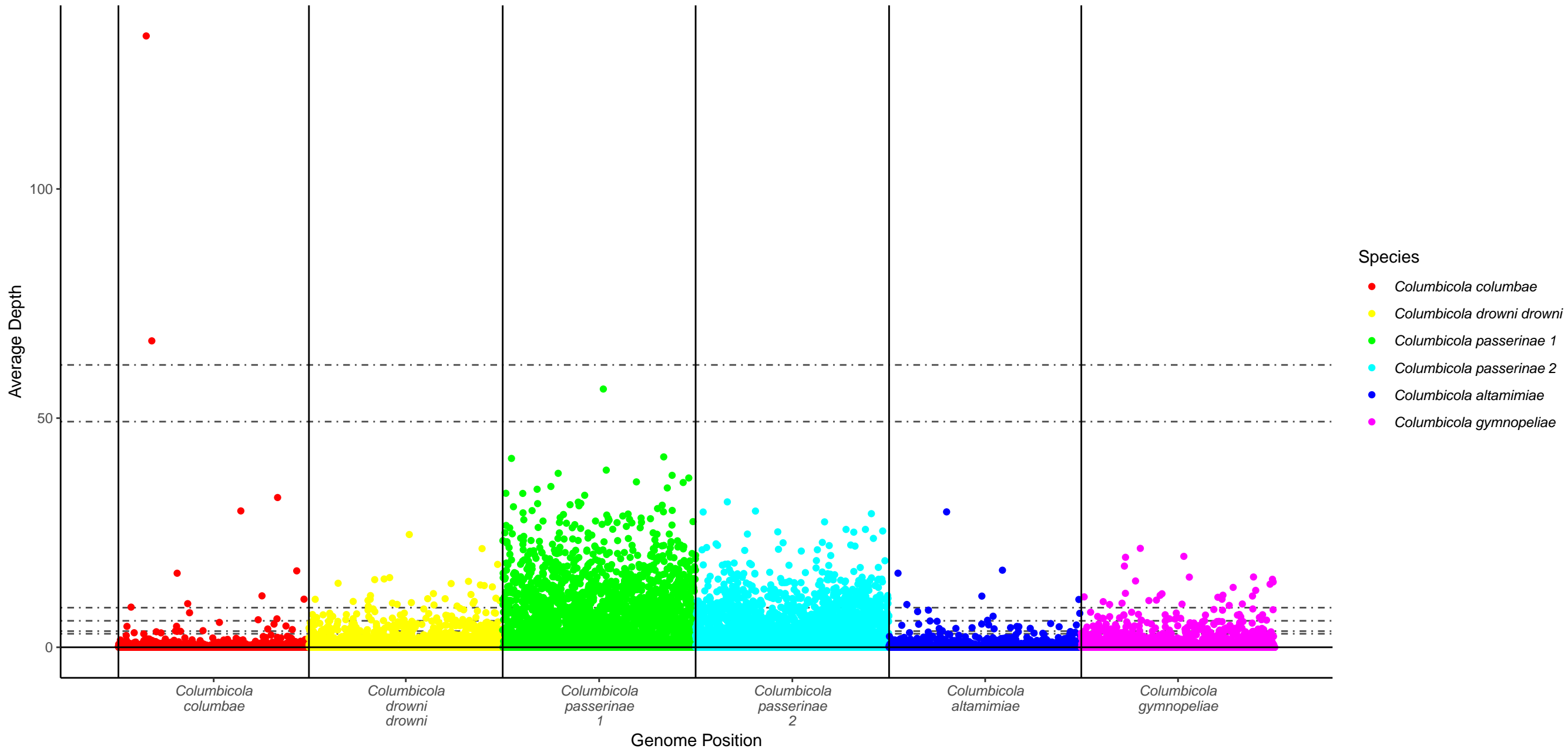
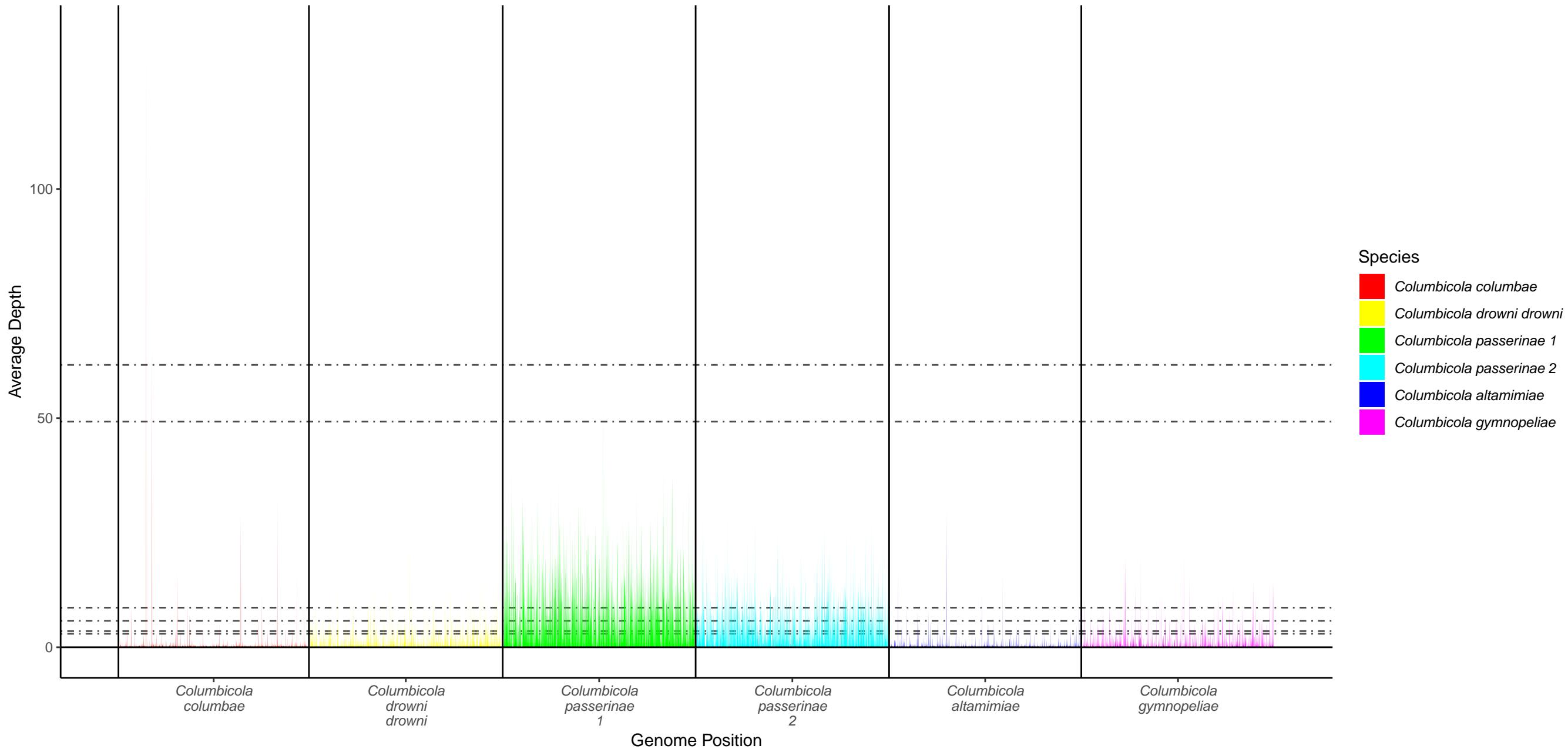


