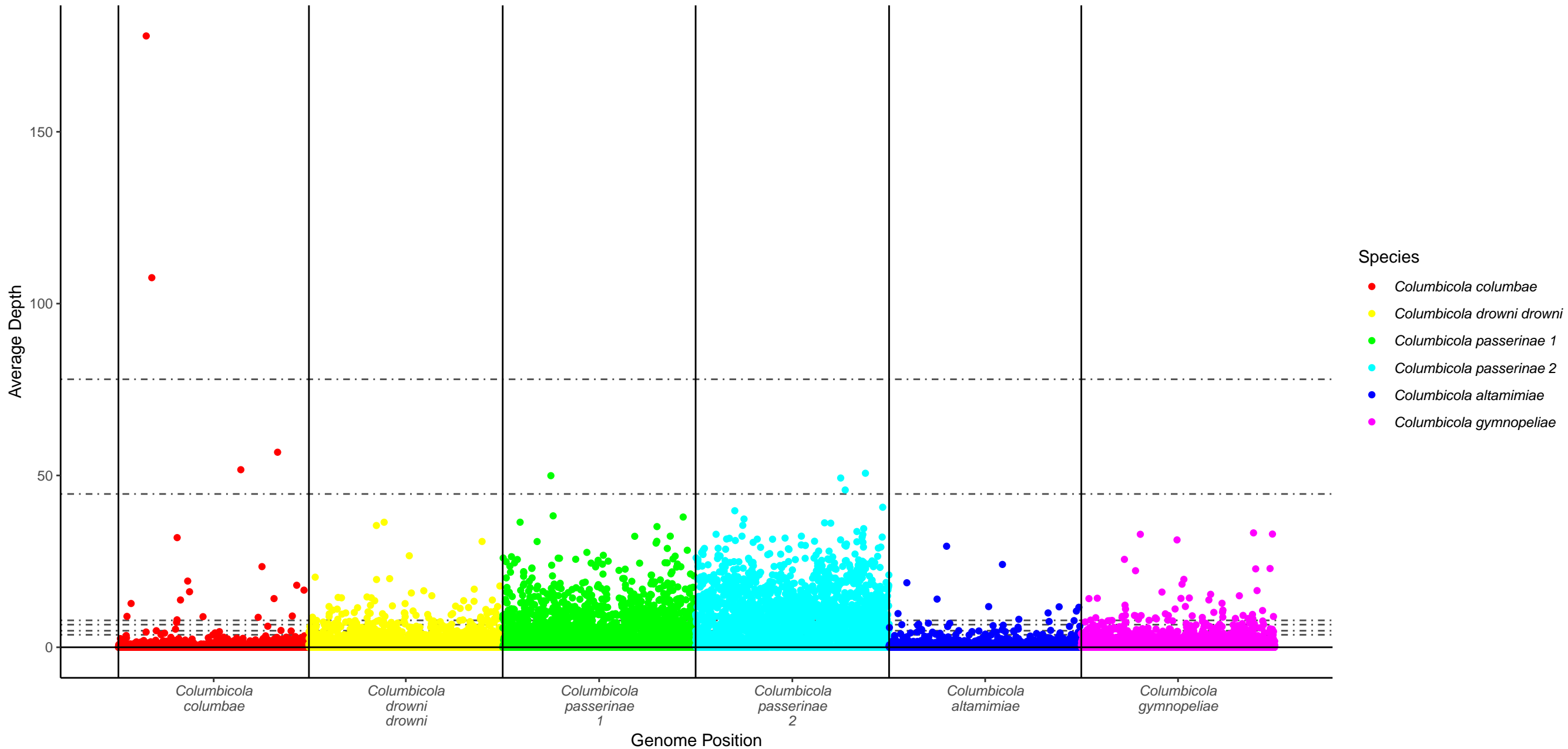
