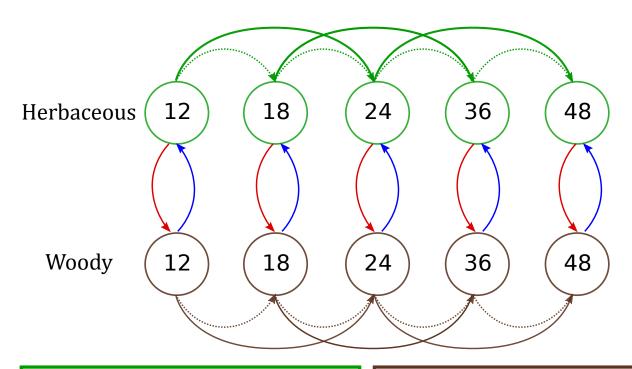
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Appendix S2. Graphical representation of the model proposed to analyze the *Solanum* data set. The *Solanum* model is a continuous time Markov Chain (Mkn model) containing 10 states, five herbaceous and five woody, each with chromosome number values 12, 18, 24, 36, and 48. It is defined via six parameters: two for chromosome doubling as a proxy of polyploidy ( $\rho_H$ ,  $\rho_w$ ), two for increase by six or 12 chromosomes as a proxy of demiploidy ( $\varepsilon_H$ ,  $\varepsilon_W$ ), and two representing the transitions between woody and herbaceous states ( $q_{HW}$ ,  $q_{WH}$ ).



 $\rho_{\rm H} = 0.139$ 

Rate polyploidy herbaceous

 $\epsilon_{H} = 0.084$ 

Rate demiploidy herbaceous

 $q_{HW} = 0.835$ 

Rate herbaceous to woody

 $\rho_{W} = 0.002$ 

Rate polyploidy woody

 $\epsilon_{\rm W} = 1.14 \times 10^{-19}$ 

Rate demiploidy woody

 $q_{WH} = 0.453$ 

Rate woody to herbaceous