OUT Avg depth of coverage



OUT Avg depth of coverage

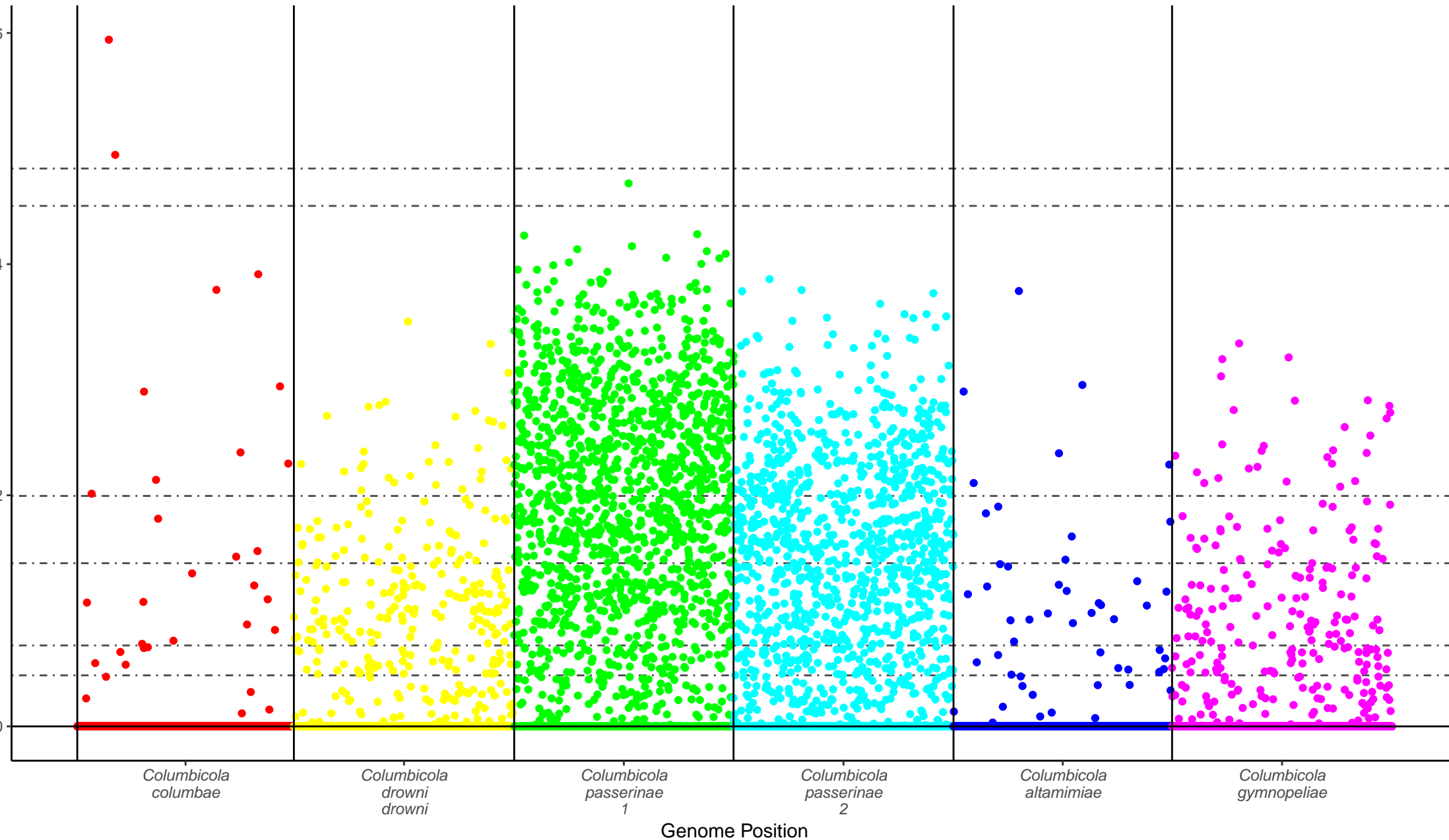


OUT Log2(Average Depth)

log2(Average Depth)

