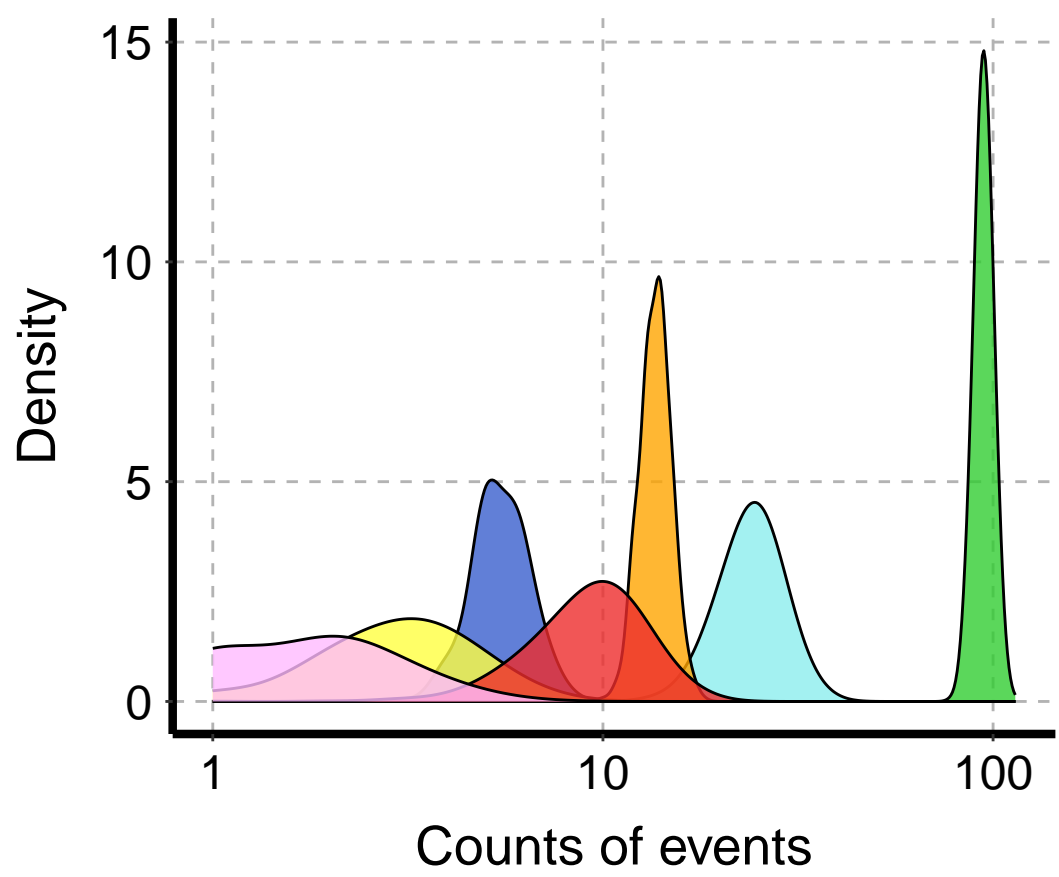
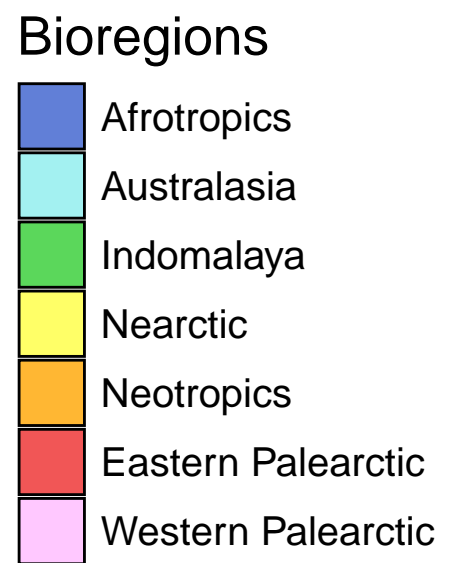
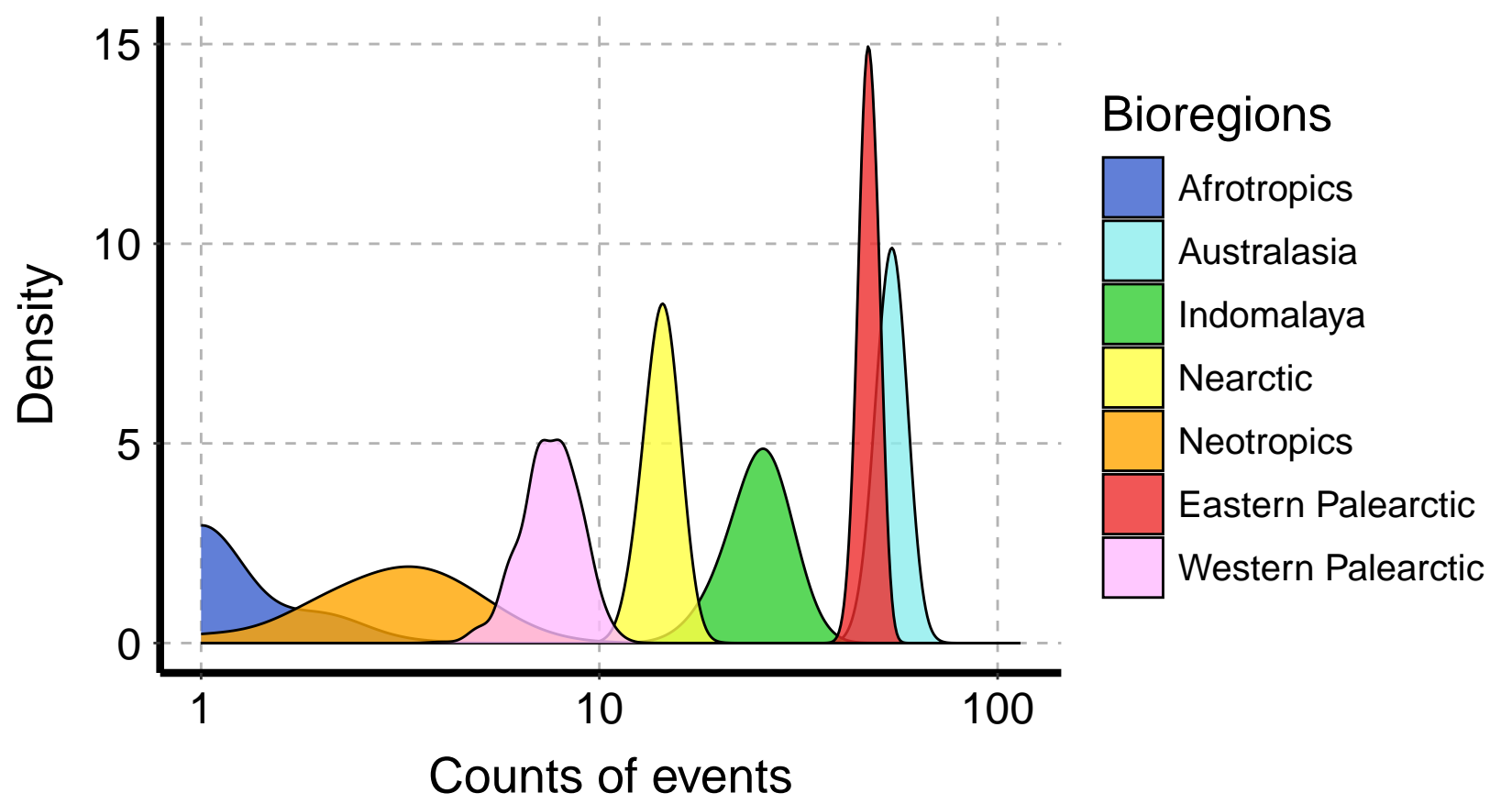


