva, Portugal (ALC)	temporary pool	1	37.7414	-7.78890	15		3' V6-V9	403	
<b><i>Eucypris pigra</i></b>	<b>466</b>	Ladik, Turkey (LAC)	temporary pool	1	35.91111	40.94694	na		3' V6-V9	403	GQ914730
<b><i>Heterocypris incongruens</i></b>	<b>HIA</b>	Wuppertal, Germany	flower pot	1	51.24	7.106275	7	KC167903	3' V6-V9	403	
<b><i>Heterocypris incongruens</i></b>	<b>HIB</b>	Wuppertal, Germany	flower pot	1	51.24	7.106275	7	KC167904	3' V6-V9	403	
<b><i>Heterocypris incongruens</i></b>	<b>HIC</b>	Wuppertal, Germany	flower pot	1	51.24	7.106275	7	KC167905, MH908934	5'-3' V1-V6	1276	<a href="#">KC168020</a>
<b><i>Heterocypris incongruens</i></b>	<b>HID</b>	Wuppertal, Germany	flower pot	1	51.24	7.106275	7	KC167906	3' V6-V9	403	KC168016
<b><i>Heterocypris incongruens</i></b>	<b>HIE</b>	Wuppertal, Germany	flower pot	1	51.24	7.106275	7	KC167907	3' V6-V9	403	XXXXXXXXXX
<b><i>Heterocypris incongruens</i></b>	<b>HIF</b>	Wuppertal, Germany	flower pot	1	51.24	7.106275	7	KC167908	3' V6-V9	403	
<b><i>Heterocypris incongruens</i></b>	<b>IS1</b>	Drongen, Belgium	flower pot	1	51.04936	3.6055139	na				
<b><i>Heterocypris incongruens</i></b>	<b>IS2</b>	Drongen, Belgium	flower pot	1	51.04936	3.6055139	1				KC168015

<i>Heterocypris incongruens</i>	IS3	Drongen, Belgium	flower pot	1	51.04936	3.6055139	1				KC168016
<i>Heterocypris incongruens</i>	IS4	Drongen, Belgium	flower pot	1	51.04936	3.6055139	4				KC168014
<i>Heterocypris incongruens</i>	IS5	Drongen, Belgium	flower pot	1	51.04936	3.6055139	na				
<i>Heterocypris incongruens</i>	IS6	Drongen, Belgium	flower pot	1	51.04936	3.6055139	na				
<i>Heterocypris incongruens</i>	Hi1A	Monte dos corvos, Portugal	temporary pool (not the same as E. virens)	1	37.43222	-7.954444	5				
<i>Heterocypris incongruens</i>	Hi1B	Monte dos corvos, Portugal	temporary pool (not the same as E. virens)	1	37.43222	-7.954444	5				
<i>Heterocypris incongruens</i>	Hi1C	Monte dos corvos, Portugal	temporary pool (not the same as E. virens)	1	37.43222	-7.954444	5				<a href="#">KC168038</a>
<i>Heterocypris incongruens</i>	Hi1E	Monte dos corvos, Portugal	temporary pool (not the same as E. virens)	1	37.43222	-7.954444	5				
<i>Heterocypris incongruens</i>	Hi1F	Monte dos corvos, Portugal	temporary pool (not the same as E. virens)	1	37.43222	-7.954444	5				<a href="#">KC168039</a>
<i>Heterocypris incongruens</i>	Hi1G	Monte dos corvos, Portugal	temporary pool (not the same as E. virens)	1	37.43222	-7.954444	5				<a href="#">KC168035</a>
<i>Heterocypris incongruens</i>	Hi1H	Monte dos corvos, Portugal	temporary pool (not the same as E. virens)	1	37.43222	-7.954444	5				<a href="#">KC168036</a>
<i>Heterocypris incongruens</i>	Hi1I	Monte dos corvos, Portugal	temporary pool (not the same as E. virens)	1	37.43222	-7.954444	5				<a href="#">KC168037</a>
<i>Heterocypris incongruens</i>	Hi1D	Monte dos corvos, Portugal	temporary pool (not the same as E. virens)	1	37.43222	-7.954444	5				
<i>Heterocypris incongruens</i>	HI2A	Naunhof, Germany	temporary pool at highway exit	1	51.291	12.626283	7				KC168023

<i>Heterocypris incongruens</i>	HI2B	Naunhof, Germany	temporary pool at highway exit	1	51.291	12.626283	7				KC168031
<i>Heterocypris incongruens</i>	HI2C	Naunhof, Germany	temporary pool at highway exit	1	51.291	12.626283	7				KC168024
<i>Heterocypris incongruens</i>	HI2D	Naunhof, Germany	temporary pool at highway exit	1	51.291	12.626283	7				KC168026
<b><i>Heterocypris incongruens</i></b>	<b>HI3A</b>	S. Manços, Portugal	pond	1	38.44361	-7.733611	6	MH908929	5' V1 & V2	376	<a href="#">KC168042</a>
<b><i>Heterocypris incongruens</i></b>	<b>HI3B</b>	S. Manços, Portugal	pond	1	38.44361	-7.733611	6				<a href="#">KC168041</a>
<b><i>Heterocypris incongruens</i></b>	<b>HI3C</b>	S. Manços, Portugal	pond	1	38.44361	-7.733611	6				
<b><i>Heterocypris incongruens</i></b>	<b>HI3D</b>	S. Manços, Portugal	pond	1	38.44361	-7.733611	6				<a href="#">KC168040</a>
<b><i>Heterocypris incongruens</i></b>	<b>HI4A</b>	Perovlades, Corfu, Greece	temporary pool	1	39.7925	19.704167	7				KC168030
<b><i>Heterocypris incongruens</i></b>	<b>HI4B</b>	Perovlades, Corfu, Greece	temporary pool	1	39.7925	19.704167	3				KC168012
<b><i>Heterocypris incongruens</i></b>	<b>HI4C</b>	Perovlades, Corfu, Greece	temporary pool	1	39.7925	19.704167	3				
<b><i>Heterocypris incongruens</i></b>	<b>HI4D</b>	Perovlades, Corfu, Greece	temporary pool	1	39.7925	19.704167	3				KC168013
<b><i>Heterocypris incongruens</i></b>	<b>HI5A</b>	Veneto, Vicenza, Italy	permanent pool	1	45.963	11.605833	7				KC168033
<b><i>Heterocypris incongruens</i></b>	<b>HI5B</b>	Veneto, Vicenza, Italy	permanent pool	1	45.963	11.605833	7	MH908931	5' V1 & V2	376	KC168032
<b><i>Heterocypris incongruens</i></b>	<b>HI5C</b>	Veneto, Vicenza, Italy	permanent pool	1	45.963	11.605833	7				KC168018
<b><i>Heterocypris incongruens</i></b>	<b>HI6A</b>	Veneto, Vicenza, Italy	permanent pool	1	45.963	11.605833	2	XXXXXXXXXX	5', 3', V1 & V2, V6-V9	376, 403	<a href="#">KC168009</a>
<b><i>Heterocypris incongruens</i></b>	<b>Hi6H</b>	Veneto, Vicenza, Italy	permanent pool	1	45.963	11.605833	2				KC168009
<i>Heterocypris incongruens</i>	CALA	Melicuccà, Calabria, Italy	temporary pool	1	38.28536	9.8466667	7				
<i>Heterocypris</i>	CALB	Melicuccà,	temporary pool	1	38.28536	9.8466667	7				KC168021