Species

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



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*

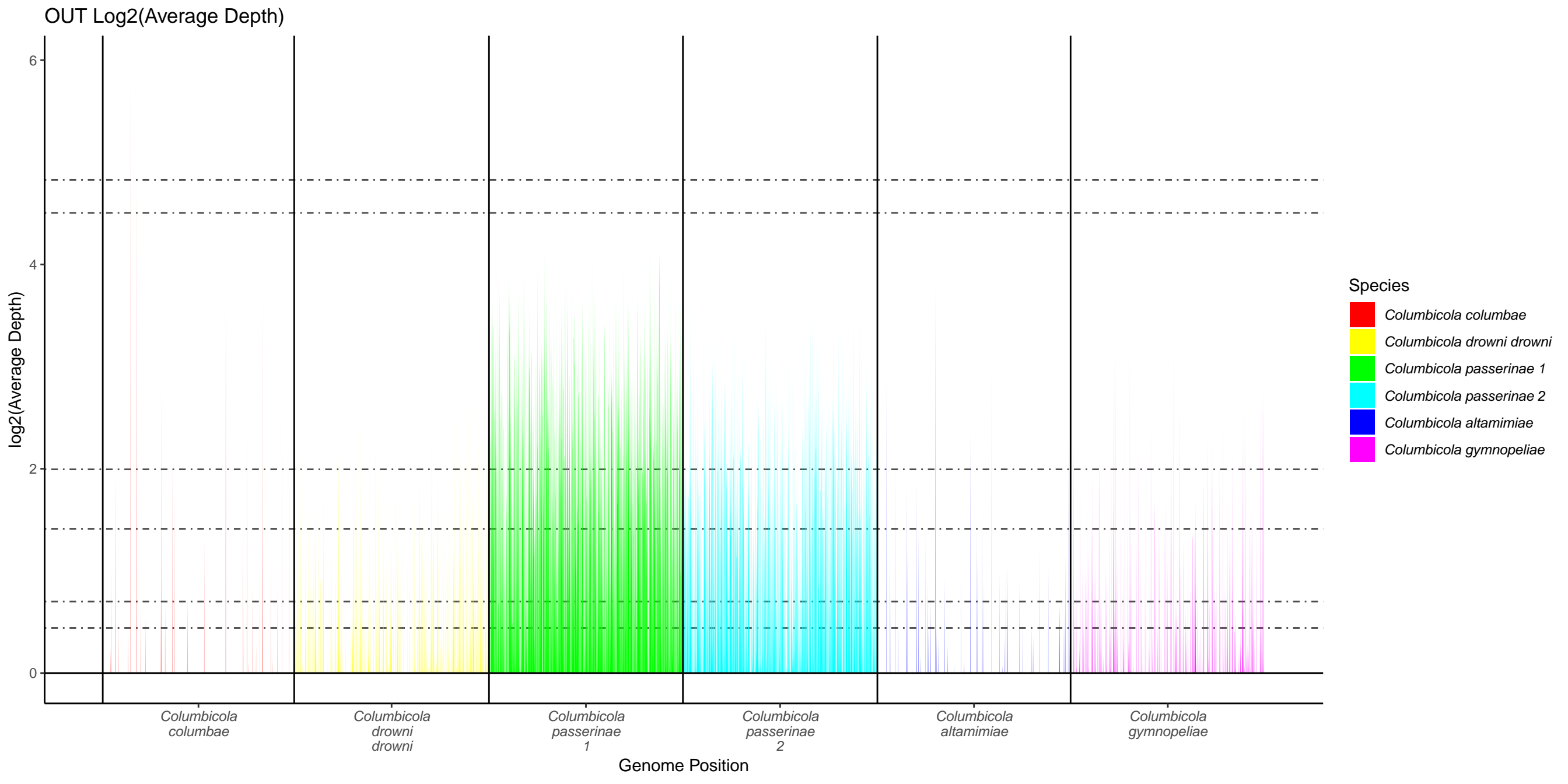
*Columbicola
passerinae
1*

*Columbicola
passerinae
2*

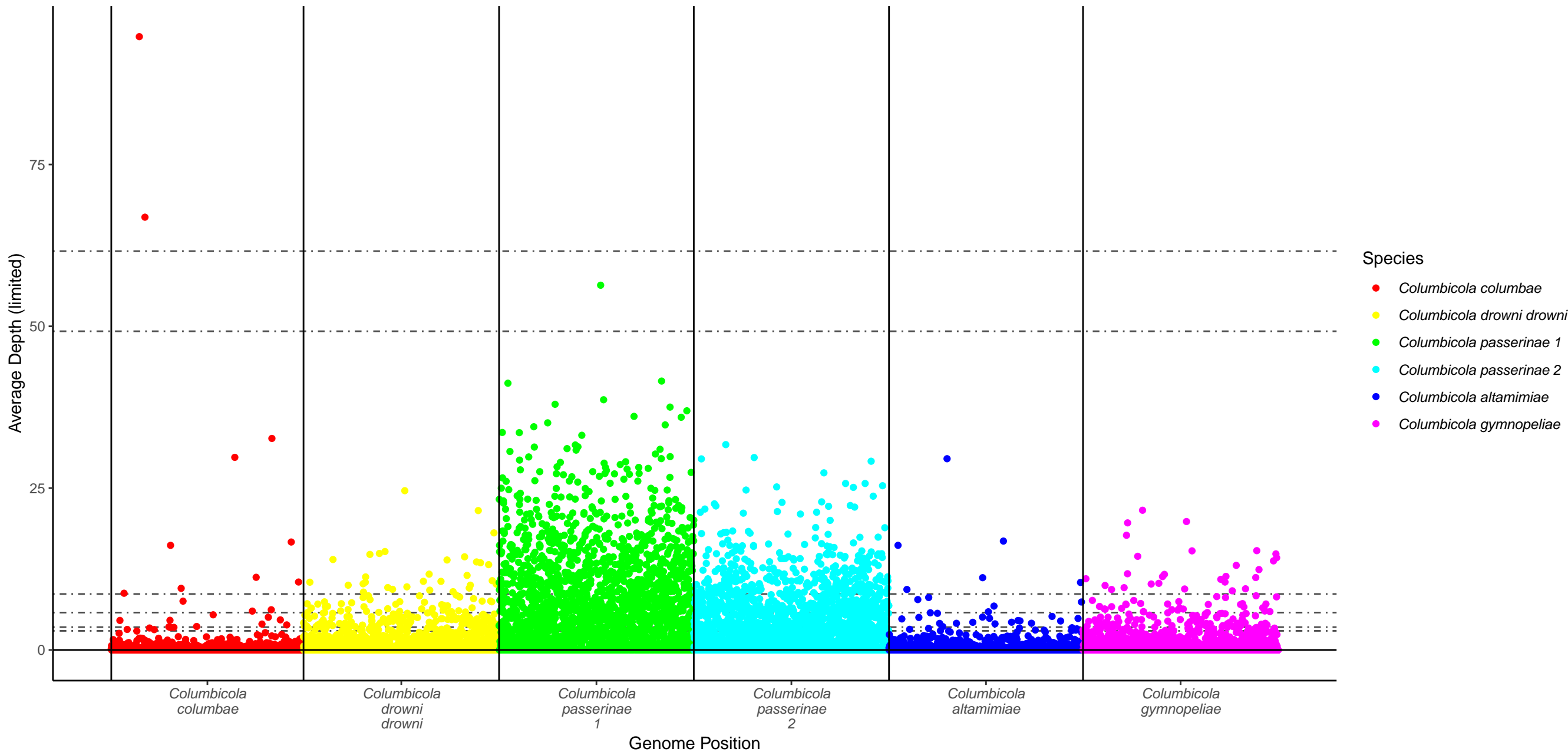
*Columbicola