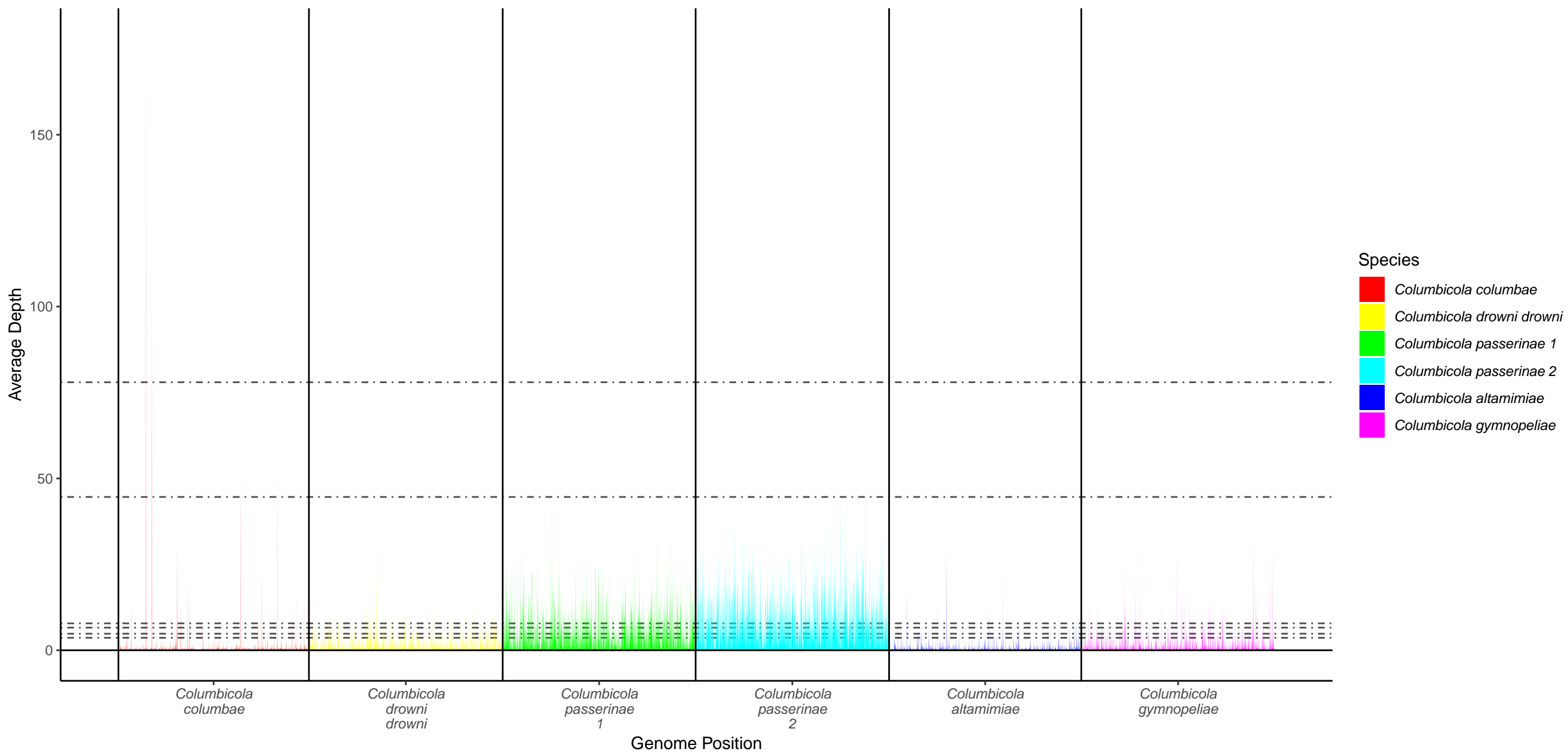


