

Supplemental material

Figure S1. Benchmarking Using Single Copy Orthologs (BUSCO) score for each of the transcriptome assemblies

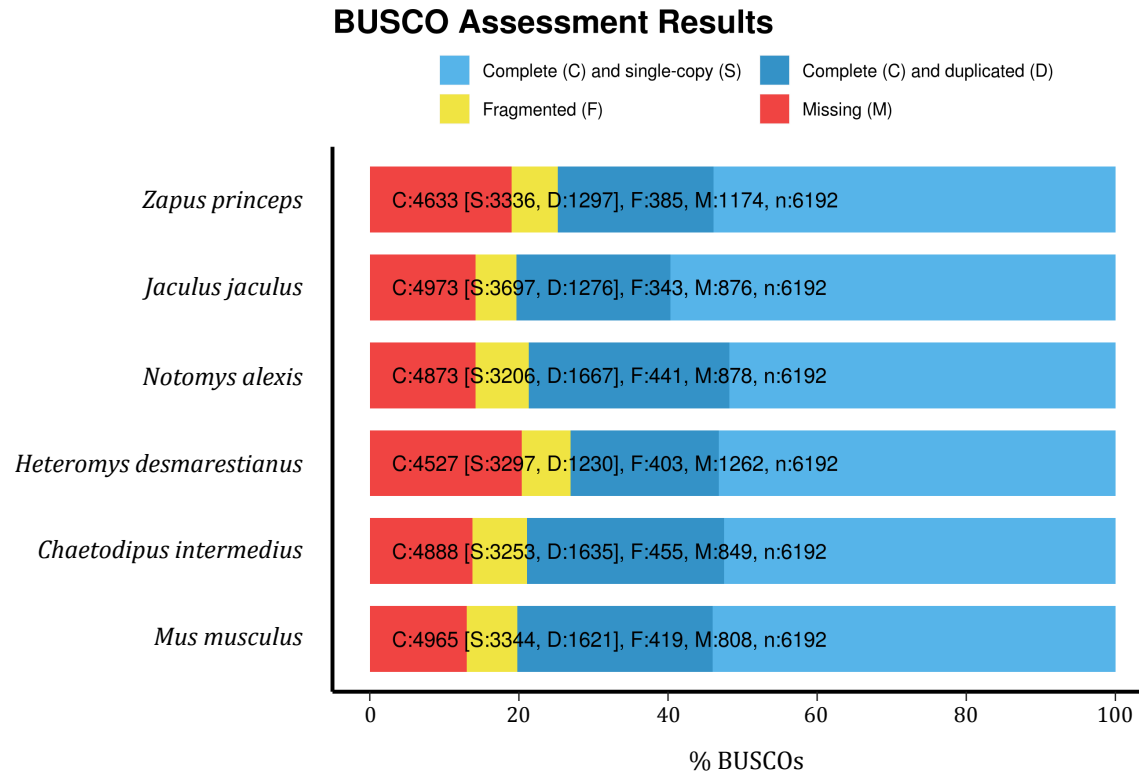


Figure S2. Genes that are differentially expressed between each desert and non-desert comparisons within each family and the overlaps among families.

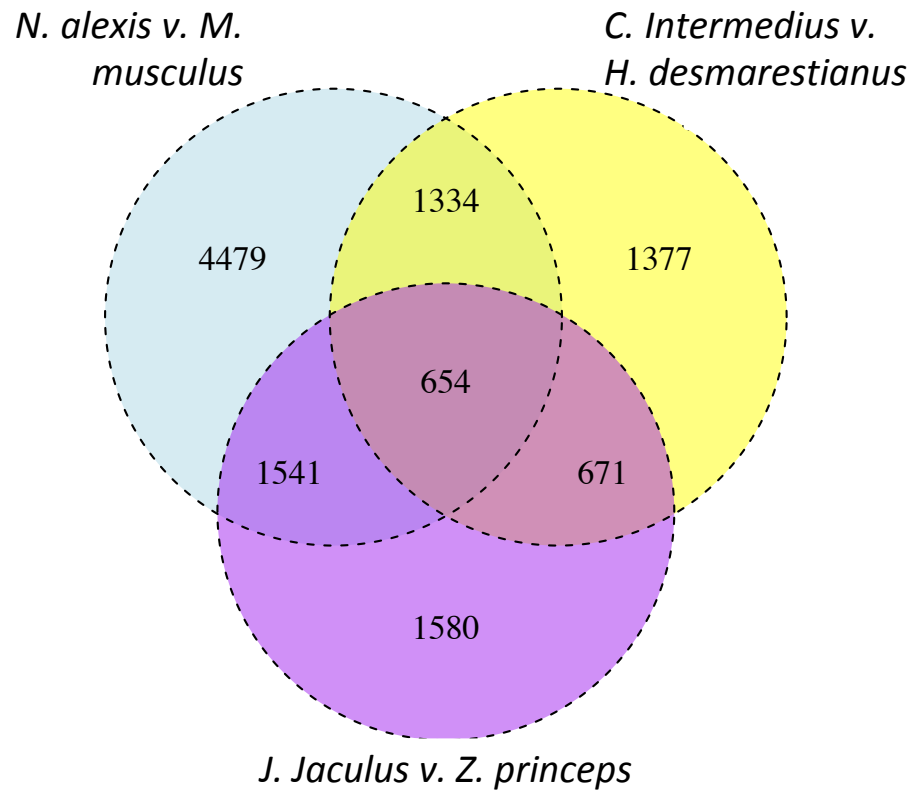


Figure S3. Magnitude of expression differences between each desert-mesic species pair