altamimiae*

*Columbicola
gymnopeliae*

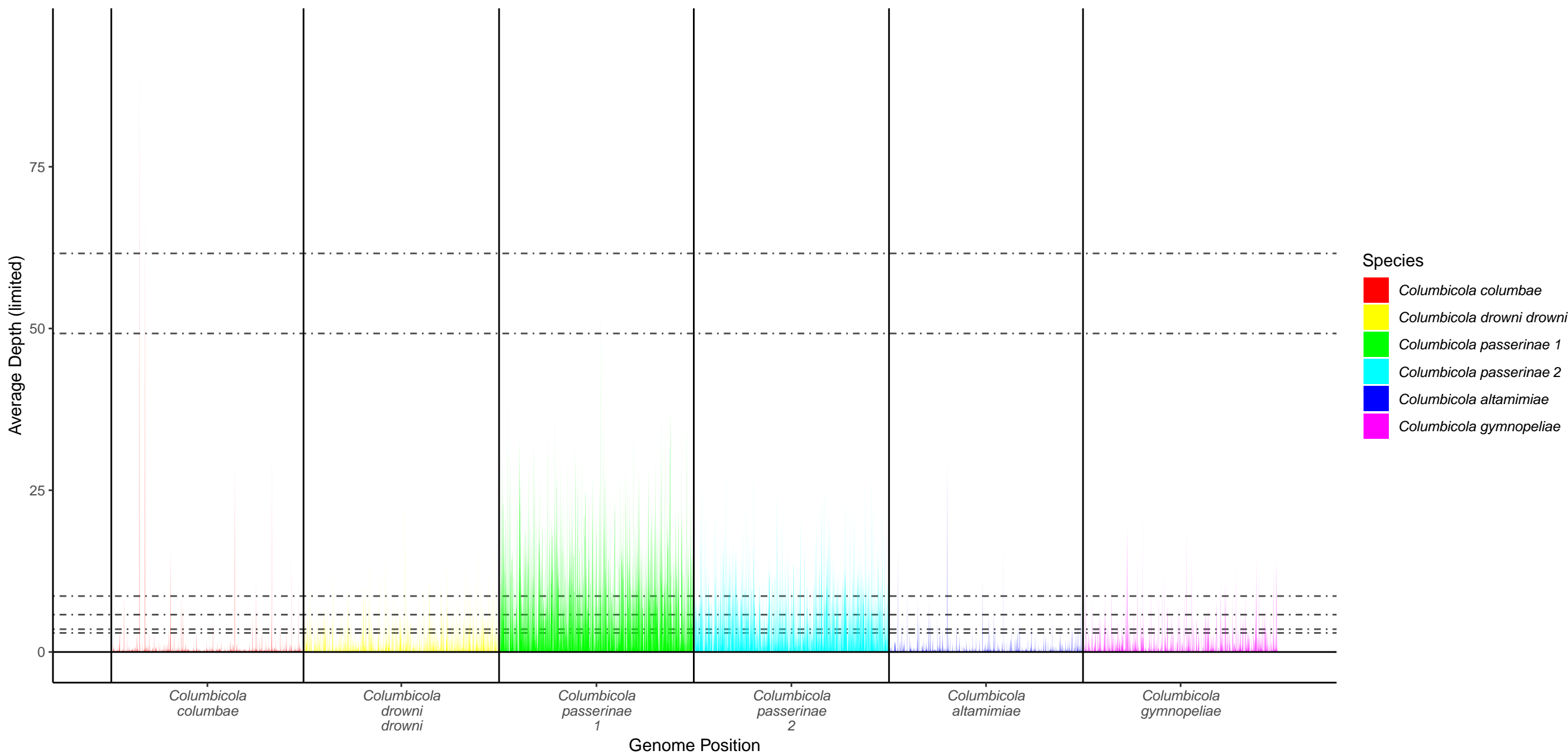
Genome Position



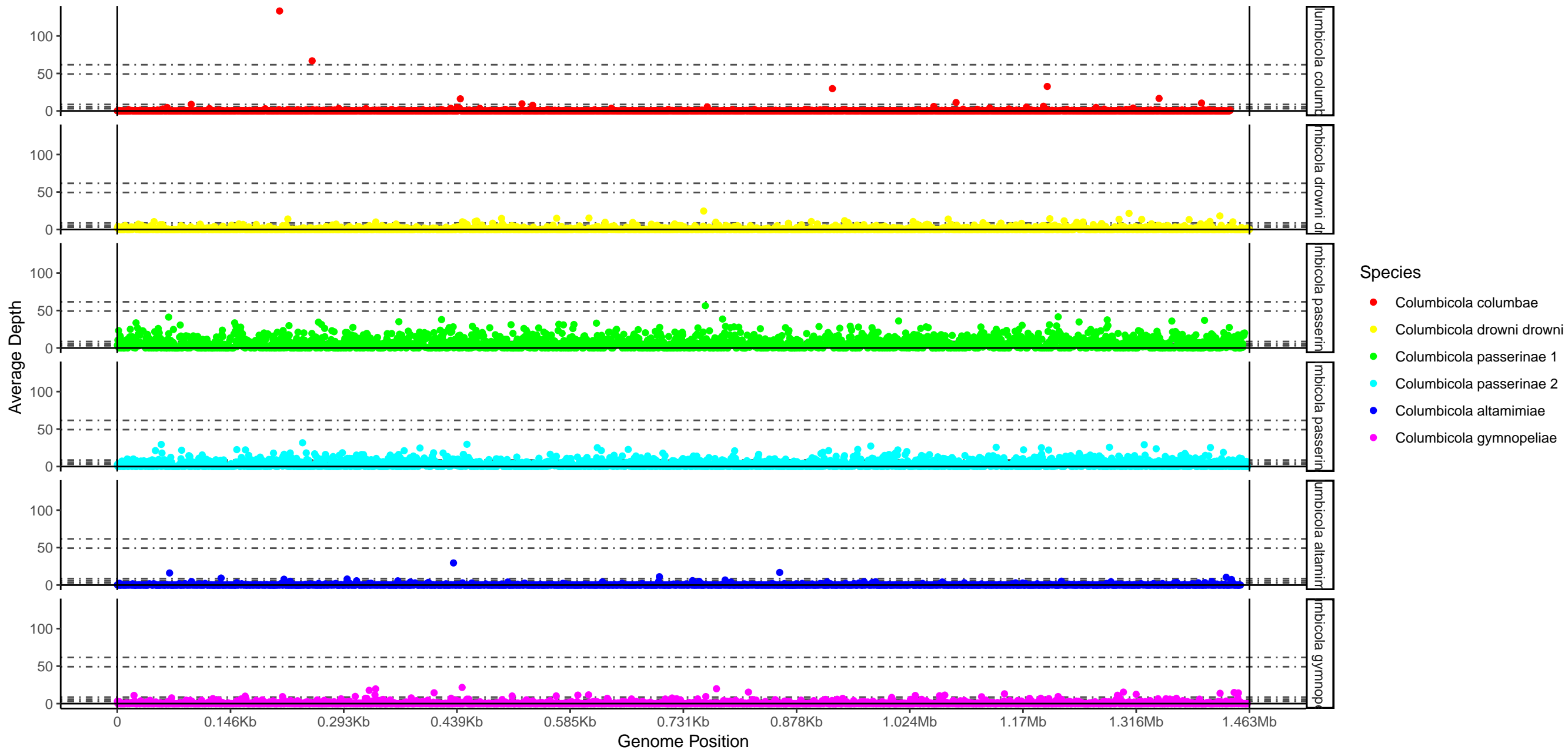
