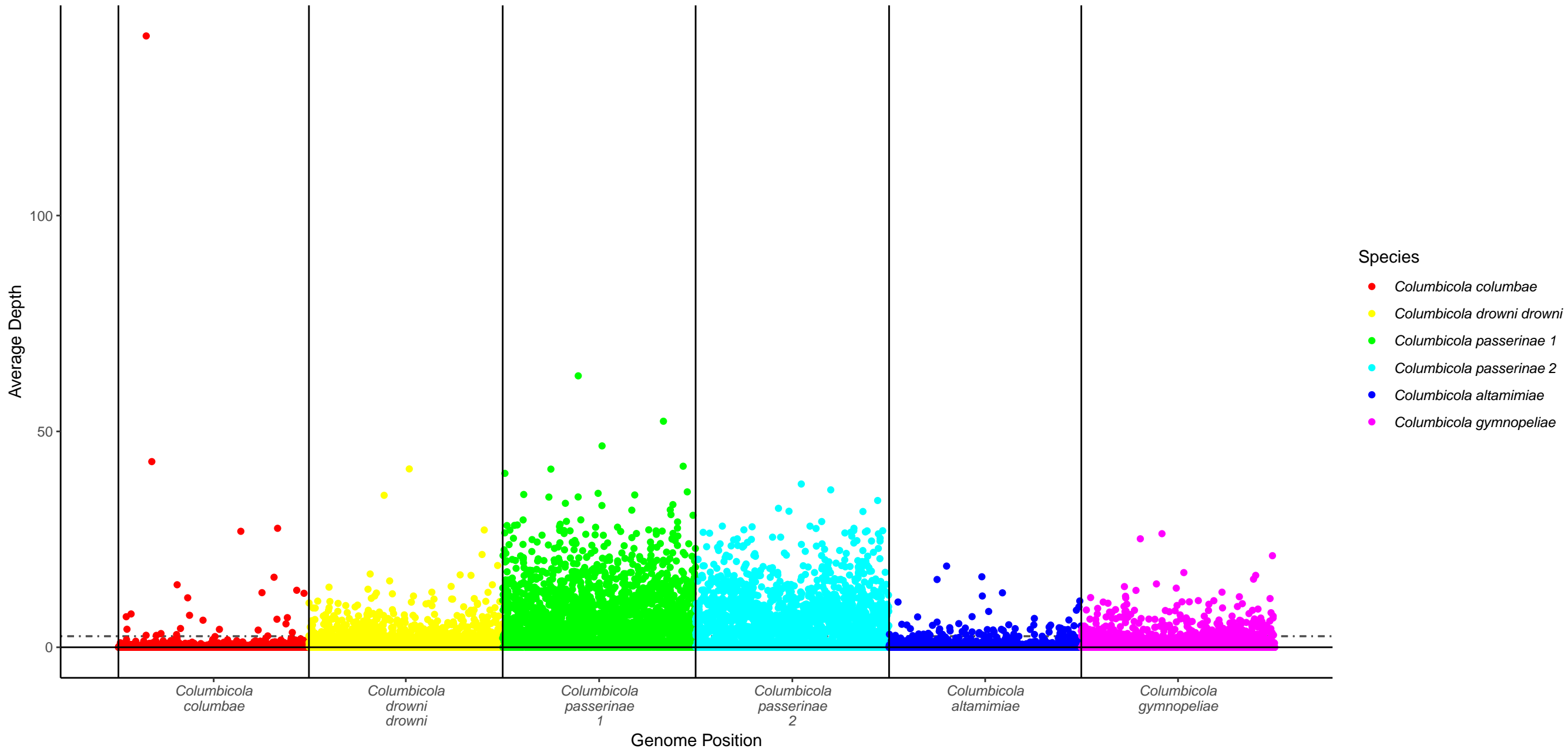
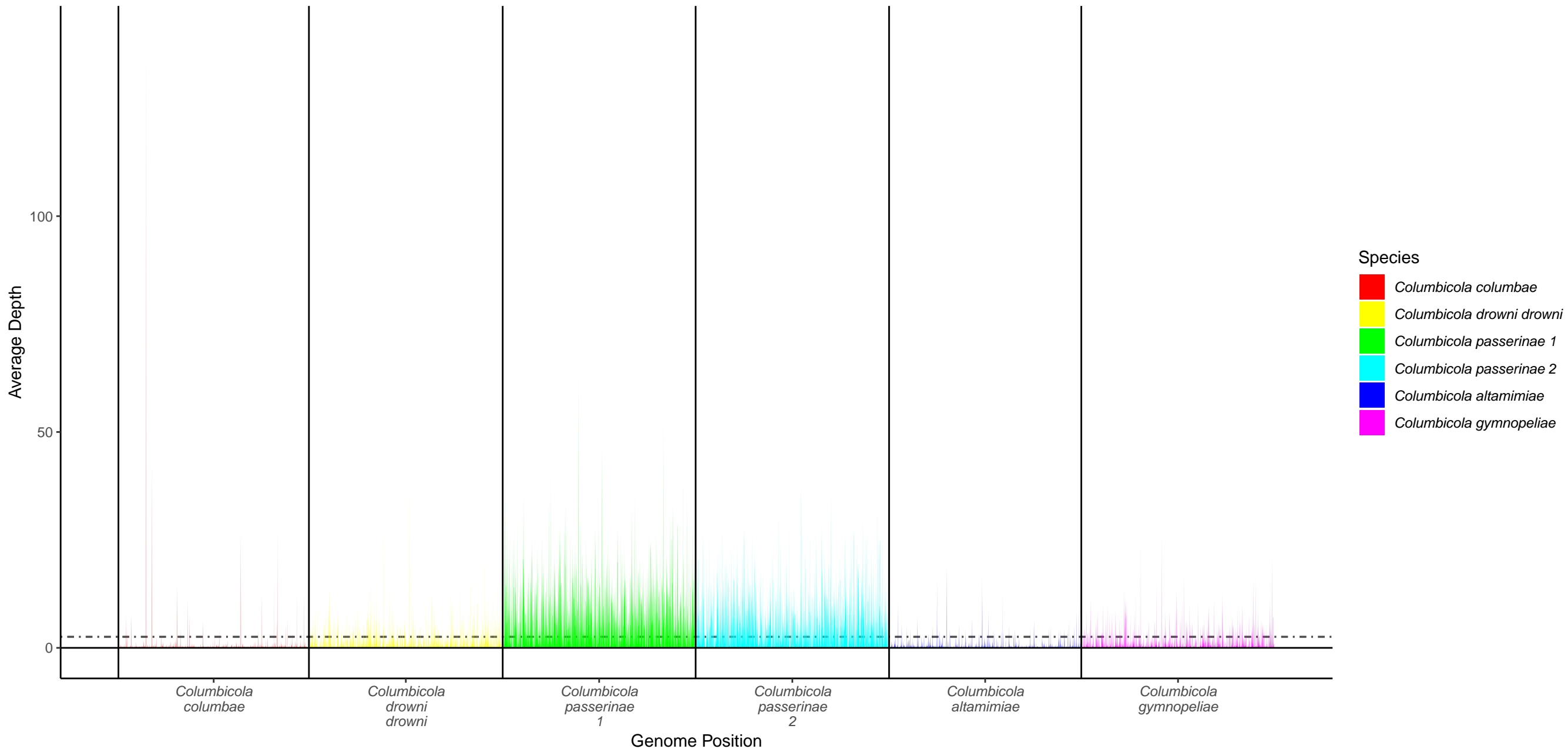