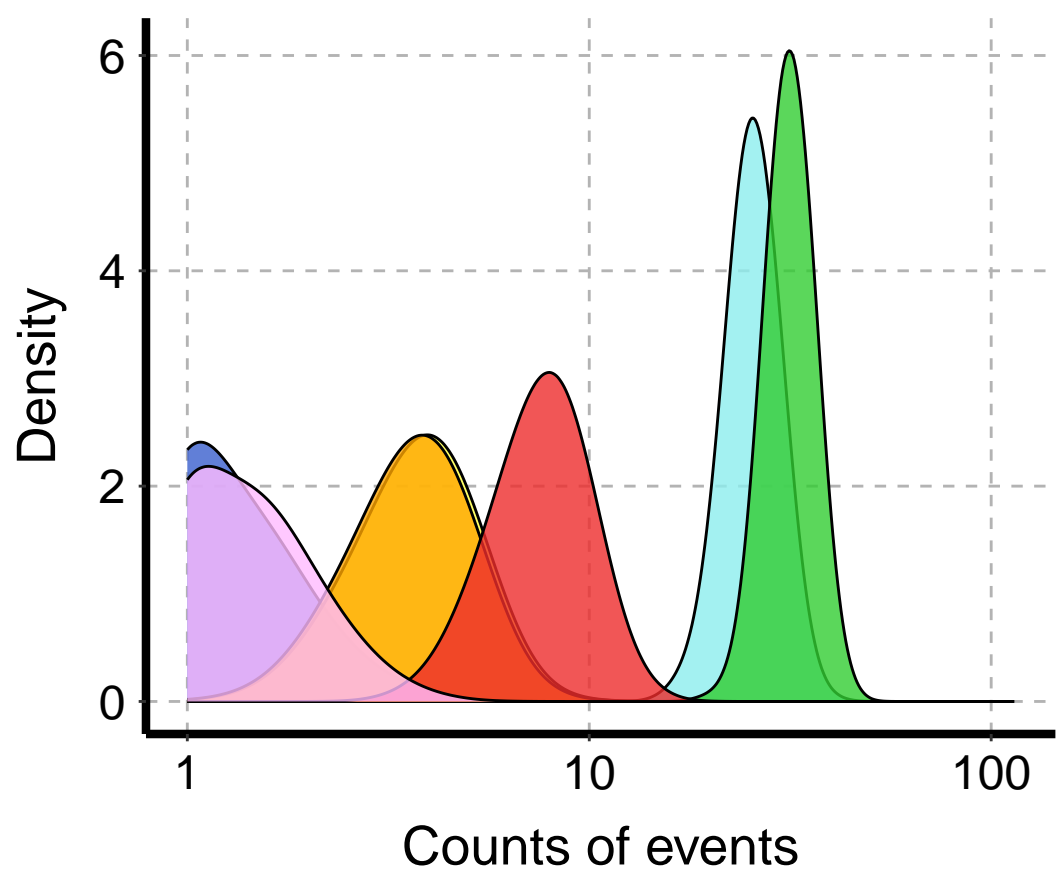
Counts of range extension (d) events
per Source bioregion
across 1000 BS maps



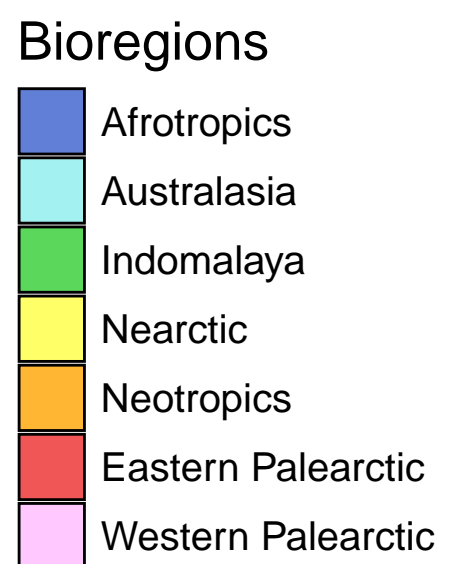
