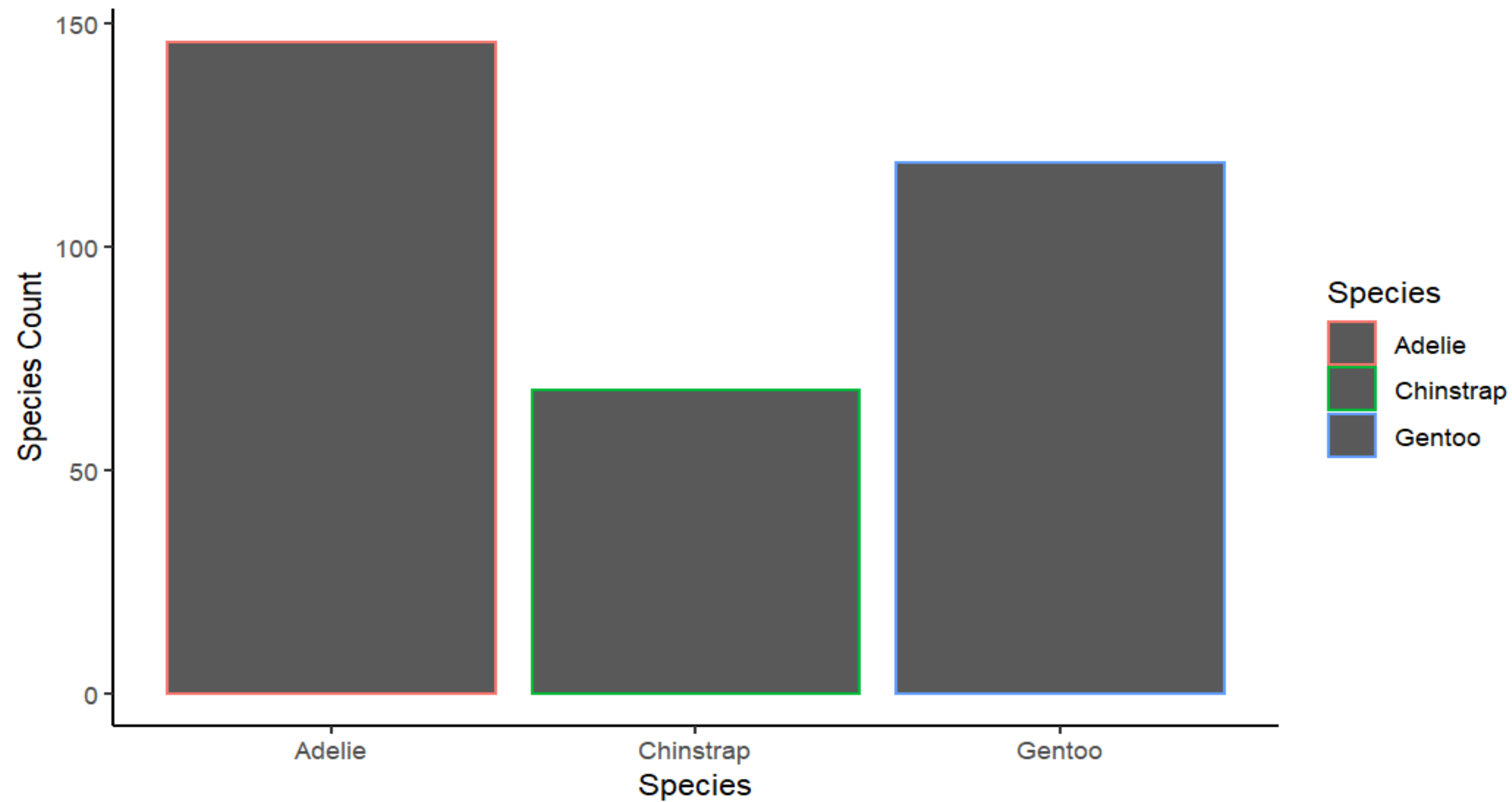
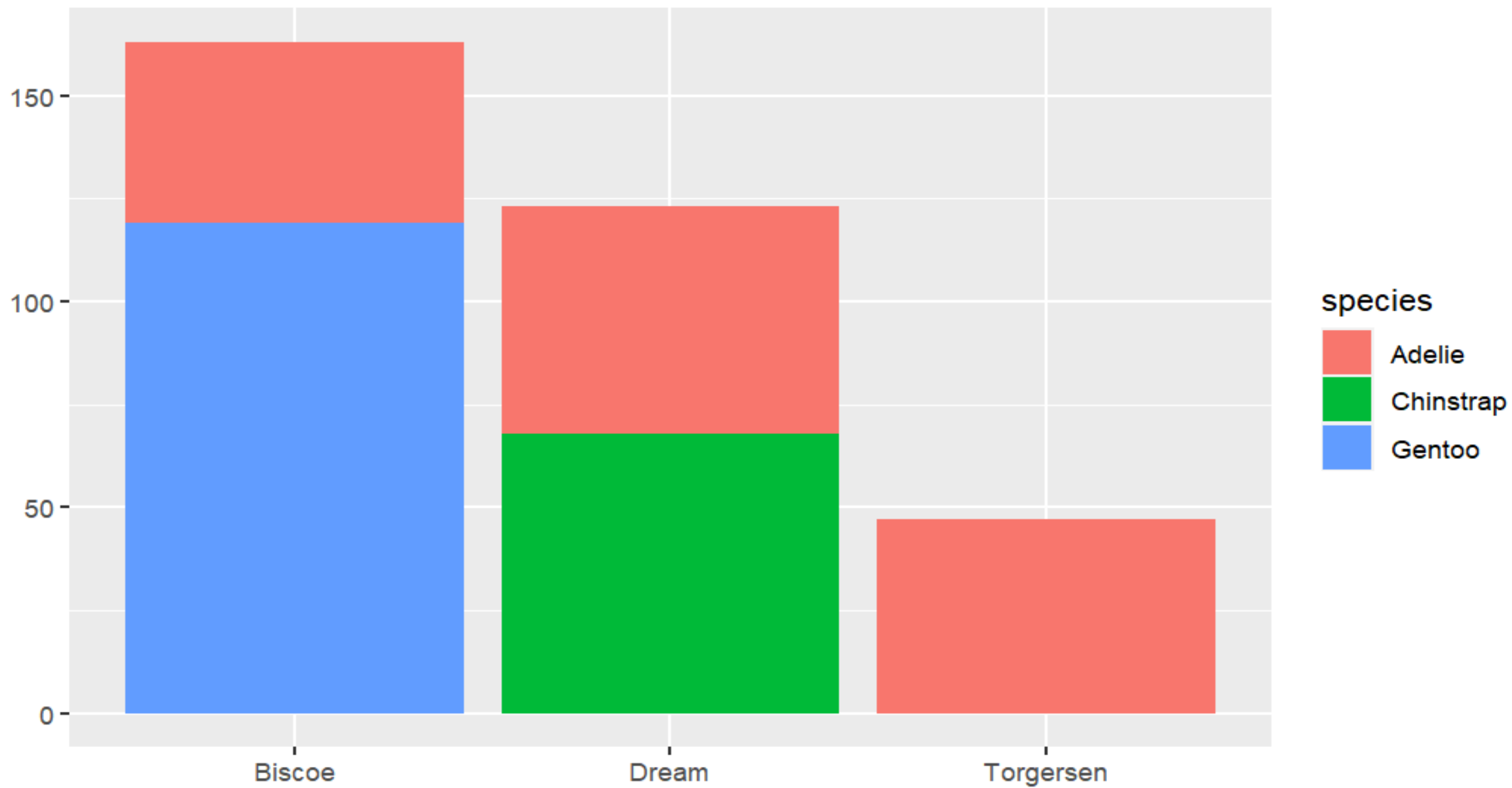


Recorded Specimen Count