OUT Avg depth of coverage



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*

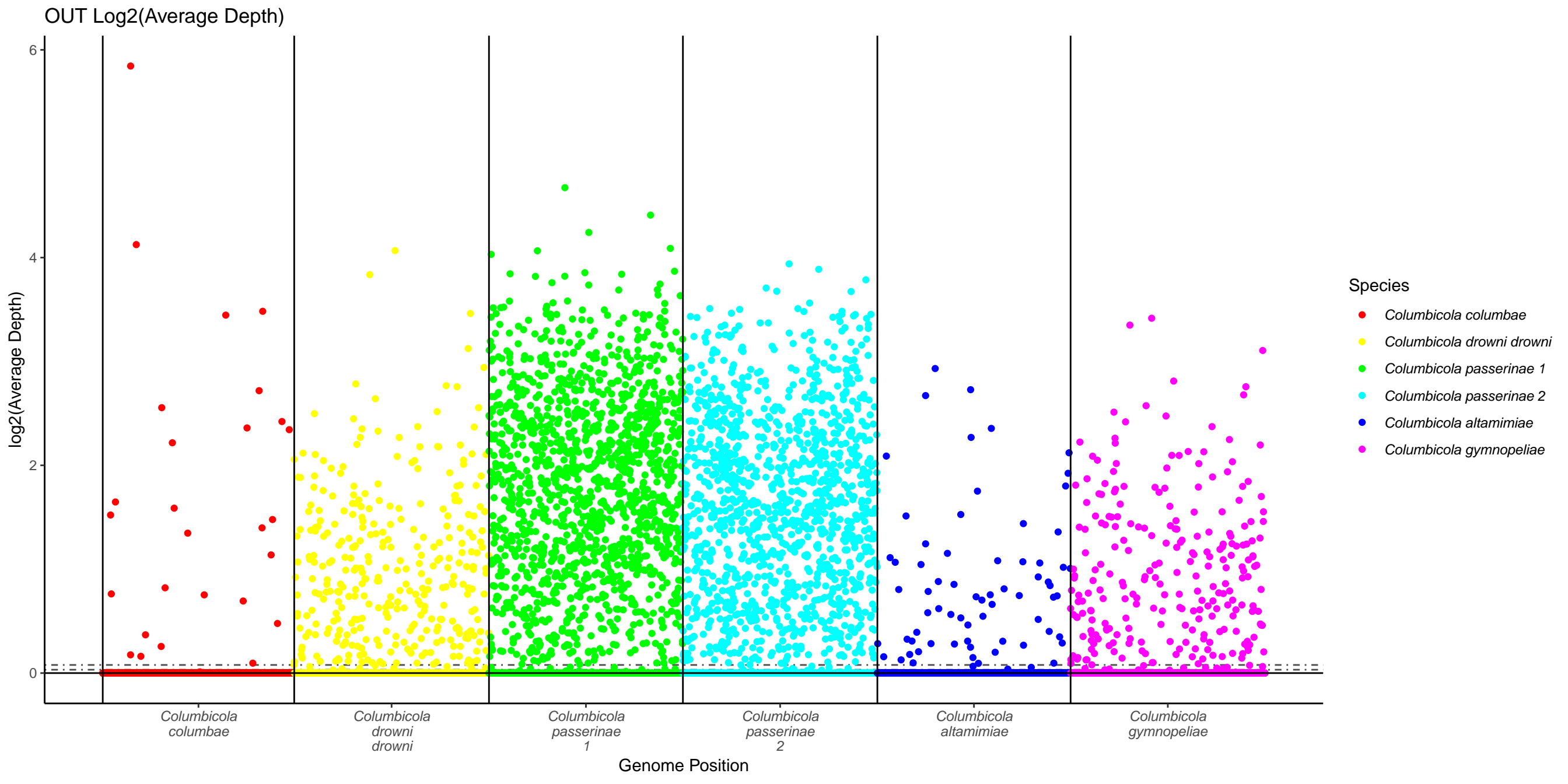
*Columbicola  
passerinae  
1*

*Columbicola  
passerinae  
2*

*Columbicola  
altamimiae*

*Columbicola  
gymnopeliae*

Genome Position

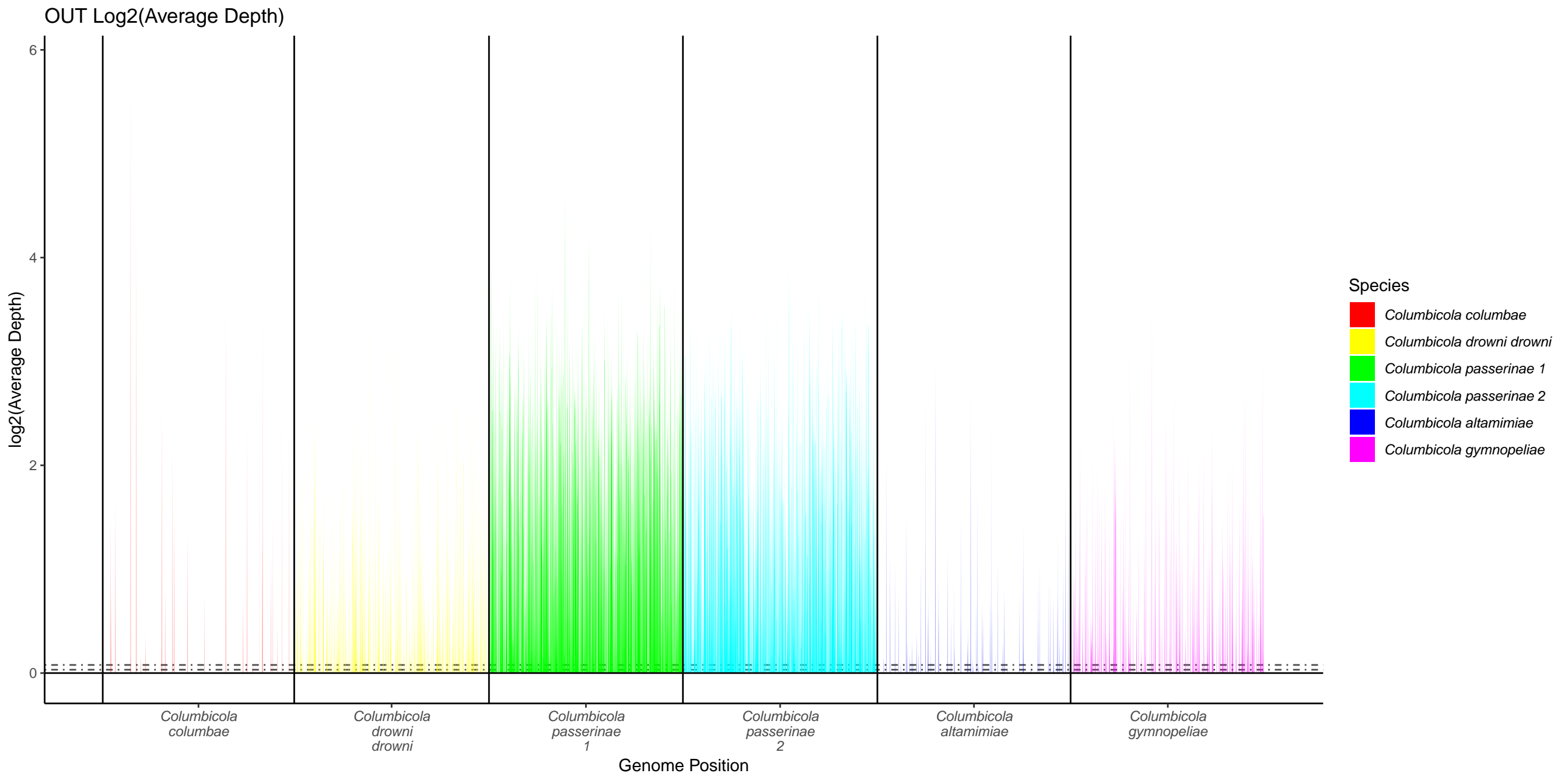


OUT Log2(Average Depth)

