

Supplementary Information

Dynamic changes in prokaryotic and eukaryotic communities and networks in a minimally managed cabbage-cultivated field soils

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Supplementary Table S1. Abundant SVs in each prokaryotic phylum and their taxa based on the SILVA database.

ID	Phylum	Class	Order	Family	Genus	Species
SV_4	Acidobacteriota	Vicinamibacteria	Vicinamibacterales	Vicinamibacteraceae	NA	NA
SV_5		Vicinamibacteria	Vicinamibacterales	Vicinamibacteraceae	NA	NA
SV_12		Subgroup_22	NA	NA	NA	uncultured_Holophaga
SV_20		Subgroup_11	NA	NA	NA	uncultured_bacterium
SV_26		Vicinamibacteria	Vicinamibacterales	Vicinamibacteraceae	NA	NA
SV_30		Blastocatellia	Blastocatellales	Blastocatellaceae	uncultured	uncultured_Acidobacteria
SV_33		Thermoanaerobaculia	Thermoanaerobacul	Thermoanaerobacul	Subgroup_10	NA
SV_42		Vicinamibacteria	Vicinamibacterales	Vicinamibacteraceae	NA	NA
SV_43		Blastocatellia	Pyrinomonadales	Pyrinomonadaceae	RB41	NA
SV_49		Blastocatellia	Blastocatellales	Blastocatellaceae	JGI_0001001-H03	uncultured_bacterium
SV_9	Actinobacteriota	Actinobacteria	Propionibacterales	Nocardiodiaceae	NA	NA
SV_19		Actinobacteria	Micrococcales	Micrococcaceae	NA	NA
SV_34		Actinobacteria	Micrococcales	Micrococcaceae	NA	NA
SV_51		Acidimicrobiia	Microtrichales	Ilumatobacteraceae	Ilumatobacter	NA
SV_75		Actinobacteria	Micrococcales	Intrasporangiaceae	NA	NA
SV_97		Acidimicrobiia	IMCC26256	NA	NA	uncultured_Actinomycetales
SV_124		Acidimicrobiia	IMCC26256	NA	NA	NA
SV_134		Actinobacteria	Micrococcales	Intrasporangiaceae	NA	NA
SV_142		Thermoleophilia	Gaiellales	Gaiellaceae	Gaiella	NA
SV_146		Actinobacteria	Frankiales	uncultured	NA	NA
SV_85	Armatimonadota	Fimbriimonadia	Fimbriimonadales	Fimbriimonadaceae	NA	uncultured_bacterium
SV_157		Fimbriimonadia	Fimbriimonadales	Fimbriimonadaceae	NA	uncultured_bacterium
SV_188		Fimbriimonadia	Fimbriimonadales	Fimbriimonadaceae	NA	uncultured_bacterium
SV_438		Fimbriimonadia	Fimbriimonadales	Fimbriimonadaceae	NA	uncultured_bacterium
SV_491		Fimbriimonadia	Fimbriimonadales	Fimbriimonadaceae	NA	NA
SV_560		Chthonomonadetes	Chthonomonadales	NA	NA	uncultured_bacterium
SV_599		Fimbriimonadia	Fimbriimonadales	Fimbriimonadaceae	NA	NA
SV_647		Fimbriimonadia	Fimbriimonadales	Fimbriimonadaceae	NA	uncultured_bacterium
SV_702		Fimbriimonadia	Fimbriimonadales	Fimbriimonadaceae	NA	NA
SV_798		Fimbriimonadia	Fimbriimonadales	Fimbriimonadaceae	NA	uncultured_bacterium
SV_8	Bacteroidota	Bacteroidia	Chitinophagales	Chitinophagaceae	uncultured	uncultured_Bacteroidetes
SV_29		Bacteroidia	Chitinophagales	Saprospiraceae	uncultured	NA
SV_58		Bacteroidia	Chitinophagales	Chitinophagaceae	uncultured	NA
SV_128		Bacteroidia	Chitinophagales	Saprospiraceae	uncultured	NA
SV_191		Bacteroidia	Cytophagales	Microscillaceae	Chryseolinea	uncultured_bacterium
SV_194		Bacteroidia	Cytophagales	Hymenobacteraceae	Adhaeribacter	uncultured_soil
SV_230		Bacteroidia	Chitinophagales	Saprospiraceae	uncultured	NA
SV_241		Bacteroidia	Chitinophagales	Chitinophagaceae	Aurantisolimonas	uncultured_bacterium
SV_251		Bacteroidia	Sphingobacterales	AKYH767	NA	metagenome
SV_257		Bacteroidia	Cytophagales	Hymenobacteraceae	Adhaeribacter	uncultured_soil
SV_442	Bdellovibrionota	Bdellovibrionia	Bdellovibrionales	Bdellovibrionaceae	OM27_clade	uncultured_bacterium
SV_499		Bdellovibrionia	Bdellovibrionales	Bdellovibrionaceae	OM27_clade	uncultured_Myxococcales
SV_1059		Bdellovibrionia	Bdellovibrionales	Bdellovibrionaceae	OM27_clade	metagenome
SV_1321		Bdellovibrionia	Bdellovibrionales	Bdellovibrionaceae	OM27_clade	metagenome
SV_1536		Bdellovibrionia	Bacteriovoracales	Bacteriovoracaceae	Peredibacter	Peredibacter_starrii
SV_1711		Oligoflexia	0319-6G20	NA	NA	NA
SV_1775		Oligoflexia	0319-6G20	NA	NA	NA
SV_1866		Bdellovibrionia	Bdellovibrionales	Bdellovibrionaceae	OM27_clade	uncultured_bacterium
SV_1903		Bdellovibrionia	Bdellovibrionales	Bdellovibrionaceae	OM27_clade	uncultured_bacterium
SV_2293		Oligoflexia	0319-6G20	NA	NA	NA
SV_2	Chloroflexi	KD4-96	NA	NA	NA	NA
SV_10		KD4-96	NA	NA	NA	NA
SV_60		KD4-96	NA	NA	NA	uncultured_bacterium
SV_79		Gitt-GS-136	NA	NA	NA	uncultured_bacterium
SV_104		OLB14	NA	NA	NA	uncultured_bacterium
SV_107		Gitt-GS-136	NA	NA	NA	uncultured_bacterium
SV_114		Anaerolineae	Anaerolineales	Anaerolineaceae	uncultured	uncultured_prokaryote
SV_120		Gitt-GS-136	NA	NA	NA	NA
SV_180		Gitt-GS-136	NA	NA	NA	NA
SV_305		Anaerolineae	Ardenticatenales	uncultured	NA	uncultured_bacterium
SV_65	Cyanobacteriia	Cyanobacteriia	Cyanobacterales	Phormidiaceae	Tychonema_CCAP_	NA
SV_190		Cyanobacteriia	Chloroplast	NA	NA	NA
SV_222		Cyanobacteriia	Cyanobacterales	Coleofasciculaceae	NA	NA
SV_376		Cyanobacteriia	Cyanobacterales	Coleofasciculaceae	Microcoleus_PCC-7	Microcoleus_sp.

SV_469	Cyanobacteria	Cyanobacteriia	Cyanobacteriales	Coleofasciculaceae	Microcoleus_SAG_1NA	
SV_518		Cyanobacteriia	Oxyphotobacteria_J	Unknown_Family	Leptolyngbya_EcFY	uncultured_bacterium
SV_800		Cyanobacteriia	Cyanobacteriales	Nostocaceae	NA	NA
SV_1132		Cyanobacteriia	Cyanobacteriales	Nostocaceae	Nostoc_PCC-7524	Nostoc_sp.
SV_1244		Cyanobacteriia	Cyanobacteriales	Phormidiaceae	uncultured	uncultured_bacterium
SV_1363		Cyanobacteriia	Cyanobacteriales	Nostocaceae	NA	NA
SV_140	Desulfobacterota	uncultured	NA	NA	NA	uncultured_bacterium
SV_300		Desulfuromonadia	Geobacterales	Geobacteraceae	uncultured	uncultured_delta
SV_356		uncultured	NA	NA	NA	NA
SV_494		Desulfuromonadia	Geobacterales	Geobacteraceae	NA	NA
SV_575		uncultured	NA	NA	NA	NA
SV_650		uncultured	NA	NA	NA	NA
SV_668		uncultured	NA	NA	NA	NA
SV_768		uncultured	NA	NA	NA	uncultured_bacterium
SV_832		Desulfuromonadia	Geobacterales	Geobacteraceae	NA	NA
SV_1057		Desulfuromonadia	Geobacterales	Geobacteraceae	NA	NA
SV_999	Elusimicrobiota	Lineage_IIa	NA	NA	NA	uncultured_bacterium
SV_1434		Lineage_IIa	NA	NA	NA	uncultured_bacterium
SV_1459		Lineage_IIb	NA	NA	NA	uncultured_soil
SV_1531		Elusimicrobia	Lineage_IV	NA	NA	uncultured_Termite
SV_1590		Lineage_IIb	NA	NA	NA	uncultured_bacterium
SV_1595		Lineage_IIa	NA	NA	NA	NA
SV_2014		Elusimicrobia	MVP-88	NA	NA	NA
SV_2146		Lineage_IIb	NA	NA	NA	uncultured_bacterium
SV_2185		Lineage_IIa	NA	NA	NA	NA
SV_2322		Elusimicrobia	Lineage_IV	NA	NA	uncultured_bacterium
SV_584	Firmicutes	Bacilli	Bacillales	Bacillaceae	Bacillus	NA
SV_826		Bacilli	Alicyclobacillales	Alicyclobacillaceae	Tumebacillus	uncultured_bacterium
SV_977		Bacilli	Bacillales	Bacillaceae	Bacillus	NA
SV_980		Bacilli	Bacillales	Bacillaceae	Bacillus	Bacillus_luciferensis
SV_981		Bacilli	Bacillales	Bacillaceae	Bacillus	NA
SV_1037		Bacilli	Bacillales	Bacillaceae	Bacillus	NA
SV_1167		Bacilli	Bacillales	Bacillaceae	Bacillus	NA
SV_1930		Clostridia	Clostridiales	Clostridiaceae	Clostridium_sensu_s	NA
SV_2278		Clostridia	Clostridiales	Clostridiaceae	Clostridium_sensu_s	NA
SV_2580		Bacilli	Alicyclobacillales	Alicyclobacillaceae	Tumebacillus	uncultured_bacterium
SV_13	Gemmatimonadota	Gemmatimonadetes	Gemmatimonadales	Gemmatimonadacea	uncultured	uncultured_soil
SV_25		Gemmatimonadetes	Gemmatimonadales	Gemmatimonadacea	uncultured	uncultured_bacterium
SV_31		Gemmatimonadetes	Gemmatimonadales	Gemmatimonadacea	uncultured	uncultured_Gemmatimonadales
SV_39		AKAU4049	NA	NA	NA	uncultured_soil
SV_46		S0134_terrestrial_group	NA	NA	NA	uncultured_bacterium
SV_47		Gemmatimonadetes	Gemmatimonadales	Gemmatimonadacea	uncultured	uncultured_bacterium
SV_53		Gemmatimonadetes	Gemmatimonadales	Gemmatimonadacea	uncultured	metagenome
SV_55		Gemmatimonadetes	Gemmatimonadales	Gemmatimonadacea	uncultured	NA
SV_61		Gemmatimonadetes	Gemmatimonadales	Gemmatimonadacea	uncultured	uncultured_bacterium
SV_66		Gemmatimonadetes	Gemmatimonadales	Gemmatimonadacea	uncultured	NA
SV_35	Latescibacterota	NA	NA	NA	NA	metagenome
SV_52		Latescibacteria	Latescibacteriales	Latescibacteraceae	NA	uncultured_bacterium
SV_73		NA	NA	NA	NA	uncultured_soil
SV_152		NA	NA	NA	NA	metagenome
SV_153		NA	NA	NA	NA	uncultured_bacterium
SV_171		NA	NA	NA	NA	NA
SV_263		NA	NA	NA	NA	metagenome
SV_280		NA	NA	NA	NA	uncultured_bacterium
SV_361		NA	NA	NA	NA	metagenome
SV_452		NA	NA	NA	NA	NA
SV_70	Methyloirabilia	Methyloirabilia	Rokubacteriales	WX65	NA	uncultured_bacterium
SV_93		Methyloirabilia	Rokubacteriales	NA	NA	uncultured_bacterium
SV_98		Methyloirabilia	Rokubacteriales	NA	NA	uncultured_bacterium
SV_130		Methyloirabilia	Rokubacteriales	NA	NA	uncultured_bacterium
SV_200		Methyloirabilia	Rokubacteriales	NA	NA	uncultured_bacterium
SV_211		Methyloirabilia	Rokubacteriales	NA	NA	uncultured_bacterium
SV_245		Methyloirabilia	Rokubacteriales	NA	NA	uncultured_bacterium
SV_287		Methyloirabilia	Rokubacteriales	NA	NA	uncultured_bacterium
SV_549		Methyloirabilia	Rokubacteriales	NA	NA	uncultured_bacterium
SV_619		Methyloirabilia	Rokubacteriales	WX65	NA	uncultured_bacterium
SV_86		Polyangia	Polyangiales	Blrii41	NA	uncultured_bacterium

SV_126		Polyangia	Haliangiales	Haliangiaceae	Haliangium	uncultured_Myxococcales
SV_164		Polyangia	Polyangiales	Polyangiaceae	Pajaroellobacter	uncultured_bacterium
SV_170		Polyangia	Polyangiales	Polyangiaceae	Pajaroellobacter	metagenome
SV_236		Polyangia	Haliangiales	Haliangiaceae	Haliangium	uncultured_bacterium
SV_269	Myxococcota	Polyangia	Polyangiales	BRii41	NA	uncultured_Sorangineae
SV_298		Polyangia	Polyangiales	Phaselicystidaceae	Phaselicystis	uncultured_Polyangiaceae
SV_366		Polyangia	Haliangiales	Haliangiaceae	Haliangium	uncultured_bacterium
SV_462		Myxococcia	Myxococcales	Myxococcaceae	Archangium	Archangium_gephyra
SV_467		Polyangia	Haliangiales	Haliangiaceae	Haliangium	uncultured_Myxococcales
SV_168		NA	NA	NA	NA	uncultured_Desulfoglaeba
SV_229		NA	NA	NA	NA	uncultured_bacterium
SV_274		NA	NA	NA	NA	uncultured_bacterium
SV_352		NA	NA	NA	NA	uncultured_Desulfoglaeba
SV_550		NA	NA	NA	NA	uncultured_Desulfoglaeba
SV_600	NB1-j	NA	NA	NA	NA	uncultured_Desulfoglaeba
SV_678		NA	NA	NA	NA	uncultured_Desulfoglaeba
SV_717		NA	NA	NA	NA	uncultured_bacterium
SV_732		NA	NA	NA	NA	NA
SV_784		NA	NA	NA	NA	uncultured_Desulfoglaeba
SV_44		Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	uncultured_soil
SV_82		Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	uncultured_Green
SV_195		Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	uncultured_Green
SV_478		Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	uncultured_Green
SV_1171	Nitrospirota	Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	uncultured_soil
SV_1414		Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	NA
SV_1537		Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	uncultured_organism
SV_1735		Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	uncultured_soil
SV_2523		Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	uncultured_Green
SV_2689		Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	NA
SV_312		ABY1	Candidatus_Magas	NA	NA	uncultured_bacterium
SV_406		Parcubacteria	NA	NA	NA	uncultured_bacterium
SV_451		Parcubacteria	NA	NA	NA	uncultured_bacterium
SV_812		Parcubacteria	Candidatus_Nomur	NA	NA	uncultured_bacterium
SV_1007	Patescibacteria	Parcubacteria	NA	NA	NA	uncultured_bacterium
SV_1022		Parcubacteria	GWA2-38-13b	NA	NA	uncultured_bacterium
SV_1146		Parcubacteria	NA	NA	NA	uncultured_bacterium
SV_1301		Parcubacteria	Candidatus_Yanof	NA	NA	uncultured_bacterium
SV_1358		Parcubacteria	Candidatus_Yanof	NA	NA	NA
SV_1429		Parcubacteria	NA	NA	NA	uncultured_bacterium
SV_37		Phycisphaerae	Tepidisphaerales	WD2101_soil_grou	NA	uncultured_soil
SV_57		Phycisphaerae	Tepidisphaerales	WD2101_soil_grou	NA	uncultured_bacterium
SV_91		Planctomycetes	Pirellulales	Pirellulaceae	Pirellula	uncultured_soil
SV_96		OM190	NA	NA	NA	uncultured_bacterium
SV_121	Planctomycetota	Planctomycetes	Planctomycetales	Rubinisphaeraceae	SH-PL14	uncultured_Planctomycetaceae
SV_129		Phycisphaerae	Tepidisphaerales	WD2101_soil_grou	NA	uncultured_bacterium
SV_135		Phycisphaerae	Tepidisphaerales	WD2101_soil_grou	NA	NA
SV_137		Phycisphaerae	Tepidisphaerales	WD2101_soil_grou	NA	uncultured_bacterium
SV_160		Phycisphaerae	Tepidisphaerales	WD2101_soil_grou	NA	uncultured_soil
SV_162		Phycisphaerae	Tepidisphaerales	WD2101_soil_grou	NA	uncultured_planctomycete
SV_1		Alphaproteobacteria	Azospirillales	uncultured	NA	uncultured_bacterium
SV_3		Gammaproteobacteria	Burkholderiales	Nitrosomonadaceae	MND1	NA
SV_6		Alphaproteobacteria	Dongiales	Dongiaceae	Dongia	NA
SV_7		Gammaproteobacteria	Burkholderiales	Nitrosomonadaceae	MND1	NA
SV_11	Proteobacteria	Alphaproteobacteria	Rhizobiales	Xanthobacteraceae	NA	NA
SV_14		Gammaproteobacteria	Burkholderiales	Nitrosomonadaceae	MND1	NA
SV_15		Gammaproteobacteria	Burkholderiales	SC-I-84	NA	uncultured_bacterium
SV_16		Gammaproteobacteria	Xanthomonadales	Xanthomonadaceae	Lysobacter	NA
SV_21		Gammaproteobacteria	Burkholderiales	Nitrosomonadaceae	Ellin6067	NA
SV_22		Gammaproteobacteria	CCD24	NA	NA	NA
SV_17		Verrucomicrobiae	Pedosphaerales	Pedosphaeraceae	uncultured	NA
SV_18		Verrucomicrobiae	Opitutales	Opitutaceae	Opitutus	NA
SV_24		Verrucomicrobiae	Pedosphaerales	Pedosphaeraceae	uncultured	metagenome
SV_28		Verrucomicrobiae	Pedosphaerales	Pedosphaeraceae	uncultured	metagenome
SV_32	Verrucomicrobiota	Verrucomicrobiae	Pedosphaerales	Pedosphaeraceae	NA	NA
SV_40		Verrucomicrobiae	Chthoniobacterales	Xiphinematobacteria	Candidatus_Xiphinei	NA
SV_69		Verrucomicrobiae	Pedosphaerales	Pedosphaeraceae	NA	uncultured_bacterium
SV_113		Verrucomicrobiae	Pedosphaerales	Pedosphaeraceae	NA	uncultured_bacterium

SV_131	Verrucomicrobiae	Pedosphaerales	Pedosphaeraceae	NA	NA
SV_149	Verrucomicrobiae	Pedosphaerales	Pedosphaeraceae	NA	uncultured_subdivision

Abundant SVs in each prokaryotic phylum and their SILVA-based taxa are shown. NA: not assigned. The top 10 SVs in abundance are indicated in the phylum. Minor and unassigned phyla-derived SVs are omitted. SVs in bold were used for Supplementary Fig. S2.