<i>incongruens</i>		Calabria, Italy									
<i>Heterocypris incongruens</i>	CALC	Melicuccà, Calabria, Italy	temporary pool	1	38.28536	9.8466667	7	MH908930	5' V1 & V2	376	KC168022
<i>Heterocypris incongruens</i>	CALD	Melicuccà, Calabria, Italy	temporary pool	1	38.28536	9.8466667	7				
<i>Heterocypris incongruens</i>	CALE	Melicuccà, Calabria, Italy	temporary pool	1	38.28536	9.8466667	7				KC168025
<i>Heterocypris incongruens</i>	CALF	Melicuccà, Calabria, Italy	temporary pool	1	38.28536	9.8466667	7				
<i>Heterocypris incongruens</i>	HI7A	Yukiari, Turkey	temporary pool	1	41.22667	36.6375	7				
<i>Heterocypris incongruens</i>	HI7B	Yukiari, Turkey	temporary pool	1	41.22667	36.6375	7				KC168029
<i>Heterocypris incongruens</i>	HI7C	Yukiari, Turkey	temporary pool	1	41.22667	36.6375	7				KC168027
<i>Heterocypris incongruens</i>	HI7D	Yukiari, Turkey	temporary pool	1	41.22667	36.6375	7				KC168028
<i>Heterocypris incongruens</i>	TPA	Trapani, Sicily, Italy	temporary pool	1	37.62688	12.635	7				KC168034
<i>Heterocypris incongruens</i>	TPB	Trapani, Sicily, Italy	temporary pool	1	37.62688	12.635					
<i>Heterocypris incongruens</i>	TPC	Trapani, Sicily, Italy	temporary pool	1	37.62688	12.635	2				KC168011
<i>Heterocypris incongruens</i>	TPD	Trapani, Sicily, Italy	temporary pool	1	37.62688	12.635					
<i>Heterocypris incongruens</i>	TPE	Trapani, Sicily, Italy	temporary pool	1	37.62688	12.635	2				KC168010
<i>Heterocypris incongruens</i>	Hi	Orkney Islands, Scotland	temporary pool	1	58.9809	-2.977261					
<i>Heterocypris incongruens</i>	Hi	Orkney Islands, Scotland	temporary pool	1	58.9809	-2.977261					
<i>Heterocypris incongruens</i>	TP_6	Trapani, Sicily, Italy	temporary pool	1	37.62688	12.635					
<i>Heterocypris salina</i>	153	Limni Keri, Greece	permanent pool	1	37.68669	12.635					

<i>Heterocypris nov. spec.1</i>	H139III	Pozza Sanguedolce, Lampedusa, Italy	hatched from sediment	1	35.5086	12.547222					MH916762
<i>Heterocypris nov. spec.1</i>	166	Taccio Vecchio 3, Lampedusa, Italy	hatched from sediment	0	35.5086	12.547222					
<i>Heterocypris nov. spec.1</i>	167	Taccio Vecchio 2, Lampedusa, Italy	hatched from sediment	0	35.5086	12.547222					
<i>Heterocypris nov. spec.1</i>	168	Taccio Vecchio 2, Lampedusa, Italy	hatched from sediment	0	35.5086	12.547222					
<i>Heterocypris nov. spec.1</i>	169	Taccio Vecchio 2, Lampedusa, Italy	hatched from sediment	1	35.5086	12.547222					
<i>Heterocypris nov. spec.1</i>	H170II	Taccio Vecchio 2, Lampedusa, Italy	hatched from sediment	0	35.5086	12.547222					MH916763
<i>Heterocypris nov. spec.1</i>	171	Taccio Vecchio 2, Lampedusa, Italy	hatched from sediment	0	35.5086	12.547222					
<i>Heterocypris nov. spec.2</i>	F2	Fogia 1, Apulia, Italy	temporary pool	1	42.11	15.480556					KC168045
<i>Heterocypris nov. spec.2</i>	F6	Fogia 1, Apulia, Italy	temporary pool	0	42.11	15.480556					KC168043
<i>Heterocypris nov. spec.2</i>	M3	Fogia 1, Apulia, Italy	temporary pool	0	42.11	15.480556					KC168044
<i>Heterocypris nov. spec.2</i>	AP_1	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_2	Fogia 2, Apulia, Italy	artificial pond	1	42.11	15.480556					

<i>Heterocypris nov. spec.2</i>	AP_3	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_4	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_5	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_6	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_7	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_8	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_9	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_10	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_11	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_12	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_1M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_2M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_3M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_4M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_5M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_6M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_7M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_8M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_9M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					



<i>Heterocypris nov. spec.2</i>	AP_10 M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Heterocypris nov. spec.2</i>	AP_11 M	Fogia 2, Apulia, Italy	artificial pond	0	42.11	15.480556					
<i>Herpetocypris chevreuxi</i>	Hc1	Drongen, Belgium	flower pot	1	51.04936	3.6055139					
<i>Herpetocypris chevreuxi</i>	Hc2	Drongen, Belgium	flower pot	1	51.04936	3.6055139					
<i>Herpetocypris chevreuxi</i>	Hc3	Drongen, Belgium	flower pot	1	51.04936	3.6055139					
<i>Herpetocypris chevreuxi</i>	Hc4	Drongen, Belgium	flower pot	1	51.04936	3.6055139					
<i>Tonnacypris lutaria</i>	T.l 1	Drongen, Belgium	flower pot	0	51.04936	3.6055139					
<i>Tonnacypris lutaria</i>	T.l 2	Drongen, Belgium	temporary pool in ditch	0	51.04936	3.6652778					
<i>Tonnacypris lutaria</i>	T.l 3	Drongen, Belgium	temporary pool in ditch	0	51.04936	3.6652778					
<i>Tonnacypris lutaria</i>	T.l 4	Drongen, Belgium	temporary pool in ditch	0	51.04936	3.6652778					
<i>Tonnacypris lutaria</i>	T.l 5	Drongen, Belgium	temporary pool in ditch	0	51.04936	3.6652778					
<i>Tonnacypris lutaria</i>	T.l 6	Drongen, Belgium	temporary pool in ditch	0	51.04936	3.6652778					
<i>Tonnacypris lutaria</i>	T.l 7	Drongen, Belgium	temporary pool in ditch	0	51.04936	3.6652778					
<i>Tonnacypris lutaria</i>	T.l 8	Drongen, Belgium	temporary pool in ditch	0	51.04936	3.6652778					
<i>Tonnacypris lutaria</i>	T.l 9	Drongen, Belgium	temporary pool in ditch	0	51.04936	3.6652778					
<i>Tonnacypris lutaria</i>	T.l 10	Drongen, Belgium	temporary pool in ditch	0	51.04936	3.6652778					
<i>Cypridopsis vidua</i>	MC183	Melsen Munte, Belgium	farm pond	1	50.96	3.7190647	2		3' V6-V9	403	
<i>Cypridopsis vidua</i>	MC090	Attenbeke, Belgium	garden pond	1	50.78417	3.9158333	1		3' V6-V9	403	MH916761

<i>Cypridopsis vidua</i>	MC110	Bornem, Belgium	forest pond	1	51.11444	4.3248111	1		3' V6-V9	403	
<i>Cypridopsis vidua</i>	MC113	Mechelen, Belgium	park pond	1	51.01889	4.4622278	1		3' V6-V9	403	MH916760
<i>Cypridopsis vidua</i>	MC003	Kortenbergh Nossegem, Belgium	industrial pond	0	50.87667	4.5045361	1				MH916759
<i>Cypridopsis vidua</i>	MC025	Tienen, Belgium	garden pool	1	50.81639	4.9064167	1		3' V6-V9	403	MH916758
<i>Cypridopsis vidua</i>	MC203	Kerkom, Belgium	duck pond	0	50.84278	4.8608611	1				MH916757
<i>Cypridopsis vidua</i>	MC208	Houwart, Belgium	garden pond	0	50.93056	4.8585639	1				MH916756
<i>Cypridopsis vidua</i>	MC211	Rotselaar, Belgium	garden pond	0	50.95806	4.7355972	2				
<b><i>Darwinula stevensoni</i></b>	<b>2A</b>	Ossiachersee, Austria	lake	0	46.6687	13.980833	5				
<b><i>Darwinula stevensoni</i></b>	<b>2B</b>	Ossiachersee, Austria	lake	0	46.6687	13.980833	5				
<b><i>Darwinula stevensoni</i></b>	<b>2C</b>	Ossiachersee, Austria	lake	0	46.6687	13.980833	5				
<i>Darwinula stevensoni</i>	Ds174	Ossiachersee, Austria	lake		46.6687	13.980833	5				AF031299
<b><i>Darwinula stevensoni</i></b>	<b>3A</b>	Keutschacher see, Austria	lake	0	46.5861	14.153333	5				
<b><i>Darwinula stevensoni</i></b>	<b>3B</b>	Keutschacher see, Austria	lake	0	46.5861	14.153333	5				
<b><i>Darwinula stevensoni</i></b>	<b>3C</b>	Keutschacher see, Austria	lake	0	46.5861	14.153333	5				
<b><i>Darwinula stevensoni</i></b>	<b>4A</b>	Mondsee, Austria	lake	0	47.8556	14.153333	5				
<i>Darwinula stevensoni</i>	Ds155	Mondsee, Austria	lake		47.8556	14.153333	5				AF031297
<b><i>Darwinula stevensoni</i></b>	<b>5A</b>	Faaker See, Austria	lake	0	46.5792	13.918611	5				
<b><i>Darwinula stevensoni</i></b>	<b>5B</b>	Faaker See, Austria	lake	0	46.5792	13.918611	5				