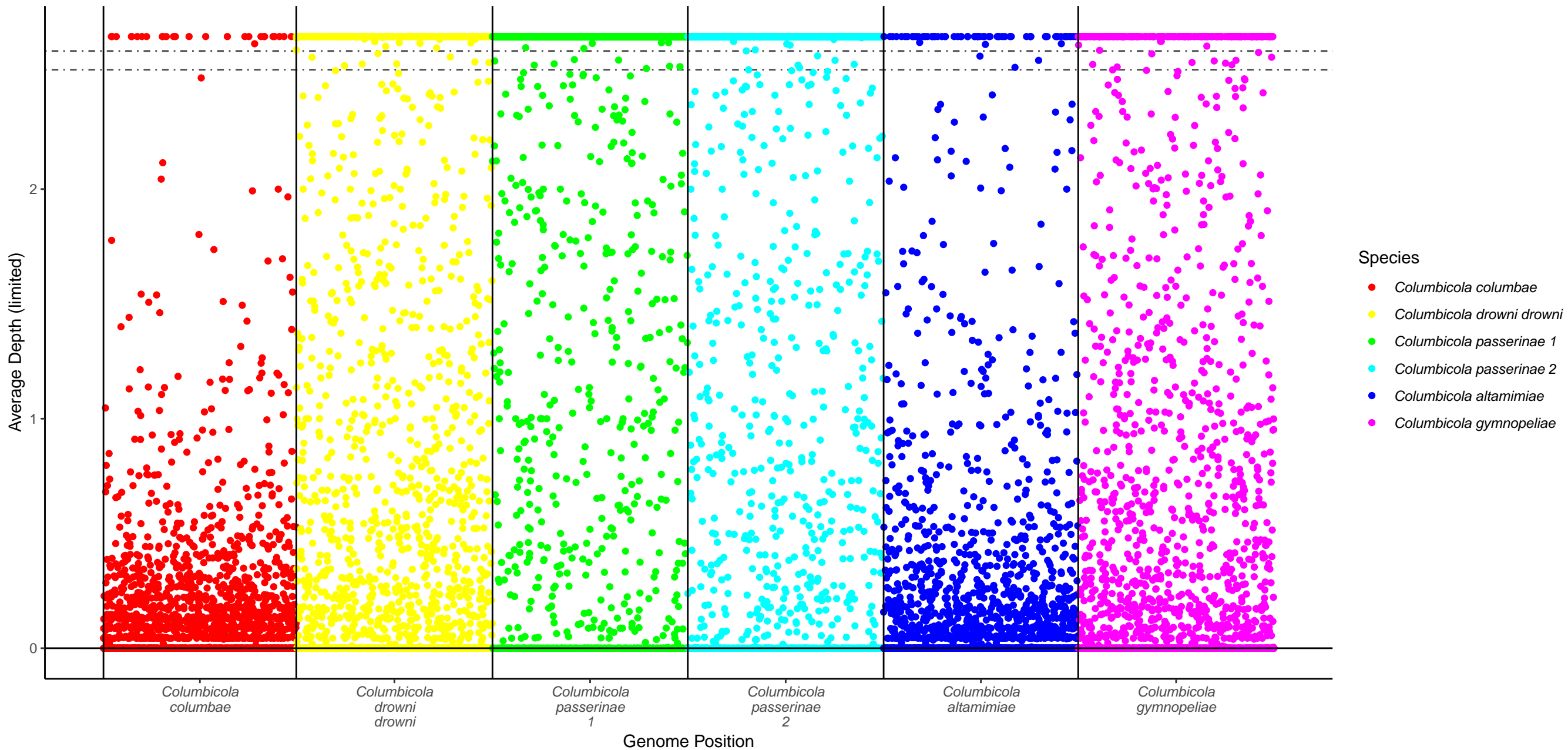
log2(Average Depth)