OUT Avg depth of coverage

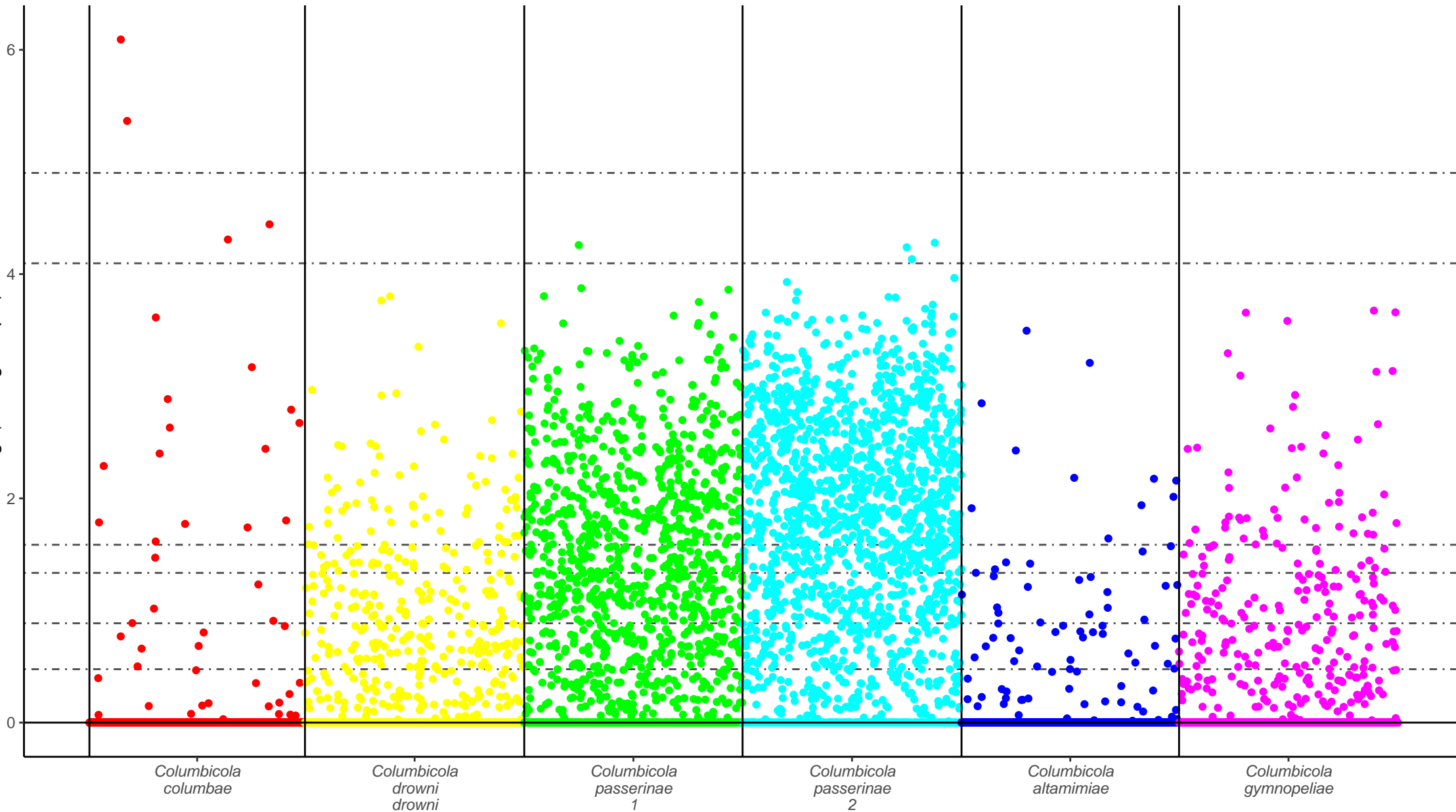


OUT Avg depth of coverage



OUT Log2(Average Depth)

log2(Average Depth)



Species

- *Columbicola columbae*
- *Columbicola drowni drowni*
- *Columbicola passerinae 1*
- *Columbicola passerinae 2*
- *Columbicola altamimiae*
- *Columbicola gymnopeliae*