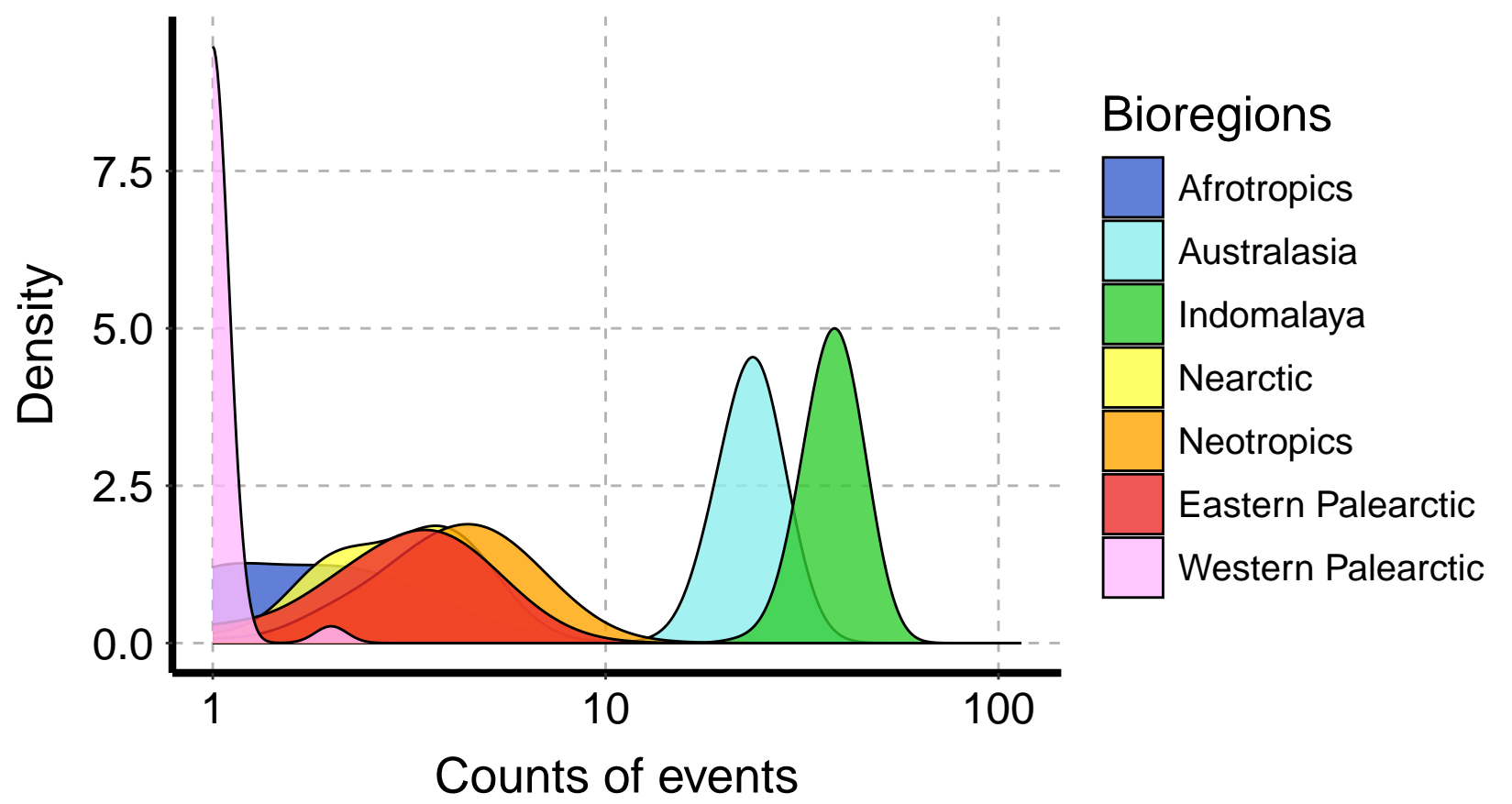
Counts of range extension (d) events
per Destination bioregion
across 1000 BS maps



