Figure 2. Maximum likelihood phylogeny from concatenated character trait alignment with the Protoss, Zerg, Tyranid and Xenomorph taxa set as outgroups.
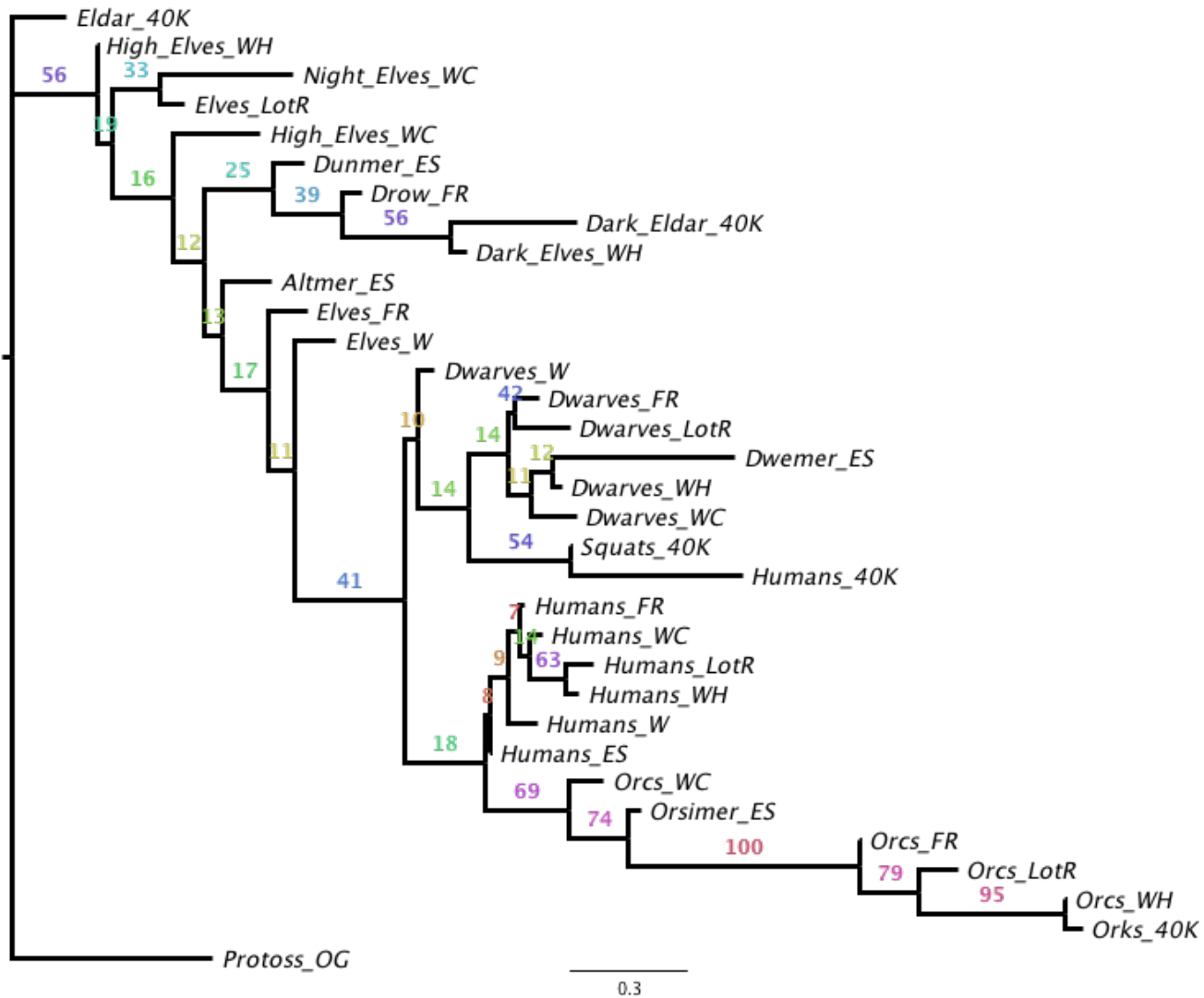
Figure 3. Maximum likelihood phylogeny from concatenated character trait alignment with the Protoss taxon set as the sole outgroup.
