Samples mostly contaminaned by the species	Contaminant species taxonomy
Number of samples Percentage of sa	mples
176	37.77 Bacteria;Firmicutes;Clostridia;Peptostreptococcales;Peptostreptococcaceae;Clostridioides;Clostridioides difficile
56 42	12.02 Bacteria;Firmicutes;Bacilli;Latctobacillaceae;Weissella;Weissella viridescens 9.01 Bacteria;Bacteroidota;Bacteroidia;Cytophagales;Microsdilaceae;Hassallia;metagenome
25	5.36 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP dade;Bilateria;Deuterostomia;Chordata;Vertebrata;Gnathostomata;Euteleostomi;Tetrapoda;Mammalia;Homo sapiens
24	5.15 Bacteria;Cyanobacteria;Cyanobacteriia;Chloroplast;seawater metagenome
15	3.22 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Ecdysozoa;Arthropoda;Chelicerata;Arachnida;Acari;Lasioerythraeus sp. OSAL 7511
9 5	1.93 Bacteria;Bacteroidota;Bacteroidia;Cytophagales;Microscillaceae;Ohtaekwangia;metagenome 1.07 Bacteria;Proteobacteria;Alphaproteobacteria;Rickettsiales;Mitochondria;Nicotiana tabacum
4	0.86 Eukaryota;Archaeplastida;Chloroplastida;Charophyta;Phragmoplastophyta;Streptophyta;Tracheophyta;Spermatophyta;Magnoliophyta;Fabales;Vigna;Vigna angularis var. angularis
4	0.86 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Deuterostomia;Chordata;Vertebrata;Gnathostomata;Euteleostomi;Actinopterygii;Neopterygii;Teleostei;Salmo salar
4	0.86 Bacteria;Proteobacteria;Gammaproteobacteria;Xanthomonadales;Xanthomonadaceae;Stenotrophomonas;[Pseudomonas] geniculata
3	0.86 Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacterales; Pasteurellaceae; Haemophilus; uncultured bacterium 0.64 Eukaryota; SAR; Rhizaria; Retaria; Polycystinea; Collodaria; Thalassicolla; Thalassicolla nucleata
3	0.64 Eukaryota;SAR;Rhizaria;Retaria;Polycystinea;Collodaria;Collosphaera;Collosphaera tuberosa
3	0.64 Eukaryota;Archaeplastida;Chloroplastida;Charophyta;Phragmoplastophyta;Streptophyta;Embryophyta;Tracheophyta;Spermatophyta;Magnoliophyta;Brassicales;Brassica;Brassica juncea var. tumida
3	0.64 Eukaryota; Archaeplastida; Chloroplastida; Charophyta; Phragmoplastophyta; Betula pendula 0.64 Eukaryota; Amorphea; Obazoa; Opisthokonta; Holozoa; Choanozoa; Metazoa; Animalia; BCP clade; Bilateria; Protostomia; Ecdysozoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Coleoptera; Conscisor a diffornicus
3	0.64 Eukaryota;Amorphea;Obazoa;Opistiokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Edysozoa;Arithropoda;Hexapoda;Insecta;Pterygota;Neoptera;Amphiesmenoptera;Lepidoptera;Ptyssoptera sp. A8-2015
3	0.64 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Ecdysozoa;Arthropoda;Crustacea;Malacostraca;Eumalacostraca;Peracarida;Proasellus parvulus
3	0.64 Bacteria;Proteobacteria;Gammaproteobacteria;Burkholderiales;Burkholderiaceae;Ralstonia;Ralstonia solanacearum
2	0.64 Bacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cyanobacteria;Cy
2	0.43 Eukaryota;Amorphea;Obazoa;Opisthokonta;Nucletmyœa;Fungi;Dikarya;Ascomycota;Saccharomycotina;Saccharomycetes;Saccharomycetales;synthetic construct
2	0.43 Eukaryota;Amorphea;Obazoa;Opisthokonta;Nucletmycea;Fungj;Dikarya;Ascomycota;Saccharomycotina;Saccharomycetes;Saccharomycetales;Incertae Sedis;Kuraishia;Kuraishia borneana
2	0.43 Eukaryota;Amorphea;Obazoa;Opisthokonta;Nucletmycea;Fungi;Dikarya;Ascomycota;Pezizomycotina;Sordariomycetes;metagenome
2 2	0.43 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Lophotrochozoa;Annelida;Polychaeta;Palpata;Phyllodocida;Pisione hartmannschroederae 0.43 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Ecdysozoa;Arthropoda;Chelicerata;Arachnida;Araneae;Leucauge blanda