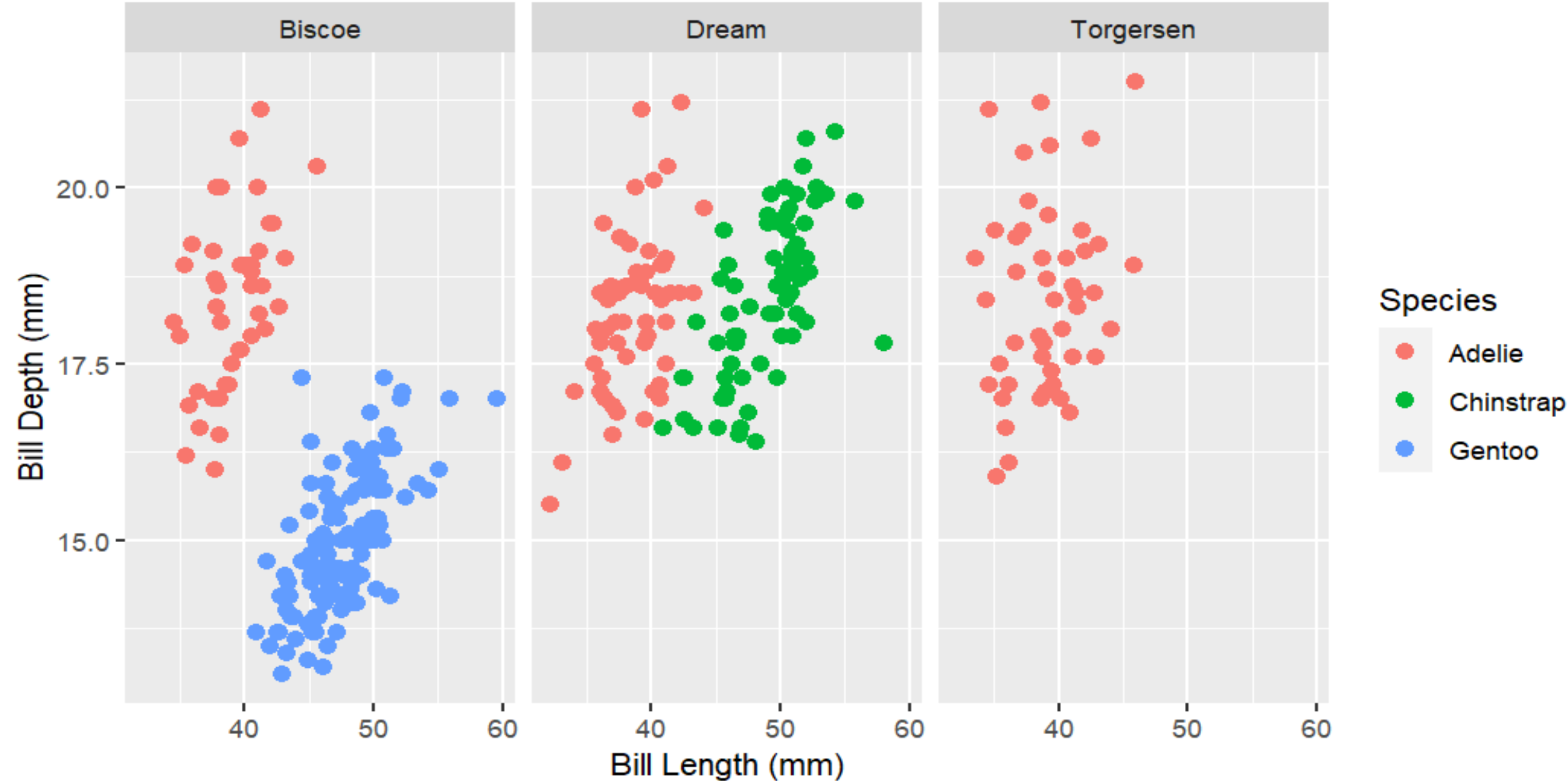


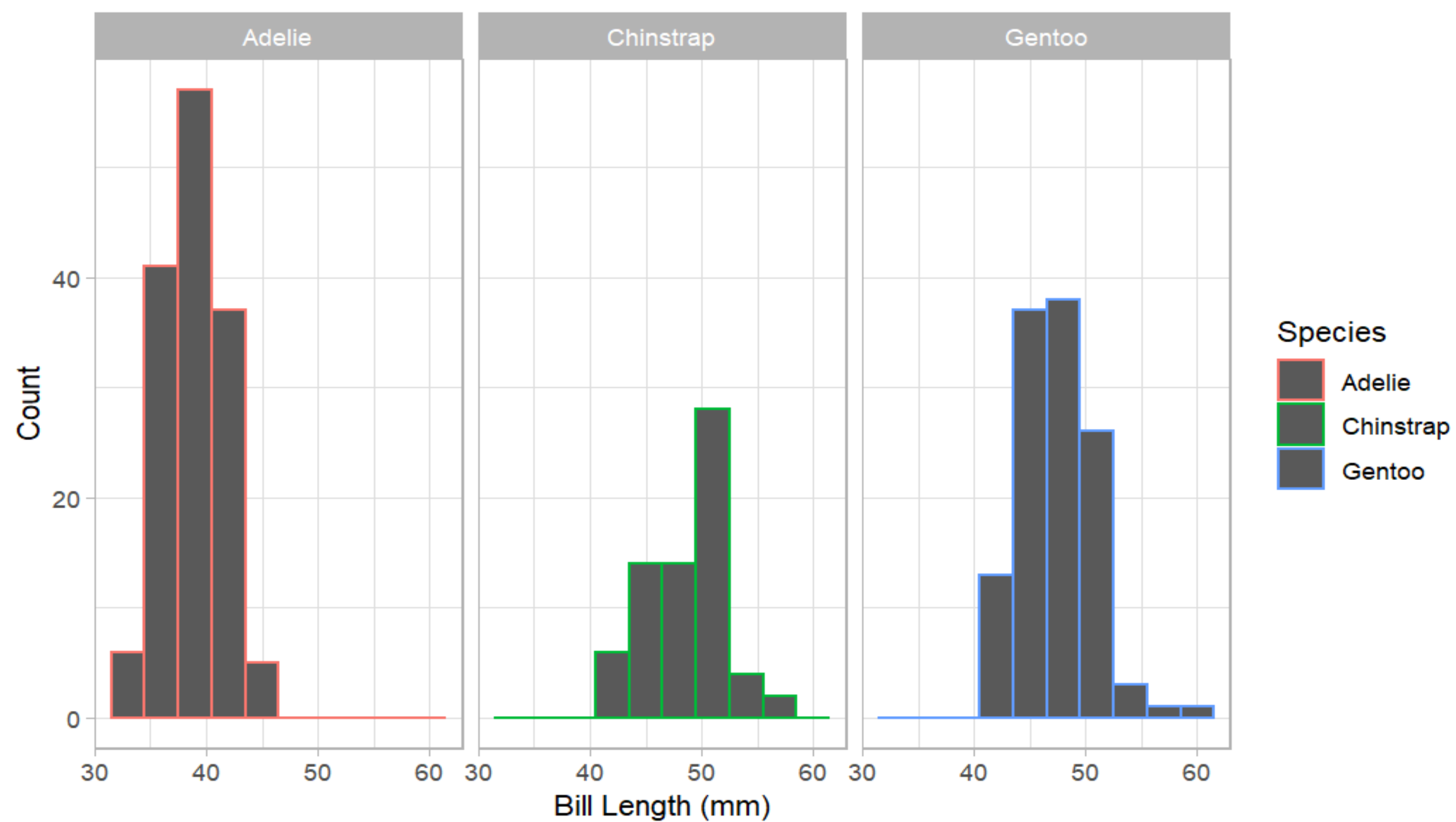
