S3 Figure. The 50 most significantly over- and under-enriched pathway components ($p < 0.001$) in the C. remanei genome (as compared to the C. elegans genome. Each value is scaled relative to the top pathway identified in each species, and pathways are grouped by biological function. C. remanei has a significant overrepresentation of pathway components involved in cellular processes, nucleotide metabolism, lipid metabolism and amino acid metabolism and a significant underrepresentation of pathway components involved in neural development and carbohydrate metabolism.