Table 5. Summary of options used for running analyses in *RaxML* per alignment/outgroup combination assessed. The model of evolution used, number of likelihood trees searched, and number of bootstrap replicates performed are indicated for each dataset.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Analysis Dataset | Outgroup(s) Used | Model Used | # of Trees Searched | # of Bootstrap Replicates |
|
| Concatenated Alignment | Zerg\_OG Tyranids\_OG Xenomorphs\_OG | MULTI8\_MK | 1000 | 1000 |
|
|
| Concatenated Alignment | Protoss\_OG Zerg\_OG Tyranids\_OG Xenomorphs\_OG | MULTI8\_MK | 1000 | 1000 |
| Concatenated Alignment | Protoss\_OG | MULTI6\_MK | 1000 | 1000 |
| Physiological Alignment | Zerg\_OG | MULTI8\_MK | 400 | 1000 |
| Behavioral Alignment | Zerg\_OG | MULTI6\_MK | 400 | 1000 |
| Archetypal Alignment | Zerg\_OG | MULTI7\_MK | 400 | 1000 |
| Physiological Alignment | Protoss\_OG | MULTI8\_MK | 400 | 1000 |
| Behavioral Alignment | Protoss\_OG | MULTI6\_MK | 400 | 1000 |
| Archetypal Alignment | Protoss\_OG | MULTI7\_MK | 400 | 1000 |