Supplementary Table S2. Abundant SVs in each eukaryotic phylum, their taxa based on the SILVA database, and the top hits identified by BLASTN search.

Fungi										
ID	Phylum	Class	Order	Family	Genus	Species	The top hits by BLASTN search	e-value	Max score	Accession No.
SV_91	Aphelidea	NA	NA	NA	uncultured	uncultured_fungus	Aphelidium parallellum	1e-174	627	LC652435.1
SV_121		NA	NA	NA	uncultured	uncultured_fungus	Aphelidium parallellum	1e-174	658	LC652435.1
SV_921		NA	NA	NA	Paraphelidium	NA	Urophycitis sp.	2e-171	616	HQ888719.1 etc.
SV_986		NA	NA	NA	uncultured	uncultured_eukaryote	Amoebophilidium occidentale	0	710	JX967274.1
SV_1150		NA	NA	NA	uncultured	uncultured_eukaryote	Aphelidium insulamus	0	699	MW186929.1
SV_1379		NA	NA	NA	uncultured	uncultured_fungus	Aphelidium insulamus	0	688	MW186929.1
SV_1471		NA	NA	NA	Paraphelidium	uncultured_eukaryote	Paraphelidium tribonematis	0	761	MK631785.1
SV_1589		NA	NA	NA	NA	NA	Amoebophilidium occidentale	0	737	JX967274.1
SV_1764		NA	NA	NA	Paraphelidium	uncultured_eukaryote	Paraphelidium tribonematis	0	728	MK631785.1
SV_1809		NA	NA	NA	uncultured	uncultured_fungus	Aphelidium parallellum	0	667	LC652435.1
SV_8	Ascomycota	Sordariomycetes	Sordariales	Chaetomiaceae	Chaetomium	NA	Chaetomium globosum/Humicola sp.	0	763	MN602645.1 etc.
SV_12		Sordariomycetes	Hypocreales	Nectriaceae	Fusarium	NA	Fusarium sp.	0	758	MN602645.1 etc.
SV_16		Sordariomycetes	Sordariales	Chaetomiaceae	Chaetomium	NA	Chaetomium globosum/Humicola sp.	0	758	MN602645.1 etc.
SV_21		Dothideomycetes	Capnodiales	Cladosporiaceae	Cladosporium	NA	Cladosporium sp.	0	767	MT644477.1 etc.
SV_27		Sordariomycetes	Sordariales	Chaetomiaceae	Chaetomium	NA	Chaetomium sp./Madurella sp./Corynascus verrucosus/Thermothelomyces sp.	0	763	MT649557.1 etc.
SV_31		Dothideomycetes	Pleosporales	Pleosporaceae	NA	NA	Bipolaris sp./Curvularia sp./Paradendryphiella arenariae	0	767	MT649586.1 etc.
SV_43		Peizomycetes	Pezizales	Rhiziniaceae	Phymatotrichopsis	uncultured_eukaryote	Ascobolus sp.	0	771	MN995515.1 etc.
SV_44		Sordariomycetes	Hypocreales	NA	NA	NA	Fusarium sp.	0	763	MT649540.1 etc.
SV_46		Dothideomycetes	Pleosporales	NA	NA	NA	Parastagonospora sp./Ophiostoma sp./Phaeosphaeria sp./Setomelanomma holmii	0	767	CP039677.1 etc.
SV_48		Dothideomycetes	Pleosporales	NA	NA	NA	Phoma sp./Boeremia sp./Ascochyta sp./Microphacopsis olivacea	0	767	LR745531.1 etc.
SV_17	Basidiomycota	Agaricomycetes	NA	NA	NA	NA	Trechispora sp.	0	713	EU909231.1 etc.
SV_37		Agaricomycetes	Agaricales	NA	NA	NA	Lyophyllum sp./Crassiosporium funariophilum/Nidula niveotomentosa/Psathyrella gracilis/Myochromella boudieri/Hebeloma angustilamellatum/Agaricus sp.	0	769	KU058570.1 etc.
SV_149		Agaricomycetes	Agaricales	NA	NA	NA	Collybia sp./Lyophyllum sp.	0	752	MT644912.1 etc.
SV_165		Agaricomycetes	Auriculariales	NA	NA	NA	Alloceidiopsis calcea	0	739	AY293130.1
SV_167		Agaricomycetes	Corticiales	Corticaceae	Sistotrema	uncultured_fungus	Mimimedusa polyspora	0	769	KC176294.1
SV_184		Agaricomycetes	Agaricales	NA	NA	NA	Hohenbuehelia sp.	0	769	JQ926736.2 etc.
SV_204		Agaricomycetes	Agaricales	NA	NA	NA	Stephanospora caroticolor	0	730	AF518591.1
SV_249		Agaricomycetes	NA	NA	NA	NA	Bjerkandera sp./Ganoderma sp./Phanerochaete chrysosporium/Lentinus crinitus/Hyphodermella sp. etc.	0	752	MT644869.1 etc.
SV_265		Agaricomycetes	uncultured	NA	NA	uncultured_Boletaceae	Serendipita vermifera	0	769	EU625993.1 etc.
SV_358		Agaricomycetes	Agaricales	NA	NA	NA	Eimeria lateralis	0	758	OP407718.1 etc.
SV_259	Blastocladiomycota	Blastocladiomycetes	Blastocladales	Physodermataceae	Paraphysoderma	Paraphysoderma sedobakerense	Paraphysoderma sedobakerense	0	747	MN203631.1 etc.
SV_663		Blastocladiomycetes	Blastocladales	Physodermataceae	Paraphysoderma	Paraphysoderma sedobakerense	Paraphysoderma sedobakerense	0	769	MN203631.1 etc.
SV_715		Blastocladiomycetes	Blastocladales	Physodermataceae	NA	uncultured_fungus	Paraphysoderma sedobakerense	0	758	MN203631.1 etc.
SV_1820		Blastocladiomycetes	Blastocladales	Physodermataceae	Paraphysoderma	Paraphysoderma sedobakerense	Paraphysoderma sedobakerense	0	747	MN203631.1 etc.
SV_1927		Blastocladiomycetes	Blastocladales	Physodermataceae	NA	NA	Rozella sp.	0	730	OQ702856.1
SV_1930		Blastocladiomycetes	Blastocladales	Physodermataceae	Paraphysoderma	Paraphysoderma sedobakerense	Paraphysoderma sedobakerense	0	706	MN203631.1 etc.
SV_2246		Blastocladiomycetes	Blastocladales	Physodermataceae	NA	uncultured_fungus	Paraphysoderma sedobakerense	0	726	MN203631.1 etc.
SV_2429		Blastocladiomycetes	Blastocladales	Physodermataceae	Paraphysoderma	Paraphysoderma sedobakerense	Paraphysoderma sedobakerense	0	747	MN203631.1 etc.
SV_2618		Blastocladiomycetes	Blastocladales	Physodermataceae	NA	NA	Paraphysoderma sedobakerense	0	715	MN203631.1 etc.
SV_3455		Blastocladiomycetes	Blastocladales	Physodermataceae	NA	NA	Paraphysoderma sedobakerense	0	712	MN203631.1 etc.
SV_13	Chytridiomycota	Chytridiomycetes	NA(Chytridiales)	NA	NA	NA	Chytridium polysiphoniae	0	695	AY032608.1
SV_47		Spizellomycetes	Olpidiaceae	Olpidium	NA	NA	Olpidium brassicae	0	763	NG_017176.1
SV_65		Chytridiomycetes	Rhizophydiales	Rhizophydiaceae	Rhizophyidium	uncultured_Rhizophydiales	Rhizophyidium patellarium/Betamycetes sp.	0	725	KF160860.1 etc.
SV_94		Chytridiomycetes	Rhizophydiales	uncultured	NA	uncultured_Chytridiomycota	Rhizophyidium sp. UGA-F16	0	752	AH009034.2
SV_102		Chytridiomycetes	NA	NA	NA	NA	Catenomyces persicius	0	763	AY635830.1
SV_134		Chytridiomycetes	Rhizophydiales	uncultured	NA	uncultured_Rhizophydiales	Paramycetes uniporus	0	656	MT731025.1 etc.
SV_138		Chytridiomycetes	Rhizophydiales	uncultured	NA	Rhizophyidium sp.	Aquamyces sp./Rhizophyidium sp.	0	747	MT730892.1 etc.
SV_139		Chytridiomycetes	NA	NA	NA	NA	Fimicolochytrium sp.	0	763	NG_062391.1 etc.
SV_141		Chytridiomycetes	Rhizophydiales	uncultured	NA	uncultured_fungus	Rhizophyidium sp.	0	769	MT730997.1 etc.
SV_150		Chytridiomycetes	Chytridiales	Incertae_Sedis	NA	Pseudorhizidium_endosporangium	Delfinachytrium mesopotamicum	0	763	MT730617.1
SV_161	Cryptomycota	Incertae_Sedis	Incertae_Sedis	Incertae_Sedis	Rozella	uncultured_fungus	Rozella sp.	0	667	OQ702932.1 etc.
SV_327		NA	NA	NA	NA	uncultured_fungus	Vinosinica sp./Bullera formosensis	6e-157	568	LC431094.1 etc.
SV_421		Incertae_Sedis	Incertae_Sedis	Incertae_Sedis	Rozella	uncultured_fungus	Rozella sp.	0	734	KX354831.1
SV_683		Incertae_Sedis	Incertae_Sedis	Incertae_Sedis	Rozella	uncultured_fungus	Rozella sp.	0	706	OQ702917.1
SV_913		Incertae_Sedis	Incertae_Sedis	Incertae_Sedis	Paramicrosporidium	uncultured_eukaryote	Paramicrosporidium sp.	1e-169	619	MW898450.1
SV_1127		Incertae_Sedis	Incertae_Sedis	Incertae_Sedis	Rozella	uncultured_fungus	Rozella sp.	0	752	OQ702856.1
SV_1265		NA	NA	NA	NA	uncultured_fungus	Vinosinica sp.	1e-158	573	LC431094.1 etc.
SV_1376		Incertae_Sedis	Incertae_Sedis	Incertae_Sedis	Rozella	uncultured_fungus	Rozella sp.	0	706	KX354831.1
SV_1411		NA	NA	NA	NA	NA	Paraphysoderma sedobakerense/Arkaya serpentina	6e-162	584	MN203631.1 etc.
SV_1465		Incertae_Sedis	Incertae_Sedis	Incertae_Sedis	Rozella	uncultured_fungus	Rozella sp.	0	695	OQ702932.1 etc.
SV_147	LKM15	NA	NA	NA	NA	uncultured_eukaryote	Podoscypha sp.	5e-148	538	KY464924.1 etc.
SV_247		NA	NA	NA	NA	metagenome	Morellospora saccaeobae	1e-178	640	MN821072.1
SV_827		NA	NA	NA	NA	NA	Morellospora saccaeobae	0	662	MN821072.1
SV_854		NA	NA	NA	NA	NA	Morellospora saccaeobae	1e-149	544	MN821072.1
SV_1420		NA	NA	NA	NA	uncultured_eukaryote	Chytrium sp. JEL341	8e-161	580	DQ536482.1
SV_1491		NA	NA	NA	NA	NA	Morellospora saccaeobae	5e-158	571	MN821072.1
SV_1770		NA	NA	NA	NA	NA	Morellospora saccaeobae	5e-163	588	MN821072.1
SV_1815		NA	NA	NA	NA	NA	Morellospora saccaeobae	0	669	MN821072.1
SV_2039		NA	NA	NA	NA	metagenome	Morellospora saccaeobae	0	737	MN821072.1
SV_2115		NA	NA	NA	NA	NA	Morellospora saccaeobae	0	710	MN821072.1
SV_9	Mucoromycota	Incertae_Sedis	Mortierellales	Mortierellaceae	Mortierella	NA	Mortierella sp./Linnemannia elongata/Entomortierella sp./Podila verticillata	0	769	MH047197.1 etc.
SV_59		Glomeromycetes	Glomerales	Glomeraceae	Funneliformis	Funneliformis_mosseae	Glomus sp./Funneliformis sp.	0	771	FR750227.1 etc.
SV_90		Glomeromycetes	Diversisporales	Gigasporaceae	NA	NA	Cetranspora gilmorei	0	763	FR773143.1 etc.
SV_124		Incertae_Sedis	Mortierellales	Mortierellaceae	Mortierella	NA	Actinomortierella wolffii	0	758	KU141329.1 etc.
SV_151		Incertae_Sedis	Mortierellales	Mortierellaceae	Mortierella	Mortierella sp.	Mortierella sp. MS-6	0	776	AY635828.1
SV_219		Glomeromycetes	Glomerales	uncultured	NA	NA	Glomus mycorrhizal symbiont of Marchantia foliacea	0	706	AJ699068.1 etc.
SV_241		Glomeromycetes	Diversisporales	NA	NA	Glomeromycotina sp.	Entrophospora sp. AS-2013	0	752	JF414174.1
SV_245		Incertae_Sedis	Mortierellales	Mortierellaceae	Mortierella	Mortierella sp.	Mortierella sp. MS-6	0	765	AY635828.1
SV_266		Glomeromycetes	uncultured	NA	NA	uncultured_fungus	Glomus macrocarpum	0	737	FR750376.1 etc.
SV_273		Glomeromycetes	Diversisporales	Acaulosporaceae	Acaulospora	Glomeromycotina sp.	Acaulospora sp.	0	706	NG_062371.1 etc.

SV_873	Neocallimastigomycota	Neocallimastigomycetes	Neocallimastigales	Neocallimastigaceae	uncultured	uncultured_fungus	Monoblepharis sp.	0	693	KJ668077.1 etc.
SV_1105		Neocallimastigomycetes	Neocallimastigales	Neocallimastigaceae	uncultured	uncultured_fungus	Olpidium brassicae	0	697	NG_017176.1
SV_2743		Neocallimastigomycetes	Neocallimastigales	Neocallimastigaceae	uncultured	uncultured_fungus	Monoblepharis sp.	0	673	KJ668077.1 etc.
SV_3667		NA	NA	Nucleariidae	Nuclearia	NA	Entrophospora etunicata	1e-174	627	Z14008.1
SV_3923	Nucleariidae_and_Fonticola_group	NA	NA	Nucleariidae	Nuclearia	metagenome	Rozella sp.	3e-165	595	OQ702932.1 etc.
SV_3924		NA	NA	Nucleariidae	Nuclearia	metagenome	Rozella sp.	1e-173	623	OQ702932.1 etc.
SV_3986		NA	NA	Nucleariidae	Nuclearia	NA	Harpochytrium sp.	8e-161	580	KJ668051.1 etc.
SV_3989		NA	NA	Nucleariidae	Nuclearia	NA	Acaulospora sp.	6e-172	617	AJ306439.1 etc.
SV_4353		NA	NA	Nucleariidae	Nuclearia	metagenome	Harpochytrium sp.	1e-163	590	KJ668051.1 etc.
SV_4893		NA	NA	Nucleariidae	Nuclearia	NA	Pompholyxophrys sp.	2e-151	549	MK547175.1 etc.
SV_5118		NA	NA	Nucleariidae	Nuclearia	NA	Rozella sp.	7e-162	584	OQ702932.1 etc.
SV_5123		NA	NA	Nucleariidae	Nuclearia	NA	Basidiascus sp. CBS 12316	2e-166	599	LT548272.1
SV_5205		NA	NA	Nucleariidae	Nuclearia	NA	Pompholyxophrys punicea	2e-142	520	MK547175.1
SV_407	Zoopagomycota	Incertae_Sedis	Kickxellales	Kickxellaceae	Ramicandelaber	Ramicandelaber_brevisporus	Ramicandelaber sp. KYK00187	0	754	AB287991.1
SV_729		Entomophthoromycetes	Entomophthorales	Ancylistaceae	Conidiobolus	Conidiobolus_brefeldianus	Conidiobolus brefeldianus	0	787	NG_065500.1
SV_1733		Entomophthoromycetes	Entomophthorales	Ancylistaceae	Conidiobolus	Conidiobolus_coronatus	Conidiobolus coronatus	0	793	FJ895303.1
SV_2038		Incertae_Sedis	Zoopagales	NA	NA	metagenome	Stylopage hadra	0	761	MN252062.1
SV_2213		Entomophthoromycetes	Entomophthorales	uncultured	NA	uncultured_fungus	Neokarlingia chitinophila	8e-171	614	HQ901766.1
SV_2454		Entomophthoromycetes	Entomophthorales	Ancylistaceae	Conidiobolus	NA	Conidiobolus sp. RSL-II	0	782	FJ895304.1
SV_2482		Incertae_Sedis	Zoopagales	NA	NA	metagenome	Stylopage hadra	0	671	MN252062.1
SV_2623		Incertae_Sedis	Kickxellales	Kickxellaceae	Ramicandelaber	Ramicandelaber_brevisporus	Ramicandelaber taiwanensis	0	756	JQ004926.1 etc.
SV_3161		Incertae_Sedis	Harpellales	NA	NA	NA	Paraphysoderma sedebokerense	1e-179	643	MN203631.1 etc.
SV_3315		Basidiobolomycetes	Basidiobolales	Basidiobolaceae	Basidiobolus	NA	Basidiobolus sp.	0	771	MW130090.1 etc.

