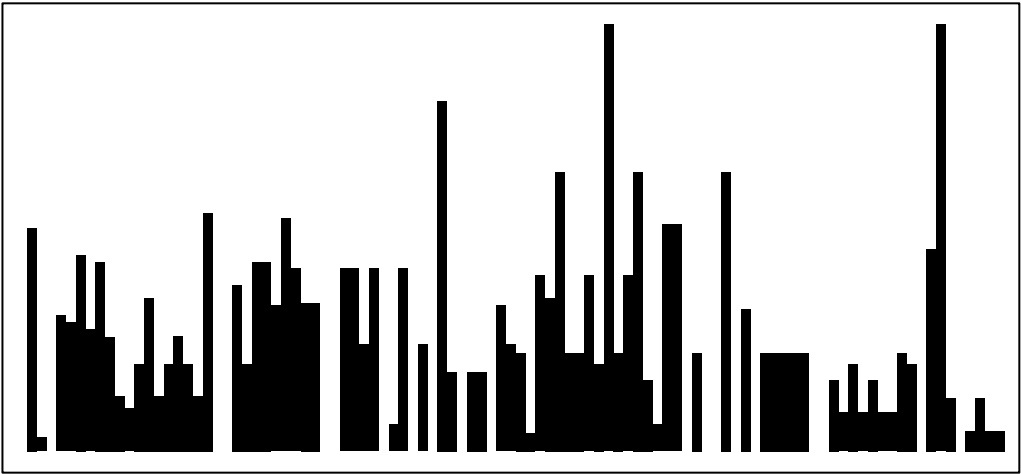


OTU_69.csv

Salina sp. BXW–2014

Paronellidae

90.48



Salina sp. BXW–2014 16S ribosomal RNA gene, partia	KM978386	90.48	28.77	1.99806e–05	Paronellidae	Salina sp. BXW–2014
Orchesellides sp. 9688–2 FZ–2013 16S ribosomal RNA	KC236217	80.65	42.47	1.99806e–05	Entomobryidae	Orchesellides sp. 9688–2 FZ–2013
Sinella cf. colorata FZ–2013 16S ribosomal RNA gen	KC236216	76.25	54.11	0.000243414	Entomobryidae	Sinella cf. colorata FZ–2013
Angela sp. MN120 16S ribosomal RNA gene, partial s	EF383381	86.36	30.14	0.000849598	Mantidae	Angela sp. MN120
Callyntrura quangdongensis 16S ribosomal RNA gene,	KM978382	86.05	29.45	0.00296539	Paronellidae	Callyntrura quangdongensis
Drepanura sp. 1 FZ–2013 16S ribosomal RNA gene, pa	KC236193	89.19	25.34	0.0103502	Entomobryidae	Drepanura sp. 1 FZ–2013
Milionia basalis gene for 16S rRNA, partial sequen	AB265396	85.71	28.77	0.0103502	Geometridae	Milionia basalis
Cyphoderus javanus 16S ribosomal RNA gene, partial	KM978390	88.89	24.66	0.0361258	Cyphoderidae	Cyphoderus javanus
Plaxiphora biramosa 16S ribosomal RNA gene, partia	KJ574069	85.37	28.08	0.0361258	Mopaliidae	Plaxiphora biramosa
Dracunculus medinensis genome assembly D_medinensi	LK978224	82.61	31.51	0.0361258	Dracunculidae	Dracunculus medinensis
Iravadiidae gen. 3 sp. 1 REG–2013 isolate IC–BOW–A	KC439885	82	34.25	0.0361258	Iravadiidae	Iravadiidae gen. 3 sp. 1 REG–2013
Tomocerus similis voucher C9674_1 16S ribosomal RN	KF591903	84.09	30.14	0.0361258	Tomoceridae	Tomocerus similis
Folsomia quadrioculata 16S ribosomal RNA gene, par	KC236199	87.18	26.71	0.0361258	Isotomidae	Folsomia quadrioculata
Epitonium ancillotoi voucher RMNH88484 16S ribosom	JF750949	82.61	31.51	0.0361258	Epitoniidae	Epitonium ancillotoi
Decellebruchus atrolineatus voucher BRU–Aa11 16S r	HQ178239	84.09	30.14	0.0361258	Chrysomelidae	Decellebruchus atrolineatus
Crenatula avicularis voucher USNM:795306 16S ribos	HQ329399	85.42	32.19	0.0361258	Isognomonidae	Crenatula avicularis
Noualhierana furtiva voucher CW 224 16S ribosomal	FJ230432	84.09	30.14	0.0361258	Reduviidae	Noualhierana furtiva
Glycine max strain Williams 82 clone GM_WBb0093C02	AC235350	82.61	31.51	0.0361258	Fabaceae	Glycine max
Candidatus Azobacteroides pseudotrichonymphae geno	AP010656	91.18	23.29	0.0361258		Candidatus Azobacteroides pseudotrichonymphae genomovar. CF
Friesea grisea mitochondrion, complete genome	EU016196	73.42	54.11	0.0361258	Neanuridae	Friesea grisea
Orchesella villosa mitochondrion, complete genome	EU016195	75.34	49.32	0.0361258	Entomobryidae	Orchesella villosa
Neogovea sp. MCZDNA101409 16S ribosomal RNA gene,	DQ825619	87.8	27.4	0.0361258	Neogoveidae	Neogovea sp. MCZDNA101409
Archiearis parthenias gene for 16S rRNA, partial s	AB265368	84.09	30.14	0.0361258	Geometridae	Archiearis parthenias
Eublaberus posticus voucher G49 16S ribosomal RNA	EF363268	88.89	30.14	0.0361258	Blaberidae	Eublaberus posticus
