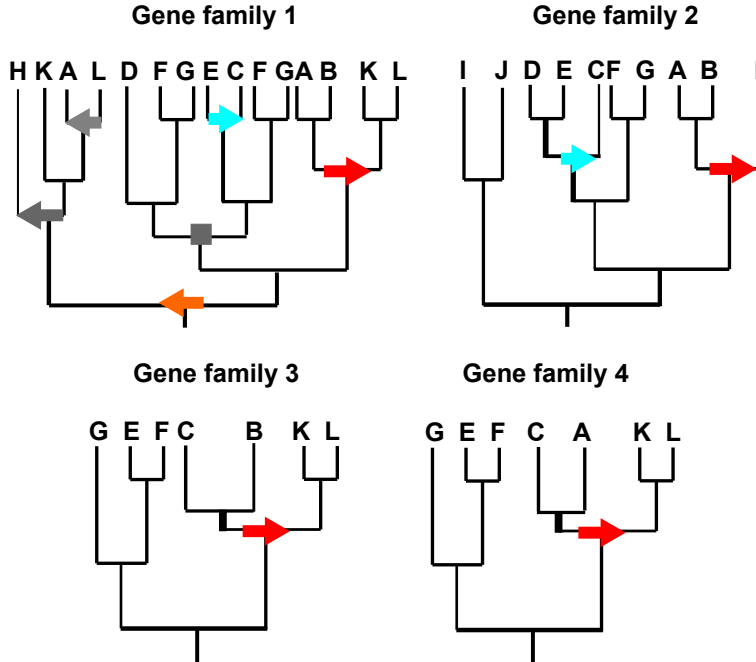
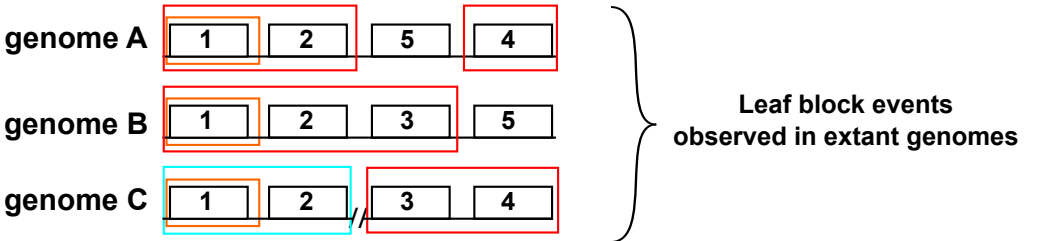
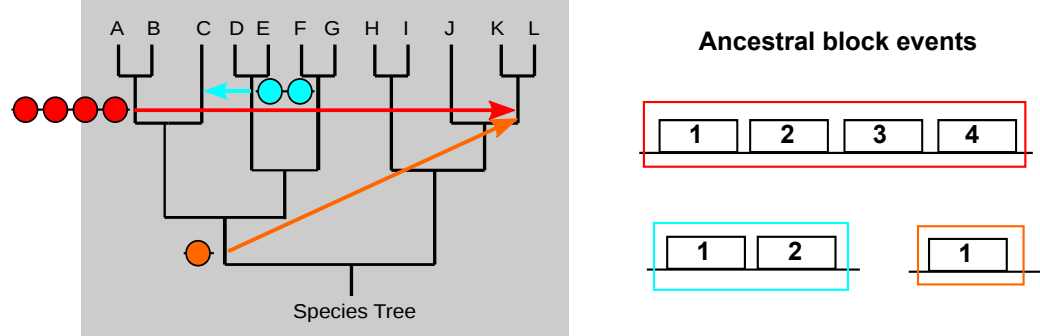
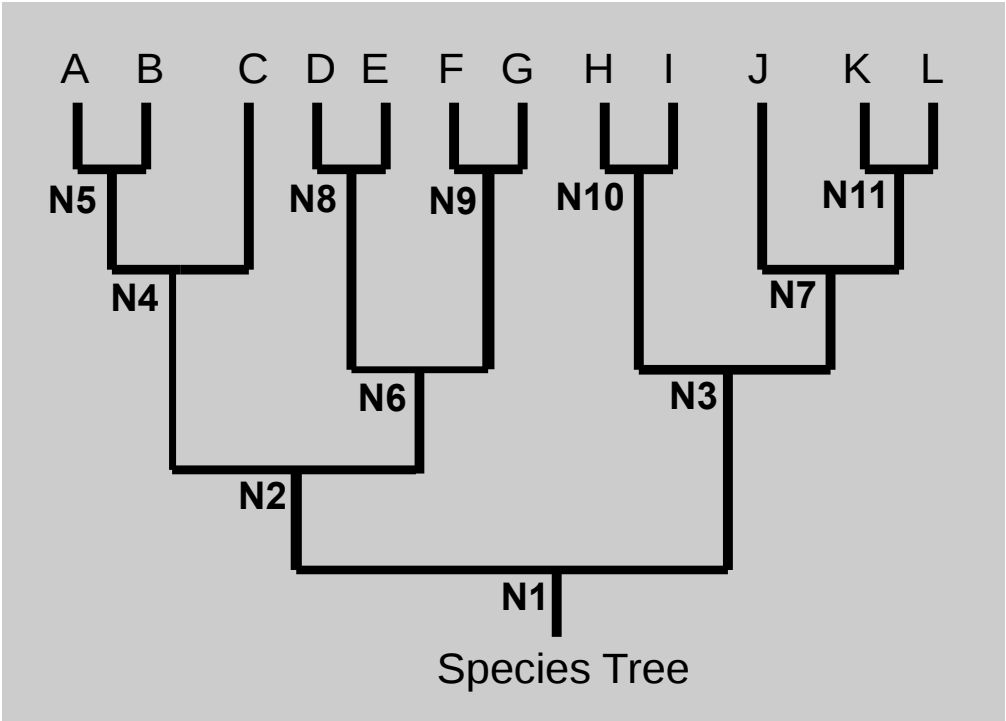
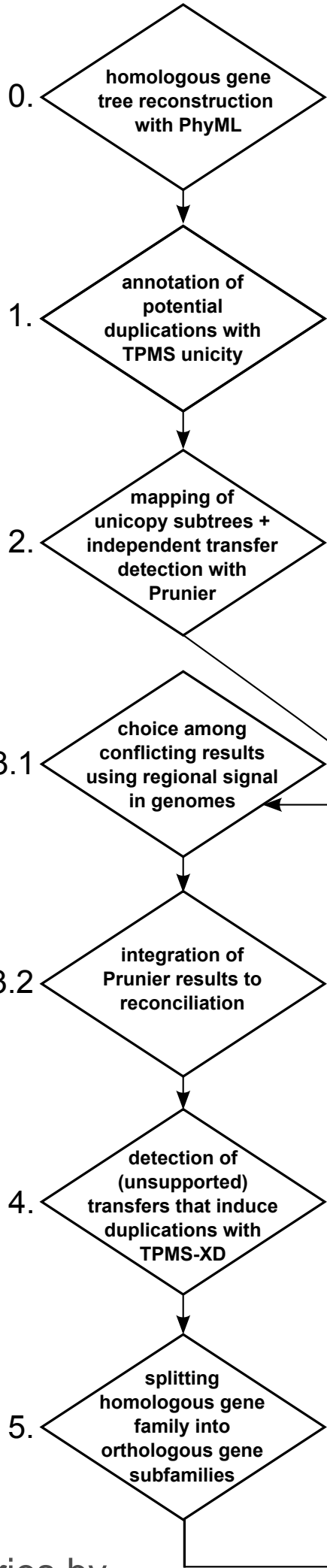