Protists

ID	Phylum	Class	Order	Family	Genus	Species	The top hits by BLASTN search	e-value	Max score	Accession No.
SV_3914	Ancyromonadida	NA	NA	NA	NA	Ancyromonas_sigmoides	Ancyromonas sp.	0	763	MW872726.1 etc.
SV_4022		NA	NA	NA	NA	Nutomonas_longa	Nutomonas longa	0	800	MW872732.1 etc.
SV_36	Apicomplexa	Conoidasida	Gregarinasina	Archigregarinorida	Selenidium	uncultured_alveolate	Eimeria tropidura	3e-160	579	AF324217.1
SV_818		Conoidasida	Gregarinasina	Eugregarinorida	Elev-18S-1089	uncultured_Eimeriidae	Coronopimeritus otoensis/Quadruspinospora mexicana	0	743	MK181532.1 etc.
SV_1364		Conoidasida	Gregarinasina	Eugregarinorida	Gregarina	NA	Gregarina sp.	2e-166	599	LR814086.1 etc.
SV_1664		Conoidasida	Gregarinasina	Eugregarinorida	uncultured	uncultured_alveolate	Coronopimeritus otoensis/Quadruspinospora mexicana	0	721	MK181532.1 etc.
SV_1835		Conoidasida	Gregarinasina	Eugregarinorida	NA	NA	Coronopimeritus otoensis/Quadruspinospora mexicana	4e-169	608	MK181532.1 etc.
SV_2481		Conoidasida	Gregarinasina	Eugregarinorida	Elev-18S-1089	NA	Coronopimeritus otoensis/Quadruspinospora mexicana	0	710	MK181532.1 etc.
SV_2637		Conoidasida	Gregarinasina	Eugregarinorida	NA	NA	Coronopimeritus otoensis/Quadruspinospora mexicana	0	682	MK181532.1 etc.
SV_2716		Conoidasida	Gregarinasina	Eugregarinorida	Gregarina	NA	Gregarina sp.	8e-161	580	LR814105.1 etc.
SV_2833		Conoidasida	Gregarinasina	Eugregarinorida	Elev-18S-1089	uncultured_Eimeriidae	Coronopimeritus otoensis/Quadruspinospora mexicana	0	710	MK181532.1 etc.
SV_3157		Conoidasida	Gregarinasina	Eugregarinorida	NA	NA	Coronopimeritus otoensis/Quadruspinospora mexicana	2e-172	619	MK181532.1 etc.
SV_680	Apusomonadidae	NA	NA	NA	Apusomonas	Apusomonas_proboscidea	Apusomonas proboscidea	0	723	OM966642.1
SV_1336		NA	NA	NA	Apusomonas	Apusomonas_proboscidea	Apusomonas proboscidea	0	758	DQ207568.1
SV_1394		NA	NA	NA	Apusomonas	Apusomonas_proboscidea	Apusomonas proboscidea	0	747	DQ207568.1
SV_1570		NA	NA	NA	Apusomonas	NA	Apusomonas proboscidea	0	713	DQ207568.1
SV_1876		NA	NA	NA	Apusomonas	Apusomonas_proboscidea	Apusomonas proboscidea	0	763	DQ207567.1
SV_1968		NA	NA	NA	Apusomonas	Apusomonas_proboscidea	Apusomonas proboscidea	0	774	DQ207567.1
SV_2236		NA	NA	NA	Apusomonas	NA	Podomonas kaiyoe/Apusomonas proboscidea	0	652	DQ207568.1 etc.
SV_2608		NA	NA	NA	Apusomonas	NA	Apusomonas australiensis	0	706	OM966637.1
SV_2625		NA	NA	NA	Apusomonas	Apusomonas_proboscidea	Apusomonas proboscidea	0	752	DQ207568.1
SV_2627		NA	NA	NA	Apusomonas	Apusomonas_proboscidea	Apusomonas proboscidea	0	723	OM966642.1
SV_307	Bicosoecida	NA	NA	NA	Bicosoeca	metagenome	Trieres chinensis	1e-153	556	MG835771.1
SV_614		NA	NA	NA	Bicosoeca	uncultured_eukaryote	Bicosoeca petiolata	0	676	AY520444.1
SV_643		NA	NA	NA	NA	NA	Adriamonas peritocrescens	0	654	AF243501.1
SV_750		uncultured	NA	NA	NA	uncultured_microeukaryote	Pseudobodo sp.	8e-156	564	LC683683.1 etc.
SV_824		NA	NA	NA	NA	NA	Nanos amicus	2e-136	499	FJ971853.1
SV_1094		NA	NA	NA	Bicosoeca	metagenome	Bicosoeca kenaiensis	2e-162	586	KM816648.1
SV_1169		NA	NA	NA	NA	NA	Adriamonas peritocrescens	0	654	AF243501.1
SV_1210		K4	NA	NA	NA	uncultured_freshwater	Adriamonas peritocrescens	0	654	AF243501.1
SV_1309		NA	NA	NA	NA	NA	Nanos amicus	2e-142	520	FJ971853.1
SV_1321		NA	NA	NA	Bicosoeca	metagenome	Bicosoeca kenaiensis	2e-157	569	KM816648.1
SV_168	Centrohelida	NA	NA	Heterophryidae	NA	metagenome	Triangulopteris lacunata	0	760	OL739463.1
SV_239		NA	NA	Heterophryidae	NA	metagenome	Khitsovia mutabilis	0	776	OP101622.1
SV_535		NA	NA	NA	NA	NA	Chlamyaster stermi	0	732	AF534709.1
SV_539		NA	NA	NA	NA	NA	Chlamyaster stermi	0	728	AF534709.1
SV_555		NA	NA	Heterophryidae	NA	metagenome	Triangulopteris lacunata	0	752	OL739463.1
SV_585		NA	NA	NA	NA	NA	Chlamyaster stermi	0	719	AF534709.1
SV_627		NA	NA	NA	NA	NA	Chlamyaster stermi	0	725	AF534709.1
SV_726		NA	NA	NA	NA	NA	Chlamyaster stermi	0	713	AF534709.1
SV_761		NA	NA	Heterophryidae	NA	metagenome	Khitsovia mutabilis	0	765	OP101622.1
SV_1104		NA	NA	NA	NA	NA	Chlamyaster stermi	0	723	AF534709.1
SV_20	Cercozoa	Thecofilosea	NA	NA	NA	NA	Rhizogostoma bowseri	0	767	MN860289.1 etc.
SV_24		Vampyrellidae	NA	NA	NA	NA	Theratomyxa weberi/Arachnula impatiens	0	713	GQ377666.1 etc.
SV_38		Vampyrellidae	NA	NA	NA	NA	Vampyrella closterii	0	701	OQ591878.1
SV_41		Vampyrellidae	NA	NA	NA	Arachnula_impatiens	Theratomyxa weberi/Arachnula impatiens	0	725	GQ377666.1 etc.
SV_52		Vampyrellidae	NA	NA	NA	Platyretia_germanica	Platyretia germanica	0	760	AY941201.1
SV_60		Cercomonadidae	NA	NA	Cercomonas	uncultured_eukaryote	Eocercomonas exploratorii	0	717	HM536154.1
SV_62		Glissomonadida	NA	NA	Heteromita	NA	Neoheteromita sp./Heteromita sp.	0	763	OR731273.1 etc.
SV_66		Vampyrellidae	NA	NA	Leptophrys	Leptophrys_vorax	Leptophrys vorax	0	710	HE609038.1 etc.
SV_67		Imbricatea	NA	Euglyphida	Euglypha	Euglypha_rotunda	Euglypha rotunda	0	761	AJ418784.1
SV_68		Vampyrellidae	NA	NA	NA	Arachnula_impatiens	Theratomyxa weberi/Arachnula impatiens	0	708	GQ377666.1 etc.
SV_111	Ciliophora	Intramacronucleata	Conthreep	Colpodea	Colpoda	metagenome	Kalometopia duplicata	0	697	KJ873050.1
SV_116		Intramacronucleata	Conthreep	Nassophorea	Obertrumia	NA	Obertrumia georgiana	0	721	X65149.1
SV_159		Intramacronucleata	Conthreep	Nassophorea	Obertrumia	NA	Obertrumia georgiana	0	726	X65149.1
SV_187		Intramacronucleata	Conthreep	Colpodea	Colpoda	uncultured_eukaryote	Colpoda sp.	0	774	MK801290.1 etc.
SV_189		Intramacronucleata	Conthreep	Colpodea	Colpoda	NA	Colpoda sp.	0	774	KJ607917.1 etc.
SV_217		Intramacronucleata	Conthreep	Colpodea	Colpoda	metagenome	Kalometopia duplicata	0	765	KJ873050.1
SV_220		Intramacronucleata	Conthreep	Oligohymenophorea	Frontonia	NA	Frontonia tchibisovae	0	675	DQ885987.1 etc.
SV_269		Intramacronucleata	Conthreep	Colpodea	NA	NA	Pseudoplatyophrya nana	0	773	AF060452.1
SV_315		Intramacronucleata	Conthreep	Colpodea	Platyophrya	NA	Platyophrya bromelicola	0	730	EU039905.1
SV_317		Intramacronucleata	Conthreep	Colpodea	Platyophrya	metagenome	Platyophrya bromelicola	0	736	EU039905.1
SV_3	Diatomea	Bacillariophyceae	NA	NA	Hantzschia	Hantzschia_amphioxys	Hantzschia amphioxys	0	773	MN696729.1 etc.
SV_18		Bacillariophyceae	NA	NA	NA	NA	Craticula subminuscula/Stauroneis acuta	0	721	KT072989.1 etc.
SV_30		Bacillariophyceae	NA	NA	NA	NA	Hantzschia amphioxys	0	761	MN696729.1 etc.
SV_103		Bacillariophyceae	NA	NA	Amphora	Amphora_montana	Halamphora montana/Amphora montana	0	767	KX120666.1 etc.
SV_106		Bacillariophyceae	NA	NA	Nitzschia	NA	Nitzschia palea	0	773	PP158997.1 etc.
SV_133		Bacillariophyceae	NA	NA	Fistulifera	Fistulifera_saprophila	Fistulifera sp.	0	774	HM997844.1 etc.
SV_213		Bacillariophyceae	NA	NA	Fistulifera	Mayamaea_atomus	Mayamaea permitis/Fistulifera pelliculosa	0	754	AM501969.1 etc.
SV_292		Bacillariophyceae	NA	NA	Fistulifera	Mayamaea_atomus	Mayamaea permitis/Fistulifera pelliculosa	0	754	AM501969.1 etc.
SV_296		Bacillariophyceae	NA	NA	Fistulifera	Mayamaea_atomus	Mayamaea permitis/Fistulifera pelliculosa	0	749	AM501969.1 etc.

SV_329		Bacillariophyceae	NA	NA	NA	NA	Craticula subminuscula/Stauroneis acuta	0	715	KT072989.1 etc.
SV_553		Dinophyceae	NA	NA	NA	NA	Aduncodinium glandula	0	767	LK934662.8
SV_1442		NA	NA	NA	NA	NA	Sourniacea diacantha/Gonyaulax whasegensis	5e-118	438	MT039425.1 etc.
SV_1667		Dinophyceae	Gymnodiniphycidae	Suessiaceae	Biecheleria	Woloszynskia_pascheri	Borghiella sp./Woloszynskia pascheri	0	761	MN604303.1 etc.
SV_2030		Dinophyceae	NA	NA	NA	NA	Esotrodinium sp.	0	734	JQ439944.1 etc.
SV_2120		Dinophyceae	Gymnodiniphycidae	Tovelliaaceae	Esotrodinium	Esotrodinium_sp.	Esotrodinium sp.	0	734	JQ439944.1 etc.
SV_2255		Dinophyceae	NA	NA	NA	NA	Pfiesteria piscicida	0	773	DQ991382.1 etc.
SV_2330		Dinophyceae	NA	NA	NA	NA	Pfiesteria sp. B112456	0	773	AF218805.2
SV_2359		Dinophyceae	NA	NA	NA	NA	Aduncusdinium glandulum	0	734	LK934662.8
SV_2680		Dinophyceae	NA	NA	NA	NA	Aduncusdinium glandulum	0	660	LK934662.8
SV_2782		Dinophyceae	NA	NA	NA	NA	Baruffeta sp./Aduncodinium sp./Stoeckeria sp./Gymnodinium sp.	0	734	MH732694.1 etc.
SV_4257		NA	NA	NA	Telaepolella	Telaepolella_tubasferens	Miligania stylosa	2e-137	503	HM640763.1 etc.
SV_4472		NA	NA	NA	Telaepolella	Telaepolella_sp.	Nudifila producta	7e-132	484	HQ121434.1
SV_4481		NA	NA	NA	Telaepolella	Telaepolella_sp.	Pterosperma sp. CCMP1384	1e-123	457	KF899835.1
SV_4658		NA	NA	NA	Telaepolella	Telaepolella_sp.	Pennarella eleganta	7e-132	484	PP10953.1
SV_4663		NA	NA	NA	Telaepolella	Telaepolella_sp.	Micromonas sp.	4e-129	475	MT117943.1 etc.
SV_5316		NA	NA	NA	Telaepolella	Telaepolella_sp.	Basidiobolus sp.	3e-111	416	MW130090.1 etc.
SV_97		NA	NA	Hyphochytriales	NA	uncultured_eukaryote	Rhizidiomyces apophysatus	0	760	AF163295.1
SV_117		NA	NA	Hyphochytriales	NA	Hyphochytrium	Hyphochytrium catenoides	0	776	X80344.1
SV_429		NA	NA	Hyphochytriales	NA	uncultured_eukaryote	Rhizidiomyces apophysatus	0	737	AF163295.1
SV_3656		NA	NA	Hyphochytriales	NA	NA	Rhizidiomyces apophysatus	1e-159	577	AF163295.1
SV_4521		NA	NA	Hyphochytriales	NA	NA	Rhizidiomyces apophysatus	3e-160	579	AF163295.1
SV_4719		NA	NA	Hyphochytriales	NA	Hyphochytrium	Hyphochytrium catenoides	1e-164	593	AF163294.1
SV_5232		NA	NA	Hyphochytriales	NA	Hyphochytrium	Hyphochytrium catenoides	1e-149	544	X80344.1
SV_803		Klebsormidiophyceae	NA	NA	NA	NA	Klebsormidium sp.	0	778	MT425960.1 etc.
SV_201		MAST-12C	NA	NA	NA	uncultured_Eimeriidae	Oblongichytrium sp./Pseudophyllomitius vesiculosus/Mastreximonas tlaamin	5e-148	538	KY980042.1 etc.
SV_409		NA	NA	NA	NA	uncultured_stramenopile	Thalassomyxa sp. MVa1x	1e-109	411	KC779515.1
SV_742		MAST-12C	NA	NA	NA	NA	Oblongichytrium sp./Pseudophyllomitius vesiculosus/Mastreximonas tlaamin	2e-146	532	KY980042.1 etc.
SV_841		MAST-12	NA	NA	NA	uncultured_Eimeriidae	Oblongichytrium sp.	1e-149	544	KY980042.1
SV_912		MAST-12C	NA	NA	NA	NA	Haptoglossa sp.	7e-147	534	KT257372.1 etc.
SV_1117		MAST-12C	NA	NA	NA	uncultured_Eimeriidae	Oblongichytrium sp./Pseudophyllomitius vesiculosus/Mastreximonas tlaamin	5e-148	538	KY980042.1 etc.
SV_3163		MAST-12C	NA	NA	NA	metagenome	Oblongichytrium sp.	2e-146	532	GQ354272.1 etc.
SV_3865		MAST-12C	NA	NA	NA	NA	Lobulomycetales sp.	2e-167	603	OQ702874.1 etc.
SV_3897		MAST-12C	NA	NA	NA	NA	Thalassomyxa sp./Vampyrella sp.	2e-141	516	KC779515.1 etc.
SV_34		NA	NA	NA	NA	NA	Globisporangium glomeratum	0	780	HQ643543.2
SV_53		NA	NA	NA	uncultured	metagenome	Myzocetiopsis sp./Lagenidium sp./Phytopythium mirpurens	0	689	KT257374.1 etc.
SV_57		NA	NA	NA	Pythium	Pythium_sp.	Pythium sp./Globisporangium sp.	0	782	MT193308.1 etc.
SV_157		NA	NA	NA	uncultured	metagenome	Pythium sp./Globisporangium sp.	0	695	KT257379.1 etc.
SV_169		NA	NA	NA	NA	NA	Thraustotheca clavata/Achlya bisexualis	0	773	KP098373.1 etc.
SV_211		NA	NA	NA	Pythium	Pythium_sp.	Pythium sp.	0	774	AY598681.2 etc.
SV_268		NA	NA	NA	Pythium	Pythium_sp.	Lagenidium sp./Lagena radicola	0	769	KJ716870.1 etc.
SV_284		NA	NA	NA	Pythium	NA	Lagenidium giganteum	0	763	KT257365.1 etc.
SV_430		NA	NA	NA	NA	NA	Pythiopsis sp./Achlya sparrowii/Leptoglena caudata/Saprolegnia sp./Protoachlya paradoxa	0	773	MK850088.1 etc.
SV_437		NA	NA	NA	Pythium	Pythium_sp.	Lagenidium sp./Lagena radicola	0	780	KJ716873.1 etc.
SV_4846		Preaxostyla	NA	NA	Paratrimastix	Paratrimastix_pyriiformis	Kinopus chlorellivorus	1e-153	556	MW694332.1
SV_928		Perkinsidae	A31	NA	NA	NA	Rana spheenocephala pathogen MJY-2007	0	662	EF675616.1
SV_2045		Perkinsidae	A31	NA	NA	uncultured_eukaryote	Parasarium dinoexitosum/Rana spheenocephala pathogen MJY-2007	8e-166	597	MZ663830.1 etc.
SV_2580		Perkinsidae	A31	NA	NA	NA	Parasarium dinoexitosum	0	651	MZ663830.1 etc.
SV_2854		Perkinsidae	A31	NA	NA	uncultured_microeukaryote	Rana spheenocephala pathogen MJY-2007	0	647	EF675616.1
SV_2902		Perkinsidae	A31	NA	NA	NA	Rana spheenocephala pathogen MJY-2007	2e-171	616	EF675616.1
SV_2983		Perkinsidae	A31	NA	NA	NA	Rana spheenocephala pathogen MJY-2007	1e-174	627	EF675616.1
SV_3106		Perkinsidae	A31	NA	NA	uncultured_microeukaryote	Rana spheenocephala pathogen MJY-2007	1e-178	640	EF675616.1
SV_3136		Perkinsidae	A31	NA	NA	uncultured_microeukaryote	Rana spheenocephala pathogen MJY-2007	0	651	EF675616.1
SV_3833		Perkinsidae	A31	NA	NA	NA	Rana spheenocephala pathogen MJY-2007	2e-157	569	EF675616.1
SV_4151		Colpodellida	NA	NA	Colpodella	uncultured_freshwater	Alphamonas edax/Olpidium brassicae	0	651	AY234843.1 etc.
SV_1306		Protosporangidae	NA	NA	Protosporangium	metagenome	Asterodiscus tamaricis	2e-52	220	NG_061222.1
SV_1371		Protosporangidae	NA	NA	Protosporangium	metagenome	Pertusaria saximontana	4e-49	209	AF113720.1
SV_1841		Protosporangidae	NA	NA	Protosporangium	metagenome	Pertusaria saximontana	4e-54	222	AF113720.1
SV_3213		Protosporangidae	NA	NA	Protosporangium	metagenome	Tilletiaria anomala/Phragmotenium dextii/Tilletia iowensis/Heterodoassansia hygrophilae/Hydnum sp.	2e-117	436	NG_061031.1 etc.
SV_1477		NA	NA	NA	NA	Micronuclearia_podoventralis	Micronuclearia_podoventralis	0	745	AY268038.1
SV_1541		NA	NA	NA	NA	Micronuclearia_podoventralis	Micronuclearia_podoventralis	0	767	AY268038.1
SV_1794		NA	NA	NA	NA	Micronuclearia_podoventralis	Micronuclearia_podoventralis	0	678	AY268038.1
SV_2129		NA	NA	NA	NA	Micronuclearia_podoventralis	Micronuclearia_podoventralis	0	712	AY268038.1
SV_4066		NA	NA	NA	NA	Micronuclearia_podoventralis	Micronuclearia_podoventralis	0	750	AY268038.1
SV_300		NA	NA	NA	Sorodiplophrys	metagenome	Sorodiplophrys stercorea	2e-126	466	KU728177.1 etc.
SV_494		NA	NA	NA	Sorodiplophrys	metagenome	Crustomastix sp.	2e-106	399	KY980137.1 etc.
SV_578		NA	NA	NA	Sorodiplophrys	uncultured_eukaryote	Sorodiplophrys stercorea	2e-137	503	KU728177.1 etc.
SV_613		NA	NA	NA	Sorodiplophrys	metagenome	Sorodiplophrys stercorea	1e-129	477	KU728177.1 etc.
SV_652		NA	NA	NA	Sorodiplophrys	metagenome	Sorodiplophrys stercorea	1e-118	440	KU728177.1 etc.
SV_709		NA	NA	NA	Sorodiplophrys	metagenome	Sorodiplophrys stercorea	4e-144	525	KU728177.1 etc.
SV_766		NA	NA	NA	Sorodiplophrys	metagenome	Nerada mexicana	4e-104	392	AY520453.1
SV_806		NA	NA	NA	Sorodiplophrys	uncultured_freshwater	Sorodiplophrys stercorea	0	675	KU728177.1 etc.
SV_908		NA	NA	NA	Sorodiplophrys	metagenome	Sorodiplophrys stercorea	4e-124	459	KU728177.1 etc.
SV_923		NA	NA	NA	Sorodiplophrys	uncultured_freshwater	Sorodiplophrys stercorea	3e-175	628	KU728177.1 etc.
SV_5161		Schizoplasmodiida	NA	NA	Phalansterium	uncultured_eukaryote	Cercomonas sp.	8e-156	564	PP174308.1 etc.