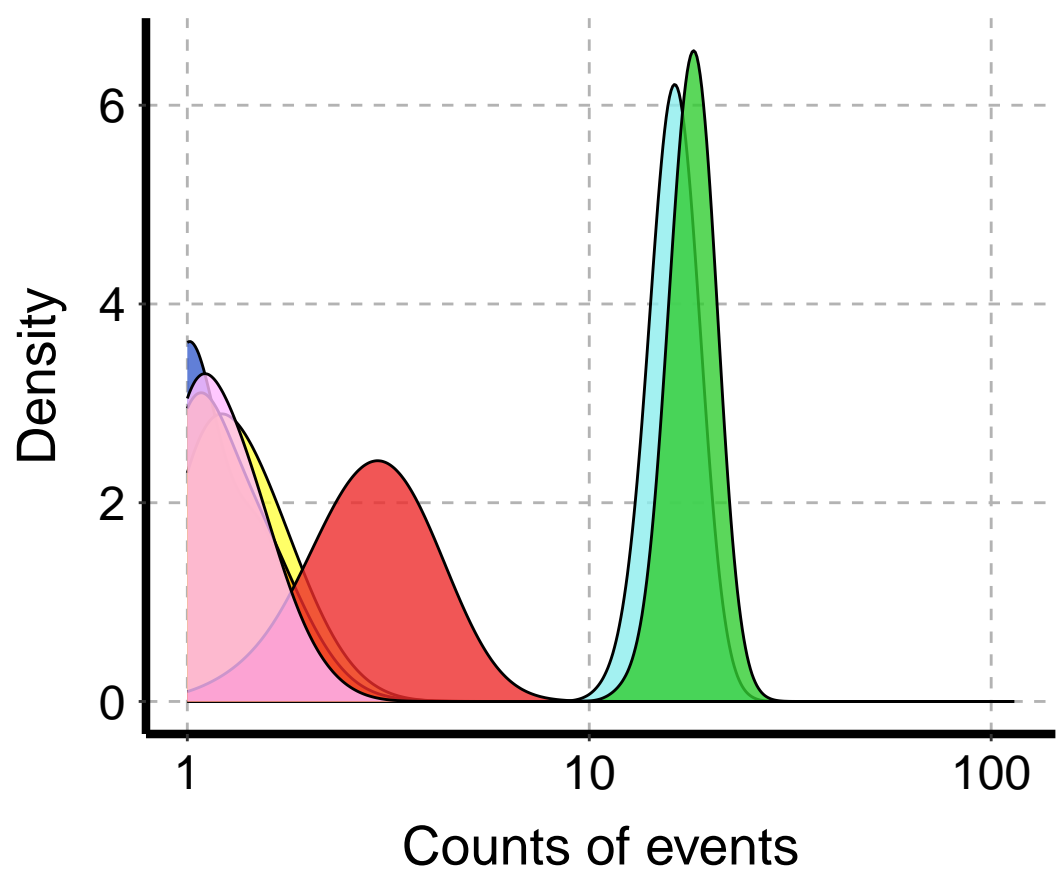
Counts of subset speciation (s) events
per Source bioregion
across 1000 BS maps



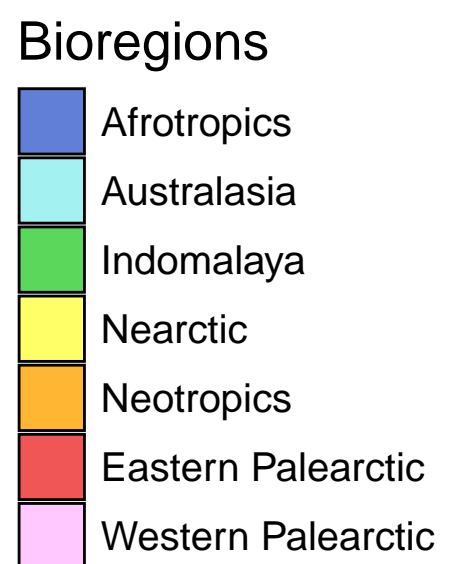