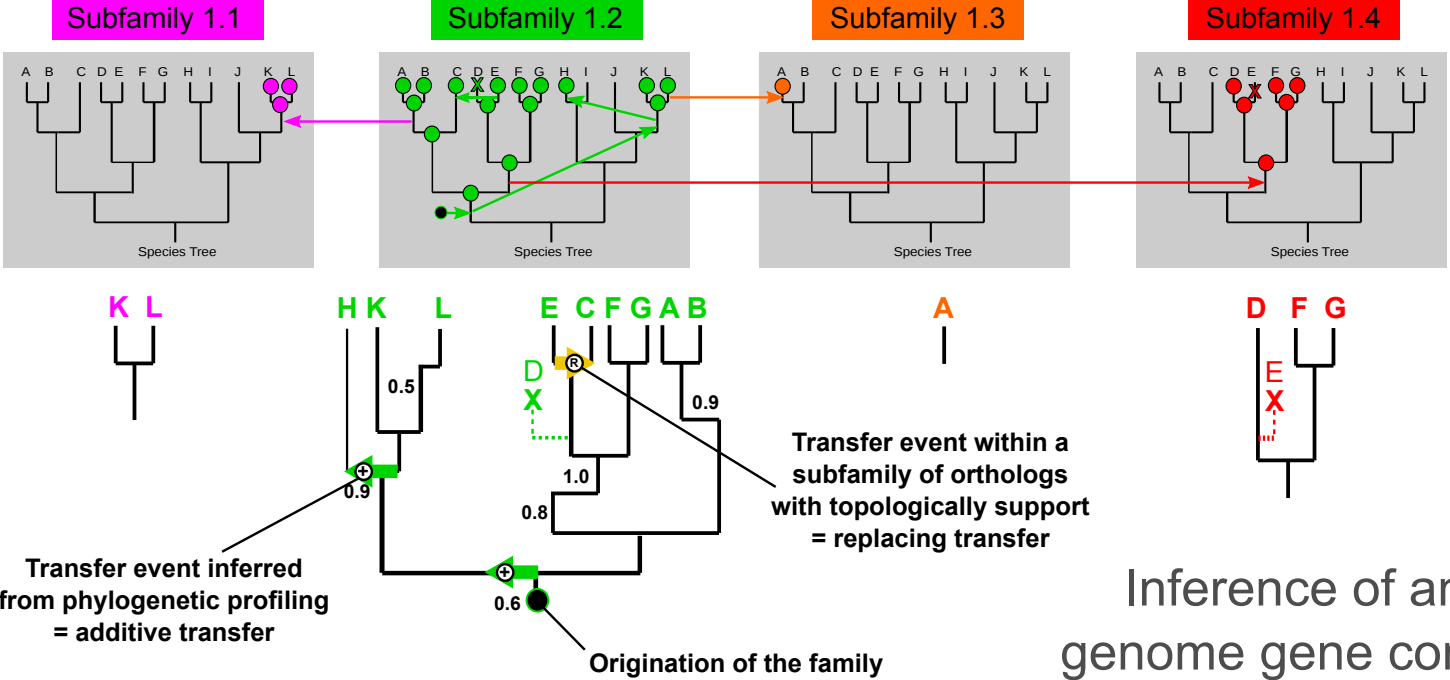


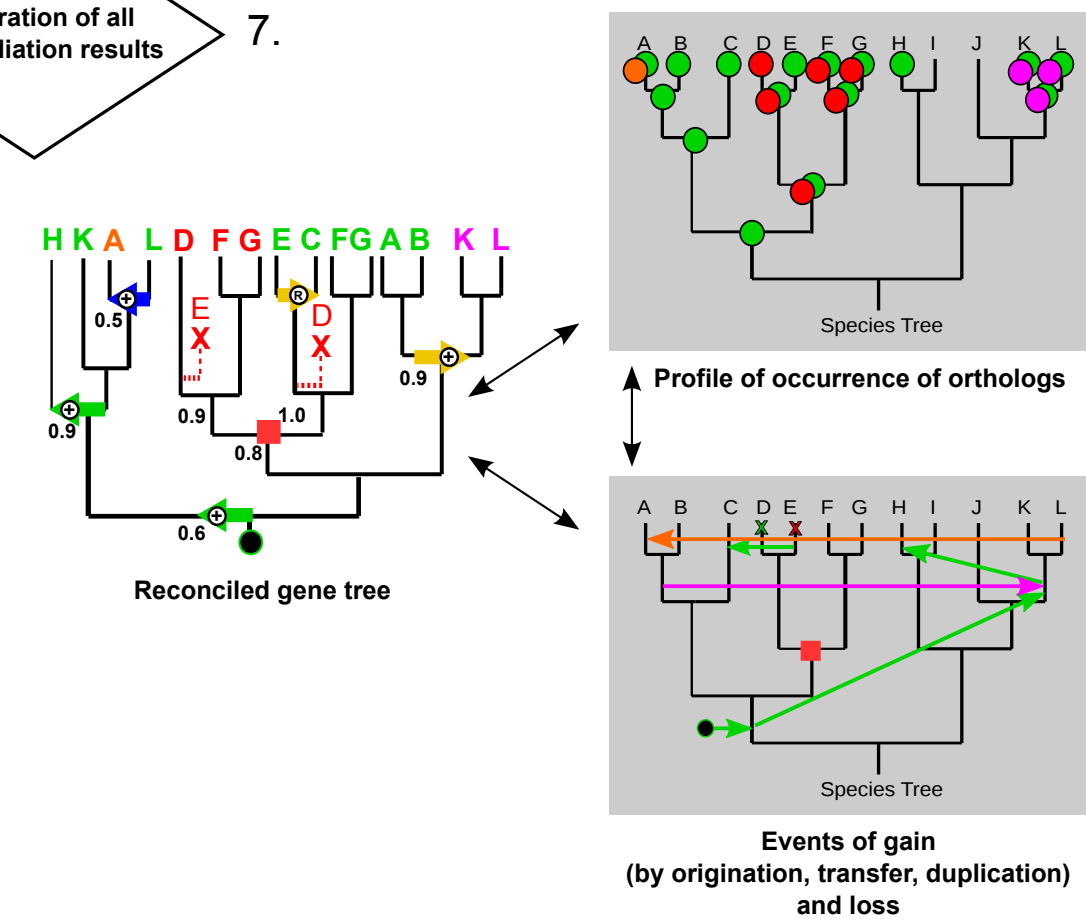
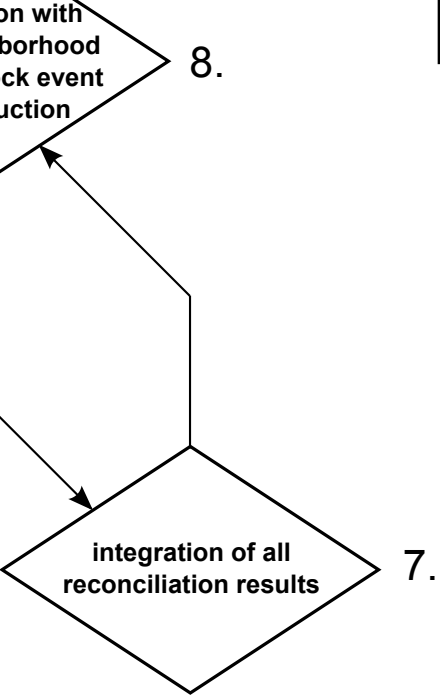
Agrogenom reconciliation pipeline



Inference of block events along genome histories by accretion of neighbouring events with congruent signatures



Inference of ancestral genome gene contents and evolutionary events from orthologous gene subfamily presence/absence profiles



Inference of events along gene histories by detection of topological incongruences and definition of orthologous gene subfamilies