Animalia

ID	Phylum	Class	Order	Family	Genus	Species	The top hits by BLASTN search	e-value	Max score	Accession No.
SV_1		Citellata	Haplotaaxida	NA	NA	NA	Enchytraeus dichaeus	0	773	MN248684.1
SV_5		Citellata	Haplotaaxida	NA	NA	NA	Enchytraeus bulbosus	0	763	GU901872.1
SV_10		Citellata	Haplotaaxida	NA	NA	NA	Achaeta sp./Hemienchytraeus sp	0	767	GU901905.1 etc.
SV_25		Citellata	Haplotaaxida	NA	NA	Amyntas_sp.	Amyntas sp.	0	771	KF179538.1 etc.
SV_64		Citellata	Haplotaaxida	NA	NA	NA	Fridericia bulboides	0	758	GU901881.1
SV_73		Citellata	Haplotaaxida	NA	NA	NA	Fridericia sp.	0	769	KX618771.1 etc.
SV_98		Citellata	Haplotaaxida	NA	NA	Aeolosoma_sp.	Aeolosoma sp.	0	750	HQ691213.1
SV_119		Citellata	Haplotaaxida	NA	NA	Rheomorpha_neiswestonovae	Rheomorpha neiswestonovae	0	756	AY527049.1
SV_171		Citellata	NA	NA	NA	NA	Helobdella sp./Oosthiuizobdella garouii/Haementeria sp.	0	665	MN312167.1 etc.
SV_199		Citellata	Haplotaaxida	NA	NA	NA	Enchytraeus bulbosus	0	769	GU901872.1
SV_14		Arachnida	NA	NA	NA	Limulidae_environmental	Alicorhagia sp.	0	773	KP325047.1 etc.
SV_28		Arachnida	NA	NA	NA	Stigmalychus_sp.	Stigmalychus sp.	0	773	KY922137.1 etc.
SV_29		Ellipura	Collembola	NA	NA	NA	Bourletiella hortensis	0	767	KY230700.1

SV_35		Arachnida	NA	NA	NA	NA	Pergamasus sp./Vulgarogamasus sp./Aclerogamasus similis/Parasitus kraepelini/Phorytocarpais fimetorum	0	758	FJ911803.1 etc.
SV_54	Arthropoda	Arachnida	NA	NA	NA	NA	Phauloppia lucorum/Oribatula tibialis	0	706	EU423198.1 etc.
SV_75		Ellipura	Collembola	NA	NA	NA	Isotomurus sp.	0	734	KY230739.1 etc.
SV_110		Arachnida	NA	NA	NA	NA	Coccopodes sp.	0	671	MN241880.1
SV_115		Ellipura	Collembola	NA	NA	Lepidocyrtus sp.	Lepidocyrtus sp.	0	765	KC236249.1
SV_142		Arachnida	NA	NA	NA	NA	Fileupodes cf. filiformis FP101	3e-155	562	MN241882.1
SV_191		Arachnida	NA	NA	NA	NA	Amblyseius sp./Neoseiulus sp./Euseius sojaensis	0	758	KP276469.1 etc.
SV_51	Gastrotricha	NA	Chaetonotida	NA	NA	NA	Neogosseia sp.	0	780	KR822111.1 etc.
SV_179		NA	Chaetonotida	NA	NA	NA	Heterolepidoderma striatum	0	732	OQ358140.1
SV_277		NA	Chaetonotida	NA	NA	Chaetonotus_cf.	Chaetonotus cf. oculifer	0	784	MN496180.1 etc.
SV_299		NA	Chaetonotida	NA	NA	Chaetonotus_cf.	Chaetonotus aff.	0	784	MN496201.1 etc.
SV_378		NA	Chaetonotida	NA	NA	NA	Heterolepidoderma striatum	0	754	OQ358140.1
SV_379		NA	Chaetonotida	NA	NA	NA	Chaetonotus aff.	0	726	MN496229.1 etc.
SV_849		NA	Chaetonotida	NA	NA	NA	Chaetonotus sp.	0	780	MN496226.1 etc.
SV_1290		NA	Chaetonotida	NA	NA	NA	Chaetonotus sp.	0	750	OM421721.1 etc.
SV_1398		NA	Chaetonotida	NA	NA	NA	Heterolepidoderma striatum	0	749	OQ358140.1
SV_1452		NA	Chaetonotida	NA	NA	NA	Bodomorpha sp.	0	763	HM536172.1
SV_433	Mollusca	Gastropoda	NA	NA	NA	NA	Coccopodes sp./Fileupodes sp.	7e-122	451	MN241880.1 etc.
SV_876		Gastropoda	Heterobranchia	NA	NA	NA	Succinea sp./Monadenia fidelis/Arion hortensis/Bradybaena similis etc.	0	773	MN186472.1 etc.
SV_3777		Gastropoda	Heterobranchia	NA	NA	NA	Oxychilus alliarius/Vitrina pellucida/Ariophanta bistrialis/Eucomulus fulvus/Geomalacus sp.	0	721	MN022707.1 etc.
SV_23		Chromadorea	Araeolaimida	NA	NA	NA	Plectus sp.	0	763	LC186814.1 etc.
SV_26	Nematodea	Chromadorea	Rhabditida	NA	NA	NA	Acrobeloides sp.	0	763	MK636581.1 etc.
SV_32		Enoplea	NA	NA	NA	NA	Dorylaimellus parvulus	0	763	AY911968.1
SV_39		Enoplea	Triplonchida	NA	NA	Prismatolaimus_intermedius	Prismatolaimus sp.	0	763	LC186851.1 etc.
SV_40		NA	NA	NA	NA	NA	Diphtherophora sp. Shahrood	1e-168	606	KY115102.1
SV_42		Enoplea	NA	NA	NA	Mononchus_truncatus	Mononchus truncatus	0	761	KJ636355.1 etc.
SV_58		Enoplea	NA	NA	NA	Mylonchulus_hawaiiensis	Mylonchulus montanus/Prionchulus oleksandri	0	765	AY919209.1 etc.
SV_84		Chromadorea	Tylenchida	NA	NA	NA	Filenchus sp.	0	767	MN542201.1 etc.
SV_87		Enoplea	NA	NA	NA	Aquatides_christei	Aquatides sp./Clavicaudoides sp.	0	769	AY552967.1 etc.
SV_122		Enoplea	NA	NA	NA	NA	Dorylaimoides sp./Tylencholaimus mirabilis/Mesodorylaimus sp.	0	752	KU662325.1 etc.
SV_71	Platyhelminthes	Catenulida	NA	Stenostomidae	NA	Stenostomum_sp.	Stenostomum sp.	0	756	FJ384833.1 etc.
SV_99		Catenulida	NA	Stenostomidae	NA	Stenostomum_sp.	Stenostomum sp.	0	739	FJ384833.1 etc.
SV_100		Catenulida	NA	Stenostomidae	NA	NA	Stenostomum leucops	0	701	KP113676.1 etc.
SV_101		Rhabditophora	Tricladida	NA	NA	Arthurdendyus_triangularis	Arthurdendyus triangularis	1e-179	643	OR797297.1
SV_107		Rhabditophora	Tricladida	NA	NA	NA	Arthurdendyus triangularis	5e-178	638	OR797297.1
SV_108		Rhabditophora	Tricladida	NA	NA	NA	Girardia sp.	0	747	MN652491.1 etc.
SV_130		Rhabditophora	Tricladida	NA	NA	NA	Arthurdendyus triangularis	5e-178	638	OR797297.1
SV_146		Rhabditophora	Rhabdocoela	NA	NA	NA	Bryoplana xerophila	0	713	KC529489.1
SV_188		Catenulida	NA	Catenulidae	NA	Catenula_turgida	Retronectes sp.	0	649	OR096307.1 etc.
SV_223		Rhabditophora	Rhabdocoela	NA	NA	Acrochordonoposthia_conica	Acrochordonoposthia conica	0	798	KC529487.1
SV_589	Rotifera	Monogononta	Ploimida	NA	NA	NA	Lecane sp.	0	760	MT025820.1 etc.
SV_1479		Monogononta	Ploimida	NA	NA	NA	Cephalodella gibba	0	726	AY218114.1
SV_1543		Monogononta	Ploimida	NA	NA	NA	Synchaeta sp./Asplanchna girodi	0	754	KY751522.1 etc.
SV_1940		Monogononta	Ploimida	NA	NA	uncultured_eukaryote	Rhinoglena frontalis/Enicentrum semiplicatum	0	732	MT522683.1 etc.
SV_1993		Monogononta	Ploimida	NA	NA	uncultured_eukaryote	Bryocella stylata	0	710	MT522631.1
SV_2285		Monogononta	Ploimida	NA	NA	NA	Enicentrum semiplicatum	0	699	MT522645.1
SV_2780		Bdelloidea	Adinetida	NA	NA	Bdelloidea_environmental	Rotaria sp./Habrotricha sp./Adineta vaga	0	761	JX494744.2 etc.
SV_2921		Monogononta	Ploimida	NA	NA	NA	Mikrocodices chlaena	0	732	MT522662.1
SV_2944		Monogononta	Ploimida	NA	NA	NA	Brachionus zahnsieri	0	721	MT522630.1
SV_3022		Monogononta	Ploimida	NA	NA	NA	Kellicottia bostoniensis	0	737	MT522652.1
SV_1468	Tardigrada	Eutardigrada	Parachela	NA	NA	NA	Doryphoribius sp.	0	761	MT872215.1
SV_3197		Eutardigrada	Parachela	NA	NA	NA	Apodibius confusus	0	769	KC582830.1

Plants										
ID	Phylum	Class	Order	Family	Genus	Species	The top hits by BLASTN search	e-value	Max score	Accession No.
SV_2	Phragmoplastophyta	Embryophyta	NA	NA	NA	Oryza_sativa	Oryza sp./Setaria viridis/Panicum hallii etc.	0	773	XR_008019782.1 etc.
SV_4		Embryophyta	NA	NA	NA	NA	Brassica napus/Arabidopsis thaliana	0	769	CP151913.1 etc.
SV_6		Embryophyta	NA	NA	NA	NA	Cerastium arcticum	0	771	EF127844.1
SV_11		Embryophyta	NA	NA	NA	NA	Bidens sp.	0	771	MN535784.1 etc.
SV_33		Embryophyta	NA	NA	NA	Triticum_aestivum	Hordeum vulgare subsp. vulgare	0	773	XR_006618434.1
SV_50		Embryophyta	NA	NA	NA	NA	Sisyrinchium campestre	0	769	MT610957.1
SV_72		Embryophyta	NA	NA	NA	NA	Silene littorea	0	767	MN325962.1
SV_76		Embryophyta	NA	NA	NA	NA	Diplostephium sp./Linochilus sp./Solidago sp./Artemisia dracunculus etc.	0	771	MT610965.1 etc.
SV_78		Embryophyta	NA	NA	NA	NA	Ptychomitrium gardneri/Ceratodon sp./Fissidens subbasilaris/Tortella tortuosa	0	778	MN307174.1 etc.
SV_81		Embryophyta	NA	NA	NA	NA	Panax sp.	0	765	MK408812.1 etc.

Abundant SVs in each eukaryotic phylum and their SILVA-based taxa are shown. The top hits by BLASTN search (May in 2024), their e-values, max scores, and accession numbers are also indicated. NA: not assigned. Accession numbers with "etc." indicate multiple hits with the same scores (these hit species are separated by forward slash). The top 10 SVs in abundance are indicated in the phylum containing more than 10 SVs, and several abundant SVs in the phylum containing less than 10 SVs. Abundant SVs in unassigned phyla are not shown. SVs in bold were used for Supplementary Figs. S6, S8, and S9.

Supplementary Table S3. Cp-values and feeding types of nematode-derived SVs.

ID	Class	Order	Species	Genus of top hits by BLASTN search	cp-value	Feeding type
SV_23	Chromadorea	Araeolaimida	NA	<i>Plectus</i>	2	3
SV_26	Chromadorea	Rhabditida	NA	<i>Acroboloides</i>	2	3
SV_32	Enoplea	NA	NA	<i>Dorylaimellus</i>	5	1
SV_39	Enoplea	Triplonchida	Prismatolaimus_intermedius	<i>Prismatolaimus</i>	3	3
SV_40	NA	NA	NA	<i>Diphtherophora</i>	3	2
SV_42	Enoplea	NA	Mononchus_truncatus	<i>Mononchus</i>	4	5
SV_58	Enoplea	NA	Mylonchulus_hawaiiensis	<i>Prionchulus</i>	4	5
SV_84	Chromadorea	Tylenchida	NA	<i>Filenchus</i>	2	2
SV_87	Enoplea	NA	Aquatides_christei	<i>Clavicaudoides/Aquatides</i>	5	5
SV_122	Enoplea	NA	NA	<i>Dorylaimoides</i>	4	8
SV_132	Enoplea	Triplonchida	Odontolaimus_sp.	<i>Odontolaimus</i>	3	3
SV_155	Enoplea	NA	NA	<i>Amblydorylaimus/Eudorylaimus/Talanema</i>	4	8
SV_175	Chromadorea	Tylenchida	NA	<i>Meloidogyne</i>	3	1
SV_198	Chromadorea	Monhysterida	Paralamyctes_environmental	<i>Eumonhystera</i>	2	3
SV_200	Enoplea	Triplonchida	Squillidae_environmental	<i>Prismatolaimus</i>	3	3
SV_205	Enoplea	NA	NA	<i>Axonchoides</i>	5	1
SV_231	Enoplea	Triplonchida	Odontolaimus_sp.	<i>Odontolaimus</i>	3	3
SV_234	Chromadorea	Tylenchida	Pratylenchus_delattrei	<i>Pratylenchus</i>	3	1
SV_237	Chromadorea	Chromadorida	Prodesmodora_circulata	<i>Achromadora</i>	3	5
SV_250	Chromadorea	Monhysterida	Paralamyctes_environmental	<i>Eumonhystera</i>	2	3
SV_276	Enoplea	NA	Mylonchulus_arenicolus	<i>Mylonchulus</i>	4	5
SV_320	Chromadorea	Tylenchida	Aphelenchus_avenae	<i>Aphelenchus</i>	2	2
SV_331	Chromadorea	Tylenchida	Filenchus_discrepans	<i>Filenchus</i>	2	2
SV_332	Enoplea	Triplonchida	Prismatolaimus_intermedius	<i>Prismatolaimus</i>	3	3
SV_360	Chromadorea	Tylenchida	Ditylenchus_sp.	<i>Ditylenchus</i>	2	2
SV_371	Chromadorea	Tylenchida	NA	<i>Ditylenchus</i>	2	2
SV_396	Enoplea	NA	NA	<i>Tylencholaimus</i>	4	2
SV_405	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_413	Enoplea	NA	Limulidae_environmental	<i>Aporcelaimellus</i>	5	8
SV_419	Chromadorea	Tylenchida	Pratylenchus_delattrei	<i>Pratylenchus</i>	3	1
SV_450	Chromadorea	Tylenchida	NA	<i>Aphelenchoides</i>	2	2
SV_454	Enoplea	Triplonchida	Prismatolaimus_intermedius	<i>Prismatolaimus</i>	3	3
SV_457	Enoplea	NA	Eudorylaimus_carteri	<i>Thonus</i>	4	5
SV_468	Chromadorea	Monhysterida	NA	<i>Sabatieria</i>	2	m
SV_492	Chromadorea	Rhabditida	NA	<i>Eucephalobus/Cephalobus</i>	2	3
SV_541	Enoplea	Triplonchida	Prismatolaimus_intermedius	<i>Prismatolaimus</i>	3	3
SV_550	Enoplea	NA	Mononchus_aquaticus	<i>Mononchus</i>	4	5
SV_579	Enoplea	Triplonchida	Odontolaimus_sp.	<i>Odontolaimus</i>	3	3
SV_599	Enoplea	NA	NA	<i>Dorylaimoides</i>	4	8
SV_600	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_646	Enoplea	NA	NA	<i>Pararhyssocolpus</i>	4	8
SV_670	Chromadorea	Chromadorida	Achromadora_cf	<i>Achromadora</i>	3	5
SV_672	Enoplea	Triplonchida	Prismatolaimus_cf.	<i>Prismatolaimus</i>	3	3
SV_694	Enoplea	NA	NA	<i>Mesodorylaimus</i>	4	8
SV_752	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_764	Chromadorea	Rhabditida	Oscheius_tipulae	<i>Oscheius</i>	1	3
SV_802	Enoplea	Triplonchida	Prismatolaimus_intermedius	<i>Prismatolaimus</i>	3	3
SV_804	Chromadorea	Tylenchida	Aphelenchoides_fujianensis	<i>Aphelenchoides</i>	2	2
SV_828	Chromadorea	Araeolaimida	Anaplectus_grandepapillatus	<i>Plectus</i>	2	3
SV_845	Enoplea	Triplonchida	NA	<i>Diphtherophora</i>	3	2
SV_846	Chromadorea	Tylenchida	NA	<i>Aphelenchoides</i>	2	2
SV_875	Chromadorea	Tylenchida	NA	<i>Meloidogyne</i>	3	1
SV_885	Chromadorea	Tylenchida	Meloidogyne_incognita	<i>Meloidogyne</i>	3	1
SV_898	Enoplea	NA	Mononchus_aquaticus	<i>Mononchus</i>	4	5
SV_902	Enoplea	NA	NA	<i>Bastiania</i>	3	3
SV_927	Chromadorea	Rhabditida	Cephalobus_cubaensis	<i>Cephalobus</i>	2	3
SV_934	Chromadorea	Tylenchida	NA	<i>Aphelenchoides</i>	2	2
SV_949	Chromadorea	Tylenchida	Irantylenchus_vicinus	<i>Irantylenchus</i>	2	1
SV_960	Chromadorea	Tylenchida	Filenchus_discrepans	<i>Filenchus</i>	2	2

