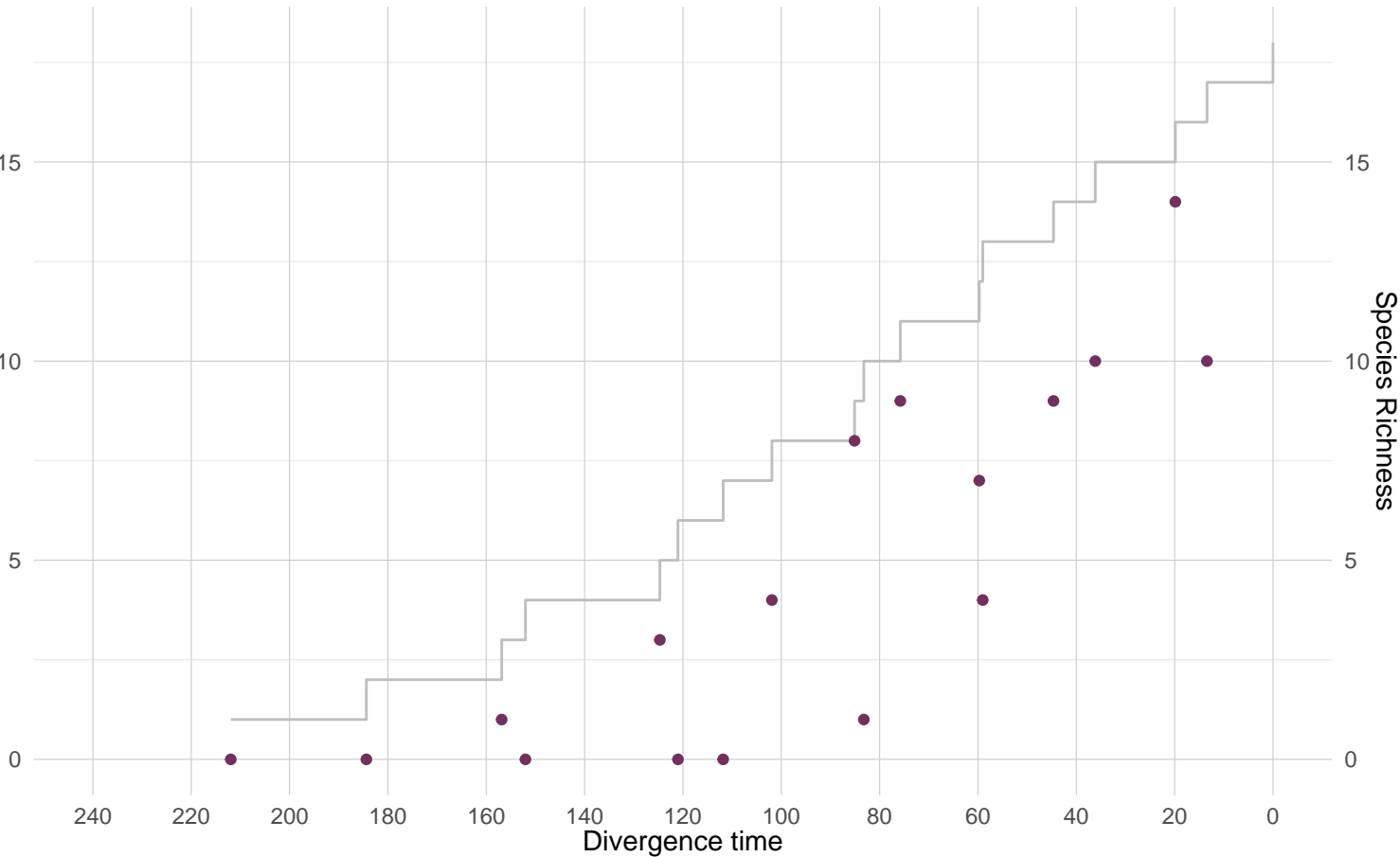


Internal nodes

Number of outlier genes present in all associated tips



Species Richness