<i>Darwinula stevensoni</i>	5E	Faaker See, Austria	lake	0	46.5792	13.918611	5				
<i>Darwinula stevensoni</i>	Ds153	Faaker See, Austria	lake		46.5792	13.918611	5				AF031296
<i>Darwinula stevensoni</i>	1A	Wörthersee, Austria	lake	0	46.6246	14.140278	5				
<i>Darwinula stevensoni</i>	1B	Wörthersee, Austria	lake	0	46.6246	14.140278	5				
<i>Darwinula stevensoni</i>	1C	Wörthersee, Austria	lake	0	46.6246	14.140278	5				
<i>Darwinula stevensoni</i>	Ds80	Wörthersee, Austria	lake		46.6246	14.140278	5				AF031295
<i>Darwinula stevensoni</i>	Ds189	Lake Sibaya, South Africa	lake		-27.3097	32.573333	3				AF031286
<i>Darwinula stevensoni</i>	6A	Lake Malawi, Malawi	lake	0	11.6701	034.96830 8391"E	3				
<i>Darwinula stevensoni</i>	6B	Lake Malawi, Malawi	lake	0	11.6701	034.96830 8391"E	3				
<i>Darwinula stevensoni</i>	Ds_A7	Semerwater, UK	lake	0	54.2802	2.1244444	5				
<i>Darwinula stevensoni</i>	Ds59	Semerwater, UK	lake		54.2802	2.1244444	5				AF031292
<i>Darwinula stevensoni</i>	<u>11</u>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778	5				
<i>Darwinula stevensoni</i>	2	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778	5				KC168007
<i>Darwinula stevensoni</i>	3	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778	5				
<i>Darwinula stevensoni</i>	4	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778	5				
<i>Darwinula stevensoni</i>	5	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778	5				

<i>Darwinula stevensoni</i>	Ds162	Hollandersga atkreek, Belgium	saline lake		51.26889	3.5352778	5					AF031290
<i>Darwinula stevensoni</i>	6	Hollandersga atkreek, Belgium	saline lake	1	51.26889	3.5352778	5					
<i>Darwinula stevensoni</i>	PIK48	Gregory Gorge, West Australia	spring	0	-21.5497	116.97083	na					
<i>Darwinula stevensoni</i>	Ds_US1	Herrick Fen, Ohio, USA	fen	0	41.21389	-81.37111	1					JX069266
<i>Darwinula stevensoni</i>	DS_US2	Herrick Fen, Ohio, USA	fen	0	41.21389	-81.37111	1					
<i>Darwinula stevensoni</i>	Ds_US5	Herrick Fen, Ohio, USA	fen	0	41.21389	-81.37111	1					JX069265
<i>Darwinula stevensoni</i>	Ds_BRA Z	Boracéia Biological Station, São Paulo, Brazil	leaf litter	0	-23.6381	-45.84028	2					JX069267
<i>Darwinula stevensoni</i>	Ds_ESP 2	La Albufera, Valencia, Spain	lake	1	39.34806	-0.324167	4	MH908933	5'-3' V1-V6	1276		
<i>Darwinula stevensoni</i>	Ds_ESP 3	La Albufera, Valencia, Spain	lake	1	39.34806	-0.324167	4	KC167886 MH908932	3' V6-V9 5' V1 & V2	403 376		AF031290
<i>Darwinula stevensoni</i>	DS30	Ulal Fosc, Spain	spring	1	39.0156	-0.090083	4					
<i>Darwinula stevensoni</i>	DS31	Ulal Fosc, Spain	spring	1	39.0156	-0.090083	4					
<i>Darwinula stevensoni</i>	DS32	Ulal Fosc, Spain	spring	1	39.0156	-0.090083	4					
<i>Darwinula stevensoni</i>	DS33	Ulal Fosc, Spain	spring	1	39.0156	-0.090083	4					
<i>Darwinula stevensoni</i>	DS34	Ulal Fosc, Spain	spring	1	39.0156	-0.090083	4					
<i>Darwinula stevensoni</i>	DS36	Ulal Fosc, Spain	spring	1	39.0156	-0.090083	4	MH910345	3' V6-V9	403		

<i>Darwinula stevensoni</i>	T5	Zaventem, Belgium	pond	1	50.8855	4.4933	5	MH910345	3' V6-V9	403	MH916755
<i>Penthesilenula aotearoa</i>	Pa1	Boracéia Biological Station, São Paulo, Brazil	leaf litter	0	-23.6381	-45.84028	3				JX069233
<i>Penthesilenula brasiliensis</i>	Pbs2_08	Campus of the University of São Paulo, Brazil	leaf litter	0	-23.5642	-46.73	4				JX069247
<i>Penthesilenula brasiliensis</i>	PbL06	Boracéia Biological Station, São Paulo, Brazil	leaf litter	0	-23.6381	-45.84028	3				
<i>Penthesilenula brasiliensis</i>	PbL06_I	Boracéia Biological Station, São Paulo, Brazil	leaf litter	0	-23.6381	-45.84028	3				
<i>Penthesilenula brasiliensis</i>	PbL2	Boracéia Biological Station, São Paulo, Brazil	leaf litter	0	-23.6381	-45.84028	3				
<i>Penthesilenula brasiliensis</i>	PbL4	Boracéia Biological Station, São Paulo, Brazil	leaf litter	0	-23.6381	-45.84028	3				JX069242
<i>Penthesilenula brasiliensis</i>	Pb_AUS41	Circular Pool Springs, Karijini, Western Australia	spring	0	-22.4767	118.56027787"E	2				JX069238
<i>Penthesilenula brasiliensis</i>	PbIRL	Lough Lickeen, Ireland	lake	0	52.96278	-9.229444	1				AJ319738
<i>Penthesilenula brasiliensis</i>	Pb236	Clue de la Fou, France	spring	0	42.7175	2.4988889	1				
<i>Microdarwinula spec.</i>	mGabA	Bolapessa, Gabon	leaf litter	0	-0.8037	12.054333					

<i>Microdarwinula spec.</i>	mGabC	Bolapessa, Gabon	leaf litter	0	-0.8037	12.054333					
<i>Microdarwinula zimmeri</i>	Micro	Clue de la Fou, France	spring		42.7175	2.4988889					AF031284
<i>Vestalenula paglioli</i>	BRAZ17_09	Salto Bandeirantes, Santa Fé, Brazil	spring	1	22.982	-51.82744		KC167883	3' V6-V9	403	
<i>Vestalenula paglioli</i>	BRAZ18_09	Salto Bandeirantes, Santa Fé, Brazil	spring	1	22.982	-51.82744		KC167884	3' V6-V9	403	KR184035
<i>Vestalenula cylindrica</i>	Vcyl 8	Lake Biwa, Biwa museum, Japan	lake	0	35.076	135.9335					
<i>Vestalenula cylindrica</i>	Vcyl 3	Lake Biwa, Biwa museum, Japan	lake	0	35.076	135.9335					
<i>Vestalenula molopoensis</i>	Vmol	Molopo Oog, South Africa	lake	0	-25.7778	25.341794					
<i>Vestalenula cornelia</i>	Vcor2	Yudomari, South Yakushima, Japan	spring	1	30.23	130.48069		KC167885	3' V6-V9	403	AJ534411
<i>Vestalenula cornelia</i>	Vcor1	Yudomari, South Yakushima, Japan	spring	1	30.23	130.48069		KC167887	3' V6-V9	403	KR184023
<i>Vestalenula nov. spec.</i>	Pb_US1	Herrick Fen, Ohio, USA	fen	0	41.21389	-81.37111				403	KR184024
<i>Alicenula serricaudata</i>	Ainv	Salto Bandeirantes, Santa Fé, Brazil	spring	0	-22.982	-51.82744					AJ534409
<i>Cypria ophthalmica</i>	<b>Cyp A</b>	Hollandersga atkreek,	saline lake	0	51.26889	3.5352778					