Species

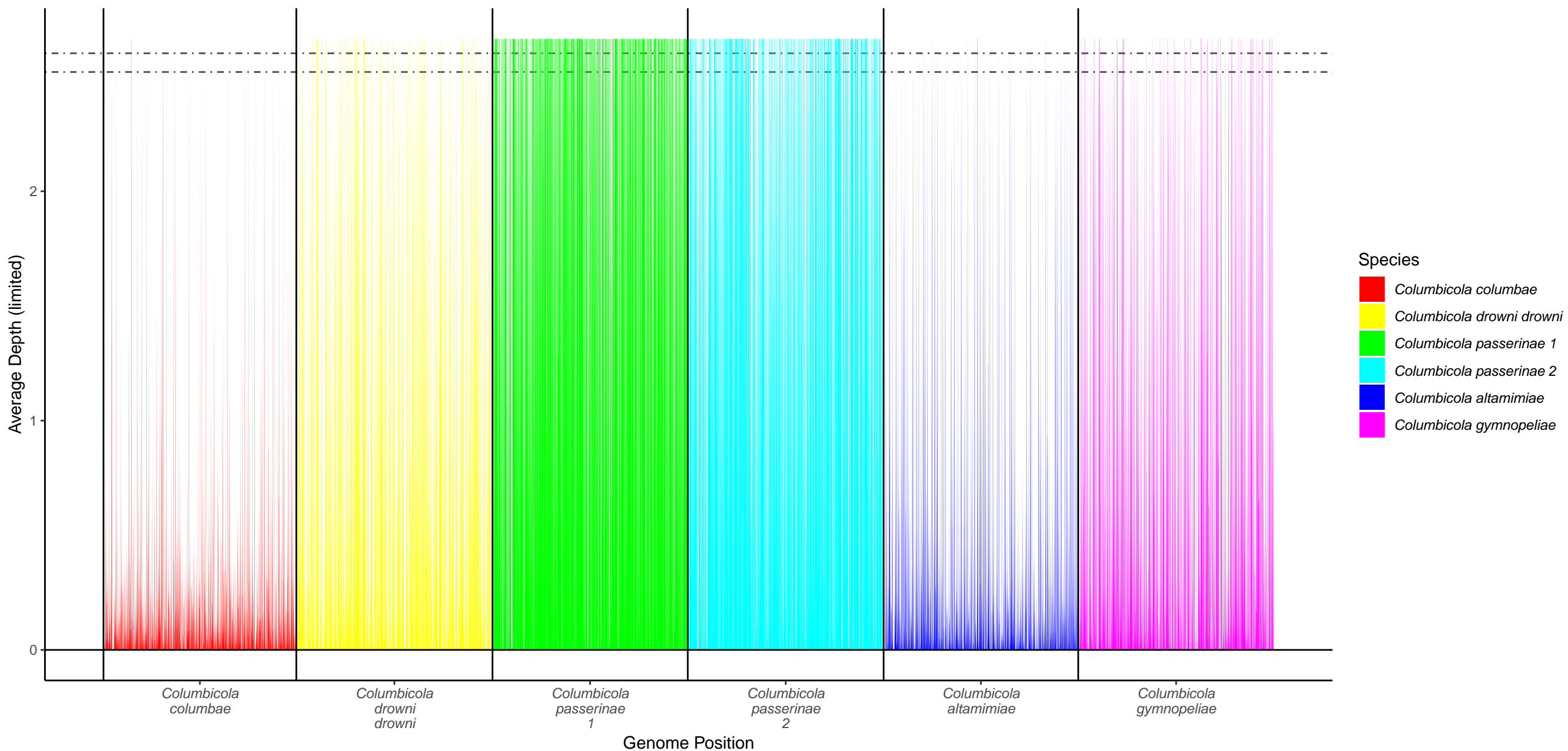
- *Columbicola columbae*
- *Columbicola drowni drowni*
- *Columbicola passerinae 1*
- *Columbicola passerinae 2*
- *Columbicola altamimiae*
- *Columbicola gymnopeliae*