Recorded Species Distribution

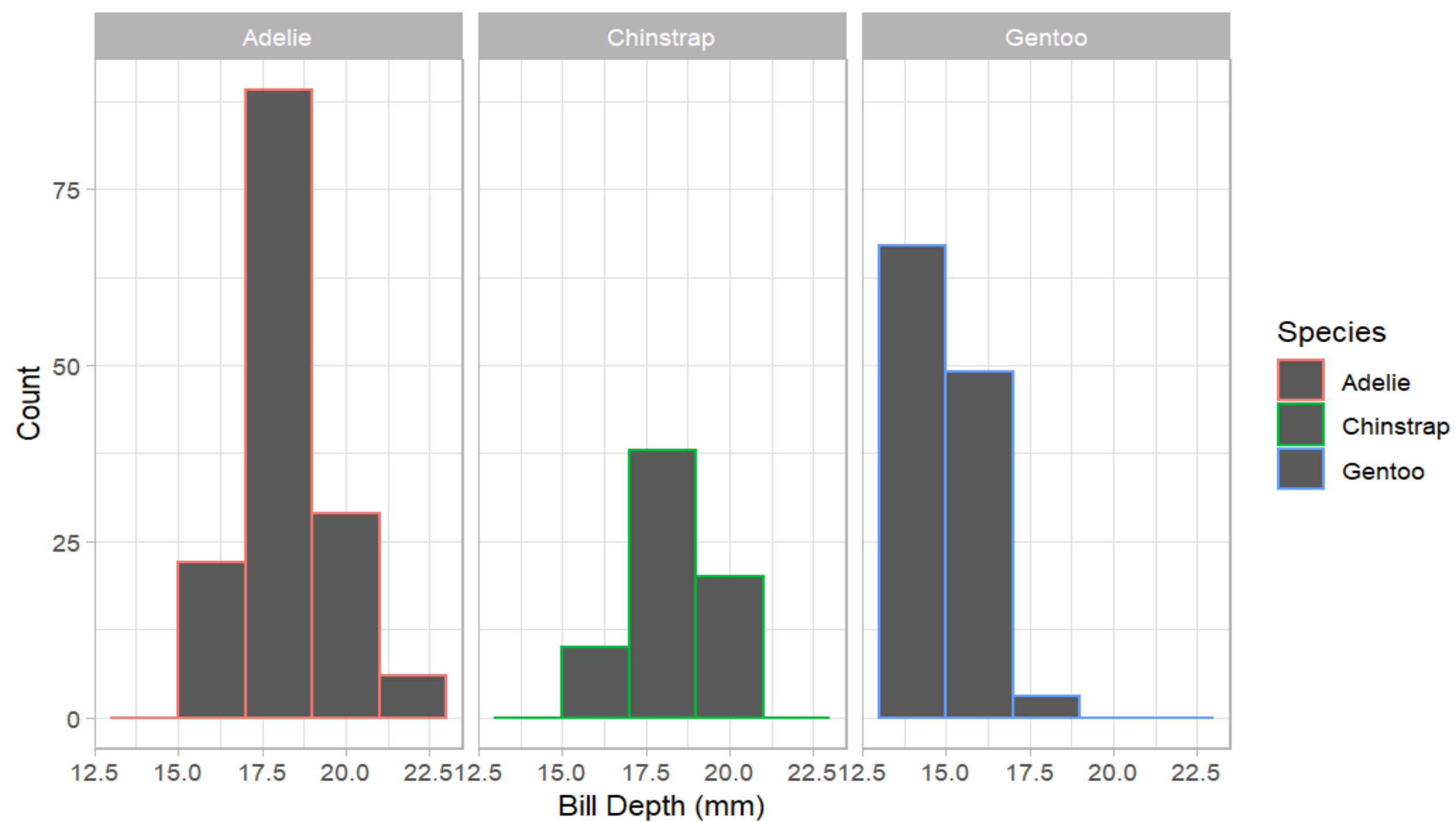


Bill Length and Depth Comparison