*Columbicola  
columbae*

*Columbicola  
drowni  
drowni*

*Columbicola  
passerinae  
1*

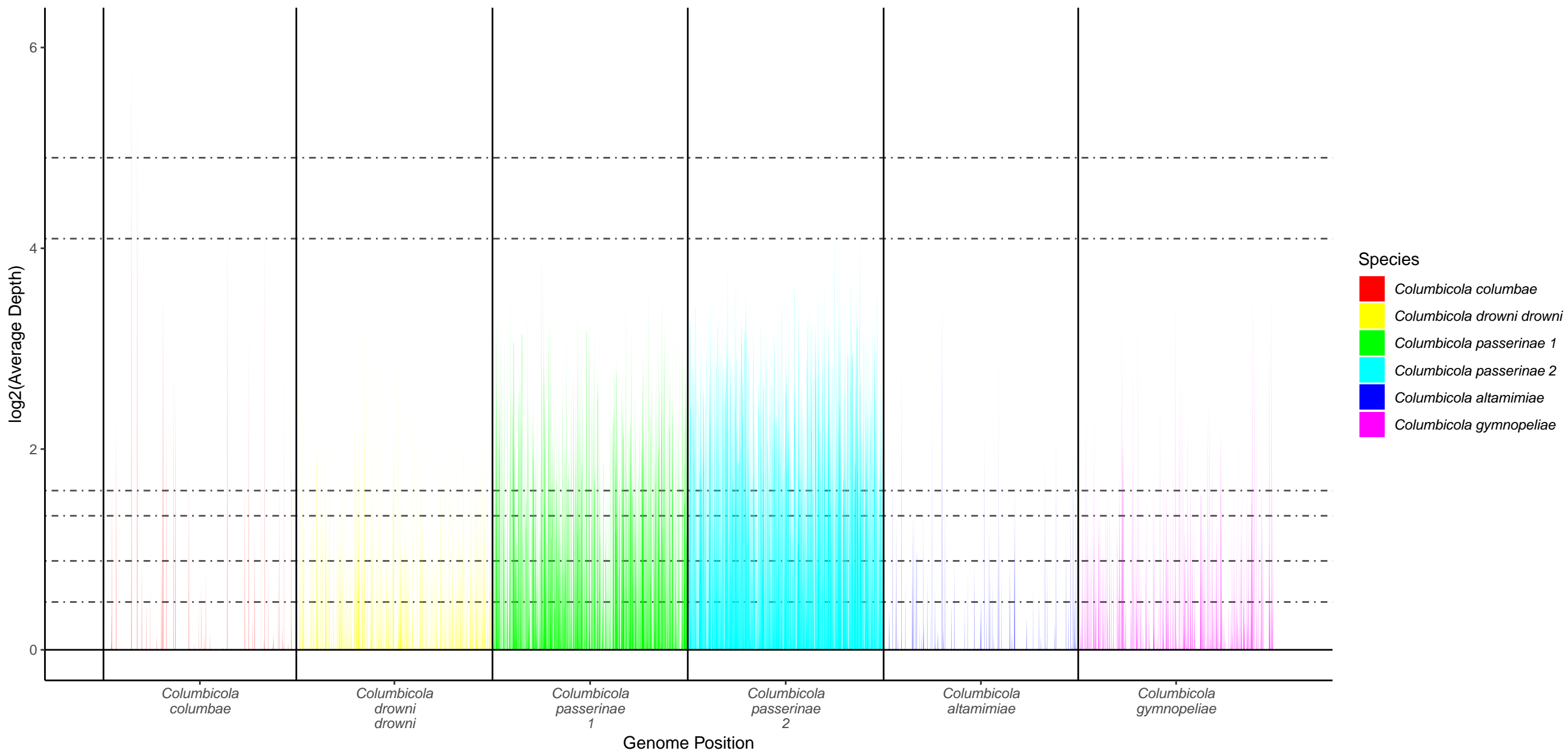
*Columbicola  
passerinae  
2*

*Columbicola  