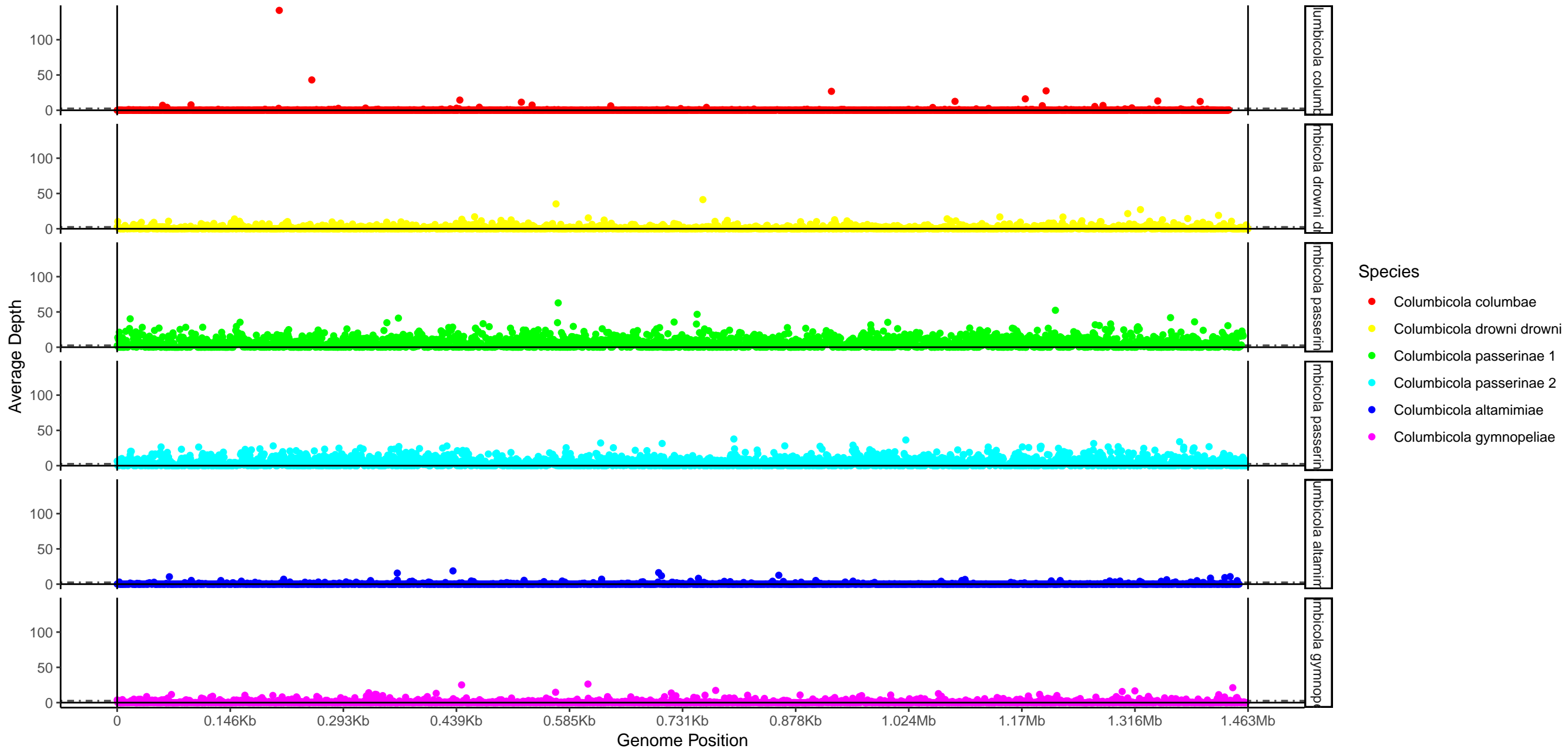
OUT Avg depth of coverage



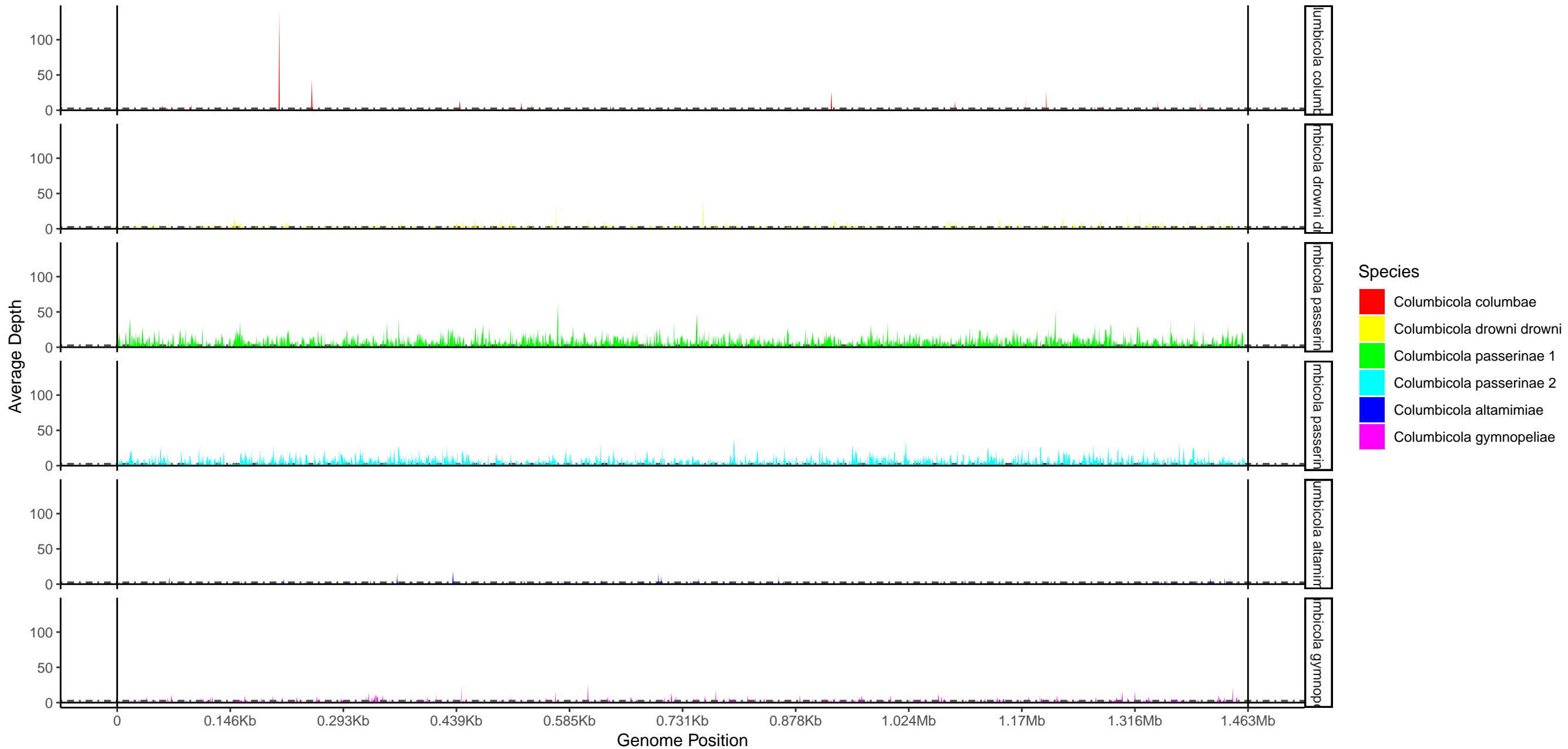