Eublaberus posticus mitochondrion 16S ribosomal RN	U17785	88.89	30.14	0.0361258	Blaberidae	Eublaberus posticus
Salina celebensis 16S ribosomal RNA gene, partial	KM978388	86.84	26.03	0.126092	Paronellidae	Salina celebensis
Ovis canadensis canadensis isolate 43U chromosome	CP011888	90.91	22.6	0.126092	Bovidae	Ovis canadensis canadensis
Enterobius vermicularis genome assembly E_vermicul	LM416355	88.57	23.97	0.126092	Oxyuridae	Enterobius vermicularis
Schistosoma mattheei genome assembly S_mattheei_De	LM156157	86.96	30.14	0.126092	Schistosomatidae	Schistosoma mattheei
Schistosoma curassoni genome assembly S_curassoni_	LM077553	86.96	30.14	0.126092	Schistosomatidae	Schistosoma curassoni
Rhodinia verecunda mitochondrial gene for 16S ribo	AB841469	78.57	38.36	0.126092	Saturniidae	Rhodinia verecunda
Rhodinia verecunda mitochondrial gene for 16S ribo	AB841468	78.57	38.36	0.126092	Saturniidae	Rhodinia verecunda
PREDICTED: Monodelphis domestica TBC1 domain famil	XM_001369738	88.57	23.97	0.126092	Didelphidae	Monodelphis domestica
PREDICTED: Monodelphis domestica TBC1 domain famil	XM_007501140	88.57	23.97	0.126092	Didelphidae	Monodelphis domestica
Folsomia candida voucher Folsomia_Candida 16S ribo	KF591920	85	27.4	0.126092	Isotomidae	Folsomia candida
PREDICTED: Saccoglossus kowalevskii uncharacterize	XM_002739036	88.57	23.97	0.126092	Harrimaniidae	Saccoglossus kowalevskii
Peripatoides sp. DNA103691 16S ribosomal RNA gene,	KC754561	76.19	43.15	0.126092	Peripatopsidae	Peripatoides sp. DNA103691
Stenygrinum quadrinotatum voucher HNAU.PP 013 16S	KF737729	81.25	32.88	0.126092	Cerambycidae	Stenygrinum quadrinotatum
Lamellicidens generosus voucher UMMZ.MC:304346 16S I	KF011263	88.57	23.97	0.126092	Unionidae	Lamellicidens generosus
Orchesellides sinensis 16S ribosomal RNA gene, par	KC236209	74.07	54.79	0.126092	Entomobryidae	Orchesellides sinensis
Folsomia candida 16S ribosomal RNA gene, partial s	KC236200	85	27.4	0.126092	Isotomidae	Folsomia candida
Peripatoides sympatrica mitochondrion, partial gen	JF800075	76.19	43.15	0.126092	Peripatopsidae	Peripatoides sympatrica
Zebrafish DNA sequence from clone DKEY–202K7 in li	CU138513	96.43	19.18	0.126092	Cypriniformes	Danio rerio
Stagmomantis sp. MN278 16S ribosomal RNA gene, par	FJ806254	83.72	29.45	0.126092	Mantodea	Stagmomantis sp. MN278
Glycine max strain Williams 82 clone GM_WBb0002A11	AC235169	76.4	57.53	0.126092	Fabales	Glycine max
Metilia brunnerii isolate MN146 16S ribosomal RNA	EF383392	83.72	29.45	0.126092	Mantodea	Metilia brunnerii
Lithobius variegatus rubriceps 16S ribosomal RNA g	AY084071	83.72	29.45	0.126092	Lithobiomorpha	Lithobius variegatus rubriceps
Homo sapiens cDNA FLJ38749 fis, clone KIDNE2012453	AK096068	77.78	41.78	0.126092	Hominidae	Homo sapiens
Gomphiocephalus hodgsoni mitochondrion, complete g	AY191995	86.84	26.03	0.126092	Hypogastruridae	Gomphiocephalus hodgsoni