SV_991	Chromadorea	Tylenchida	NA	<i>Basiria</i>	2	1
SV_1022	Chromadorea	Tylenchida	NA	<i>Basiria</i>	2	1
SV_1046	Chromadorea	Araeolaimida	NA	<i>Plectus</i>	2	3
SV_1047	Chromadorea	Tylenchida	<i>Aphelenchus_avenae</i>	<i>Aphelenchus</i>	2	2
SV_1054	Chromadorea	Tylenchida	NA	<i>Lelenchus</i>	2	1
SV_1055	Chromadorea	Tylenchida	NA	<i>Meloidogyne</i>	3	1
SV_1059	Enoplea	NA	NA	<i>Microdorylaimus</i>	4	8
SV_1084	Chromadorea	Tylenchida	NA	<i>Aphelenchoides</i>	2	2
SV_1100	Chromadorea	Chromadorida	<i>Prodesmodora_circulata</i>	<i>Achromadora</i>	3	5
SV_1134	Chromadorea	Tylenchida	<i>Basiria_sp.</i>	<i>Basiria</i>	2	1
SV_1162	Chromadorea	Tylenchida	<i>Aphelenchoides_bicaudatus</i>	<i>Aphelenchoides</i>	2	2
SV_1174	Enoplea	Triplonchida	<i>Odontolaimus_sp.</i>	<i>Diphtherophora</i>	3	2
SV_1175	Chromadorea	Monhysterida	<i>Paralamyctes_environmental</i>	<i>Eumonhystera</i>	2	3
SV_1192	Enoplea	Triplonchida	NA	<i>Odontolaimus</i>	3	3
SV_1199	Enoplea	NA	<i>Mylonchulus_brachyuris</i>	<i>Mylonchulus</i>	4	5
SV_1207	Enoplea	NA	NA	<i>Mesodorylaimus</i>	4	8
SV_1211	Chromadorea	Tylenchida	NA	<i>Lelenchus</i>	2	1
SV_1220	Enoplea	NA	NA	<i>Discolaimus</i>	4	8
SV_1256	Chromadorea	Tylenchida	NA	<i>Lelenchus</i>	2	1
SV_1304	Enoplea	NA	NA	<i>Aporcelaimellus</i>	5	8
SV_1307	Chromadorea	Tylenchida	<i>Pratylenchus_penetrans</i>	<i>Pratylenchus</i>	3	1
SV_1335	Chromadorea	Tylenchida	<i>Filenchus_discrepans</i>	<i>Filenchus</i>	2	2
SV_1404	Chromadorea	Tylenchida	NA	<i>Geocenamus</i>	3	1
SV_1433	Enoplea	NA	NA	<i>Tylencholaimellus</i>	4	2
SV_1562	Enoplea	NA	<i>Eudorylaimus_carteri</i>	<i>Eudorylaimus</i>	4	8
SV_1574	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_1597	Enoplea	Triplonchida	NA	<i>Diphtherophora</i>	3	2
SV_1613	Chromadorea	Tylenchida	<i>Aphelenchoides_fujianensis</i>	<i>Aphelenchoides</i>	2	2
SV_1622	Chromadorea	Tylenchida	<i>Basiria_sp.</i>	<i>Basiria</i>	2	1
SV_1635	Chromadorea	Chromadorida	<i>Prodesmodora_circulata</i>	<i>Achromadora</i>	3	5
SV_1652	Enoplea	NA	<i>Mylonchulus_brachyuris</i>	<i>Mylonchulus</i>	4	5
SV_1657	Chromadorea	Tylenchida	NA	<i>Tylenchus/Filenchus</i>	2	1, 2
SV_1695	Enoplea	NA	NA	<i>Clavicaudoides</i>	5	5
SV_1753	Chromadorea	Tylenchida	<i>Malenchus_sexlineatus</i>	<i>Malenchus</i>	2	1
SV_1760	Enoplea	NA	<i>Nematoda_sp.</i>	<i>Gastromermis</i>	0	ap
SV_1785	Chromadorea	Tylenchida	<i>Filenchus_misellus</i>	<i>Filenchus</i>	2	2
SV_1810	Chromadorea	Rhabditida	<i>Poikilolaimus_oxycercus</i>	<i>Poikilolaimus</i>	1	3
SV_1860	Chromadorea	Chromadorida	<i>Achromadora_cf</i>	<i>Achromadora</i>	3	5
SV_1879	Chromadorea	Tylenchida	<i>Pratylenchus_delattrei</i>	<i>Pratylenchus</i>	3	1
SV_1922	Enoplea	Triplonchida	<i>Prismatolaimus_intermedius</i>	<i>Prismatolaimus</i>	3	3
SV_1986	Chromadorea	Tylenchida	NA	<i>Lelenchus</i>	2	1
SV_1992	Enoplea	NA	NA	<i>Nematimermis</i>	0	ND
SV_2013	Chromadorea	Tylenchida	<i>Aphelenchoides_bicaudatus</i>	<i>Aphelenchoides</i>	2	2
SV_2021	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_2036	Enoplea	NA	NA	<i>Paractinolaimus</i>	5	5
SV_2043	Chromadorea	Tylenchida	NA	<i>Ditylenchus</i>	2	2
SV_2055	Enoplea	NA	<i>Eudorylaimus_carteri</i>	<i>Prodorylaimus</i>	4	8
SV_2087	Chromadorea	Tylenchida	NA	<i>Lelenchus</i>	2	1
SV_2155	Enoplea	NA	NA	<i>cf. Tylencholaimus/cf. Calodorylaimus</i>	4	2
SV_2170	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_2174	Chromadorea	Araeolaimida	NA	<i>Plectus</i>	2	3
SV_2202	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_2212	Chromadorea	Tylenchida	NA	<i>Nothotylenchus</i>	2	2
SV_2242	Enoplea	NA	NA	<i>Microdorylaimus</i>	4	8
SV_2271	Chromadorea	Araeolaimida	<i>Anaplectus_grandepapillatus</i>	<i>Anaplectus</i>	2	3
SV_2301	Enoplea	NA	NA	<i>Tylencholaimus</i>	4	2
SV_2304	Chromadorea	Tylenchida	NA	<i>Lelenchus</i>	2	1
SV_2348	Chromadorea	Tylenchida	<i>Aphelenchoides_sp.</i>	<i>Aphelenchoides</i>	2	2
SV_2356	Chromadorea	Chromadorida	<i>Achromadora_cf</i>	<i>Achromadora</i>	3	5
SV_2423	Chromadorea	Tylenchida	NA	<i>Miculenchus</i>	2	1
SV_2470	Enoplea	NA	NA	<i>Laimydorus</i>	4	8
SV_2478	Chromadorea	Tylenchida	<i>Ditylenchus_halictus</i>	<i>Ditylenchus</i>	2	2

SV_2510	Chromadorea	Tylenchida	Aphelenchoides_bicaudatus	<i>Aphelenchoides</i>	2	2
SV_2518	Chromadorea	Tylenchida	Ditylenchus_halictus	<i>Ditylenchus</i>	2	2
SV_2538	Chromadorea	Monhysterida	uncultured_eukaryote	<i>Eumonhystera</i>	2	3
SV_2629	Enoplea	NA	NA	<i>Prionchulus</i>	4	5
SV_2643	Chromadorea	Tylenchida	NA	<i>Filenchus</i>	2	2
SV_2648	Chromadorea	Monhysterida	Nematoda_environmental	<i>Eumonhystera</i>	2	3
SV_2658	Chromadorea	Chromadorida	NA	<i>Achromadora</i>	3	5
SV_2662	Enoplea	NA	NA	<i>Heterodorus/Longidorella/Microdorylaimus/Allodorylaimus/Eudorylaimus</i>	4	8
SV_2686	Chromadorea	Tylenchida	NA	<i>Discopersicus/Basiria</i>	2	1
SV_2758	Enoplea	Triplonchida	NA	<i>Trischistoma</i>	3	5
SV_2760	Enoplea	NA	NA	<i>Amblydorylaimus/Eudorylaimus/Talanema</i>	4	8
SV_2781	Chromadorea	Tylenchida	NA	<i>Psilenchus</i>	2	1
SV_2812	Enoplea	NA	NA	<i>Tylencholaimus</i>	4	2
SV_2858	Chromadorea	Tylenchida	NA	<i>Ficophagus/Schistonchus</i>	2	1
SV_2864	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_2867	Chromadorea	Tylenchida	NA	<i>Meloidogyne</i>	3	1
SV_2923	Chromadorea	Tylenchida	NA	<i>Boleodorus</i>	2	1
SV_2933	Chromadorea	Rhabditida	NA	<i>Cephalobus</i>	2	3
SV_2934	Chromadorea	Araeolaimida	Anaplectus_grandepapillatus	<i>Anaplectus</i>	2	3
SV_2939	Enoplea	Triplonchida	Tripylina_sp.	<i>Tripylina</i>	3	5
SV_2977	Chromadorea	Tylenchida	NA	<i>Meloidogyne</i>	3	1
SV_3050	Chromadorea	Chromadorida	NA	<i>Achromadora</i>	3	5
SV_3056	Chromadorea	Chromadorida	NA	<i>Achromadora</i>	3	5
SV_3090	Chromadorea	Rhabditida	NA	<i>Distolabrellus</i>	1	3
SV_3093	Chromadorea	Tylenchida	Pratylenchus_delattrei	<i>Pratylenchus</i>	3	1
SV_3104	Chromadorea	Tylenchida	NA	<i>Psilenchus</i>	2	1
SV_3112	Chromadorea	Tylenchida	Filenchus_misellus	<i>Filenchus</i>	2	2
SV_3201	Chromadorea	Rhabditida	NA	<i>Panagrolaimus</i>	1	3
SV_3233	Enoplea	NA	NA	<i>Axonchium</i>	5	1
SV_3305	Chromadorea	Tylenchida	NA	<i>Labrys</i>	2	1
SV_3311	Chromadorea	Tylenchida	Pratylenchus_delattrei	<i>Pratylenchus</i>	3	1
SV_3317	Chromadorea	Rhabditida	Mesorhabditis_sp.	<i>Mesorhabditis</i>	1	3
SV_3380	Enoplea	Triplonchida	Trischistoma_taignensis	<i>Trischistoma</i>	3	5
SV_3382	Chromadorea	Rhabditida	Mesorhabditis_sp.	<i>Mesorhabditis</i>	1	3
SV_3397	Chromadorea	Tylenchida	NA	<i>Pratylenchus</i>	3	1
SV_3399	Chromadorea	Araeolaimida	NA	<i>Plectus</i>	2	3
SV_3414	Chromadorea	Tylenchida	NA	<i>Pratylenchus</i>	3	1
SV_3502	Chromadorea	Rhabditida	NA	<i>Cervidellus/Heterocephalobus</i>	2	3
SV_3512	Chromadorea	Tylenchida	NA	<i>Labrys</i>	2	1
SV_3566	Chromadorea	Araeolaimida	Anaplectus_grandepapillatus	<i>Anaplectus</i>	2	3
SV_3627	Chromadorea	Rhabditida	NA	<i>Helicotylenchus</i>	3	1
SV_3635	Chromadorea	Tylenchida	NA	<i>Pratylenchus</i>	3	1
SV_3650	Chromadorea	Rhabditida	Propanagrolaimus_sp.	<i>Panagrolaimus</i>	1	3
SV_3684	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_3748	Chromadorea	Tylenchida	NA	<i>Malenchus</i>	2	1
SV_3797	Enoplea	NA	NA	<i>Clarkus/Actus</i>	4	5
SV_3838	Chromadorea	Monhysterida	Eumonhystera_cf.	<i>Eumonhystera</i>	2	3
SV_3948	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_3960	Chromadorea	Tylenchida	Filenchus_discrepans	<i>Sakia</i>	2	1
SV_4048	Enoplea	NA	NA	<i>Xiphinemella</i>	4	2
SV_4127	Chromadorea	Monhysterida	NA	<i>Eumonhystera</i>	2	3
SV_4247	Chromadorea	Tylenchida	Filenchus_vulgaris	<i>Myionchulus</i>	4	5
SV_4260	Chromadorea	Rhabditida	NA	<i>Acrobeles</i>	2	3
SV_4288	Chromadorea	Rhabditida	NA	<i>Acrobeles</i>	2	3
SV_4322	Chromadorea	Monhysterida	uncultured_eukaryote	<i>Eumonhystera</i>	2	3
SV_4507	Chromadorea	Tylenchida	Aphelenchus_avenae	<i>Aphelenchus</i>	2	2
SV_4790	Chromadorea	Tylenchida	NA	<i>Meloidogyne</i>	3	1
SV_4795	Chromadorea	Tylenchida	Filenchus_vulgaris	<i>Filenchus</i>	2	2
SV_4810	Chromadorea	Tylenchida	NA	<i>Meloidogyne</i>	3	1
SV_4811	Chromadorea	Tylenchida	NA	<i>Meloidogyne</i>	3	1
SV_4812	Chromadorea	Tylenchida	NA	<i>Meloidogyne</i>	3	1

SV_4829	Enoplea	NA	NA	<i>Aporcella/Dorylaimellus</i>	5	1, 8
SV_4853	Chromadorea	Rhabditida	Cephalobus_cubaensis	<i>Cephalobus</i>	2	3
SV_4854	Chromadorea	Tylenchida	NA	<i>Filenchus</i>	2	2
SV_4875	Enoplea	NA	NA	<i>Clavicaudoides</i>	5	5
SV_4918	Chromadorea	NA	NA	<i>Achromadora</i>	3	5
SV_5078	Enoplea	NA	NA	<i>Microdorylaimus/Eudorylaimus</i>	4	8
SV_5101	Chromadorea	Rhabditida	Cephalobus_cubaensis	<i>Cephalobus</i>	2	3

Class, order, and species of nematode-derived SVs based on the SILVA database. The genus of the hit with the lowest e-value by BLASTN search (May 2024) is shown. The genera derived from the hits with the same lowest e-values are separated by forward slash. The cp-value and feeding type were determined based on the genus of top hit by BLASTN search in the Nematode Ecophysiological Parameter Search at the Nemaplex homepage of UC Davis, USA (<http://nemaplex.ucdavis.edu/Ecology/EcophysiologyParms/EcoParameterMenu.html>). NA: not assigned. ND: no data. ap: animal-parasitic nematodes. m: marine nematode. The feeding type of the family was used if the genera are absent in nematode genus list (i.e., cp-value is zero).

Supplementary Table S4. Parameter values of prokaryotic and eukaryotic networks of top 150 abundant SVs in each sample.