2	0.43 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Deuterostomia;Chordata;Vertebrata;Gnathostomata;Euteleostomi;Callithrix jacchus
2	0.43 Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Enterobacterisceae;Enterobacteria;Dateriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteriaceae;Enterobacteria
2	0.43 Bacteria;Patescibacteria;Microgenomatia;Candidatus Levybacteria,Candidatus Levybacteria bacterium GW2011_GWC1_40_19 0.43 Bacteria;Bacteroidota;Bacteroidia;Flavobacteriales;Flavobacteriaceae;Flavivirga;Flavobacteriaceae bacterium TID809
1	0.21 Eukaryota;SAR;Stramenopiles;Ochrophyta;Diatomea;Bacillariophytina;Bacillariophyceae;Placoneis;Placoneis hambergii
1	0.21 Eukaryota;SAR;Rhizaria;Retaria;Polycystinea;Collodaria;Thalassicolla;Thalassicolla pellucida
1	0.21 Eukaryota;SAR;Alveolata;Dinoflagellata;Dinophyceae;Peridiniphycidae;Gonyaulacales;Gonyaulax;uncultured dinoflagellate
1	0.21 Eukaryota;Cryptophyceae;Cryptomonadales;Cryptomonas;Cryptomonas borealis 0.21 Eukaryota;Archaeplastida;Chloroplastida;Chlorophyta;Phragmoplastophyta;Pinus taeda
1	0.21 Eukaryota;Archaeplastida;Chloroplastida;Charophyta;Phragmoplastophyta;Picea abies
1	0.21 Eukaryota;Archaeplastida;Chloroplastida;Charophyta;Phragmoplastophyta;Nicotiana sylvestris
1	0.21 Eukaryota;Archaeplastida;Chloroplastida;Chlorophyta;Phragmoplastophyta;Alosa alosa 0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Nudetmycea;Fungi;Dikarya;Basidiomycota;Agaricomycotina;Tremellomycetes;Filobasidiales;Filobasidiaceae;Heterocephalacria;uncultured Tremellaceae
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Porifera;Demospongiae;Poecilosderida;Crambe crambe
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Lophotrochozoa;Platyhelminthes;Rhabditophora;Seriata;Tridadida;Maricola;Ectoplana limuli
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Lophotrochozoa;Platyhelminthes;Cestoda;Eucestoda;Cyclophyllidea;Raillietina mitchelli 0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Lophotrochozoa;Platyhelminthes;Cestoda;Eucestoda;Cyclophyllidea;Hymenolepis nana
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Lophotrochozoa;Mollusca;Gastropoda;Diaulula sandiegensis
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Ecdysozoa;Nematozoa;Nematoda;Enoplea;Dorylaimia;Trichocephalida;Trichinella murrelli
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Ecdysozoa;Nematozoa;Nematozoa;Nematodoa;Enoplea;Dorylaimia;Trichinella pseudospiralis 0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Ecdysozoa;Arthropoda;Hexapoda;Insecta;Pterygota;Palaeoptera;Ephemeroptera;Platybaetis probus
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Protostomia;Ecdysozoa;Arthropoda;Chelicerata;Arachnida;Opiliones;Cimex hemipterus
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Deuterostomia;Chordata;Vertebrata;Gnathostomata;Euteleostomi;Tetrapoda;Mammalia;Monodelphis domestica
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Deuterostomia;Chordata;Vertebrata;Gnathostomata;Euteleostomi;Tetrapoda;Mammalia;Malademys terrapin terrapin 0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Deuterostomia;Chordata;Vertebrata;Gnathostomata;Euteleostomi;Tetrapoda;Aves;Gallus gallus
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Deuterostomia;Chordata;Vertebrata;Gnathostomata;Euteleostomi;Oryctolagus cuniculus