		Belgium								
<i>Cypria ophthalmica</i>	<b>Cyp B</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cypria ophthalmica</i>	<b>Cyp 3C</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cypria ophthalmica</i>	<b>Cyp D</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cypria ophthalmica</i>	<b>Cyp E</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cypria ophthalmica</i>	<b>Cyp F</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cyprideis torosa</i>	<b>Ct A</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cyprideis torosa</i>	<b>Ct B</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cyprideis torosa</i>	<b>Ct C</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cyprideis torosa</i>	<b>Ct D</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cyprideis torosa</i>	<b>Ct E</b>	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cyprideis torosa</i>	CT6	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				
<i>Cyprideis torosa</i>	CT7	Hollandersga atkreek, Belgium	saline lake	0	51.26889	3.5352778				

<i>Romecytheridea ampla</i>	Amp 21	Lake Tanganyika, Katoto, Zambia	lake	0	-8.8108	31.028889					
<i>Romecytheridea ampla</i>	Amp 22	Lake Tanganyika, Katoto, Zambia	lake	0	-8.8108	31.028889					
<i>Romecytheridea ampla</i>	Amp 23	Lake Tanganyika, Katoto, Zambia	lake	0	-8.8108	31.028889					
<i>Romecytheridea ampla</i>	Amp 24	Lake Tanganyika, Katoto, Zambia	lake	0	-8.8108	31.028889					
<i>Romecytheridea ampla</i>	Amp 25	Lake Tanganyika, Katoto, Zambia	lake	0	-8.8108	31.028889					
<i>Romecytheridea ampla</i>	Amp 26	Lake Tanganyika, Katoto, Zambia	lake	0	-8.8108	31.028889					
<i>Romecytheridea ampla</i>	Amp 27	Lake Tanganyika, Katoto, Zambia	lake	0	-8.8108	31.028889					
<i>Romecytheridea ampla</i>	Amp 28	Lake Tanganyika, Katoto, Zambia	lake	0	-8.8108	31.028889					
<i>Romecytheridea ampla</i>	Amp 29	Lake Tanganyika, Katoto, Zambia	lake	0	-8.8108	31.028889					
<i>Romecytheridea ampla</i>	Amp 30	Lake Tanganyika,	lake	0	-8.8108	31.028889					



		Katoto, Zambia								
<i>Cytherissa lacustris</i>	CL 4 II	Semerwater, UK	lake	1	54.2802	2.1244444			3' V6-V9	403
<i>Cytherissa lacustris</i>	CL5 I	Semerwater, UK	lake	1	54.2802	2.1244444			3' V6-V9	403
<i>Cytherissa lacustris</i>	CL3 I	Semerwater, UK	lake	1	54.2802	2.1244444			3' V6-V9	403
<i>Cytherissa lacustris</i>	CL1 II	Semerwater, UK	lake	1	54.2802	2.1244444			3' V6-V9	403

**Table S1B. Origin and identity of 16S *Cardinium* and non-*Cardinium* sequences used for phylogenetic reconstructions, comparisons of genetic diversity, and EPA mapping.** #= number. If no reference is provided, the sequence was directly submitted to GenBank. “short” indicates that this sequence was included in the alignment of 403 basepairs, “long” in the alignments comprising 1276, 1638 and 1777 basepairs, respectively.

<i>Cardinium</i> sequences						
GenBank #	Host genus	Host family - order	Host class	short	long	Reference
AB001518	Ixodes	Ixodida	Acari-Arachnida	x	x	Kurtti et al. 1996
AB116514	Brevipalpus	Trombidiformes	Acari-Arachnida		x	
AB116515	Brevipalpus	Trombidiformes	Acari-Arachnida	x	x	
AB241129	Eotetranychus	Acarina	Acari-Arachnida		x	
AB241130	Oligonychus	Trombidiformis	Acari-Arachnida		x	
AB241131	Amphitetranynchus	Acarina	Acari-Arachnida		x	
AB241132	Tetranychus	Acarina	Acari-Arachnida		x	
AB241135	Tetranychus	Acarina	Acari-Arachnida		x	
AB506773	Harmalia	Delphacidae-Hemiptera	Insecta		x	Nakamaru et al. 2009
AB506774	Sogatella	Fulgoromorpha-Hemiptera	Insecta	x	x	Nakamura et al. 2009
AB506775	Euides	Fulgoromorpha-Hemiptera	Insecta	x	x	Nakamura et al. 2009
AB506776- AB506778	Culicoides	Diptera	Insecta		x	Nakamura et al. 2009
AB506779	Culicoides	Diptera	Insecta	x	x	Nakamura et al. 2009
AF350221	Brevipalpus	Trombidiformes	Acari-Arachnida		x	
AM042540	Scaphoideus	Auchorryncha-Hemiptera	Insecta	x	x	Marzorati et al. 2006
AY279401	Brevipalpus	Trombidiformes	Acari-Arachnida	x	x	Weeks et al. 2003
AY279410	Petrobia	Trombidiformes	Acari-Arachnida	x	x	Weeks et al. 2003
AY279412	Brevipalpus	Trombidiformes	Acari-Arachnida		x	Weeks et al. 2003
AY279413	Metaseiulus	Phytoseiidae	Acari-Arachnida		x	Weeks et al. 2003
AY279414	Opiella	Oribatei	Acari-Arachnida		x	
AY279415	Dicantropis	Delphacidae-Hemiptera	Insecta		x	Weeks et al. 2003
AY327472	Plagiomerus	Encyrtidae-Hymenoptera	Insecta	x	x	Zchori-Fein & Perlman 2004
AY635291	Metaseiulus	Phytoseiidae	Acari-Arachnida		x	Jeyaprakash & Hoy 2004
AY753169- AY753170	Metaseiulus	Phytoseiidae	Acari-Arachnida		x	Jeyaprakash & Hoy 2004