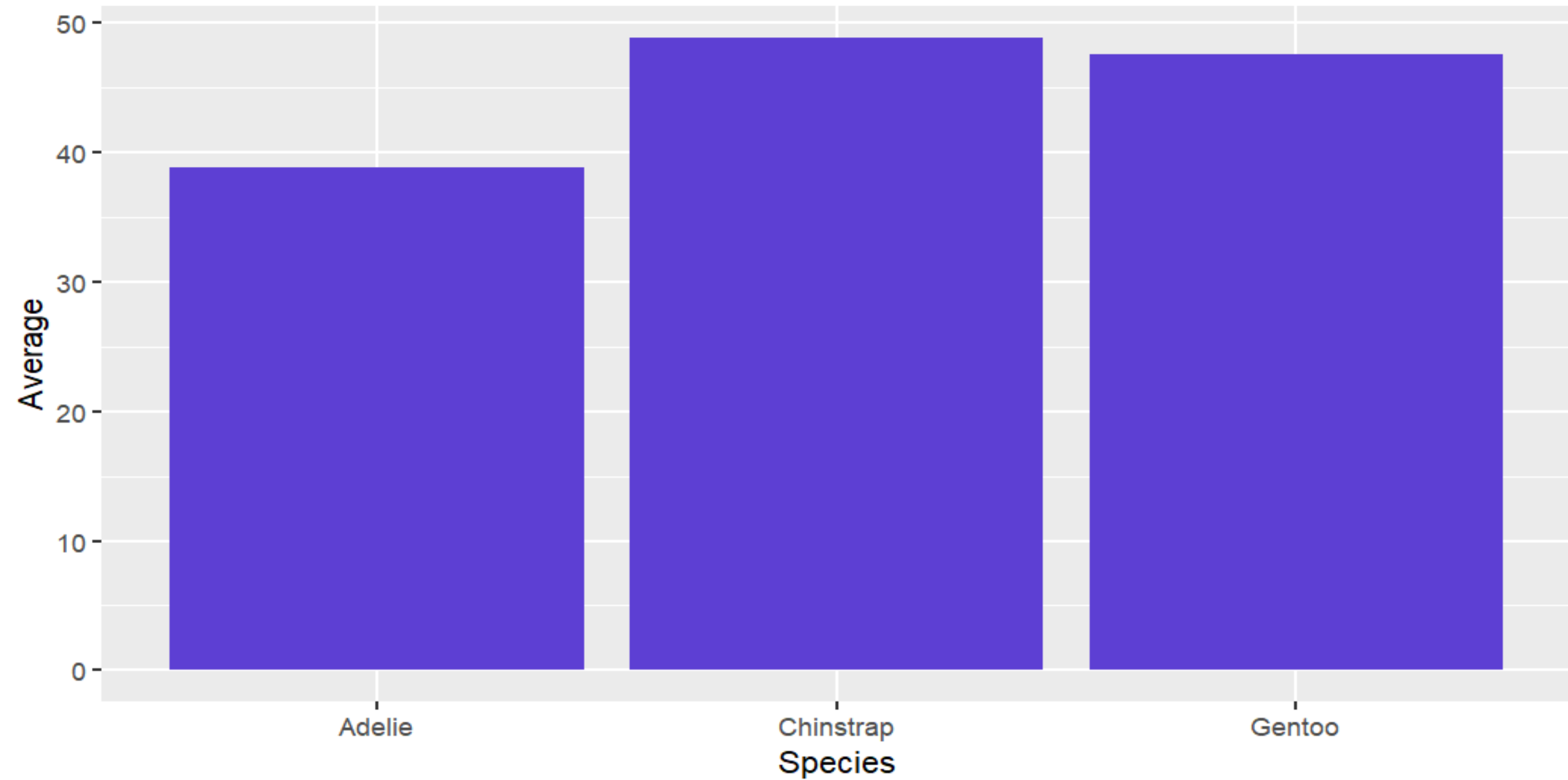
Comparison of Bill Length and Bill Depth of