altamimiae*

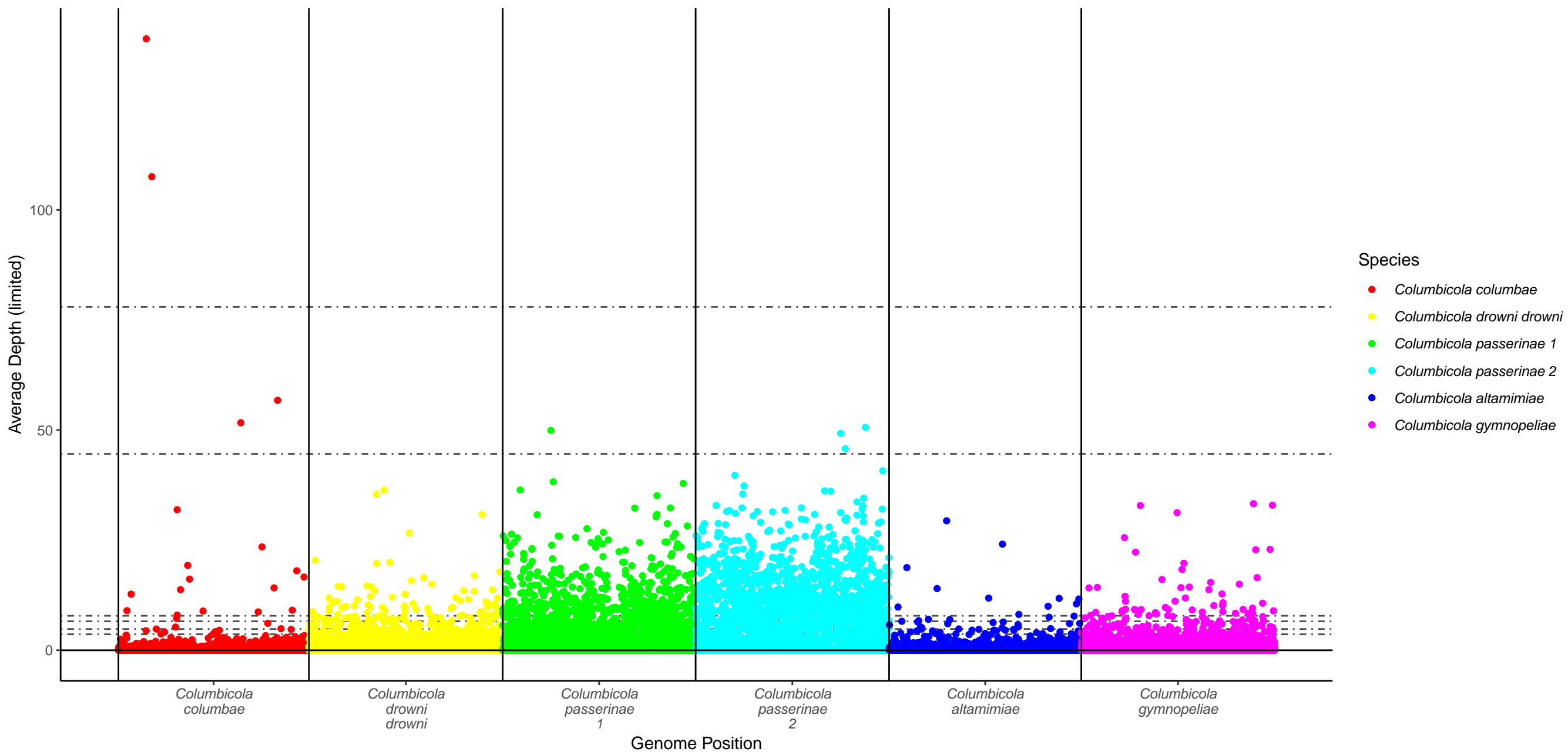
*Columbicola  
gymnopeliae*

Genome Position

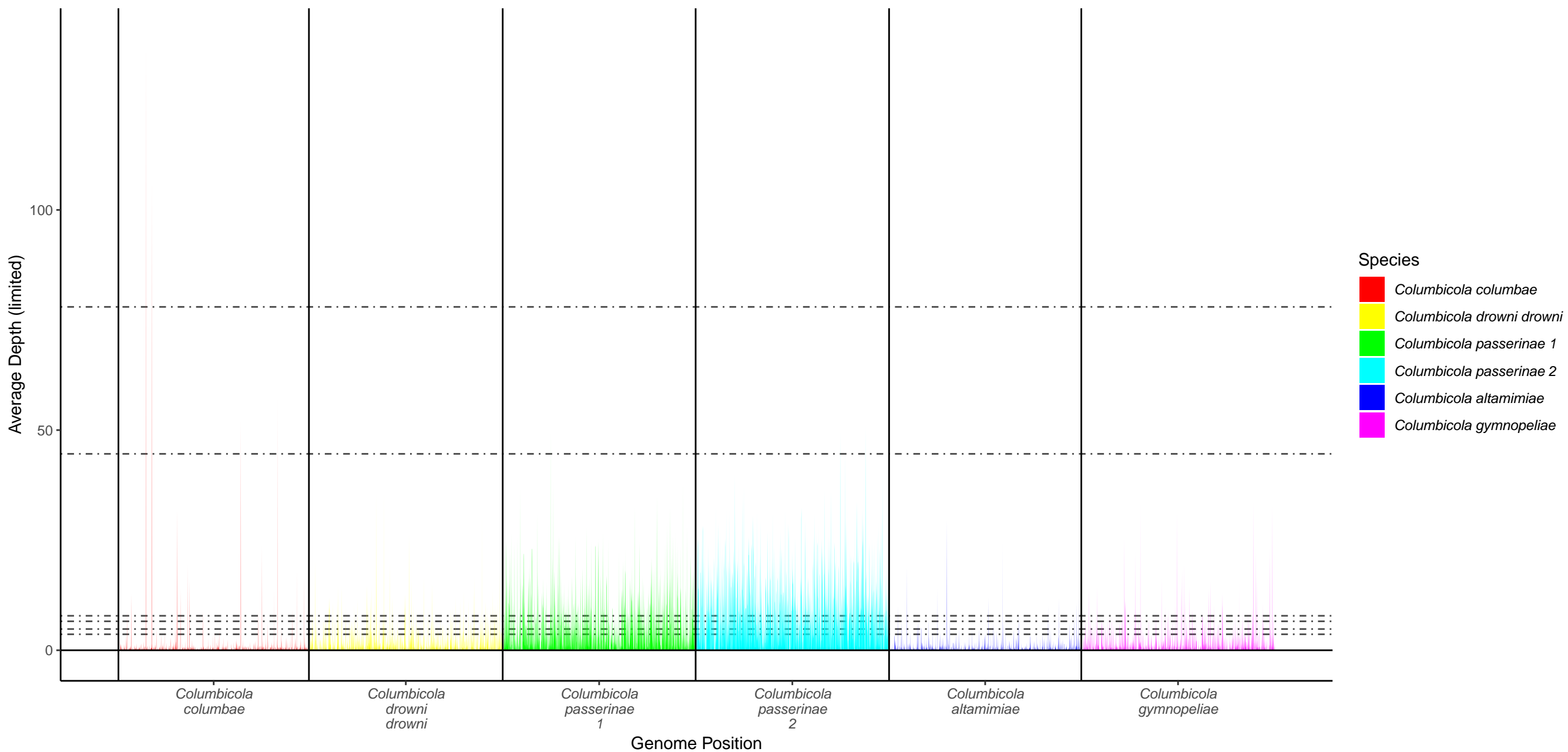
OUT Log2(Average Depth)