OUT Avg depth of coverage



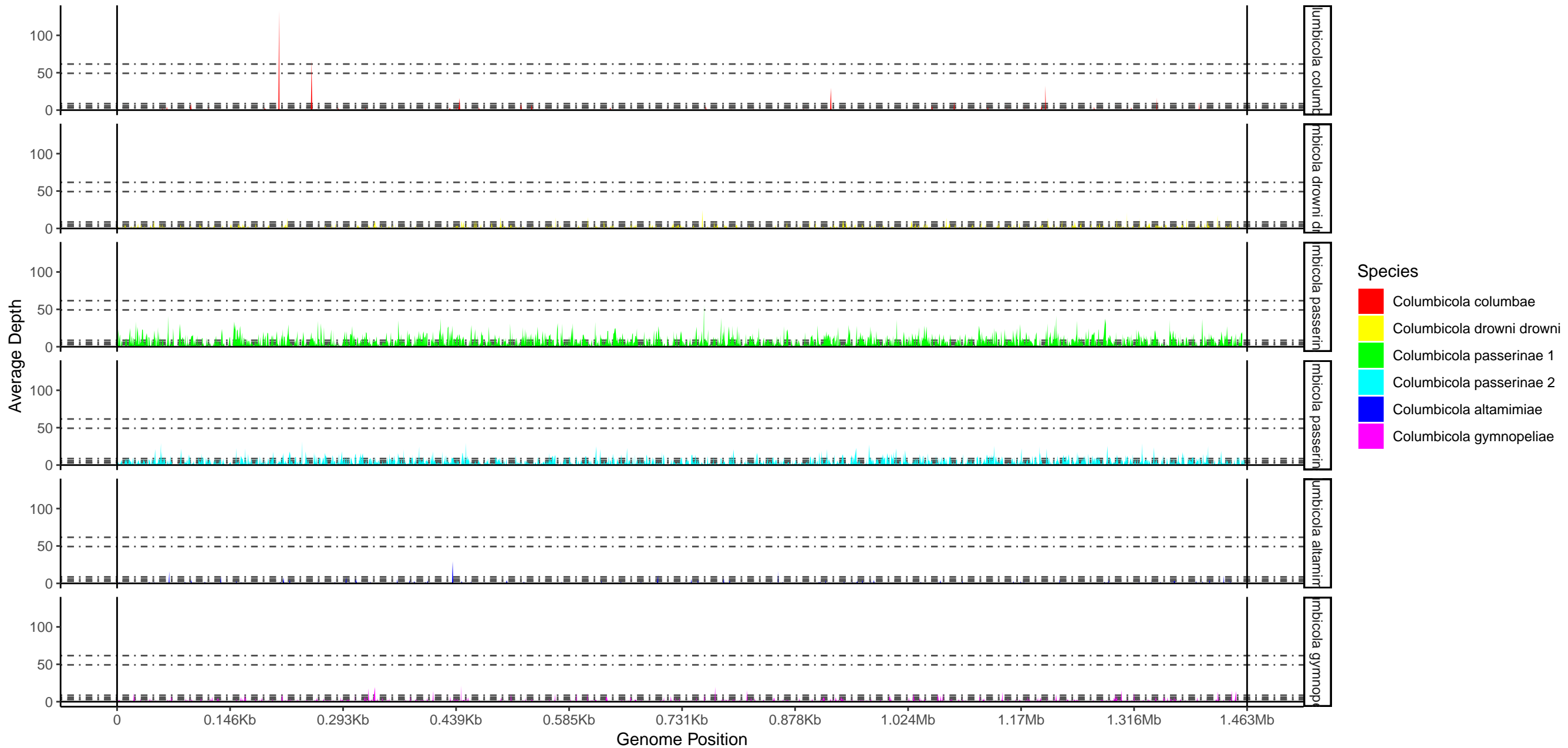
OUT Avg depth of coverage



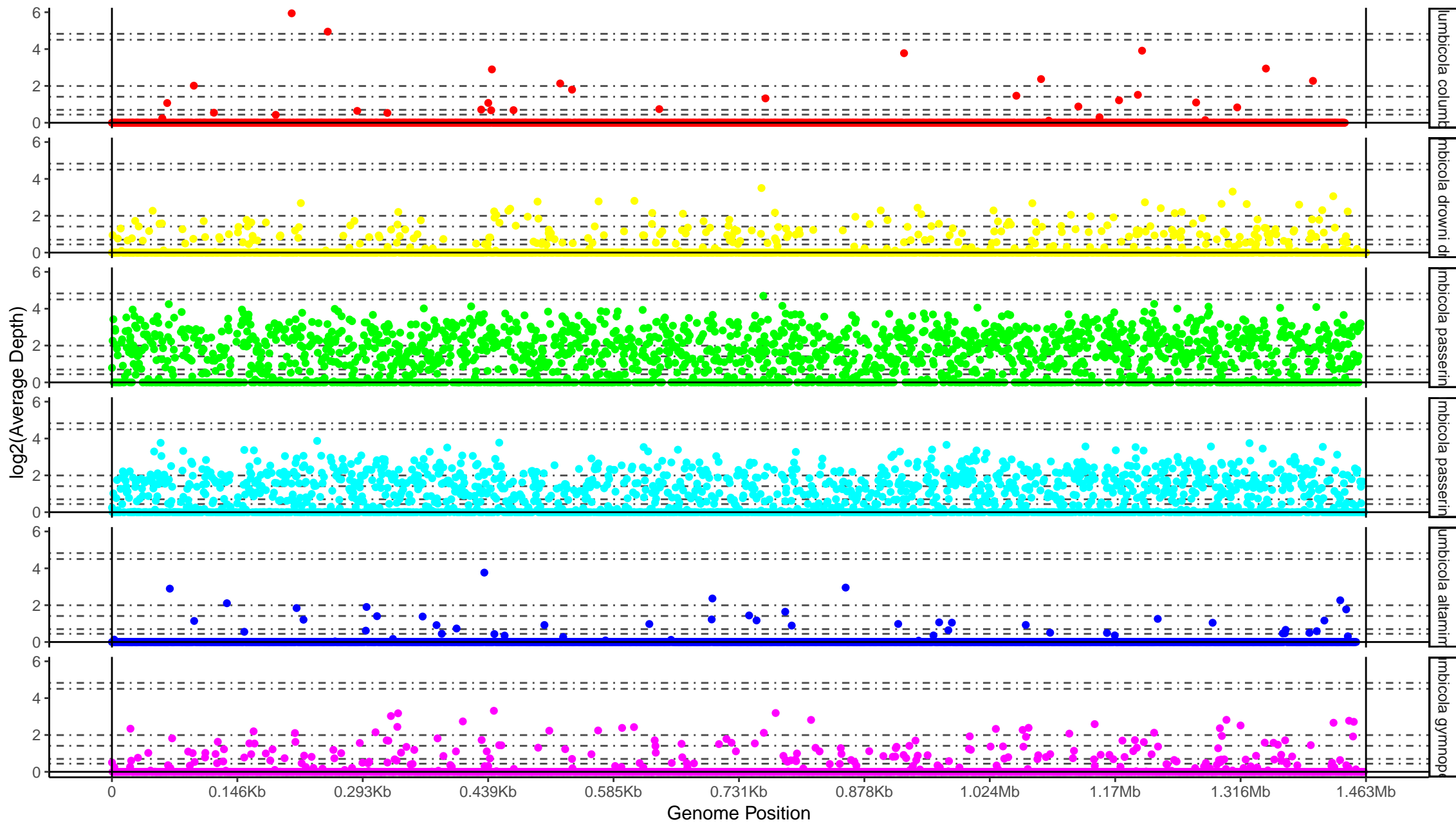