DQ314214	Heterodera	Tylenchida	Nematoda	x		Noel & Atibalentja 2006
DQ369961- DQ369965	Tetranychus	Acarina	Acari-Arachnida		x	Liu et al. 2006
DQ449047	Tetranychus	Acarina	Acari-Arachnida		x	Liu et al. 2006
DQ854713	Encarsia	Chalcidoidea -Hymenoptera	Insecta	x	x	
EU333926	Cyclosa	Araneae	Arachnida	x		Duron et al. 2008
EU333927	Alopecosa	Araneae	Arachnida	x		Duron et al. 2008
EU333928	Evarcha	Araneae	Arachnida	x		Duron et al. 2008
EU333929	Pachygnatha	Araneae	Arachnida	x		Duron et al. 2008
EU333930	Holocnemus	Araneae	Arachnida	x		Duron et al. 2008
EU333931	Linyphia	Araneae	Arachnida	x		Duron et al. 2008
FJ766335	Bemisia	Aleyrodidae-Hemiptera	Insecta	x		Gueguen et al. 2010
FJ766337- FJ766340	Bemisia	Aleyrodidae-Hemiptera	Insecta	x		Gueguen et al. 2010
FJ766341	Bemisia	Aleyrodidae-Hemiptera	Insecta	x	x	Gueguen et al. 2010
GQ206320	Sogatella	Fulgoromorpha-Hemiptera	Insecta	x	x	
GQ455411	Abgrallaspis	Diaspididae-Hemiptera	Insecta	x		
GQ455412	Unaspis	Diaspididae-Hemiptera	Insecta	x		
GQ455413	Howardia	Diaspididae-Hemiptera	Insecta	x		
GQ455414	Leucaspis	Diaspididae-Hemiptera	Insecta	x		
GQ455415	Hemiberlesia	Diaspididae-Hemiptera	Insecta	x		
GQ455416	Pseudoparlatoria	Diaspididae-Hemiptera	Insecta	x		
GQ455417	Melanaspis	Diaspididae-Hemiptera	Insecta	x		
GQ455418	Prodigiaspis	Diaspididae-Hemiptera	Insecta	x		
GQ455419	Aonidomytilus	Diaspididae-Hemiptera	Insecta	x		
GQ455420	Protargionia	Diaspididae-Hemiptera	Insecta	x		
GQ455421	Lepidosaphes	Sternorrhyncha-Hemiptera	Insecta	x		
GQ455422	Oceanaspidiotus	Coccoidea-Hemiptera	Insecta	x		
GQ455423- GQ455424	Palinaspis	Diaspididae-Hemiptera	Insecta	x		
GQ455426	Chionaspis	Diaspididae-Hemiptera	Insecta	x		
GQ455427	Aspidiotus	Diaspididae-Hemiptera	Insecta	x		
GQ455428	Poliaspis	Diaspididae-Hemiptera	Insecta	x	x	
GQ455429	Pallulaspis	Diaspididae-Hemiptera	Insecta	x		
GQ455430- GQ455438	Aspidiotus	Diaspididae-Hemiptera	Insecta	x		

GU451191	Sogatella	Delphacidae-Hemiptera	Insecta		x	
GU451193	Sogatella	Delphacidae-Hemiptera	Insecta		x	
GU451207	Sogatella	Delphacidae-Hemiptera	Insecta		x	
GU731426	Tetranychus	Acarina	Acari-Arachnida		x	
HE983995	Encarsia	Chalcidoidea-Hymenoptera	Insecta	x	x	Penz et al. 2012
JN166961	Culicoides	Diptera	Insecta		x	Morag et al. 2012
JN166962	Culicoides	Diptera	Insecta	x	x	Morag et al. 2012
JN204479	Bemisia	Aleyrodidae-Hemiptera	Insecta	x		Singh et al. 2012
JN204479- JN204482	Bemisia	Aleyrodidae-Hemiptera	Insecta		x	Sing et al. 2012
JN236327	Dermatophagoides	Aracina	Acari-Arachnida		x	
JN236332	Dermatophagoides	Aracina	Acari-Arachnida		x	
JN236335- JN236336	Dermatophagoides	Aracina	Acari-Arachnida		x	
JN236341	Dermatophagoides	Aracina	Acari-Arachnida		x	
JN236354	Dermatophagoides	Aracina	Acari-Arachnida		x	
JN236355	Dermatophagoides	Aracina	Acari-Arachnida	x	x	
JN236357	Dermatophagoides	Aracina	Acari-Arachnida		x	
JN236359	Dermatophagoides	Aracina	Acari-Arachnida		x	
JQ406682- JQ406683	Dermatophagoides	Aracina	Acari-Arachnida	x		Kopecky et al. 2013
JQ406686	Dermatophagoides	Aracina	Acari-Arachnida	x		Kopecky et al. 2013
JQ406690	Dermatophagoides	Aracina	Acari-Arachnida	x		Kopecky et al. 2013
JQ406719	Glycyphagus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406727	Tyrophagus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406730	Tyrophagus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406732	Tyrophagus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406734	Tyrophagus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406741	Acarus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406755	Tyroborus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406810- JQ406812	Tyrophagus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013

JQ406814	Tyrophagus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406816	Tyrophagus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406830	Acarus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406831	Acarus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406834	Lepidoglyphus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406838	Lepidoglyphus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JQ406841	Lepidoglyphus	Sarcoptiformes	Acari-Arachnida	x		Kopecky et al. 2013
JX001272	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	
JX064607	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	
JX064623- JX064624	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	
JX064626	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	
KC677578	Phalangium	Opiliones	Arachnida	x		Duron 2013
KC677579	Dermanyssus	Mestostigmata	Acari-Arachnida	x		Duron 2013
KC677580	Tetranychus	Acarina	Acari-Arachnida	x		Duron 2013
KC677581	Acari sp.	Acarina	Acari-Arachnida	x		Duron 2013
KC677582	Encarsia	Chalcidoidea-Hymenoptera	Insecta	x		Duron 2013
KC677583	Scaphoideus	Auchorryncha-Hemiptera	Insecta	x		Duron 2013
KC677584	Bemisia	Aleyrodidae-Hemiptera	Insecta	x		Duron 2013
KC677585	Aspidiotus	Diaspididae-Hemiptera	Insecta	x		Duron 2013
KF111022- KF111026	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111028- KF111030	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111033	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111035- KF111038	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111059	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111065	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111069	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111078- KF111080	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111086	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111090	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111092 - KF111093	Cheyletus	Trombidiformes	Acari-Arachnida	x		

KF111109	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111118	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KF111140	Cheyletus	Trombidiformes	Acari-Arachnida	x		
KM464155	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	Kopecky et al. 2014
KM464157	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	Kopecky et al. 2014
KM464160	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	Kopecky et al. 2014
KM464167	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	Kopecky et al. 2014
KM464187- KM464188	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	Kopecky et al. 2014
KM464203	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	Kopecky et al. 2014
KM464281	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	Kopecky et al. 2014
KR026907	Culicoides	Diptera	Insecta		x	Mee et al. 2015
KR026910- KR026912	Culicoides	Diptera	Insecta		x	Mee et al. 2015
KR026915	Culicoides	Diptera	Insecta		x	Mee et al. 2015
KR026920- KR026923	Culicoides	Diptera	Insecta		x	Mee et al. 2015
KX022130	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	
KX022134	Tyrophagus	Sarcoptiformes	Acari-Arachnida		x	
KY660634- KY660637	Ornithodoros	Ixodida	Acari-Arachnida	x		
KY660638	Ixodes	Ixodida	Acari-Arachnida	x		
LC159289	Bemisia	Aleyrodidae-Hemiptera	Insecta		x	
LN829672- LN829682	Bemisia	Aleyrodidae-Hemiptera	Insecta		x	
LN829684- LN829685	Bemisia	Aleyrodidae-Hemiptera	Insecta		x	
LN829688- LN829689	Bemisia	Aleyrodidae-Hemiptera	Insecta		x	
MF188893- MF188900	Culicoides	Diptera	Insecta	x		Pages et al. 2017
MF669559	Achipteria	Sarcoptiformes	Acari-Arachnida		x	

MH057586- MH057593	Dermatophagoides	Aracina	Acari-Arachnida	x		
MH057587	Dermatophagoides	Aracina	Acari-Arachnida		x	
MH057596	Dermatophagoides	Aracina	Acari-Arachnida	x		
MH057597	Dermatophagoides	Aracina	Acari-Arachnida		x	
MH057598- MH057604	Dermatophagoides	Aracina	Acari-Arachnida	x		
MH057601- MH057602	Dermatophagoides	Aracina	Acari-Arachnida		x	
MH057605	Dermatophagoides	Aracina	Acari-Arachnida		x	
MH057610	Dermatophagoides	Aracina	Acari-Arachnida	x		
MH057611	Dermatophagoides	Aracina	Acari-Arachnida		x	
MH057612	Dermatophagoides	Aracina	Acari-Arachnida	x		
MH057614	Dermatophagoides	Aracina	Acari-Arachnida	x	x	
MH057615	Dermatophagoides	Aracina	Acari-Arachnida	x		

<b>Non-Cardinium sequences</b>			
<b>GenBank #</b>	<b>Bacterial identity</b>	<b>long</b>	<b>Reference</b>
AB078038	Flammeovirgaceae; Flexithrix	X	Nakagawa et al. 2002.
AB547639	Firmicutes; Tissierellia	X	Sakamoto & Ohkuma 2010
AY695840	Opiritutus, Opiritutaceae	X	
DQ16669	Chryseobacterium, Flavobacteriaceae	X	
EU016445	Cellulomonadaceae, Micrococcales	X	
EU705634	Acinetobacter, Pseudomonadales	X	
FR821218	Pseudorhodobacter, Rhodobacteraceae	X	
HQ425320	Paracoccus, Rhodobacteraceae	X	
JN934492	Bacteriovoracaceae; Bacteriovorax	X	
JQ435729	Pseudoxanthomonas, Xanthomonadaceae	X	
JQ977513	Pseudomonas, Pseudomonadaceae	X	
KF054791	Sphingobacterium, Sphingobacteriaceae;	X	
KJ957195	Flavobacterium, Flavobacteriaceae	X	Park et al. 2015





**Table S2: Overview on PCR primers used to amplify 16S of *Cardinium* from non-marine ostracod hosts.**

The full length of 16S was assembled from sequences of three different PCR products from non-marine ostracods with the three different primer combinations detailed below. Rev comp= reverse complementary.