Prokaryotic network (top150 abundant SVs, $r = 0.8$, $p = 0.01$)

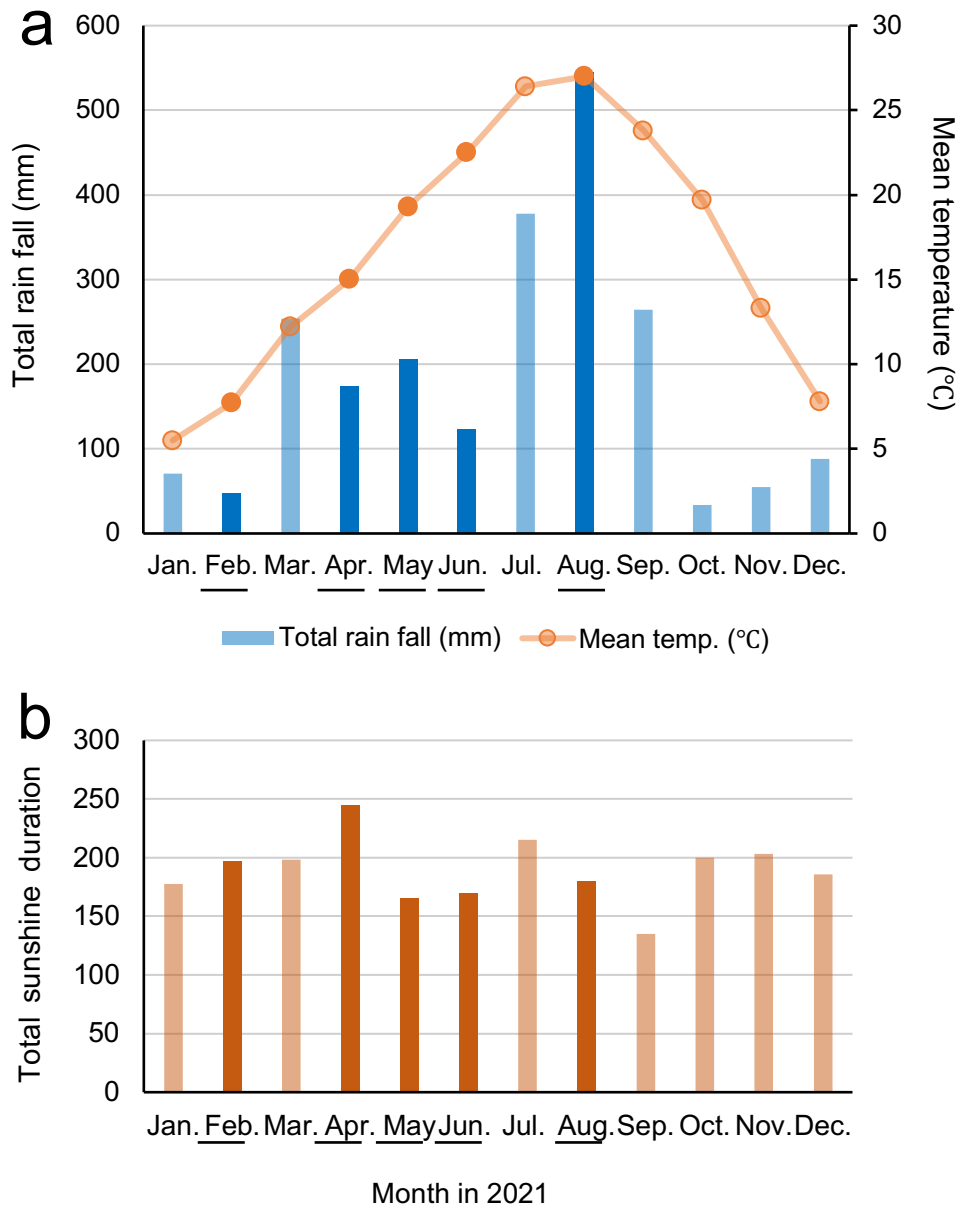
Papameters	February	April	May	June	August
Number of links	316	346	262	344	298
Number of positive links	268	314	165	247	246
Number of negative links	48	32	97	97	52
Number of vertices	132	133	131	129	139
Connectance (link_density)	0.037	0.039	0.031	0.042	0.031
Average degree (Average K)	4.788	5.203	4.000	5.333	4.288
Average path length	6.032	5.204	4.807	6.432	5.715
Diameter	16.024	14.967	13.131	22.467	15.958
Mean clustering coefficient (Average.CC)	0.541	0.577	0.646	0.600	0.548
Number of clusters	12	10	13	13	10
Centralization degree	0.070	0.059	0.077	0.099	0.049
Centralization betweenness	0.228	0.119	0.087	0.177	0.121
Centralization closeness	1.371	1.347	1.231	1.346	1.385
Relative modularity (RM)	0.727	0.975	0.931	0.702	0.732
The number of keystone nodes	1	5	1	2	1

Eukaryotic network (top150 abundant SVs, $r = 0.7$, $p = 0.01$)

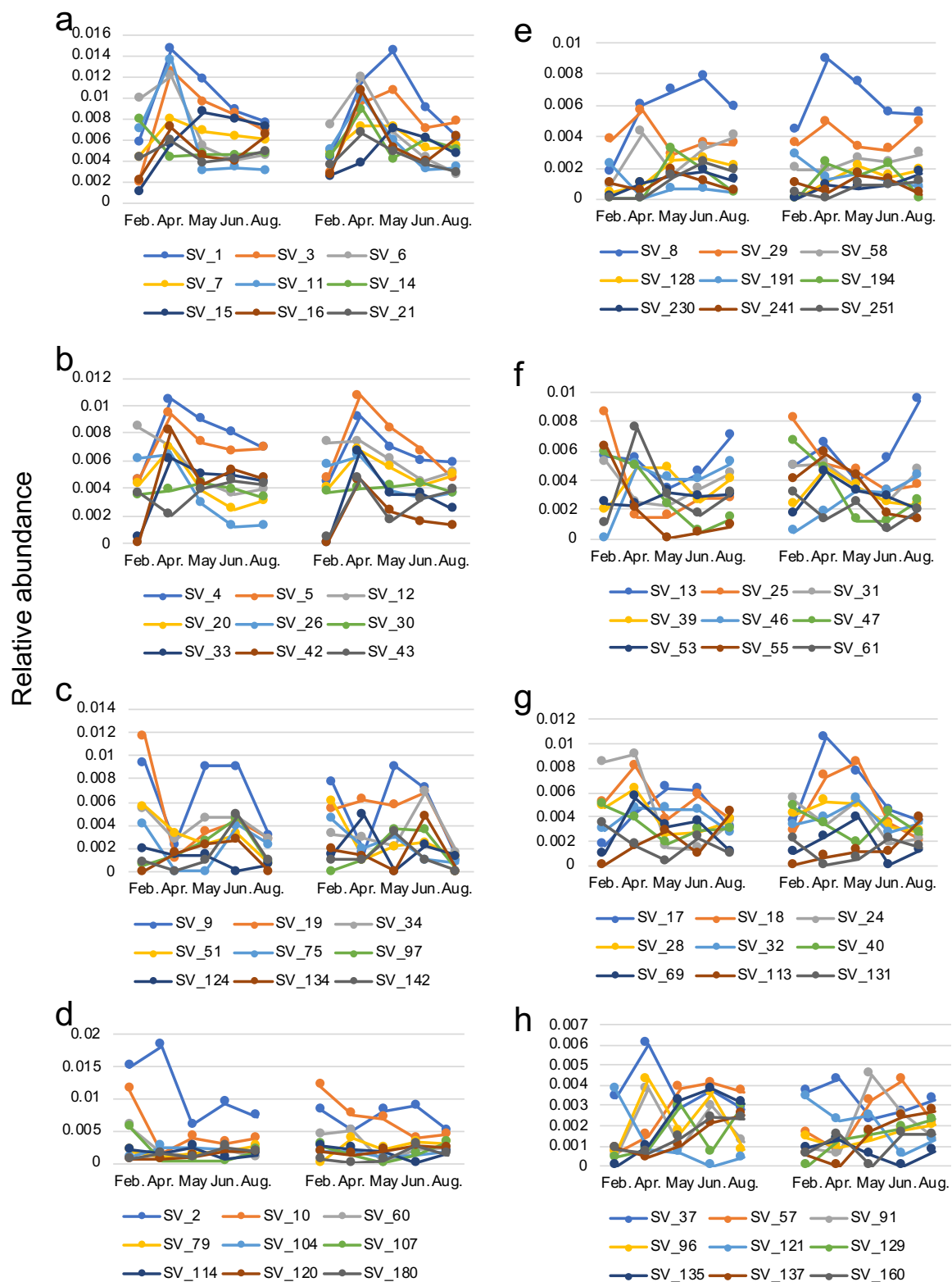
Papameters	February	April	May	June	August
Number of links	288	325	485	266	314
Number of positive links	208	302	478	246	222
Number of negative links	80	23	7	20	92
Number of vertices	135	137	135	126	137
Connectance (link_density)	0.032	0.035	0.054	0.034	0.034
Average degree (Average K)	4.267	4.745	7.185	4.222	4.584
Average path length	5.720	5.081	4.281	5.779	6.617
Diameter	14.075	12.279	11.318	15.033	17.843
Mean clustering coefficient (Average.CC)	0.518	0.643	0.573	0.628	0.550
Number of clusters	9	12	10	11	9
Centralization degree	0.080	0.046	0.088	0.078	0.099
Centralization betweenness	0.242	0.182	0.091	0.218	0.252
Centralization closeness	1.420	1.275	1.329	1.319	1.439
Relative modularity (RM)	0.896	1.087	1.142	0.993	0.909
The number of keystone nodes	1	4	4	0	0

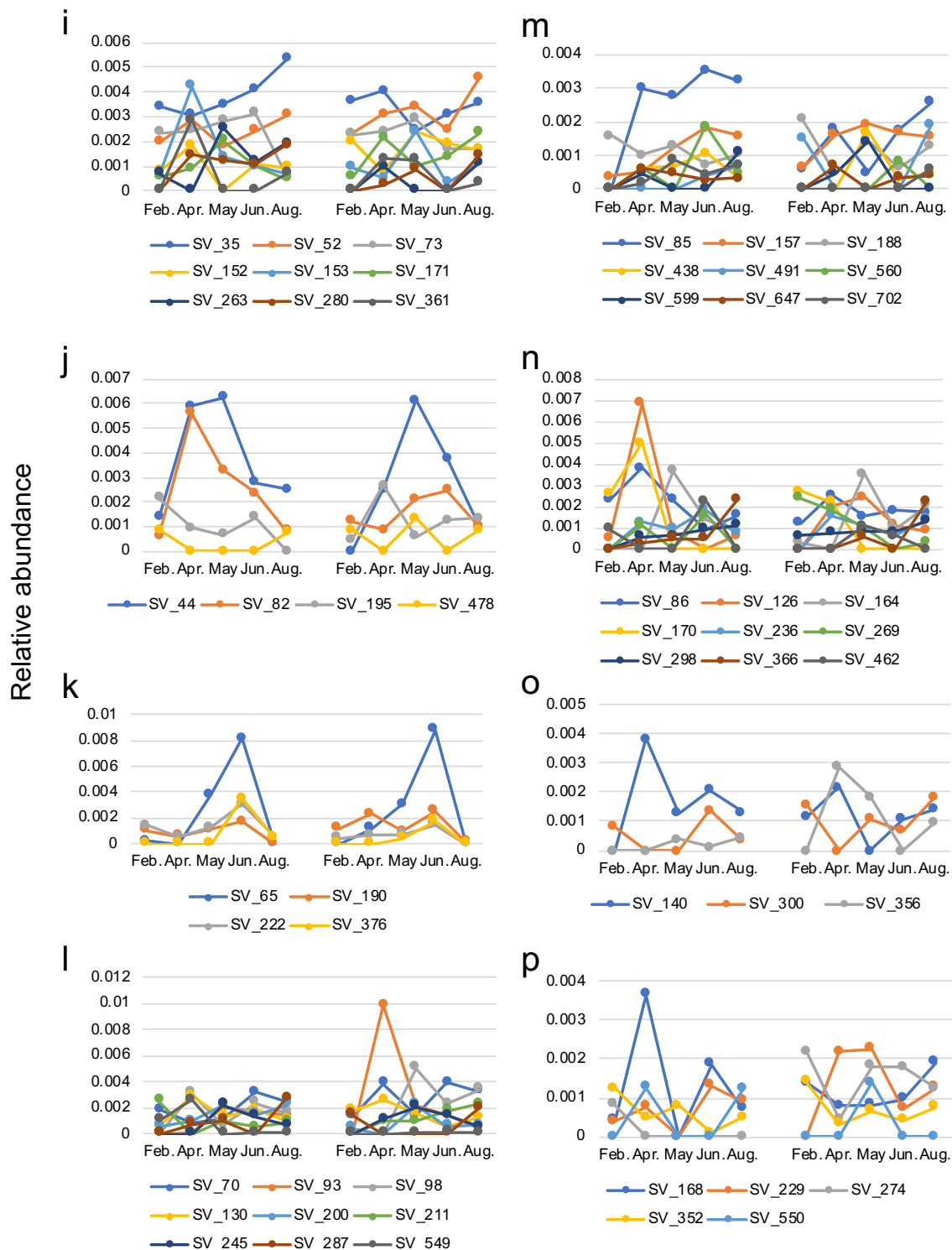
The values of the parameters of the prokaryotic and eukaryotic co-occurrence networks of top 150 abundant SVs are shown in each sample. Networks were prepared as described in the Methods section.

Supplementary Figures

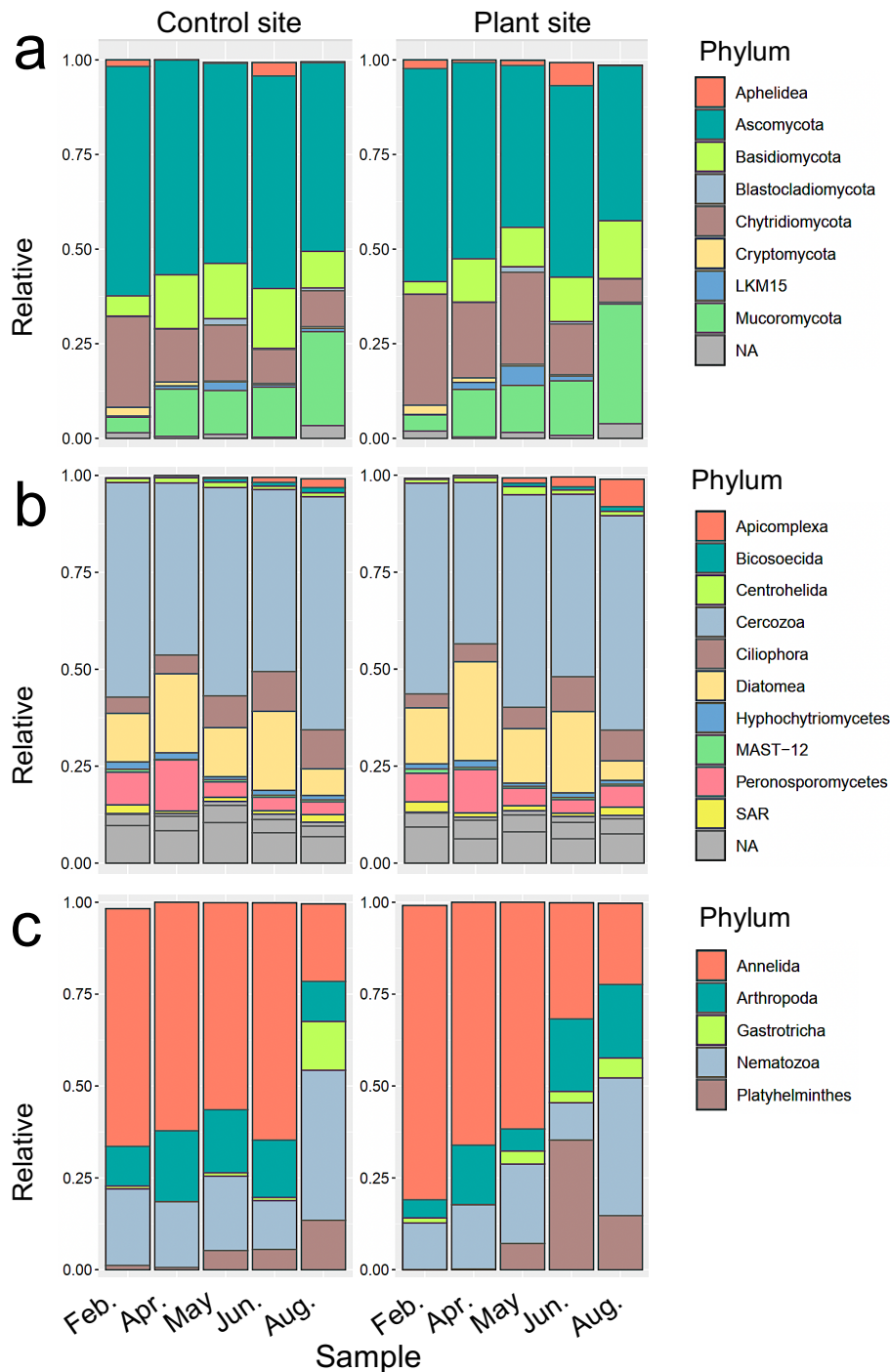


Supplementary Figure S1. Total rainfall and mean temperature (a) and total sunshine duration (b) per month at Toyohashi city in 2021. The total rainfall (mm) and total sunshine duration (h), and mean temperature (°C) per month in Toyohashi city in 2021 are shown by bar and line graphs in panel (a) and (b), respectively. Data was obtained from home page of Japan Meteorological Agency (https://www.data.jma.go.jp/obd/stats/etrn/view/monthly_a1.php?prec_no=51&block_no=0470&year=2021&month=&day=&view=p1). Note: Total sunshine durations of January to March were measured by different type of sunshine meter.

