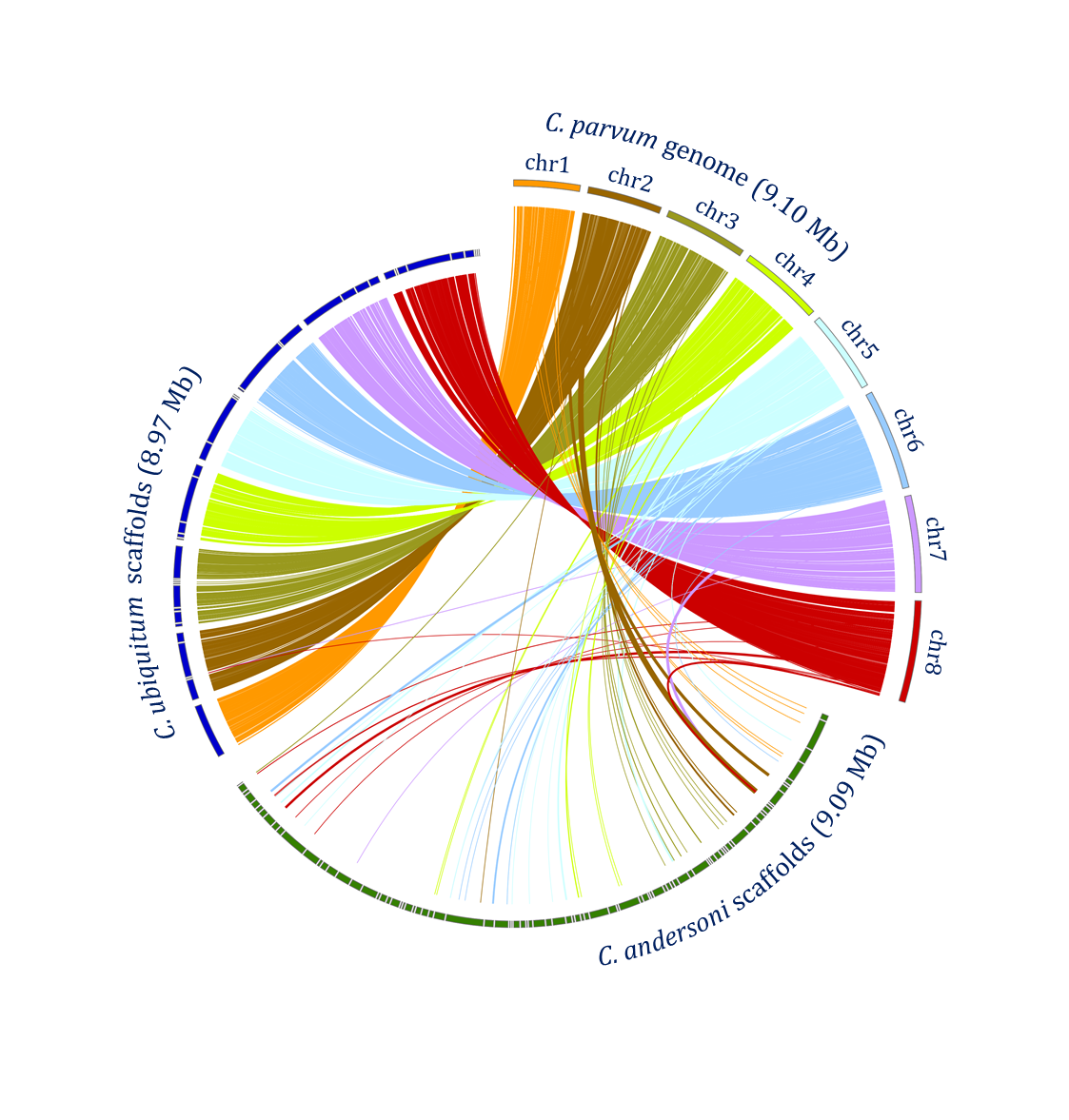
**Additional file 3: Figure S2:** A) Syntenic relationship between the genomes of *Cryptosporidium parvum* and *C. ubiquitum* or *C. andersoni*. Syntenic sequences (identity >75%) are connected with lines. The colors of lines represent different chromosomes of *C. parvum*. B) Venn diagram of shared orthologs and species-specific genes among four *Cryptosporidium* species. Because of the use of different gene prediction approaches in genome annotation, species-specific genes are generally over-estimated.



**A**

**B**