<b>Aim</b>	<b>Primer forward</b>	<b>Primer reverse</b>	<b>Part of 16S</b>	<b>Sequence length in basepairs</b>
Presence of <i>Cardinium</i> ?	CLO-f1 (5'-GGA ACC TTA CCT GGG CTA GAA TGT ATT) (Gotoh et al., 2007)	CLO-r1 (5'-GCC ACT GTC TTC AAG CTC TAC CAA C) (Gotoh et al., 2007)	V1-V2	376
Phylogeny	16S_Bact_fd1 (Weisburg et al., 1991)	16S_Bact_926r (Weisburg et al., 1991)	V6-V9	403
	rev comp of CLO-r1 (5'- G TTG GTA GAG CTT GAA GAC AGT GGC)	rev comp of 16S_Bact_fd1 (5'-CTG AGC CAG GAT CAA ACT CTG TTG TCG ACG AAT TCG G)	V2-V6	500

**Table S3: Results of DADA2 analyses with different parameters for filtering and trimming.**

mEE=maximum expected error; trim64\_64= left hand trimming of 64 bp, including primers, barcodes and Illumina adaptors; maxmismatch=maximum number of mismatches allowed in the overlap region when merging read pairs. truncq=truncation quality score; trim160\_100=trimming 160 bp of forward and 100 bp of reverse reads, respectively; minlength= minimum sequence length kept after trimming and merging read pairs; bimeras= removed *de novo* chimeras; average reads are calculated per sample; OTUs=number of ASVs used to classify bacteria.

parameters	total reads used	paired reads	average reads	bimeras	OTUs	comments	reads in	reads out
mEE5, trim64_64,maxmismatch3,truncq3, minlength75	436680	39452	28069.67	1068	463	none filtered		
mEE2, trim64_64,maxmismatch3, truncQ3, trim160_100, minlength75	436680	0	0	0	0	all filtered		
mEE5, trim64_64,maxmismatch3,truncq3, minlength75	436680	39452	28069.67	1068	463	none filtered		
mEE5, trim64_64,maxmismatch3,truncq2, minlength75	436680	39452	28069.67	1068	463	none filtered		
mEE5, trim64_64,maxmismatch1,truncq4, minlength75	436680	30285	24092.62	902	545	Hi.yes2 omitted		
mEE3, trim64_64,maxmismatch3,truncq2, minlength75	190042	18469	13288.56	647	354	Ds.no Ds.yes Hi.yes1 Hi.yes2 Hi.no Cv.yes Ev.no1 Ev.no2 Ev.yes	52550 49428 442389 34908 47951 46189 47554 57682 56129	22147 20155 21382 14961 17444 21456 20835 27081 24581
mEE4, trim64_64,maxmismatch3,truncq2, minlength75	328777	30092	21749.78	870	417	Ds.no Ds.yes Hi.yes1 Hi.yes2 Hi.no Cv.yes Ev.no1 Ev.no2 Ev.yes	52550 49428 44289 34908 47951 46189 47554 57682 56129	38940 36413 34902 26180 34113 35417 35919 44691 42202











	Ds.no	Ds.yes	Hi.yes1	Hi.yes2	Hi.no	Cv.yes	Ev.no1	Ev.yes	Ev.no2	Kingdom	Phylum	Class	Order	Family	Genus	Species	EDPL in EPA	RaxML weight in EPA	mapped to branch	internal	external	DNA#	ID			
seq375	0	0	9	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.08	0.99		173	internal			Ostracoda		
seq376	0	0	0	0	9	0	0	0	0	0	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Rhizobiaceae		0.99	0.99		219	internal	HQ425320		Paracoccus, Rhodobacteraceae		
seq377	0	0	0	0	0	0	9	0	0	0	Bacteria	Proteobacteria	Gammaproteobacteria	Betaproteobacteriales			0.02	0.96		223	external	JO977513		Pseudomonas, Pseudomonadaceae		
seq378	0	0	0	0	0	0	0	9	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.33	0.99		175	external	Hi_1C		H. incongruens, Ostracoda		
seq379	0	0	0	0	0	0	0	9	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.33	1		173	internal			Ostracoda		
seq380	0	0	7	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa	Candidatus_Cardinium		0.05	1		173	internal			Ostracoda	
seq381	0	0	0	0	0	0	7	0	0	0	Bacteria	Proteobacteria	Gammaproteobacteria	Betaproteobacteriales	Burkholderiaceae	Polaromonas		0.31	0.97		3	internal	JO435729		Pseudoxanthomonas, Xanthomonadaceae	
seq382	0	6	0	0	0	0	0	0	0	0	Bacteria	Firmicutes	Bacilli	Bacillales	Staphylococcaceae	Staphylococcus		0.19	0.98		214	external	EU016445		Cellulomonadaceae, Micrococcales	
seq383	0	0	6	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.06	1		173	internal			Ostracoda		
seq384	0	0	0	0	0	0	0	6	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.33	1		173	internal			Ostracoda		
seq385	0	0	5	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.08	0.99		173	internal			Ostracoda		
seq386	0	0	5	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.04	0.96		174	external	454		E. virens, Ostracods		
seq387	0	0	0	0	5	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Chitinophagales	Chitinophagaceae	Parasegetibacter	luojensis		0.32	1		223	external	KF054791		Sphingobacteriaceae; Sphingobacterium
seq388	0	0	0	0	0	0	5	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Flavobacteriales	Weeksellaceae		0.11	1		210	external	DO279361		Chryseobacterium, Flavobacteriaceae		
seq389	0	0	0	0	0	0	5	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Flavobacteriales	Weeksellaceae		0.18	1		210	external	DO279361		Chryseobacterium, Flavobacteriaceae		
seq390	0	0	0	0	0	0	0	5	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.33	0.99		173	internal			Ostracoda		
seq391	0	0	0	0	0	0	0	5	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa	Candidatus_Cardinium		0.33	1		173	internal			Ostracoda	
seq392	0	0	4	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.07	0.99		173	internal			Ostracoda		
seq393	0	0	4	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.06	1		173	internal			Ostracoda		
seq394	0	0	4	0	0	0	0	0	0	0	Bacteria	Proteobacteria	Gammaproteobacteria	Betaproteobacteriales	Burkholderiaceae		0.02	0.97		223	external	JO977513		Pseudomonas, Pseudomonadaceae		
seq395	0	0	4	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa	Candidatus_Cardinium		0.06	1		173	internal			Ostracoda	
seq396	0	0	4	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa	Candidatus_Cardinium		0.96	0.96		175	external	Hi_1C		H. incongruens, Ostracoda	
seq397	0	0	0	0	0	4	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Flavobacteriales	Weeksellaceae		0.05	1		210	external	DO279361		Chryseobacterium, Flavobacteriaceae		
seq398	0	0	0	0	0	0	4	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.32	0.97		173	external	Hi_1C		H. incongruens, Ostracoda		
seq399	0	0	0	0	0	0	0	4	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.33	1		173	internal			Ostracoda		
seq400	0	0	0	0	0	0	0	0	4	0	Bacteria	Bacteroidetes	Bacteroidia	Flavobacteriales	Flavobacteriaceae	Flavobacterium		1	1		208	external	KJ057195		Flavobacterium, Flavobacteriaceae	
seq401	3	0	0	0	0	0	0	0	0	0	Bacteria	Proteobacteria	Gammaproteobacteria	Betaproteobacteriales	Burkholderiaceae		0.27	0.97		2	internal	JO435729		Pseudoxanthomonas, Xanthomonadaceae		
seq402	0	0	0	0	0	0	0	3	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.32	0.97		175	external	Hi_1C		H. incongruens, Ostracoda		
seq403	0	0	0	0	0	0	0	3	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.33	1		173	internal			Ostracoda		
seq404	0	0	0	0	0	0	0	3	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.33	1		173	internal			Ostracoda		
seq405	2	0	0	0	0	0	0	0	0	0	Bacteria	Proteobacteria	Gammaproteobacteria	Betaproteobacteriales	Burkholderiaceae		0.23	0.96		2	internal	JO435729		Pseudoxanthomonas, Xanthomonadaceae		
seq406	0	0	2	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa	Candidatus_Cardinium		0.03	0.97		174	external	454		E. virens, Ostracods	
seq407	0	0	2	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa	Candidatus_Cardinium		0.03	0.97		174	external	454		E. virens, Ostracods	
seq408	0	0	0	2	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Flavobacteriales	Flavobacteriaceae		0.07	0.98		223	external	KF054791		Sphingobacteriaceae; Sphingobacterium		
seq409	0	0	0	0	0	0	2	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Flavobacteriales	Weeksellaceae		0.04	0.97		209	internal	DO16669		Chryseobacterium, Flavobacteriaceae		
seq410	0	0	0	0	0	0	0	2	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.33	0.99		173	internal			Ostracoda		
seq411	0	0	0	0	0	0	0	2	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.32	0.99		175	external	Hi_1C		H. incongruens, Ostracoda		
seq412	0	0	0	0	0	0	0	2	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.33	1		173	internal			Ostracoda		
seq413	0	1	0	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa	Candidatus_Cardinium		1	1		172	external	DS_ESP		D. stivensoni, Ostracoda	
seq414	0	0	1	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.04	0.96		174	external	454		E. virens, Ostracods		
seq415	0	0	1	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa	Candidatus_Cardinium		0.96	0.96		175	external	Hi_1C		H. incongruens, Ostracoda	
seq416	0	0	1	0	0	0	0	0	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.08	1		173	internal			Ostracoda		
seq417	0	0	0	0	0	0	0	1	0	0	Bacteria	Bacteroidetes	Bacteroidia	Cytophagales	Amoebozoa		0.32	0.99		175	external	Hi_1C		H. incongruens, Ostracoda		



