

Supplementary Table 2 – Number of species sampled for BAMM analysis and percentage relative to the total number of species reported for tribes Tulbaghieae, Gilliesieae (divided in subtribes Leucocoryninae and Gilliesiinae) and Allieae (divided in the three *Allium* evolutionary lineages)

Taxa	Number of sampled species in BAMM analysis	% of sampled species for BAMM analysis
Allieae		
<i>Allium</i> 1st evolutionary lineage	23	16%
<i>Allium</i> 2nd evolutionary lineage	55	30%
<i>Allium</i> 3rd evolutionary lineage	73	14%
Gilliesieae		
Gilliesiinae	6	27%
Leucocoryninae	30	45%
Tulbaghieae	3	14%