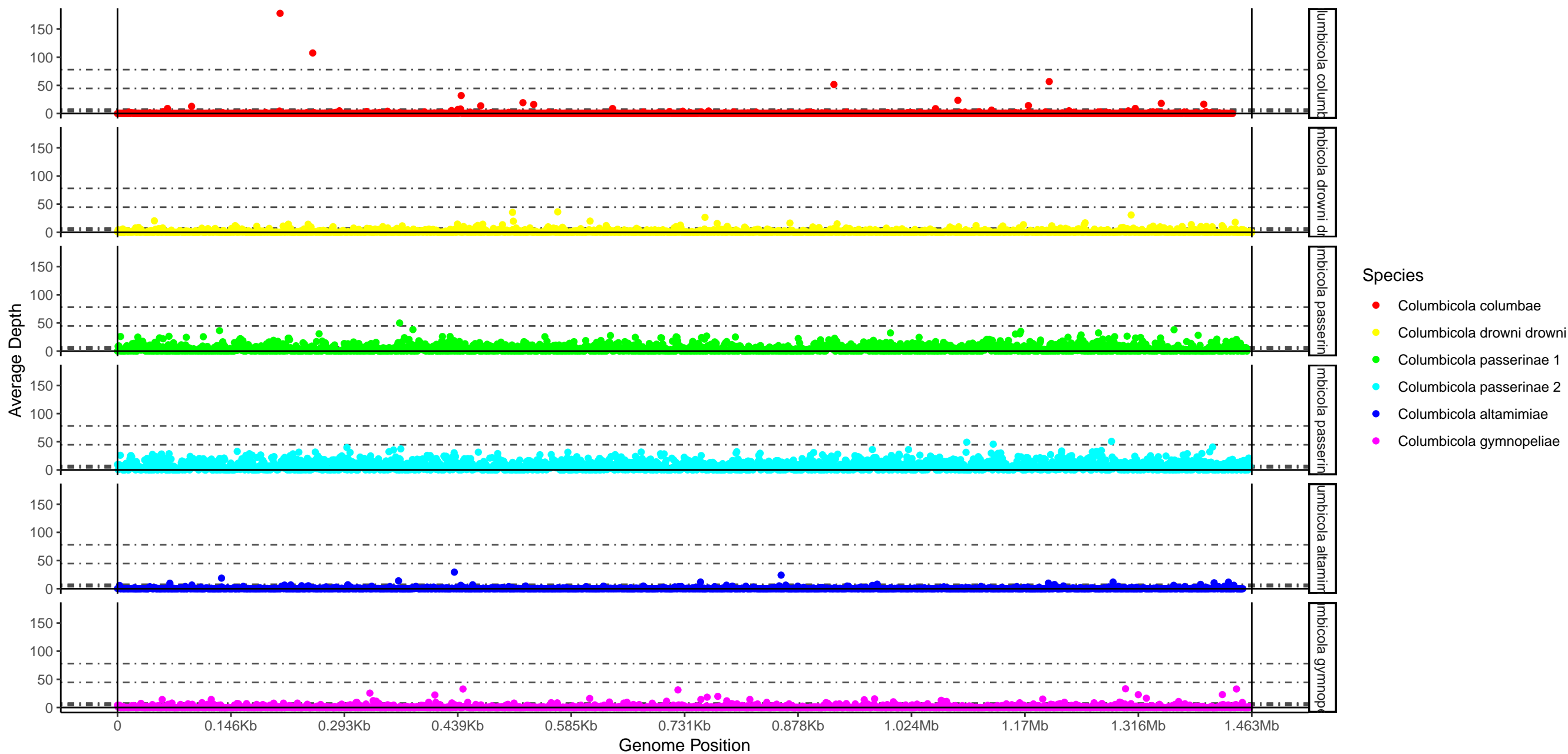
OUT Avg depth of coverage



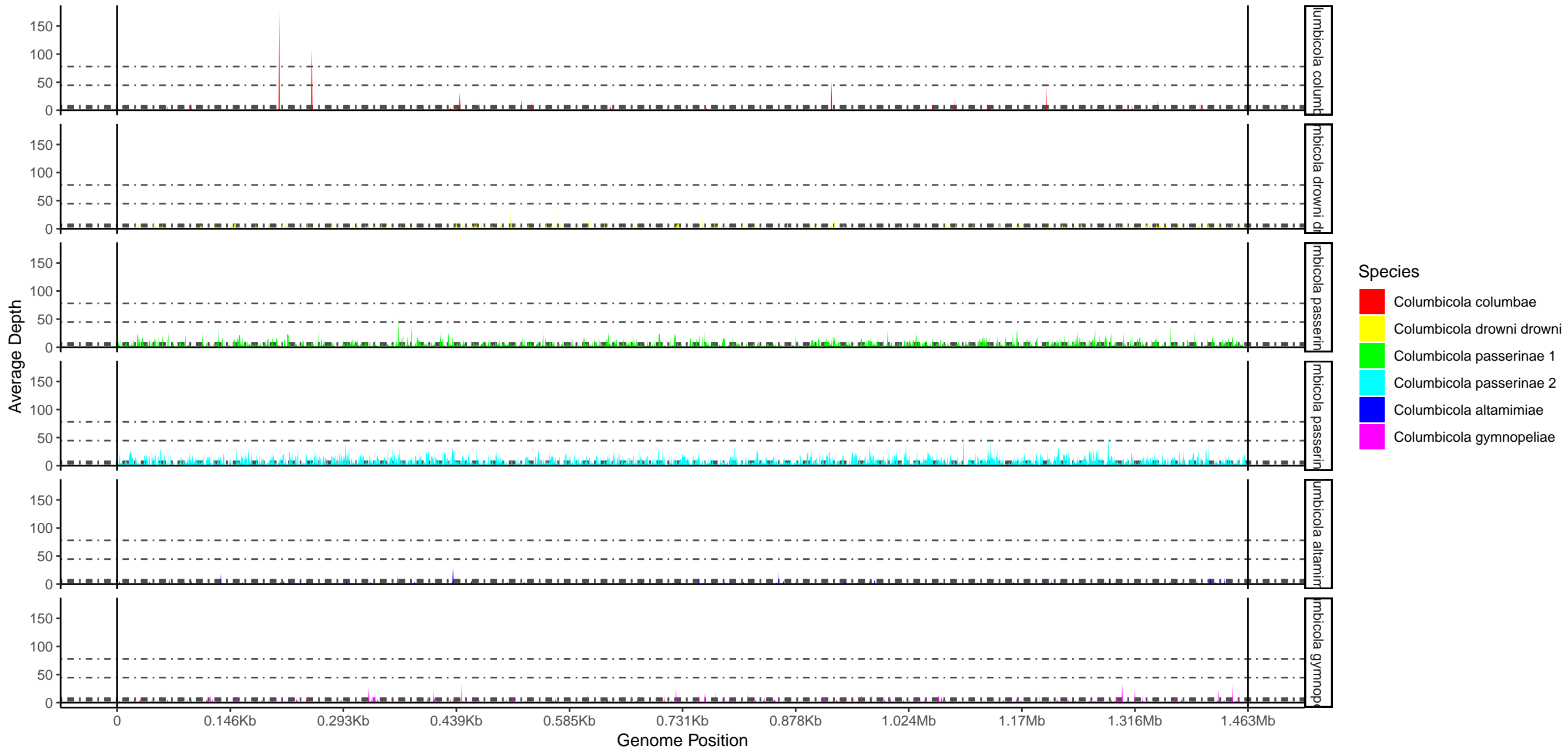
OUT Avg depth of coverage