**Table S4B: Results of mapping the shorter 5' and 3' 16S sequences of *Cardinium* from ostracod hosts on the RAxML reference tree with EPA.**

The RAxML tree was based on an alignment of 1638 basepairs of 16S from various bacteria, including *Cardinium* from different hosts. More details on the identity of ostracod DNA numbers is provided in Table S1A, on the 16S reference sequences in Table S1B. Branch numbers refer to the RAxML reference tree, of which the relevant section is shown in Figure S2B. Green indicates that 16S sequences are placed on external or internal branches of *Cardinium* from non-marine ostracods. EDPL= Expected Distance between Placement Locations (Matsen et al., 2010). Estimated values for EDPL and RAxML weight can range between 0 and 1. Here, only the highest RAxML weights and corresponding EDPL values are shown. The complete EPA results are available from the first author on request.

DNA#	ID	16S region	EDPL in EPA	RAxML weight in EPA	mapped to branch	internal_external	ID branch
143	E. virens	5'	0.09	1	40	external	1C, H. incongruens
151	E. virens	3'	0.33	1	40	external	1C, H. incongruens
198	E. virens	3'	0.33	1	41	external	454, E. virens
207	E. virens	5'	0.09	1	40	external	1C, H. incongruens
217	E. virens	3'	0.33	1	40	external	1C, H. incongruens
218	E. virens	3'	0.33	1	41	external	454, E. virens
225	E. virens	5'	0.2	1	39	internal	Ostracoda
262	E. virens	3'	0.33	1	40	external	1C, H. incongruens
324	E. virens	3'	0.33	1	41	external	454, E. virens
349	E. virens	3'	0.33	1	40	external	1C, H. incongruens
351	E. virens	3'	0.33	1	40	external	1C, H. incongruens
361	E. virens	3'	0.33	1	40	external	1C, H. incongruens
362	E. virens	3'	0.33	1	40	external	1C, H. incongruens
363	E. virens	3'	0.33	1	40	external	1C, H. incongruens
364	E. virens	3'	0.33	1	41	external	454, E. virens
386	E. virens	3'	0.33	1	40	external	1C, H. incongruens
391	E. virens	3'	0.33	1	40	external	1C, H. incongruens
450	E. virens	5'	0.2	1	39	internal	Ostracoda
451	E. virens	3'	0.33	1	40	external	1C, H. incongruens
466	E. virens	3'	0.33	1	40	external	1C, H. incongruens
CALC	H. incongruens	5'	1	1	40	external	1C, H. incongruens
Hi_A	H. incongruens	3'	1	1	40	external	1C, H. incongruens
Hi_B	H. incongruens	3'	1	1	40	external	1C, H. incongruens
Hi_C	H. incongruens	3'	1	1	40	external	1C, H. incongruens
Hi_D	H. incongruens	3'	1	1	40	external	1C, H. incongruens
Hi_E	H. incongruens	3'	1	1	40	external	1C, H. incongruens
Hi_F	H. incongruens	3'	1	1	40	external	1C, H. incongruens
Hi_6A	H. incongruens	5'	1	1	40	external	1C, H. incongruens
3A	H. incongruens	5'	1	1	40	external	1C, H. incongruens
5B	H. incongruens	5'	1	1	40	external	1C, H. incongruens

DNA#	ID	16S region	EDPL in EPA	RAXML weight in EPA	mapped to branch	internal_external	ID branch
MC025	<i>C. vidua</i>	3'	0.12	1	41	external	454, <i>E. virens</i>
MC090	<i>C. vidua</i>	3'	0.12	1	41	external	454, <i>E. virens</i>
MC110	<i>C. vidua</i>	3'	0.12	1	41	external	454, <i>E. virens</i>
MC113	<i>C. vidua</i>	3'	0.12	1	41	external	454, <i>E. virens</i>
MC183	<i>C. vidua</i>	3'	0.12	1	41	external	454, <i>E. virens</i>
pbBR0917	<i>V. paglioli</i>	3'	0.16	1	38	internal	Ostracoda
pbBR0918	<i>V. paglioli</i>	3'	0.16	1	38	internal	Ostracoda
Vjp1c	<i>V. cornelia</i>	3'	0.13	1	38	internal	Ostracoda
VC309	<i>V. cornelia</i>	3'	1	1	42	external	<i>D. stevensoni</i>
DsESP11	<i>D. stevensoni</i>	3'	1	1	42	external	<i>D. stevensoni</i>
DsESP3	<i>D. stevensoni</i>	5'	1	1	42	external	<i>D. stevensoni</i>
T5	<i>D. stevensoni</i>	3'	1	1	42	external	<i>D. stevensoni</i>
Ds36	<i>D. stevensoni</i>	3'	1	1	42	external	<i>D. stevensoni</i>
CI1	<i>C. lacustris</i>	3'	0.07	0.96	42	external	<i>D. stevensoni</i>
CI3	<i>C. lacustris</i>	3'	0.07	0.96	42	external	<i>D. stevensoni</i>
CI4	<i>C. lacustris</i>	3'	0.06	0.97	42	external	<i>D. stevensoni</i>
CI5	<i>C. lacustris</i>	3'	0.06	0.97	42	external	<i>D. stevensoni</i>