Each Recorded Specimen



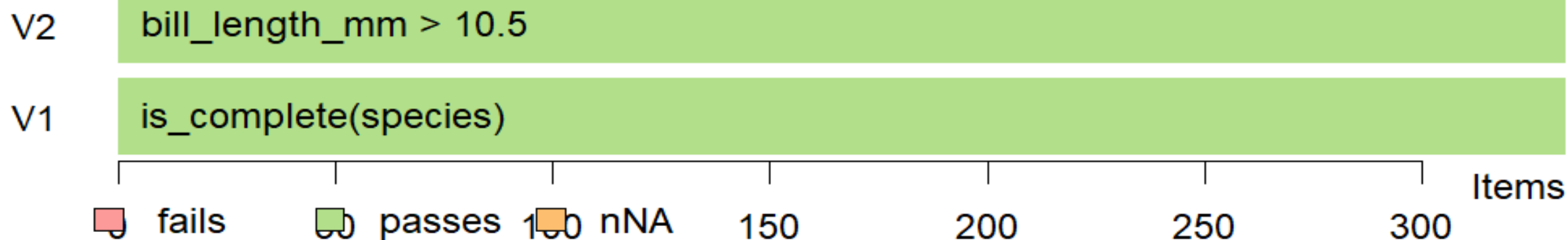




Average Bill Length
Of Each Observed Species



confront(dat = penguins, x = pTest)



confront(dat = penguins, x = pTest)

