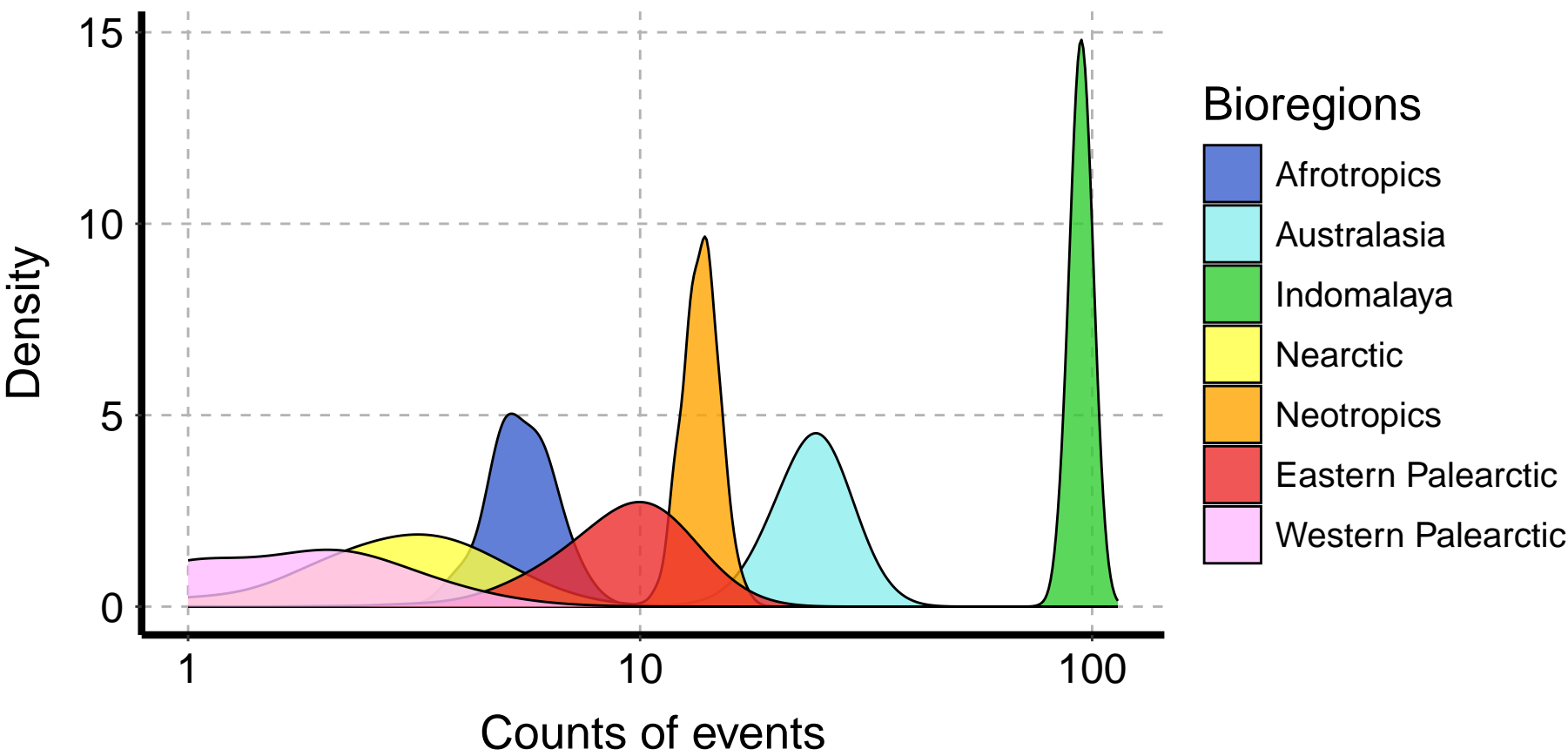
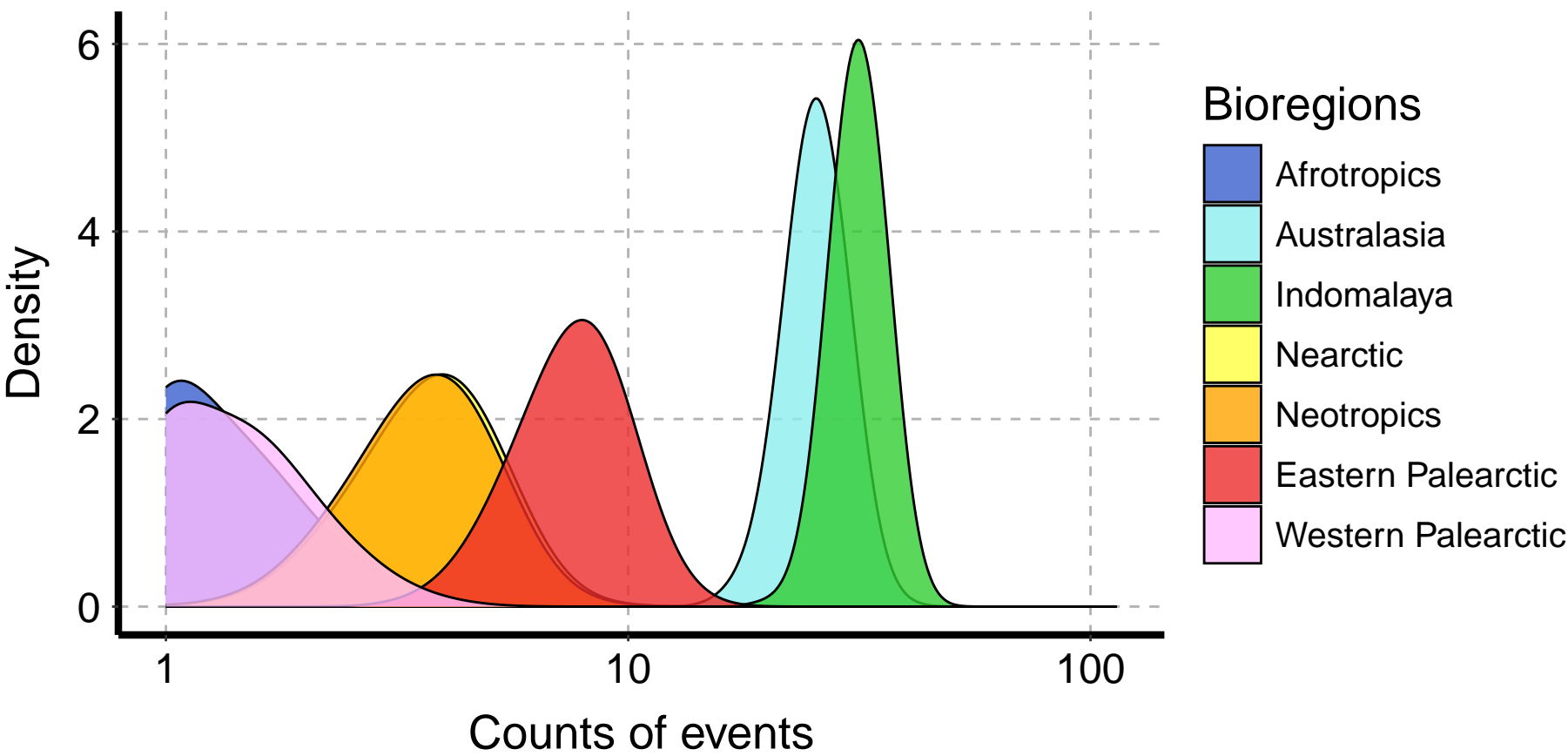


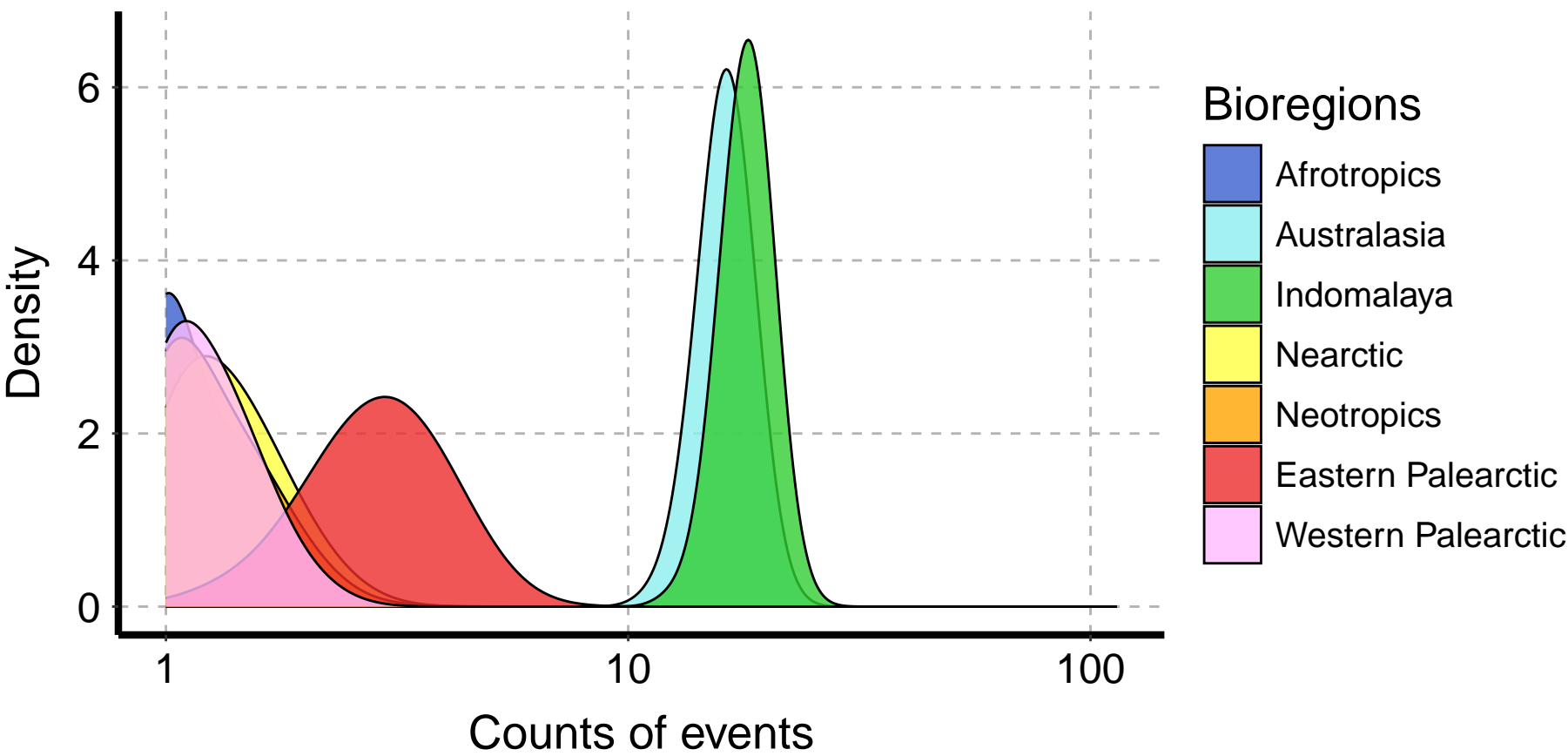
Counts of range extension (d) events
per Source bioregion
across 1000 BS maps



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

