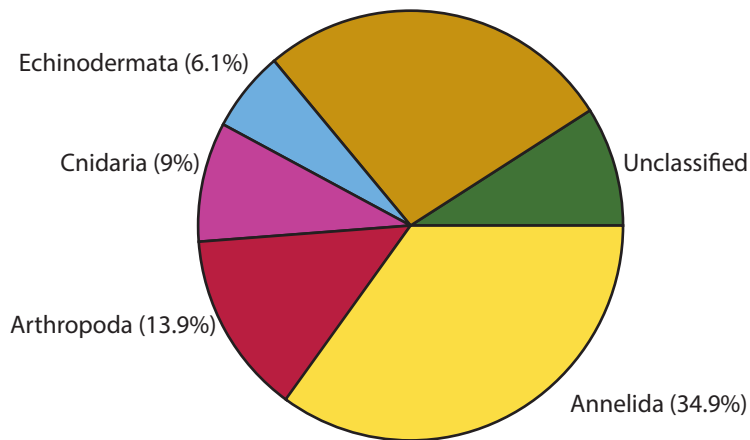


a

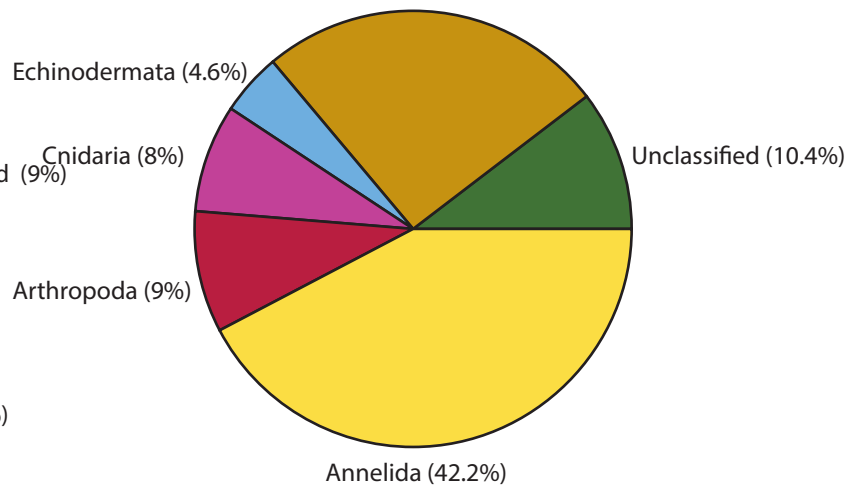
Mollusca (27.1%)



Phyla identified with Illumina
MiSeq metabarcoding

b

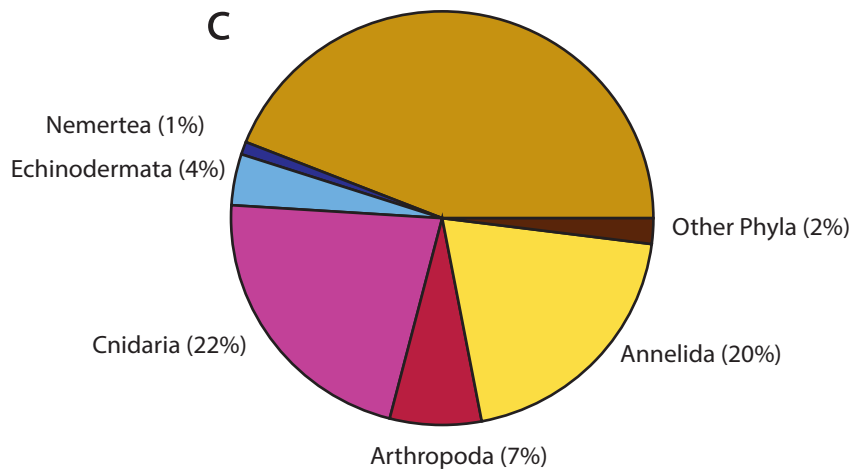
Mollusca (25.7%)



Phyla identified with Nanopore
metabarcoding

c

Mollusca (44 %)



Phyla identified with Novaseq
metagenomics