OUT Avg depth of coverage



# OUT Avg depth of coverage

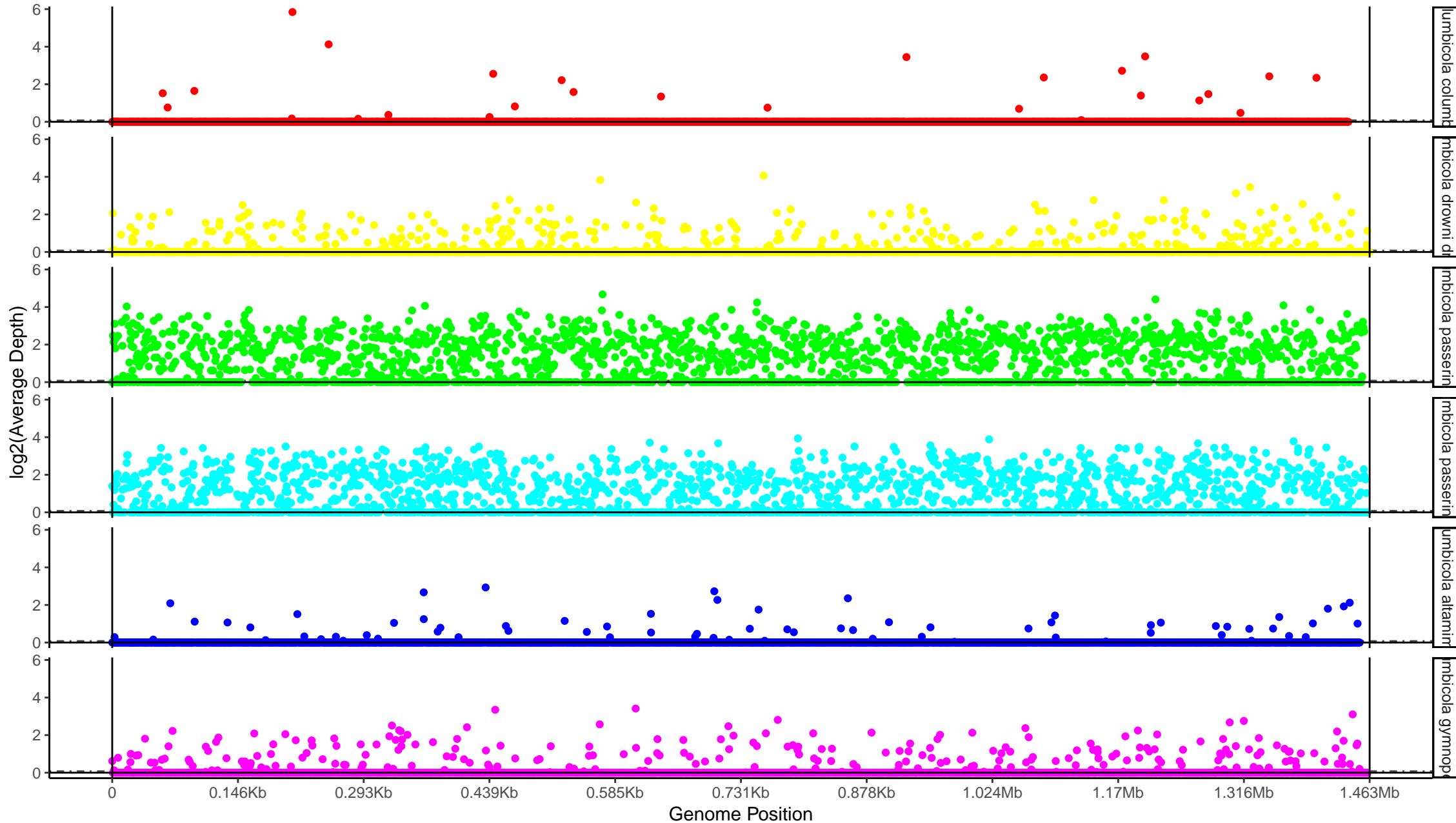


