

README for *Parhyale hawaiiensis* 454 transcriptome sequences.

Please contact [Vzeng@oeb.harvard.edu](mailto:Vzeng@oeb.harvard.edu) with any questions.

Note: This is the plain fasta file for all sequences assembled from the high-throughput 454 *P. hawaiiensis* transcriptome. The accession for each sequence is unique, and are used to identify the specific sequences in literature. There will soon be an interactive website available for browsing these sequences including their annotation.