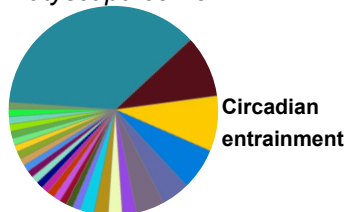
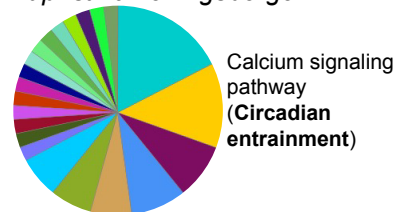


Figure 3: KEGG network enrichment analysis of genes near TEs inserted at 31-35 Mya.
 In the pie charts, the regions with different colour represented the merged group. Yellow regions were groups leading by Ciradian entrainment pathway or by other terms which contained Ciradian entrainment pathway.

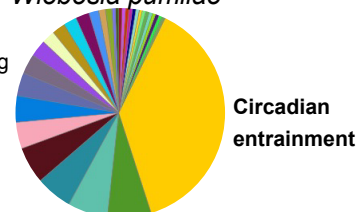
Platyscapa corneri



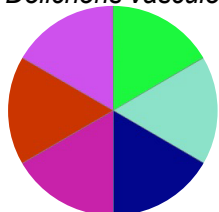
Eupristina koningsbergeri



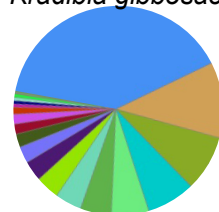
Wiebesia pumilae



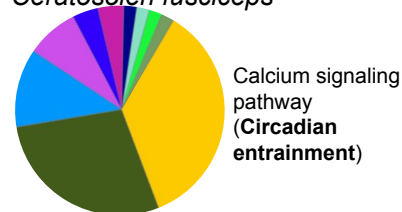
Dolichoris vasculosae



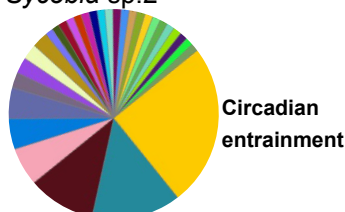
Kradibia gibbosae



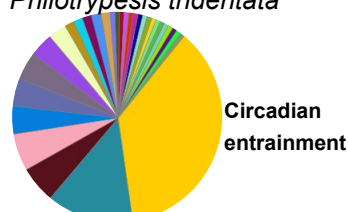
Ceratosolen fusciceps



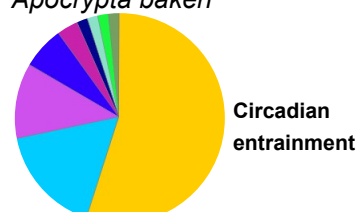
Sycobia sp.2



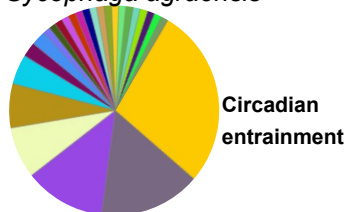
Philotrypesis tridentata



Apocrypta bakeri



Sycophaga agraensis



Sycophila sp.2