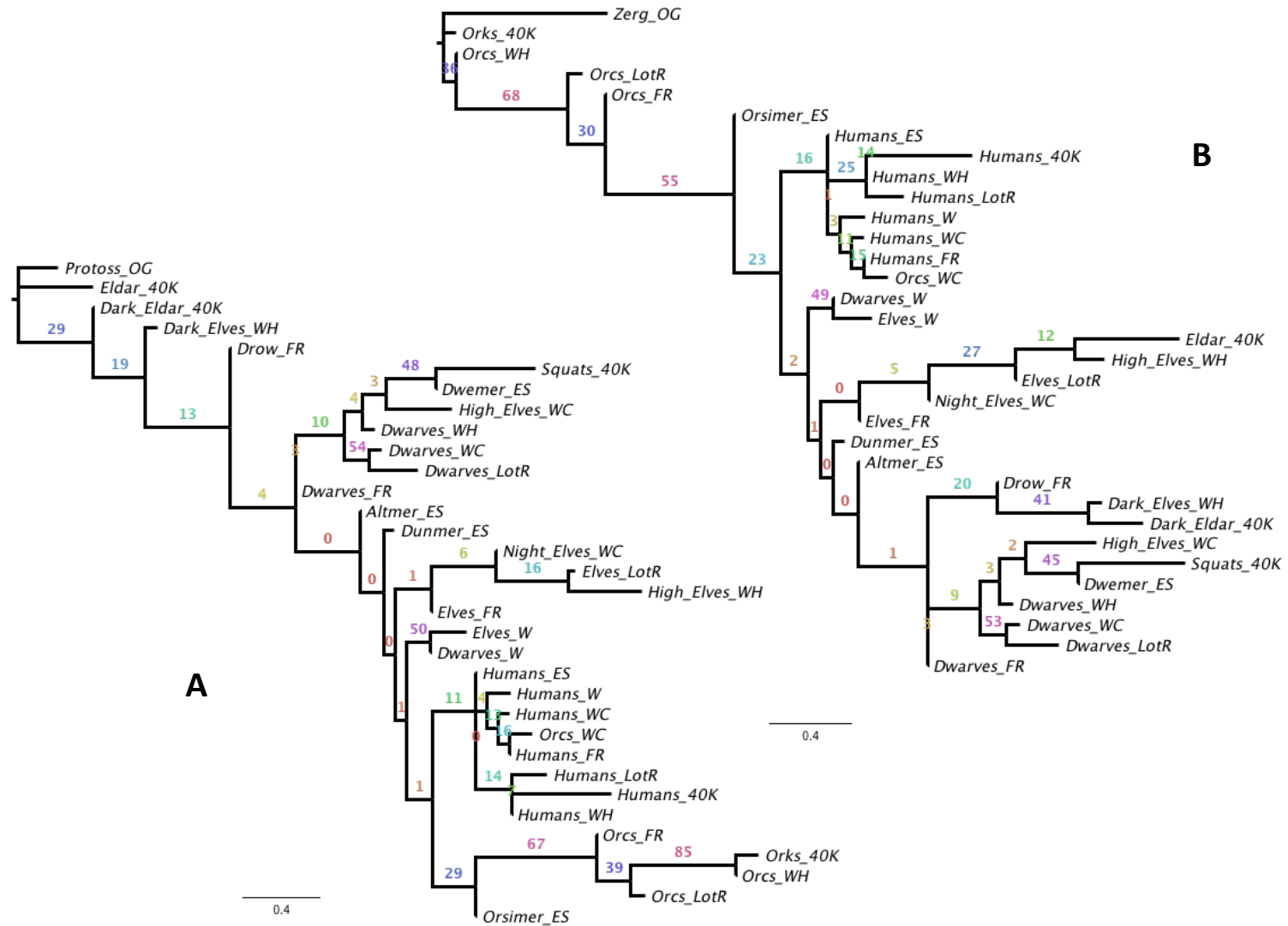
Figure 4. Maximum likelihood phylogenies from archetypal trait alignments; A. Protoss taxon set as outgroup; B. Zerg taxon set as outgroup.