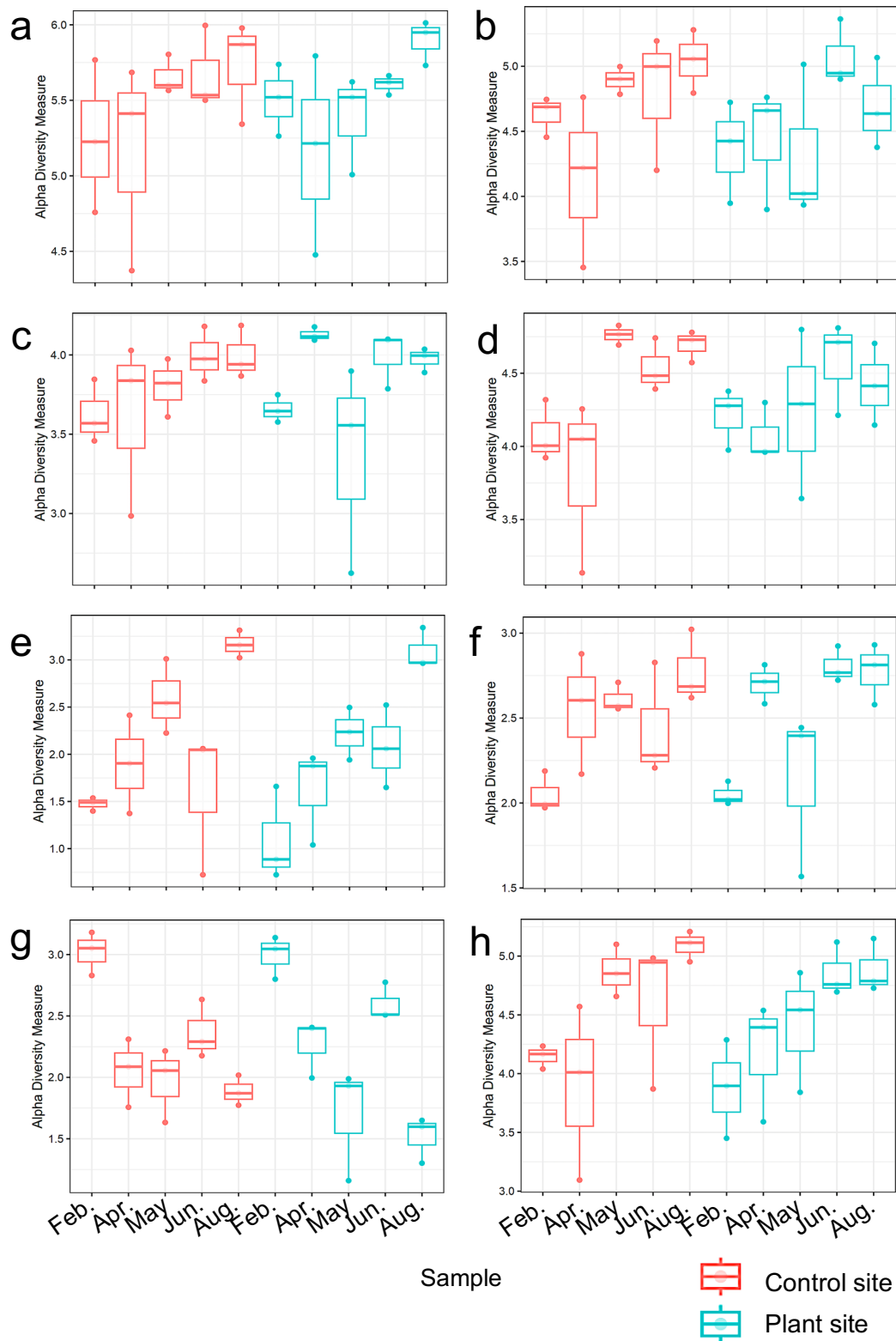




Supplementary Figure S2. Changes in the relative abundances of abundant prokaryotic SVs in each phylum. The relative abundances of abundant SVs shown in the bottom of the panels of Proteobacteria (a), Acidobacteriota (b), Actinobacteriota (c), Chloroflexi (d), Bacteroidota (e), Gemmatimonadota (f), Verrucomicrobiota (g), Planctomycetota (h), Latescibacterota (i), Nitrospirota (j), Cyanobacteria (k), Methylomirabilota (l), Armatimonadota (m), Myxococcota (n), Desulfobacterota (o), and NB1-j (p), are indicated during the cultivation period (see Supplementary Table S1). The left and right line graphs indicate the results from control and plant sites, respectively.

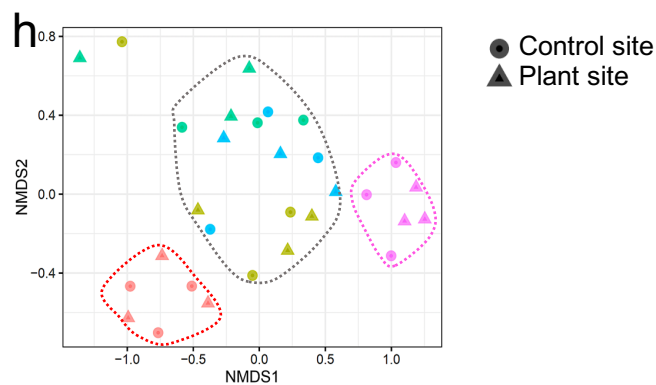
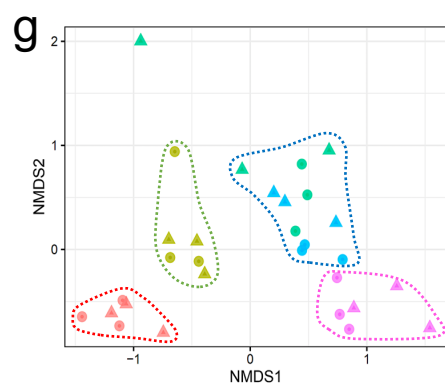
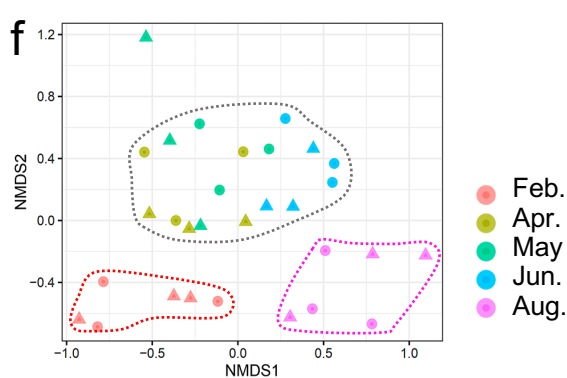
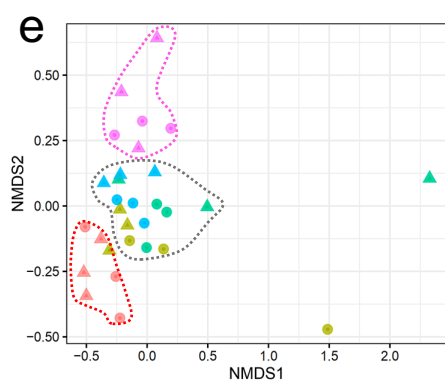
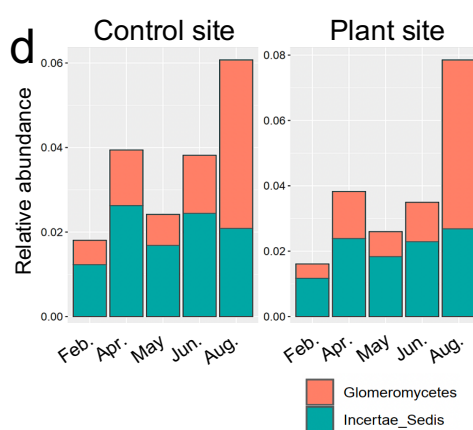
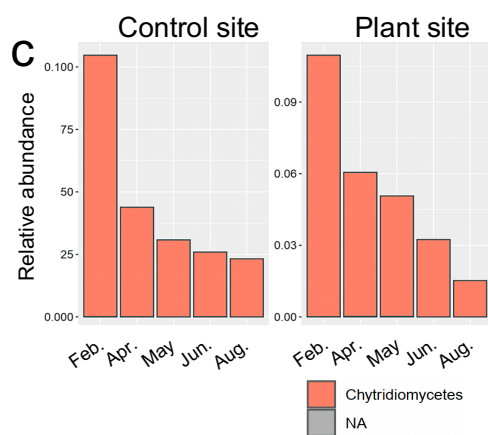
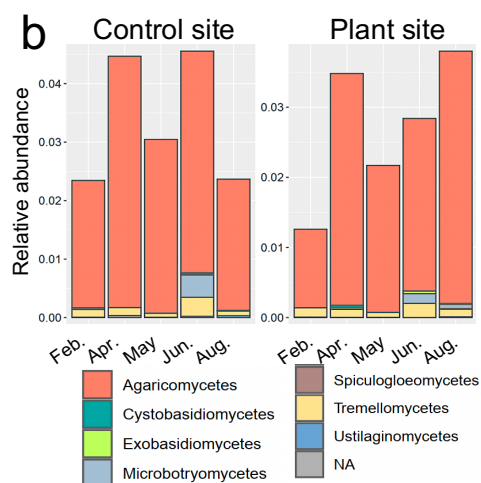
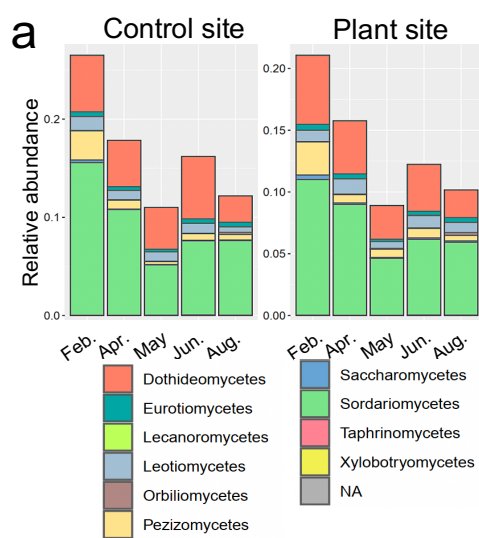


Supplementary Figure S3. Phylum compositions of fungi- (a), protists- (b), and animal-derived (c) SVs across the cultivation period at control (left) and plant (right) sites. The SILVA-based taxa are indicated by the colors shown on the right of the figures. The relative abundances of the SVs at phylum levels in each sample are shown by bar charts: kingdom Fungi (a), protists (i.e., kingdoms Alveolata, Amoebozoa, Rhizaria, and Stramenopiles, phyla Nucleariidae_and_Fonticula_group, and Preaxostyla, and classes Choanoflagellida, Filasterea, and Ichthyospora) (b), and kingdom Animalia (c), respectively with a relative abundance more than 0.5%. NA, not assigned phylum.

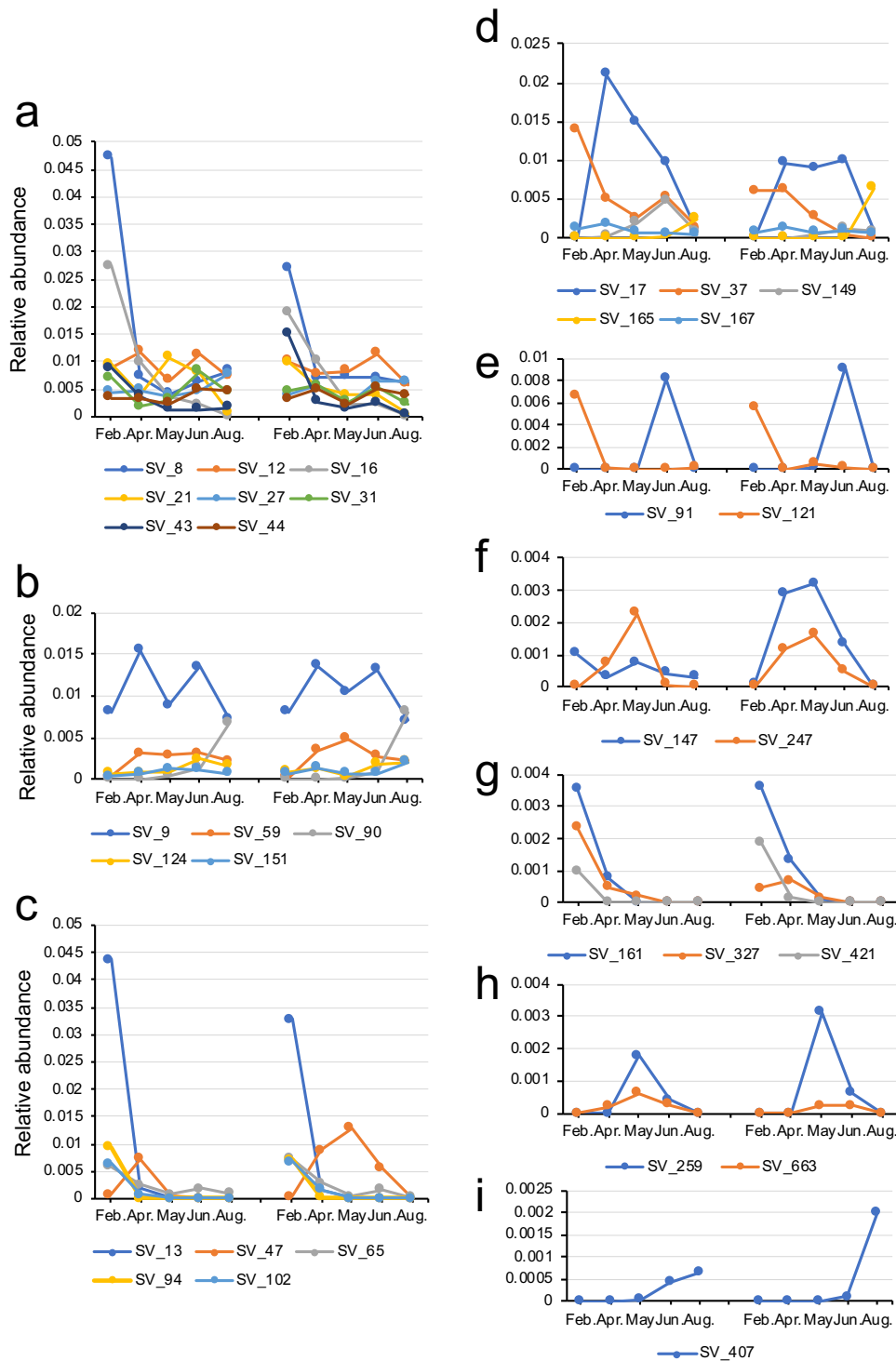


Supplementary Figure S4. Boxplots showing the α -diversities by the Shannon index across the five samples. The α -diversities of prokaryote (a), eukaryote (b), fungi (c), protists (d), animals (e), nematodes (f), plants (g), and eukaryotes without plants (h) across the cultivation period (month) are indicated in each sample derived from control (left) and plant (right) sites by boxplots. The taxa of fungi, protists,

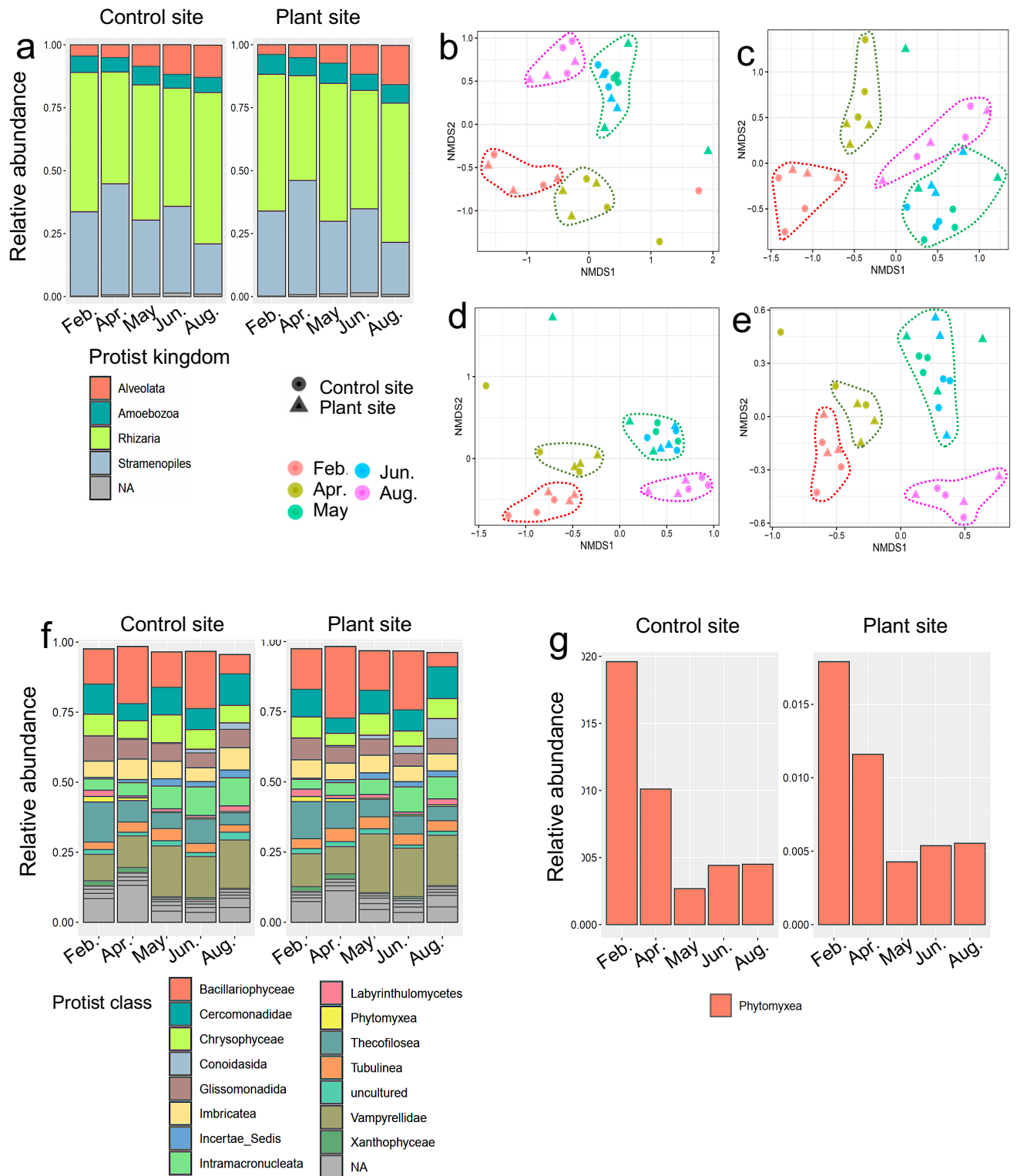
and animals are referred to in the legend for Supplementary Figure S3, and those of nematodes and plants represent the SVs derived from phyla Nematozoa and Phragmoplastophyta, respectively.



Supplementary Figure S5. Taxonomic compositions and nonmetric multidimensional scaling (NMDS) ordinations of four fungal phyla-derived SVs in each sample isolated from control and plant sites. The relative abundances of fungal SVs at the class level derived from phyla Ascomycota (a), Basidiomycota (b), Chytridiomycota (c), and Mucoromycota (d) are shown by bar charts in each sample from control (left) and plant (right) sites. Classes based on the SILVA database are indicated by the colors shown on the bottom of the bar graphs. NA and Incertae_Sedis: unassigned class. NMDS ordinations showing differences in β -diversity, based on the Bray–Curtis dissimilarity in fungal SVs derived from phyla Ascomycota (e), Basidiomycota (f), Chytridiomycota (g), and Mucoromycota (h) in the samples. The samples designated as Feb., Apr., May, Jun. and Aug., which were isolated on February, April, May, June and August in 2021, are shown by colored symbols on the right of figures. Samples derived from control (circles) and plant (triangles) sites are also shown. Clusters of samples are surrounded by dotted lines in the figures.