OUT Avg depth of coverage



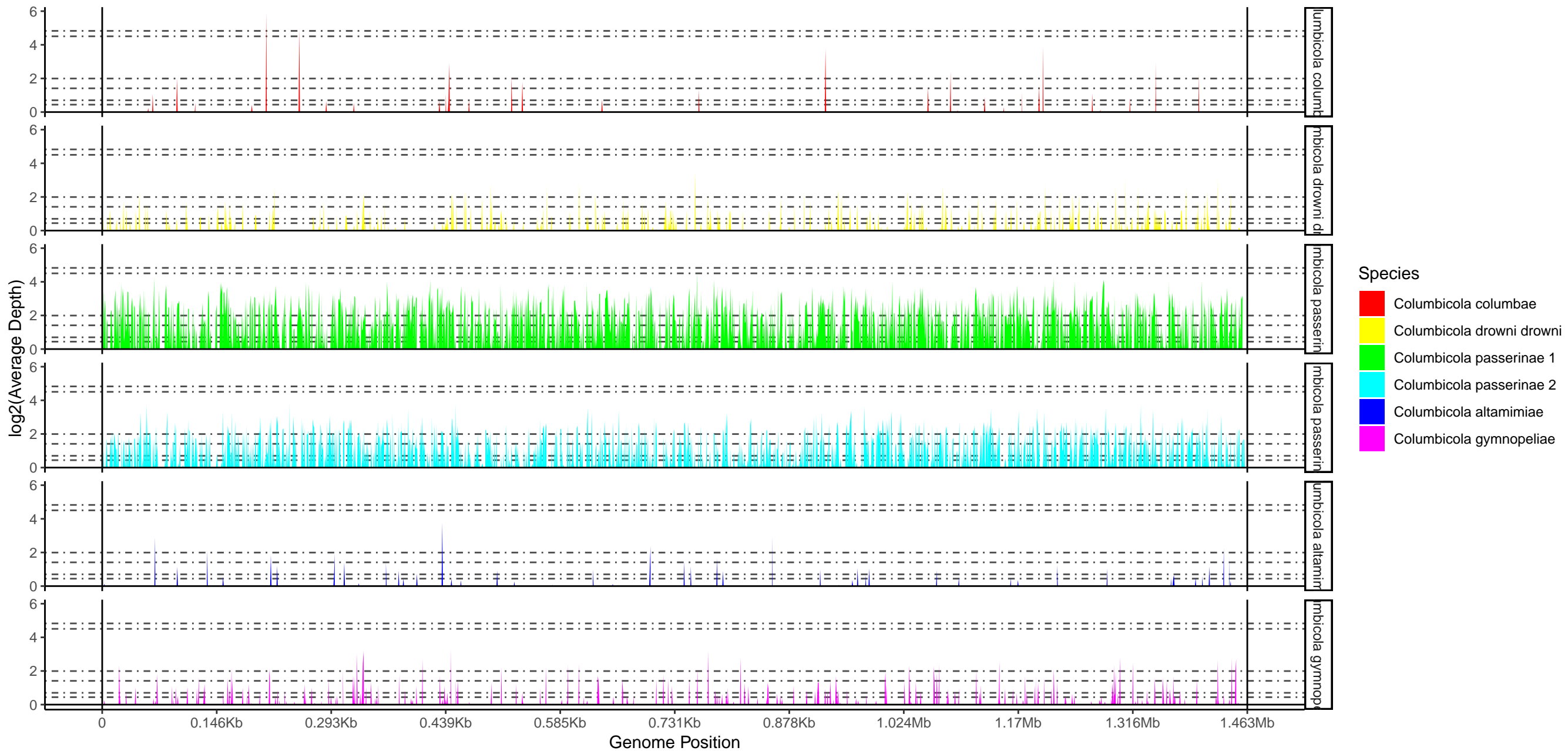
OUT Avg depth of coverage



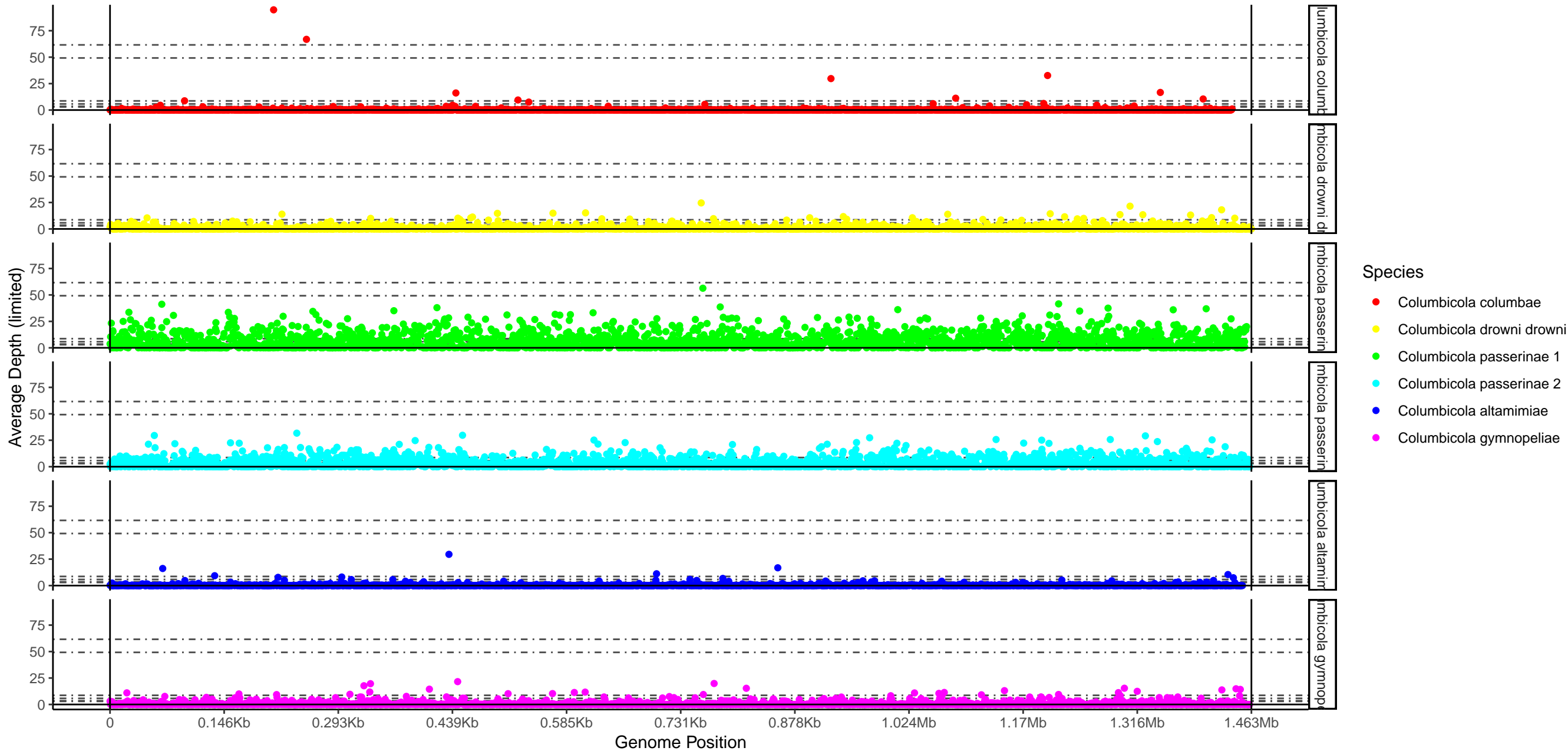
OUT Log2(Avg depth of coverage)



OUT Log2(Avg depth of coverage)



OUT Avg depth of coverage



OUT Avg depth of coverage

