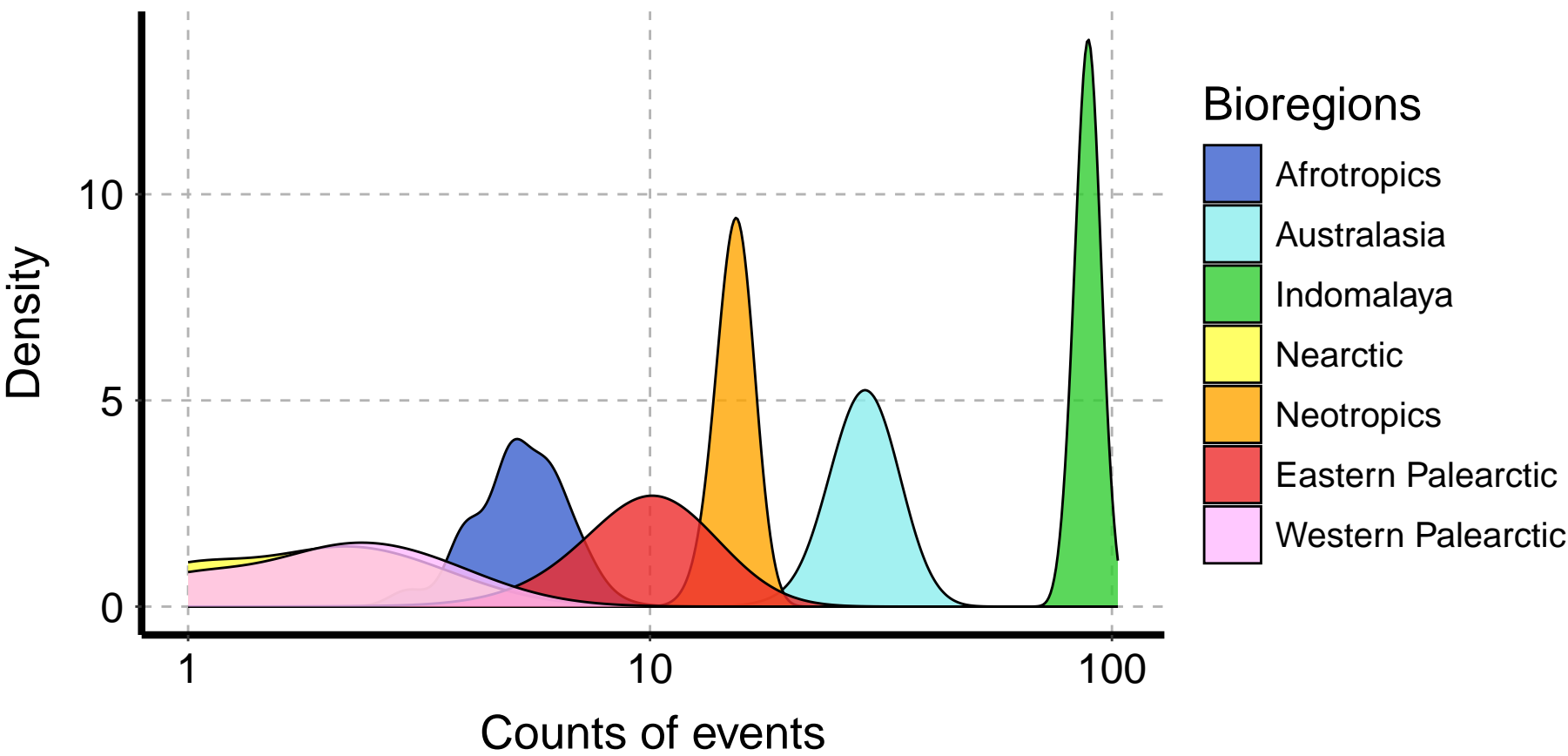
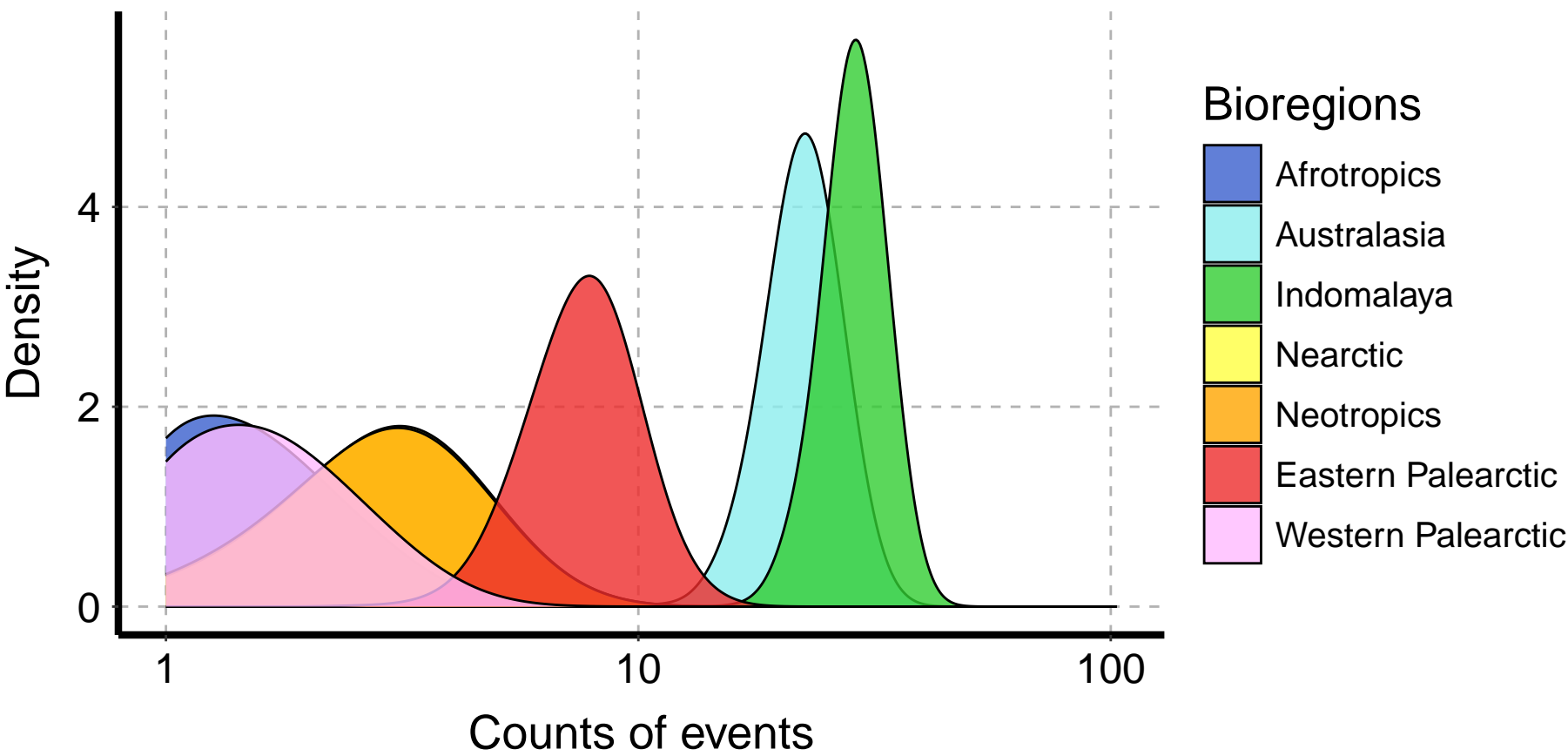


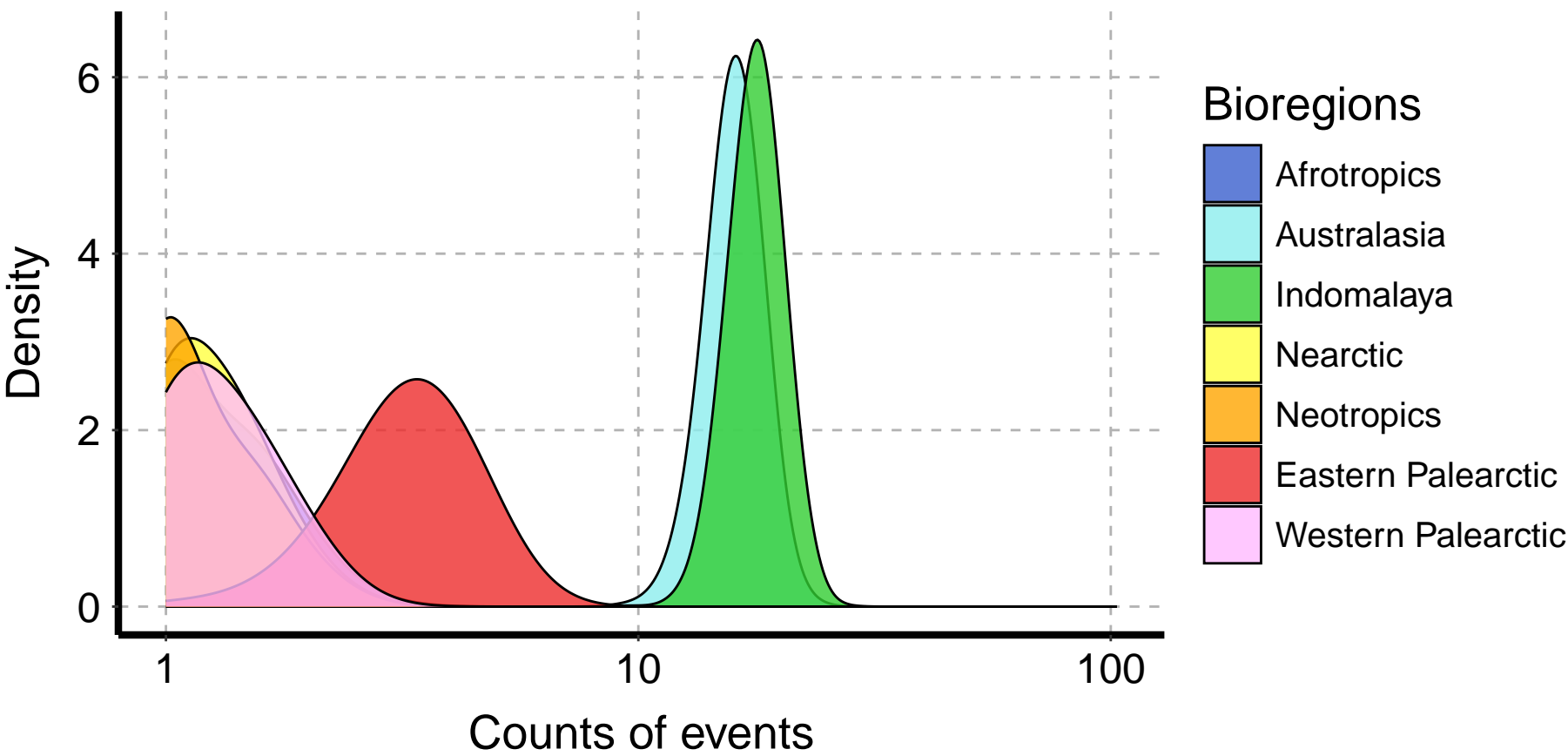
Counts of range extension (d) events  
per Source bioregion  
across 1000 BS maps



Counts of subset speciation (s) events  
per Source bioregion  
across 1000 BS maps



Counts of vicariance (v) events  
per Source bioregion  
across 1000 BS maps



Counts of jump-dispersal (j) events  
per Source bioregion  
across 1000 BS maps

