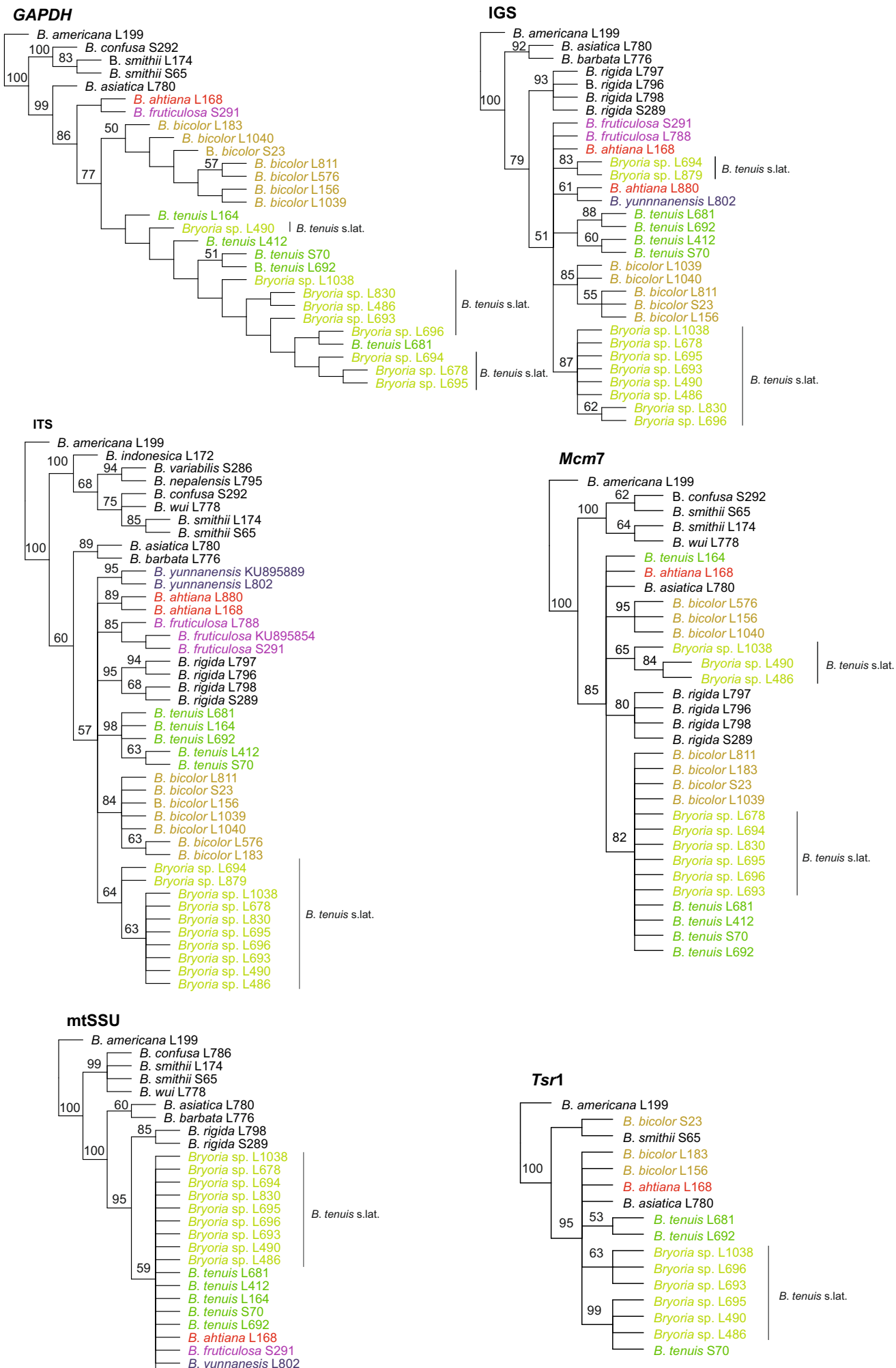

Fig S2. Separate *Divaricatae* phylogenies obtained from TNT analyses. Bootstrap values ≥ 50% are shown at nodes.