OUT Avg depth of coverage





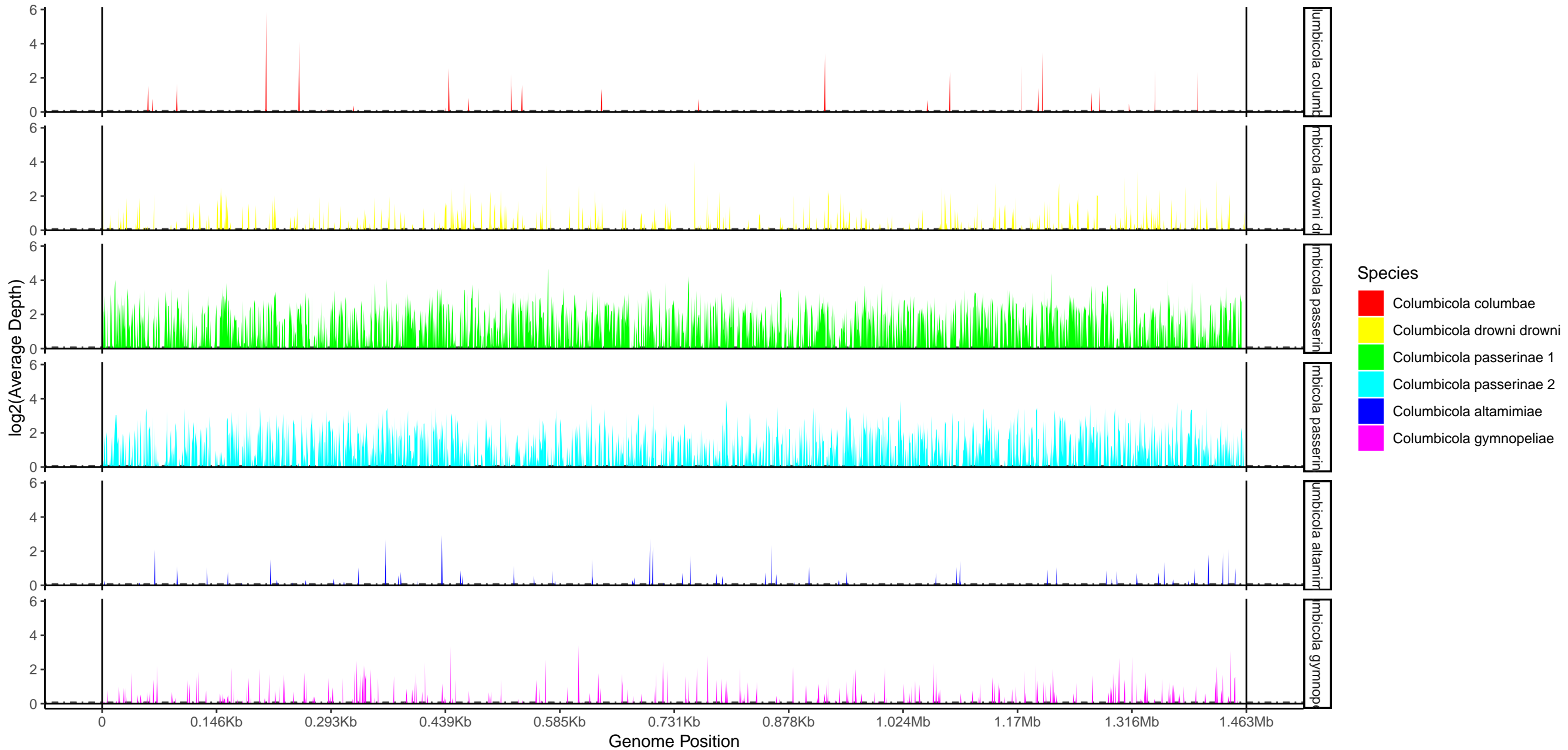
OUT Log2(Avg depth of coverage)