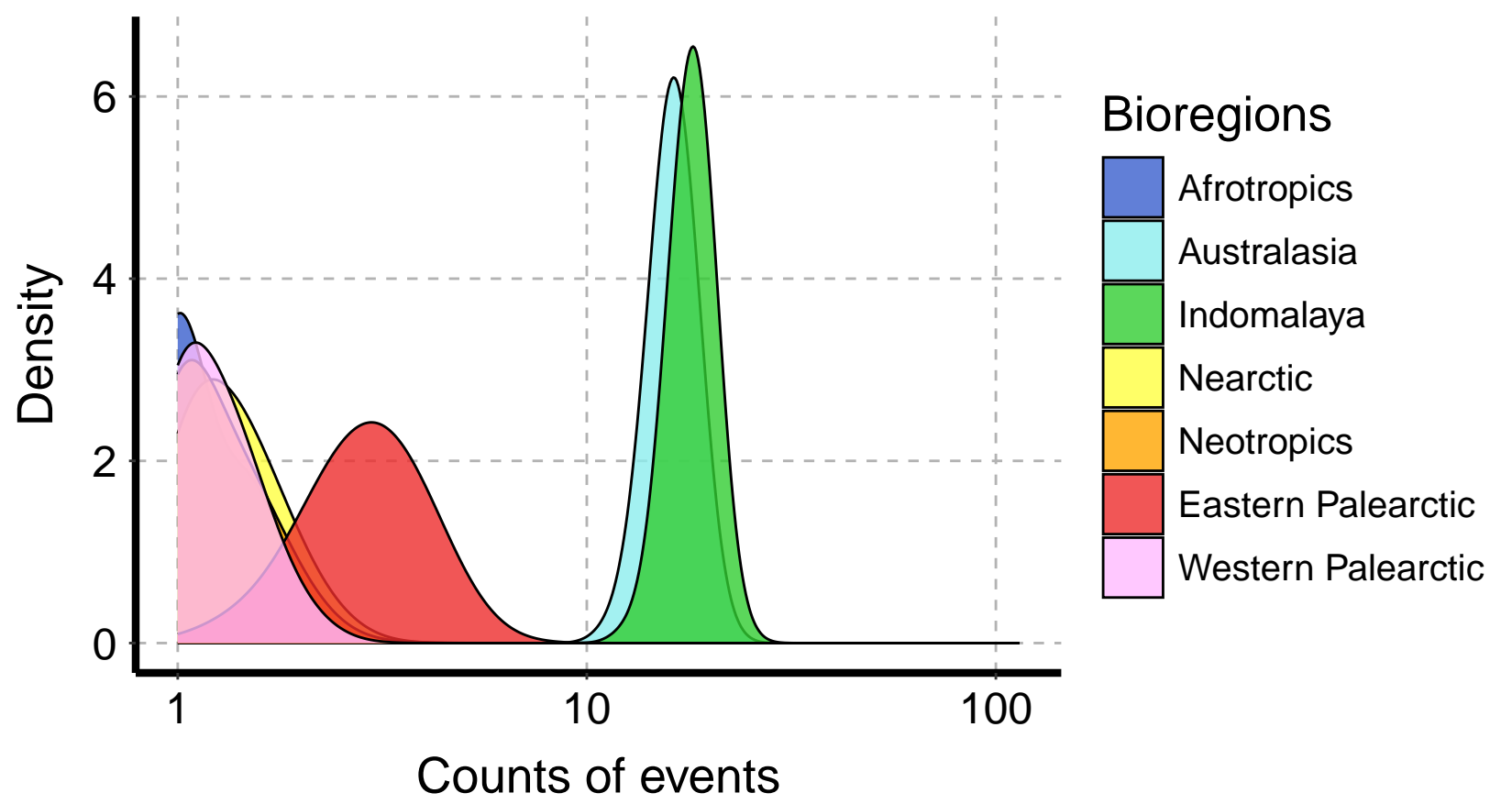
Counts of subset speciation (s) events
per Destination bioregion
across 1000 BS maps



