

Results_RADSeqMicronbiome

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RADSeq Microbiome Results

The majority of the sequences found matched with *Arthropoda* database references, but also another seven problematic phyla were present and filtered.

Table 1: Total matches with 90% similarity

	Actinobacteria	Annelida	Apicomplexa	Arthropoda	Artverviricota
Denovo_95	0	0	3	645	1
Denovo_97	0	0	2	630	0
DPond_95	1	1	0	277	0
DPond_97	0	1	0	252	0
DRef_97	0	0	2	84	0
GRef_97	0	0	0	33	0

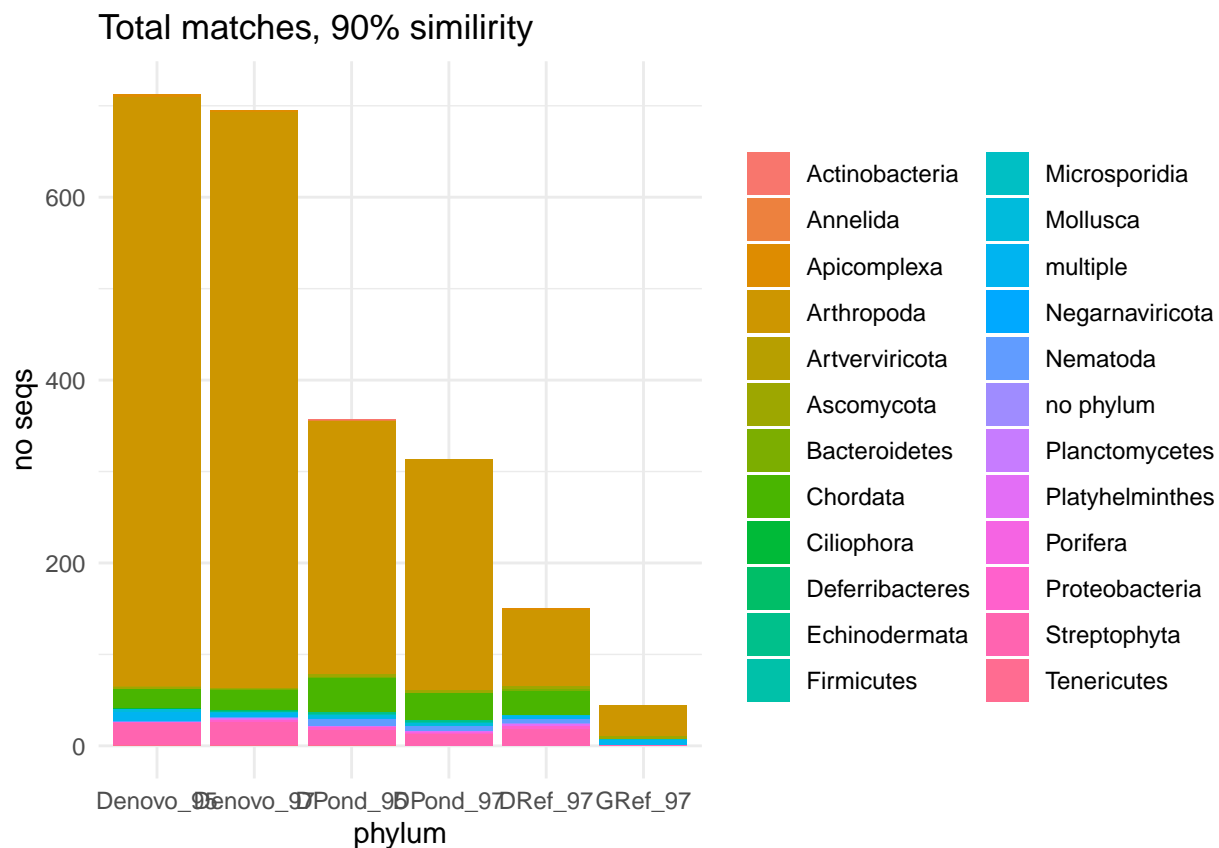
	Ascomycota	Bacteroidetes	Chordata	Ciliophora	Deferribacteres
Denovo_95	2	0	21	1	0
Denovo_97	1	1	22	0	1
DPond_95	3	1	37	0	0
DPond_97	3	0	30	0	0
DRef_97	3	2	26	0	0
GRef_97	3	0	1	0	0

	Echinodermata	Firmicutes	Microsporidia	Mollusca	multiple
Denovo_95	0	0	0	3	10
Denovo_97	1	0	0	1	4
DPond_95	2	1	0	4	1
DPond_97	2	1	0	4	0
DRef_97	0	0	0	1	4
GRef_97	0	0	1	0	5

	Negarnaviricota	Nematoda	no phylum	Planctomycetes	Platyhelminthes
Denovo_95	0	1	0	0	0
Denovo_97	1	1	0	1	1
DPond_95	0	7	0	0	1
DPond_97	0	5	0	0	1
DRef_97	0	4	1	0	2
GRef_97	0	0	0	0	0

	Porifera	Proteobacteria	Streptophyta	Tenericutes
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Denovo_95	0	2	23	1
Denovo_97	0	2	25	1
DPond_95	1	3	16	1
DPond_97	1	1	13	0
DRef_97	0	4	18	0
GRef_97	0	0	1	0



Overall, the most common phyla found were *Ascomycota* and *Proteobacteria*. Furthermore the most promising assembly method used in this survey was the use of an insect reference genome as filter on the subset of the reference specie samples (*Dendroctonus ponderosae*) with a clustering threshold of 95%.

Table 2: Filtered matches with 90% similarity

	Actinobacteria	Annelida	Apicomplexa	Artverviricota	Ascomycota
Denovo_95	0	0	3	1	2
Denovo_97	0	0	2	0	1
DPond_95	1	1	0	0	3
DPond_97	0	1	0	0	3
DRef_97	0	0	2	0	3
GRef_97	0	0	0	0	3

	Bacteroidetes	Ciliophora	Deferribacteres	Firmicutes	Microsporidia
Denovo_95	0	1	0	0	0
Denovo_97	1	0	1	0	0
DPond_95	1	0	0	1	0

DPond_97	0	0	0	1	0
DRef_97	2	0	0	0	0
GRef_97	0	0	0	0	1

	Negarnaviricota	Nematoda	no phylum	Planctomycetes	Proteobacteria
Denovo_95	0	1	0	0	2
Denovo_97	1	1	0	1	2
DPond_95	0	7	0	0	3
DPond_97	0	5	0	0	1
DRef_97	0	4	1	0	4
GRef_97	0	0	0	0	0

	Tenericutes
Denovo_95	1
Denovo_97	1
DPond_95	1
DPond_97	0
DRef_97	0
GRef_97	0