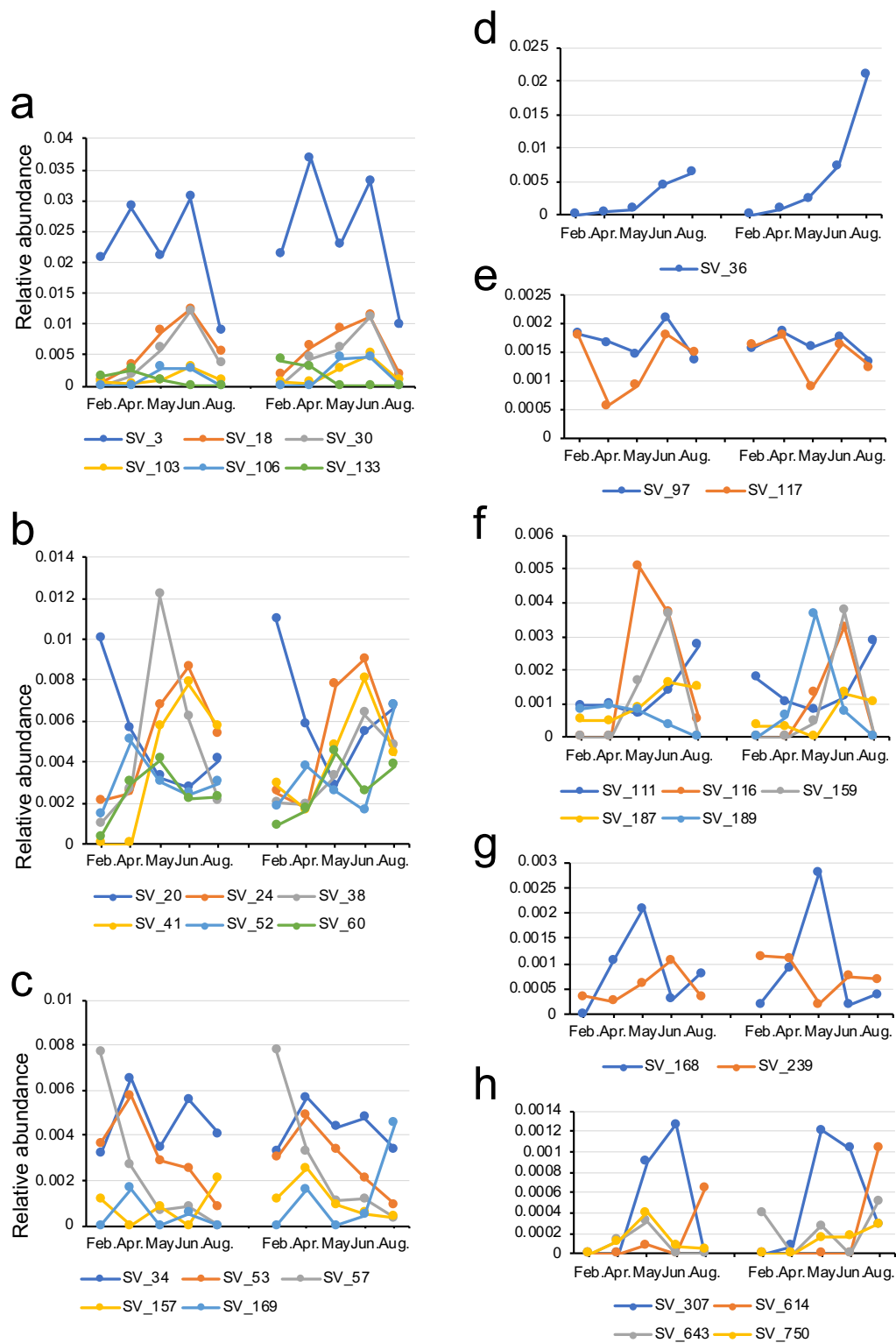


Supplementary Figure S6. Changes in the relative abundances of abundant SVs in each fungal phylum. The relative abundances of abundant fungal SVs (shown in the bottom of the panels) in total eukaryotic SVs, which are derived from Ascomycota (a), Mucoromycota (b), Chytridiomycota (c), Basidiomycota (d), Aphelidea (e), LKM15 (f), Cryptomycota (g), Blastocladiomycota (h), and Zoopagomycota (i), are indicated in the samples (see Supplementary Table S2). The left and right line graphs indicate the relative abundances of SVs at control and plant sites, respectively.

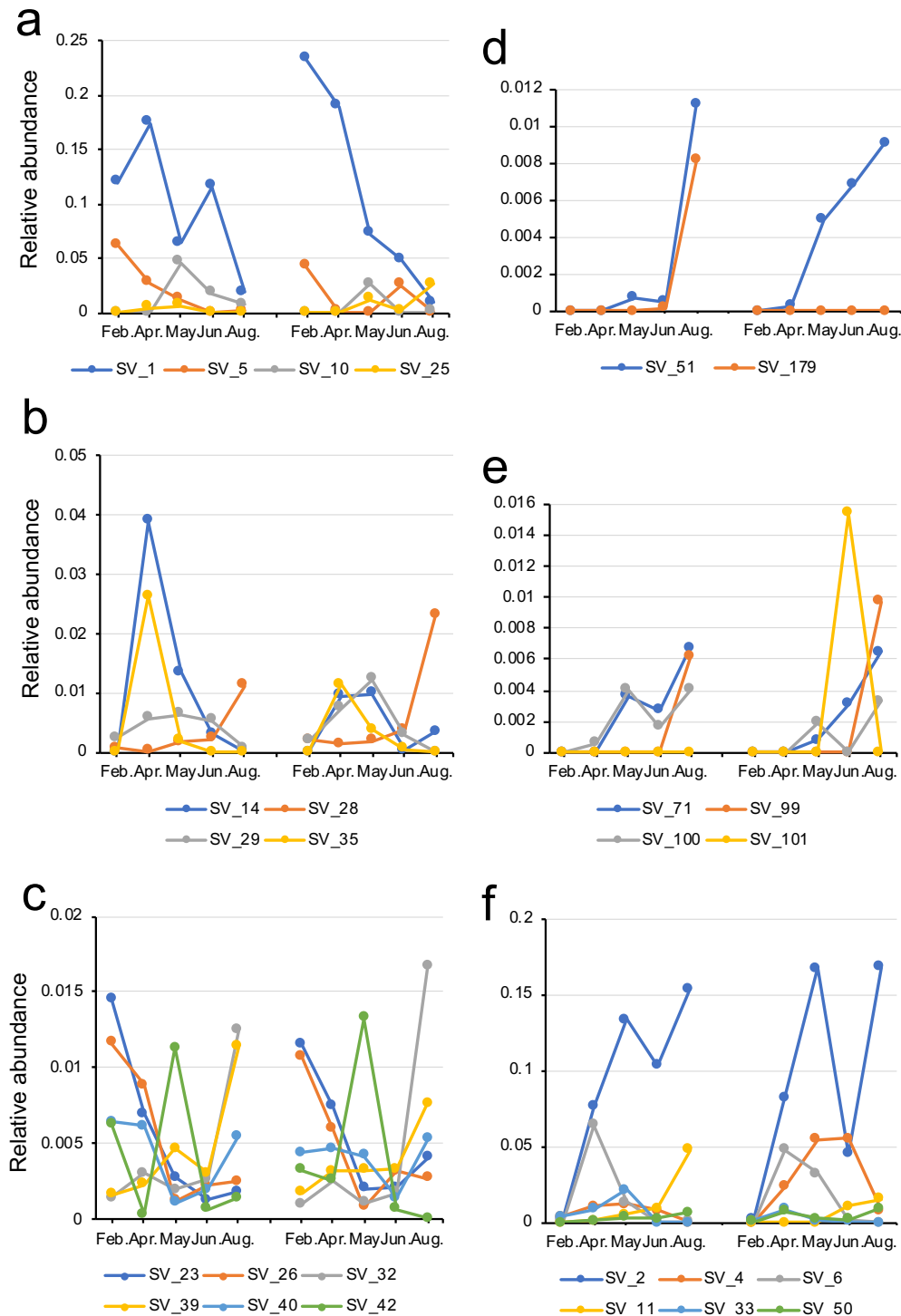


Supplementary Figure S7. Taxonomic compositions and Bray–Curtis dissimilarity plots of protist-derived SVs in the samples. (a) The relative abundances of kingdoms-derived SVs in total protists-derived SVs are shown in the samples. The taxa based on the SILVA database are indicated by the colors shown on the right of the bar graphs. NA: unassigned protist clades. The Bray–Curtis dissimilarity plots of SVs assigned to kingdoms Alveolata (b), Amoebozoa (c), Rhizaria (d), and Stramenopiles (e) are

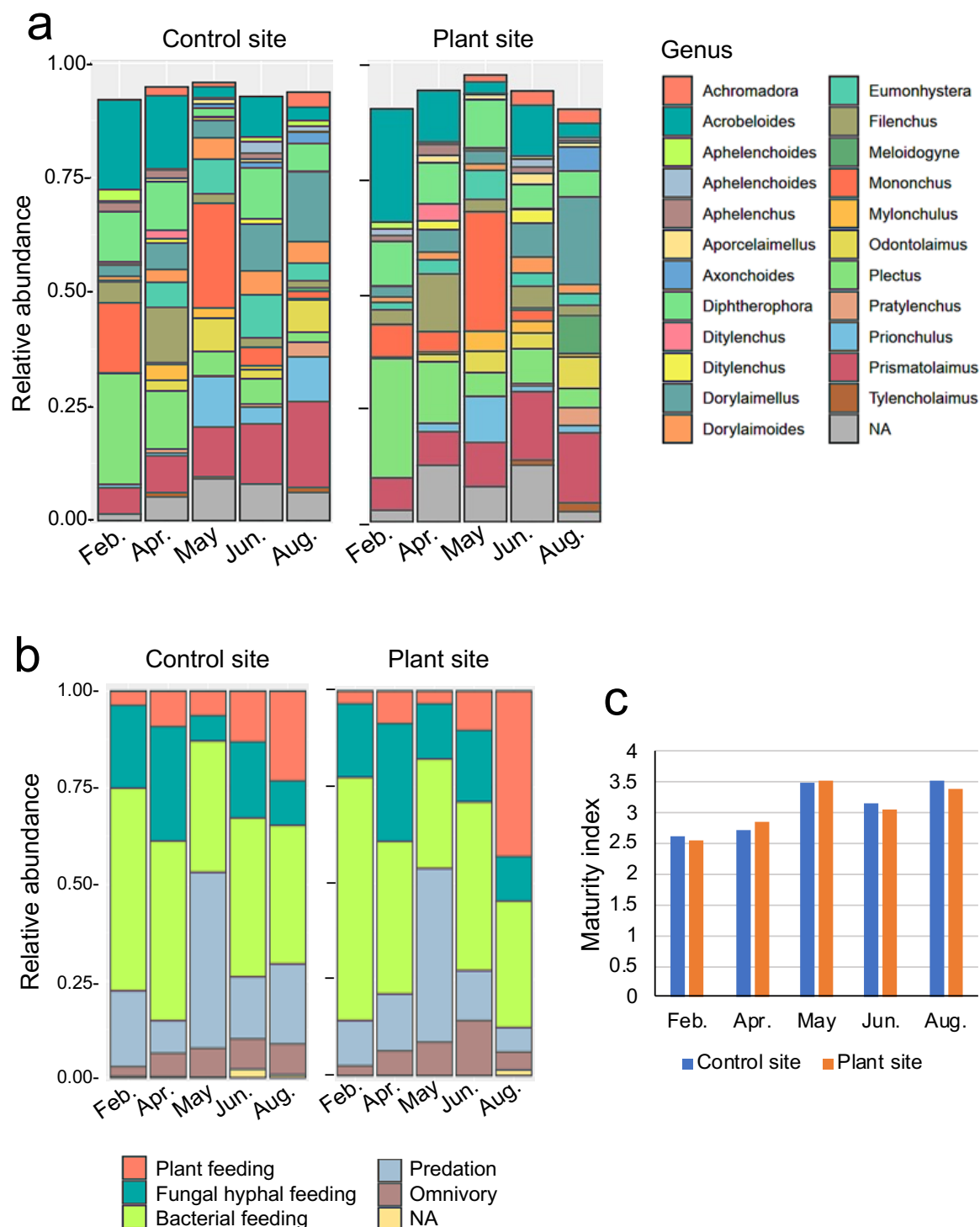
shown. Each sample and sampling site are indicated by colors and symbols (circles: control site; triangles: plant site) as shown on the left side of the figures. The compositions of protists class-derived SVs (f) in total protists SVs, and the relative abundances of Phytomyxea-derived SVs in total eukaryotic SVs (g) are indicated by bar charts in the samples. Bar charts of the samples derived from control and plant sites are shown on the left and right, respectively. NA and Incertae_Sedis: unassigned protist classes. uncultured: uncultured protists.



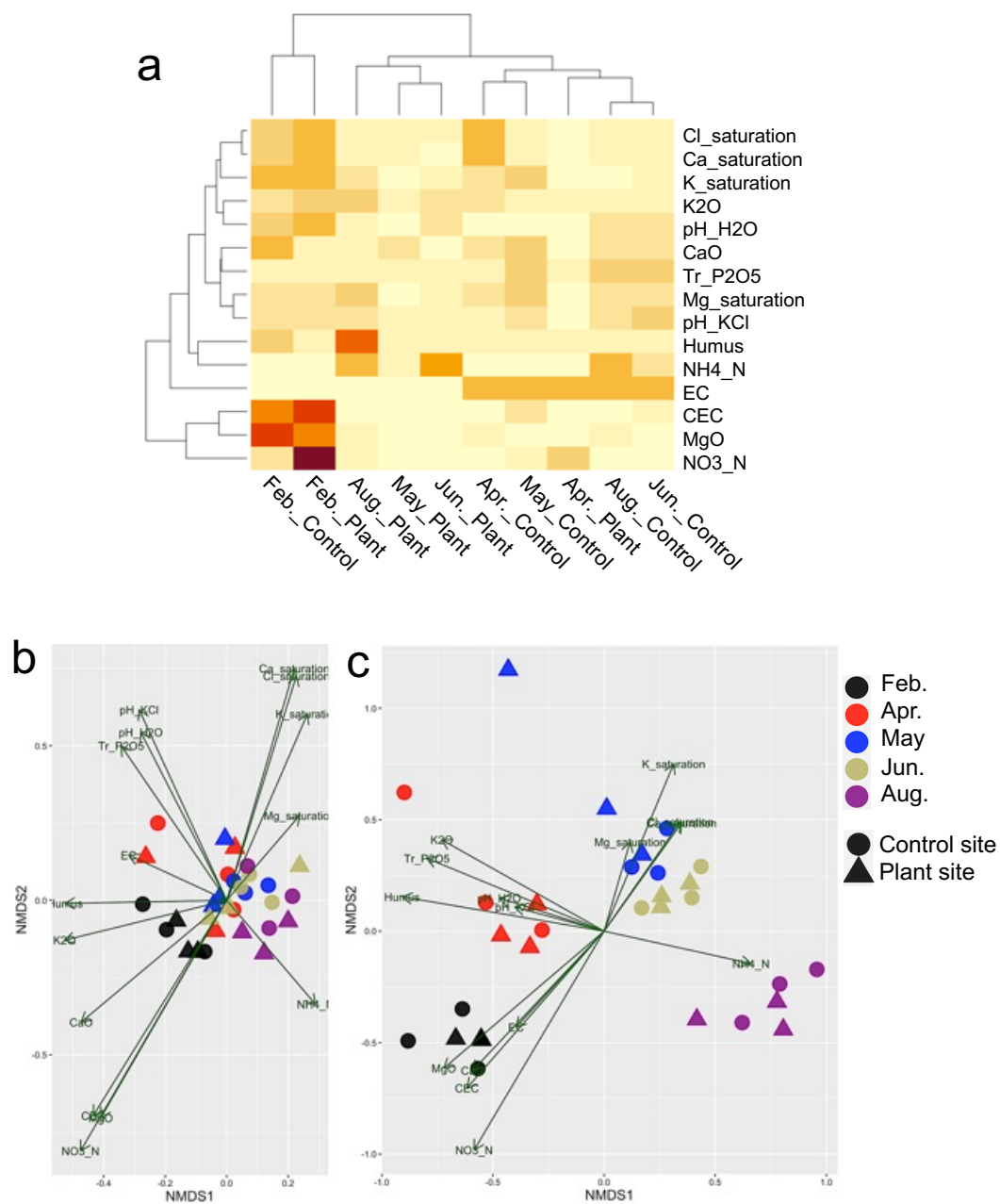
Supplementary Figure S8. Changes in the relative abundances of abundant SVs in each protistian phylum. The relative abundances of abundant SVs shown in the bottom of the panels of Diatomea (a), Cercozoa (b), Peronosporomycetes (c), Apicomplexa (d), Hyphochytriomycetes (e), Ciliophora (f), Centrohelida (g), and Bicosoecida (h), in total eukaryotic SVs are indicated by line graphs in the samples (see Supplementary Table S2). The left and right graphs indicate the results from control and plant sites, respectively.



Supplementary Figure S9. Changes in the relative abundances of animal- and plant-derived abundant SVs in each phylum. The relative abundances of abundant SVs shown on the bottom of the panels of Annelida (a), Arthropoda (b), Nematode (c), Gastrotrocha (d), Platyhelminthes (e), and Phragmoplastophyta (plants) (f), in total eukaryotic SVs are indicated by line graphs in the samples (see Supplementary Table S2). The left and right graphs indicate the results from control and plant sites, respectively.



Supplementary Figure S10. Compositions of nematode genus-derived SVs (a) and feeding habitat-classified SVs (b) in total Nematozoa-derived SVs in the samples. The results from the control and plant sites are shown on the left and right, respectively. The SILVA database-based genera are shown on the right of (a). NA: unassigned nematode genus. Feeding types of SVs are derived from the Supplementary Table S3 as shown on the bottom of (b). NA: unassigned feeding habitat. (c) Maturity index of nematode-derived SVs are indicated by bar charts in the samples derived from control and plant sites.



Supplementary Figure S11. (a) Heatmap of soil chemical parameters in the samples. Samples are clustered by the relative contents of chemical parameters based on the measured values (see Supplementary Data S3). Each parameter and the sample with its corresponding sampling site are indicated on the right and bottom of the map, respectively. Abbreviations of chemical parameters: cation-exchange capacity (CEC), chloride saturation (Cl_saturation), calcium saturation (Ca_saturation), potassium saturation (K_saturation), magnesium saturation (Mg_saturation), pH in water and KCl solution (pH_H2O, pH_KCl), nitrate nitrogen (NO3_N), ammonium nitrogen (NH4_N), exchangeable potassium (K2O), exchangeable magnesium (MgO), exchangeable calcium (CaO), electric conductivity, available phosphorus (Tr_P2O5), and humus content (Humus). (b) Redundancy analysis (RDA) of prokaryotic (b) and eukaryotic (c) communities in the samples showing soil chemical properties as

vectors. The colors and symbols for samples and sampling sites are shown on the right of the figure. Longer arrows indicate a stronger association.