Mainwaringia rhizophila 16S ribosomal RNA gene, mi	U46829	85	27.4	0.126092	Littorinidae	Mainwaringia rhizophila
Candidatus Westeberhardia cardiocondylae genome as	LN774881	84.62	26.71	0.440103	Enterobacteriales	Candidatus Westeberhardia cardiocondylae
PREDICTED: Microcebus murinus pleiomorphic adenoma	XM_012774084	80.85	32.19	0.440103	Primates	Microcebus murinus
Bryopsis plumosa chloroplast complete genome, stra	LN810504	88.24	23.29	0.440103	Bryopsidales	Bryopsis plumosa
Caulibugula pearsei voucher BRBRY–Cau2 16S ribosom	KF714807	87.18	26.71	0.440103	Hemiptera	Caulibugula pearsei
Protopolystoma xenopodis genome assembly P_xenopod	LM861591	93.1	19.86	0.440103	Polypisthocotylea	Protopolystoma xenopodis
Schistosoma margrebowiei genome assembly S_margreb	LL888452	84.62	26.71	0.440103	Strigeidida	Schistosoma margrebowiei
Schistosoma margrebowiei genome assembly S_margreb	LL883872	84.62	26.71	0.440103	Strigeidida	Schistosoma margrebowiei
Gongylonema pulchrum genome assembly G_pulchrum_Ho	LL748524	88.24	23.29	0.440103	Spirurida	Gongylonema pulchrum
Brugia pahangi genome assembly B_pahangi_Glasgow ,	LK964536	84.09	30.14	0.440103	Spirurida	Brugia pahangi
Angiostrongylus costaricensis genome assembly A_co	LK939637	100	16.44	0.440103	Rhabditida	Angiostrongylus costaricensis
Schistosoma mattheei genome assembly S_mattheei_De	LM180696	84.62	26.71	0.440103	Strigeidida	Schistosoma mattheei
Strongyloides ratti genome assembly S_ratti_ED321	LN609528	88.24	23.29	0.440103	Rhabditida	Strongyloides ratti
Plasmodium chabaudi chabaudi genome assembly PCHAS	LK022885	93.1	19.86	0.440103	Haemosporida	Plasmodium chabaudi chabaudi
Emmelinea monodactyla mitochondrion, partial genome	KJ508063	83.33	28.77	0.440103	Pterophoridae	Emmelinea monodactyla
Monochamus alternatus mitochondrion, complete geno	KJ809086	81.25	43.15	0.440103	Coleoptera	Monochamus alternatus
PREDICTED: Nasonia vitripennis uncharacterized LOC	XM_008217494	90.62	21.92	0.440103	Hymenoptera	Nasonia vitripennis
PREDICTED: Nasonia vitripennis uncharacterized LOC	XM_003424965	90.62	21.92	0.440103	Hymenoptera	Nasonia vitripennis
Coecobrya communis 16S ribosomal RNA gene, partial	KC236218	80	34.25	0.440103	Entomobryidae	Coecobrya communis
Entomobrya sp. 2 FZ–2013 16S ribosomal RNA gene, p	KC236198	84.62	26.71	0.440103	Entomobryidae	Entomobrya sp. 2 FZ–2013
Coecobrya tenebricosa 16S ribosomal RNA gene, part	KC236191	80	34.25	0.440103	Entomobryidae	Coecobrya tenebricosa
Phaedrotoma rudis isolate CVA4259 16S ribosomal RN	JQ736266	71.13	60.27	0.440103	Braconidae	Phaedrotoma rudis
Ciona intestinalis clone 4H2_9L22 genomic sequence	JF357717	93.1	19.86	0.440103	Cionidae	Ciona intestinalis
Hypogastrura vernalis 16S ribosomal RNA gene, part	JN970945	73.2	64.38	0.440103	Hypogastruridae	Hypogastrura vernalis
Eublaberus posticus voucher ZMUC:JD–879 16S riboso	JN615276	86.67	30.14	0.440103	Blaberidae	Eublaberus posticus