OUT Avg depth of coverage

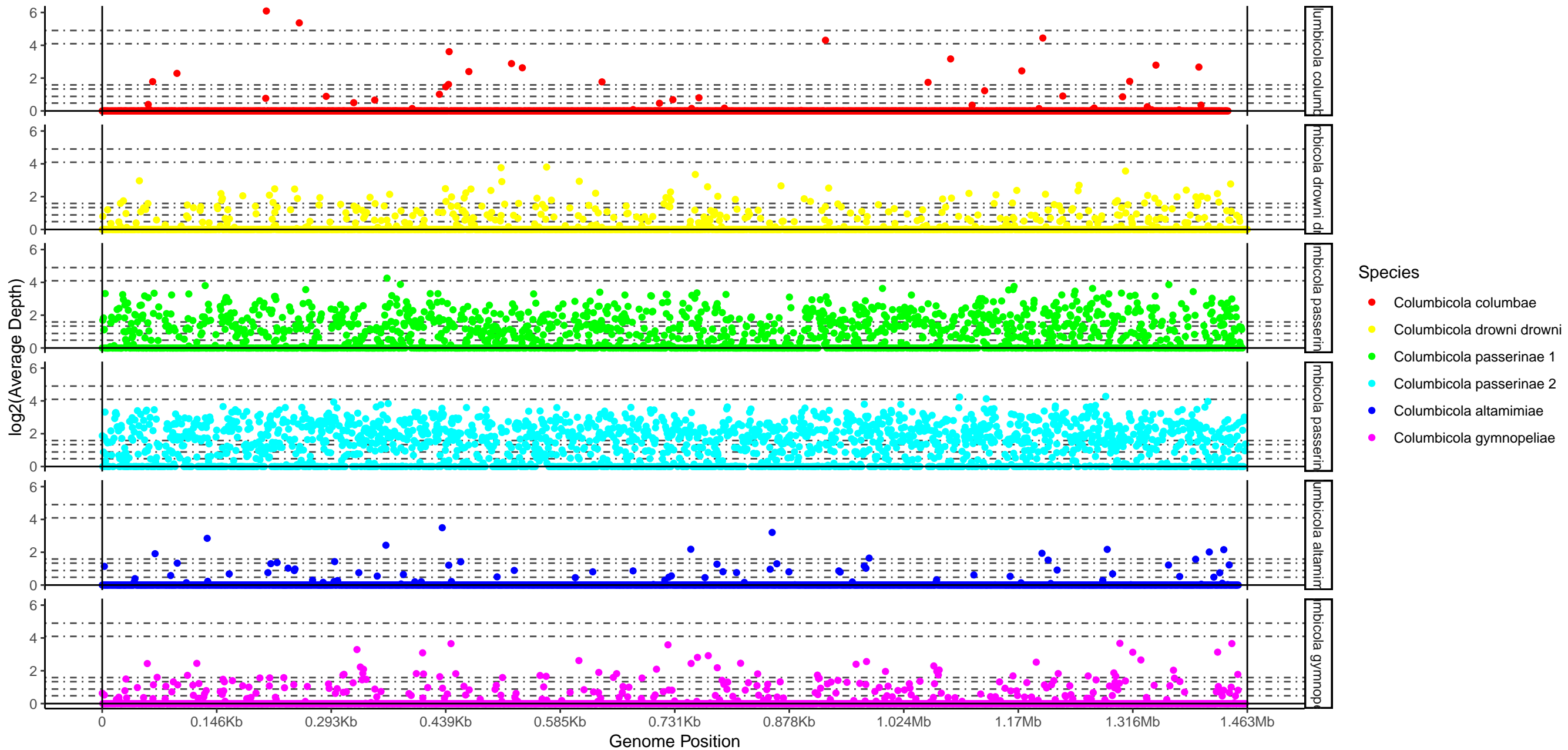


OUT Avg depth of coverage