## References of the supplementary material

- Berger, S.A., Krompaß, D., Stamatakis, A. 2011. Performance, accuracy and web-server for evolutionary placement of short sequence reads under maximum likelihood. *Syst. Biol.* 60, 291-302.
- Bode, S.N.S., Adolfsson, S., Lamatsch, D.K., Martins, M.J.F., Schmit, O., Vandekerkhove, J., Mezquita, F., Namiotko, T., Rossetti, G., Schön, I., Butlin, R.K., Martens, K., 2010. Exceptional cryptic diversity and multiple origins of parthenogenesis in a freshwater ostracod. *Mol. Phyl. Evol.* 54, 542-552.
- Callahan, B.J., McMurdie, P.J., Rosen, M.J., Han, A.W., Johnson, A.J., Holmes, S.P. 2016. DADA2: High-resolution sample inference from Illumina amplicon data. *Nat. Methods* 13, 581-583.
- Duron, O. 2013. Lateral transfers of insertion sequences between *Wolbachia*, *Cardinium* and *Rickettsia* bacterial endosymbionts. *Heredity* 111, 330-337.
- Duron, O., Hurst, G.D., Hornett, E.A., Josling, J.A., Engelstadter, J. 2008. High incidence of the maternally inherited bacterium *Cardinium* in spiders. *Mol. Ecol.* 17, 1427-1437.
- Gotoh, T., Noda, H., Ito, S. 2007. *Cardinium* symbionts cause cytoplasmic incompatibility in spider mites. *Heredity* 98, 13-20.
- Gueguen, G., Vavre, F., Gnankine, O., Peterschmitt, M., Charif, D., Chiel, E., Gottlieb, Y., Ghanim, M., Zchori-Fein, E., Fleury, F. 2010. Endosymbiont metacommunities, mtDNA diversity and the evolution of the *Bemisia tabaci* (Hemiptera: Aleyrodidae) species complex. *Mol. Ecol.* 19, 4365-4376.
- Guindon, S., Gascuel, O., 2003. A simple, fast, and accurate algorithm to estimate large phylogenies by maximum likelihood. *Syst. Biol.* 52, 696-704.
- Kopecky, J., Perotti, M.A., Nesvorna, M., Erban, T., Hubert, J. 2013. *Cardinium* endosymbionts are widespread in synanthropic mite species (Acari: Astigmata). *J. Invertebr. Pathol.* 112, 20-23.
- Kopecky, J., Nesvorna, M., Mareckova-Sagova, M., Hubert, J. 2014. The effect of antibiotics on associated bacterial community of stored product mites. *PLoS ONE* 9, E112919.
- Kurtti, T.J., Munderloh, U.G., Andreadis, T.G., Magnarelli, L.A., Mather, T.N. 1996. Tick cell culture isolation of an intracellular prokaryote from the tick *Ixodes scapularis*. *J. Invertebr. Pathol.* 67, 318-321.
- Liu, Y., Miao, H., Hong, X.-Y. 2006. Distribution of the endosymbiotic bacterium *Cardinium* in Chinese populations of the carmine spider mite *Tetranychus cinnabarinus* (Acari: Tetranychidae). *J. Appl. Entomol.* 130, 523-529.

- Marzorati, M., Alma, A., Sacchi, L., Pajoro, M., Palermo, S., Brusetti, L., Raddadi, N., Balloi, A., Tedeschi, R., Clementi, E., Corona, S., Quaglino, F., Bianco, P.A., Beninati, T., Bandi, C., Daffonchio, D. 2006. A novel Bacteroidetes symbiont is localized in *Scaphoideus titanus*, the insect vector of *Flavescence doree* in *Vitis vinifera*. *Appl. Environ. Microbiol.* 72, 1467-1475.
- Matsen, F.A., Kodner, R.B., Armbrust, E.V. 2010. pplacer: linear time maximum-likelihood and Bayesian phylogenetic placement of sequences onto a fixed reference tree. *BMC Bioinformatics*, 11:538.
- McMurdie, P.J., Holmes, S. 2013. phyloseq: An R package for reproducible interactive analysis and graphics of microbiome census data. *PLoS ONE*. 8, e61217
- Mee, P.T., Weeks, A.R., Walker, P.J., Hoffmann, A.A., Duchemin, J.B. 2015. Novel detection of low level *Cardinium* and *Wolbachia* infections in Culicoides. *Appl. Environ. Microbiol.* 81, 6177-6188.
- Morag, N., Klement, E., Saroya, Y., Lensky, I., Gottlieb, Y. 2012. Prevalence of the symbiont *Cardinium* in Culicoides (Diptera: Ceratopogonidae) vector species is associated with land surface temperature. *FASEB J.* 26, 4025-4034.
- Pages, N., Munoz-Munoz, F., Verdun, M., Pujol, N., Talavera, S. 2017. First detection of *Wolbachia*-infected Culicoides (Diptera: Ceratopogonidae) in Europe: *Wolbachia* and *Cardinium* infection across Culicoides communities revealed in Spain. *Parasit. Vectors* 10, 582.
- Penz, T., Schmitz-Esser, S., Kelly, S.E., Cass, B.N., Muller, A., Woyke, T., Malfatti, S.A., Hunter, M.S., Horn, M. 2012. Comparative genomics suggests an independent origin of cytoplasmic incompatibility in *Cardinium hertigii*. *PLoS Genet.* 8, E1003012.
- Nakagawa, Y., Sakane, T., Suzuki, M. and Hatano, K. 2002. Phylogenetic structure of the genera *Flexibacter*, *Flexithrix*, and *Microscilla* deduced from 16S rRNA sequence analysis. *J. Gen. Appl. Microbiol.* 48, 155-165.
- Nakamura, Y., Kawai, S., Yukuhiro, F., Ito, S., Gotoh, T., Kisimoto, R., Yanase, T., Matsumoto, Y., Kageyama, D., Noda, H. 2009. Prevalence of *Cardinium* bacteria in planthoppers and spider mites and taxonomic revision of 'Candidatus *Cardinium hertigii*' based on detection of a new *Cardinium* group from biting midges. *Appl. Environ. Microbiol.* 75, 6757-6763.
- Noel, G.R., Atibalentja, N. 2006. 'Candidatus *Paenicardinium endonii*', an endosymbiont of the plant-parasitic nematode *Heterodera glycines* (Nemata: Tylenchida), affiliated to the phylum Bacteroidetes. *Int. J. Syst. Evol. Microbiol.* 56, 1697-1702.
- Park, S.H., Kim, J.Y., Kim, Y.J. and Heo, M.S. 2015. *Flavobacterium jejuensis* sp. nov., isolated from marine brown alga *Ecklonia cava*. *J. Microbiol.* 53, 756-761.
- Sakamoto, M., Ohkuma, M. 2010. Usefulness of the hsp60 gene for the identification and classification of Gram-negative anaerobic rods. *J. Med. Microbiol.* 59, 1293-1302.

- Schön, I., Pinto, R.L., Halse, S., Smith, A.J., Martens, K., Birky, C.W., 2012. Cryptic species in putative ancient asexual darwinulids (Crustacea, Ostracoda). Plos ONE 7, e39844.
- Singh, S.T., Priya, N.G., Kumar, J., Rana, V.S., Ellango, R., Joshi, A., Priyadarshini, G., Asokan, R., Rajagopal, R. 2012. Diversity and phylogenetic analysis of endosymbiotic bacteria from field caught *Bemisia tabaci* from different locations of North India based on 16S rDNA library screening. Infect. Genet. Evol. 12, 411-419.
- Weeks, A.R., Velten, R., Stouthamer, R. 2003. Incidence of a new sex-ratio-distorting endosymbiotic bacterium among arthropods. Proc. R. Soc. Lond. B Biol. Sci. 270, 1857-1865.
- Weisburg, W.G., Barns, S.M., Pelletier, D.A., Lane, D.J. 1991. 16S ribosomal DNA amplification for phylogenetic study. J. Bact. 173, 697-703.
- Wickham, H. 2016. ggplot2: Elegant graphics for data analysis. Springer-Verlag New York.
- Zchori-Fein, E., Perlman, S.J. 2004. Distribution of the bacterial symbiont Cardinium in arthropods. Mol. Ecol. 13, 2009-2016.