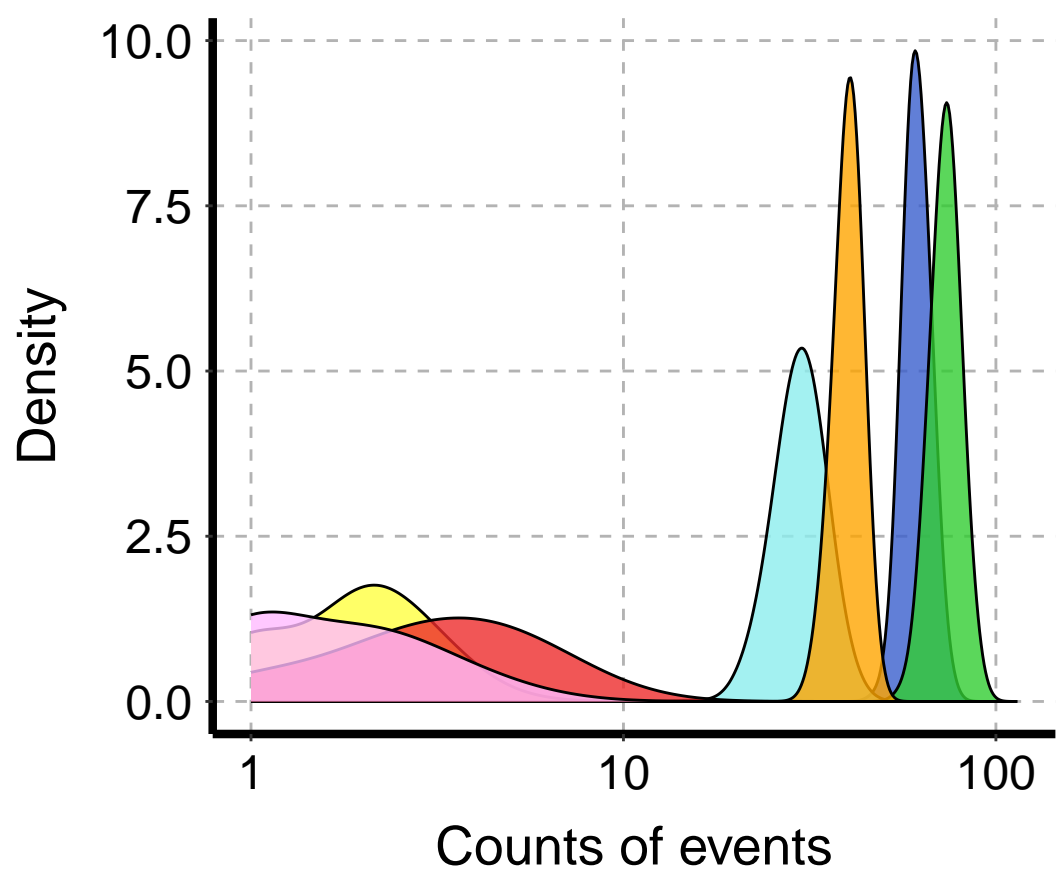
Counts of vicariance (v) events
per Source bioregion
across 1000 BS maps



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



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



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