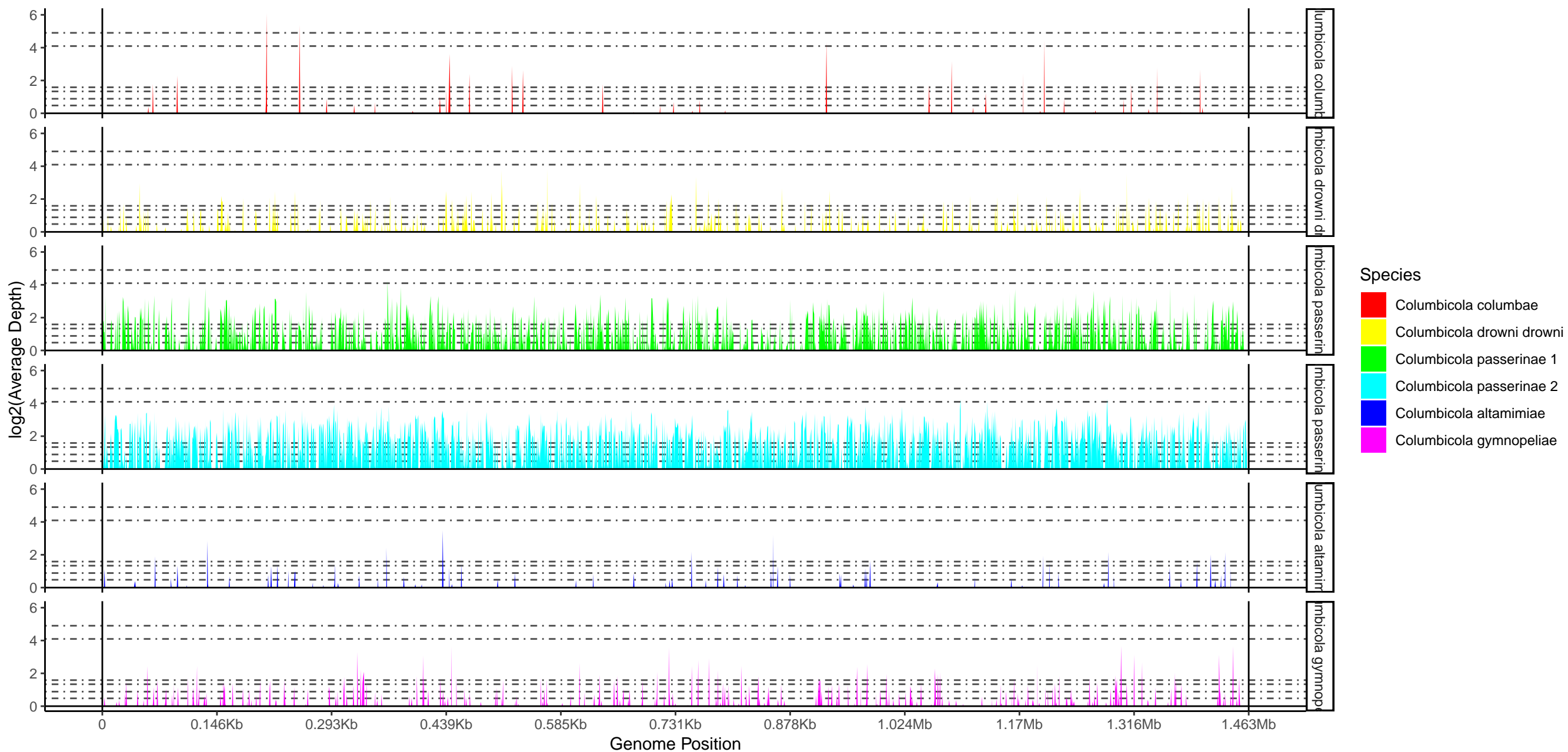




OUT Log2(Avg depth of coverage)



OUT Log2(Avg depth of coverage)



# OUT Avg depth of coverage