Dichromacalles dioctletianus bio–material ZFMK–DNA	GU988590	76.19	43.15	0.440103	Coleoptera	Dichromacalles dioctletianus
Armatoglyptes habei isolate Acro–sp–13 16S ribosom	HQ148947	84.62	26.71	0.440103	Pygophora	Armatoglyptes habei
Armatoglyptes habei isolate Acro–sp–10 16S ribosom	HQ148946	84.62	26.71	0.440103	Pygophora	Armatoglyptes habei
Armatoglyptes habei isolate Acro–sp–8 16S ribosoma	HQ148945	84.62	26.71	0.440103	Pygophora	Armatoglyptes habei
Armatoglyptes habei isolate Acro–sp–6 16S ribosoma	HQ148944	84.62	26.71	0.440103	Pygophora	Armatoglyptes habei
Armatoglyptes habei isolate Acro–sp–3 16S ribosoma	HQ148943	84.62	26.71	0.440103	Pygophora	Armatoglyptes habei
Aphengoecus multiserratus isolate AMDAR03 16S ribo	GQ289726	79.37	42.47	0.440103	Coleoptera	Aphengoecus multiserratus
Aphengoecus multiserratus isolate AMDAR02 16S ribo	GQ289725	79.37	42.47	0.440103	Coleoptera	Aphengoecus multiserratus
Xanthomantis mantispoides isolate MN335 16S riboso	FJ806306	83.33	28.77	0.440103	Mantodea	Xanthomantis mantispoides
Empusa sp. MN131 16S ribosomal RNA gene, partial s	FJ806202	81.82	30.14	0.440103	Mantodea	Empusa sp. MN131
Parastagmatoptera sp. MN028 16S ribosomal RNA gene	FJ806159	84.09	30.14	0.440103	Mantidae	Parastagmatoptera sp. MN028
Psychobiella sordida 16S ribosomal RNA gene, parti	EU734896	81.82	30.14	0.440103	Neuroptera	Psychobiella sordida
Notocrypta curvifascia 16S ribosomal RNA gene, par	DQ445109	83.33	28.77	0.440103	Lepidoptera	Notocrypta curvifascia
Brotia verbecki voucher ZMB 200.118b 16S ribosomal	AY330780	81.82	30.14	0.440103	Pachychilidae	Brotia verbecki
Brotia verbecki voucher ZMB 200.118 16S ribosomal	AY330779	81.82	30.14	0.440103	Pachychilidae	Brotia verbecki
Podura aquatica mitochondrion, partial genome	AY639939	84.62	26.71	0.440103	Poduridae	Podura aquatica
Parastagmatoptera sp. MN028 16S ribosomal RNA gene	AY491117	84.09	30.14	0.440103	Mantidae	Parastagmatoptera sp. MN028
Pachysoma bennigseni 16S ribosomal RNA gene, parti	AF499698	77.78	43.15	0.440103	Scarabaeidae	Pachysoma bennigseni
Zebrafish DNA sequence from clone RP71–4M17 in lin	BX296520	89.47	24.66	0.440103	Cypriniformes	Danio rerio
Zebrafish DNA sequence from clone CH211–132F13 in	BX005096	100	16.44	0.440103	Cypriniformes	Danio rerio
Folsomia candida 16S ribosomal RNA gene, partial s	JN970944	82.5	27.4	1.53611	Isotomidae	Folsomia candida
Monochamus alternatus endai mitochondrial gene for	AB533599	79.69	43.15	1.53611	Cerambycidae	Monochamus alternatus endai
Epactoides mangabeensis voucher Ale.AND24.32.4 16S	EU030499	80.95	28.77	1.53611	Scarabaeidae	Epactoides mangabeensis
Folsomia candida 16S ribosomal RNA gene, partial s	AY555551	82.5	27.4	1.53611	Isotomidae	Folsomia candida
Epactoides mangabeensis voucher Ale.AND24.32.2 16S	EU030498	80.95	28.77	5.36155	Scarabaeidae	Epactoides mangabeensis
Epactoides mangabeensis voucher Ale.AND24.32.1 16S	EU030497	80.95	28.77	5.36155	Scarabaeidae	Epactoides mangabeensis