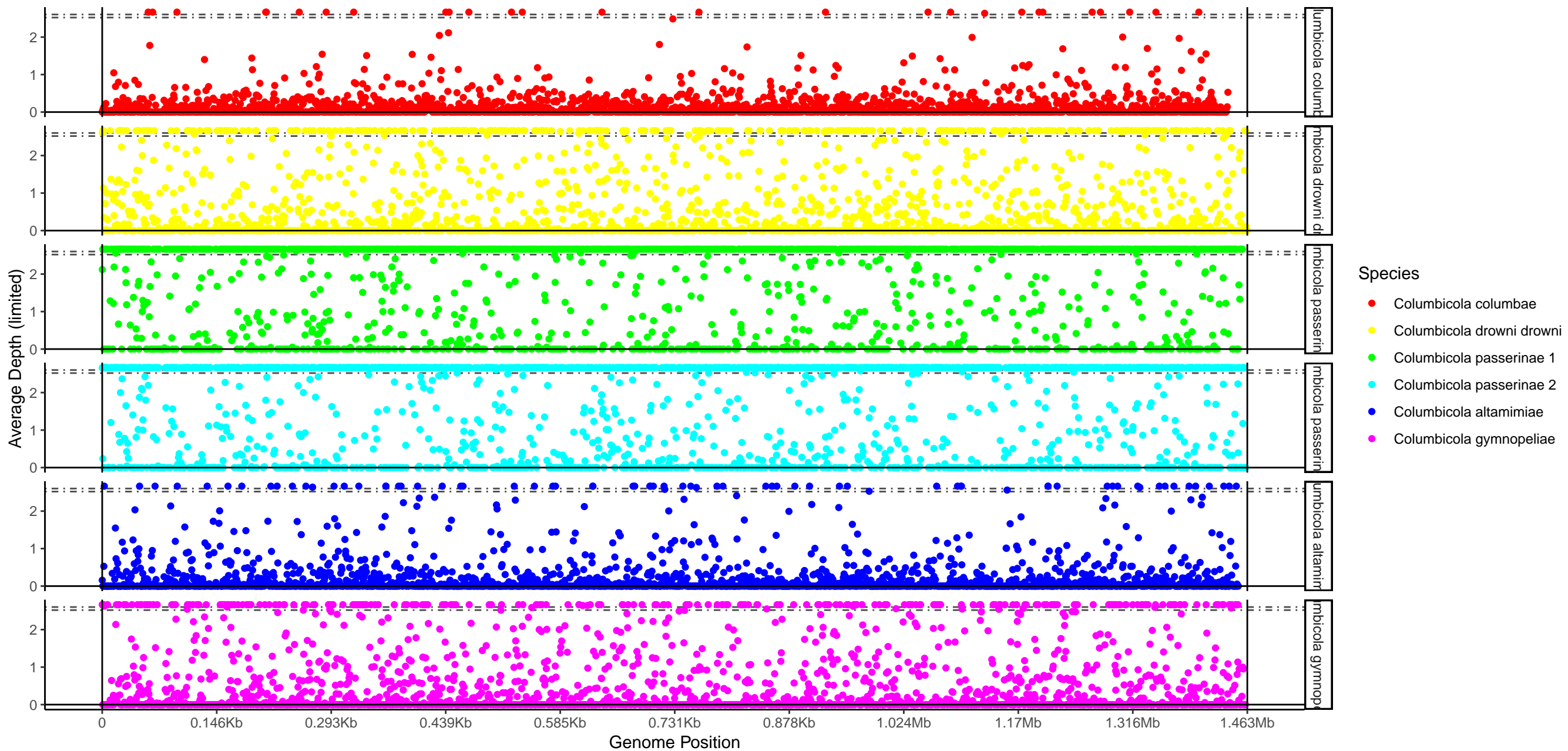
Species

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

OUT Log2(Avg depth of coverage)



OUT Avg depth of coverage



OUT Avg depth of coverage