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Deuterostomia;Chordata;Vertebrata;Gnathostomata;Euteleostomi;Lates calcarifer
1	0.21 Eukaryota;Amorphea;Obazoa;Opisthokonta;Holozoa;Choanozoa;Metazoa;Animalia;BCP clade;Bilateria;Deuterostomia;Chordata;Vertebrata;Gnathostomata;Euteleostomi;Homo sapiens 0.21 Bacteria;Verrucomicrobiota;Verrucomicrobiae;Opitutales;Puniceicoccaceae;marine metagenome
1	0.21 Bacteria;Spirochaetota;Spirochaetia;Spirochaetales;Spirochaetaceae;Treponema;Clostridiales bacterium Firm_16
1	0.21 Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Sponglibacteraceae;uncultured gamma proteobacterium HF4000_36I10
1	0.21 Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales; Pseudomonadaceae; Pseudomonas; uncultured bacterium 0.21 Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales; Pseudomonadaceae; Pseudomonas; uncultured bacterium
1	0.21 Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales; Pseudomonadales; Pseudomonas; Pseudomonas linyingensis
1	0.21 Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Moraxellaceae;Acinetobacter;Acinetobacter sp. AR2-3
1	0.21 Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacterale; Morganellaceae; endosymbionts, gamma proteobacterium symbiont of Odoiporus longicollis 0.21 Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacterales; Enterobacteriaceae; Escherichia-Shigella; Escherichia coli
1	0.21 Bacteria; Proteobacteria; Gammaproteobacteria; Coxiellales; Coxiella; uncultured bacterium
1	0.21 Bacteria;Proteobacteria;Alphaproteobacteria;Sphingomonadales;Sphingomonadaceae;Sphingomonas koreensis
1	0.21 Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales; Mitochondria; Cryptomonas curvata 0.21 Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales; Devosia œae; Devosia œae; Devosia exae; Devos
1	0.21 Bacteria;Proteobacteria;Alphaproteobacteria;Rhizobiales;D05-2;metagenome
1	0.21 Bacteria;Patescibacteria;Microgenomatia;Candidatus Levybacteria;uncultured bacterium
1	0.21 Bacteria;Patescibacteria;Microgenomatia;Candidatus Gottesmanbacteria;uncultured bacterium 0.21 Bacteria;Firmicutes;Clostridia;Peptostreptococcales-Tissierellales;Peptostreptococcaeae;Clostridioides;Clostridioides difficile
1	0.21 Bacteria;Firmicutes;Clostridia;Peptostreptococcales-lissierellales;Peptostreptococcaceae;Clostridioides;Clostridioides difficile 0.21 Bacteria;Firmicutes;Clostridia;Eubacteriales;Eubacteriaceae;Eubacterium;Eubacterium callanderi
1	0.21 Bacteria; Firmicutes; Bacilli; Staphylococcales; Staphylococcaceae; Staphylococcus aureus
1	0.21 Bacteria;Firmicutes;Bacilli;tactobacillales;Lactobacillaceae;Limosilactobacillus;Lactobacillus fermentum
1	0.21 Bacteria;Chloroffexi;Anaerolineae;Anaerolineaeles;Anaerolineaceae;Thermomarinilinea;uncultured bacterium 0.21 Bacteria;Actinobacteriota;Actinobacteria;Micrococcaeles;Micrococcaee;Rothia;Rothia mucilaginosa ATCC 25296
1	0.21 Bacteria;Actinobacteriota;Actinobacteria;Frankiales;Sporidhthyaceae;Bgd dade;uncultured actinobacterium
1	0.21 Bacteria;Actinobacteriota;Actimicrobiia;Actinomarinales;uncultured;uncultured bacterium
1	0.21 Bacteria;Addobacteriota;Thermoanaerobaculia;Thermoanaerobaculales;Thermoanaerobaculaceae;Subgroup 10; uncultured bacterium 0.21 Archaea;Halobacterota;Methanosarcinia;Methanosarciniales;Syntrophoarchaeaceae;uncultured archaeon
1	U.21 Archaea;Halobacterota;Methanosaronia;Methanosaroniales;Syntropnoarchaeaceae;uncultured archaeon U.21 Archaea;Asgardarchaeota;Lokiarchaeia;uncultured crenarchaeote