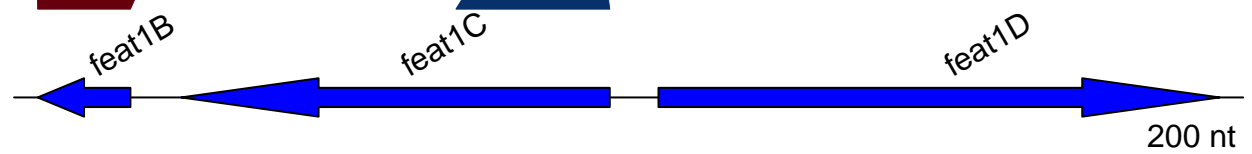
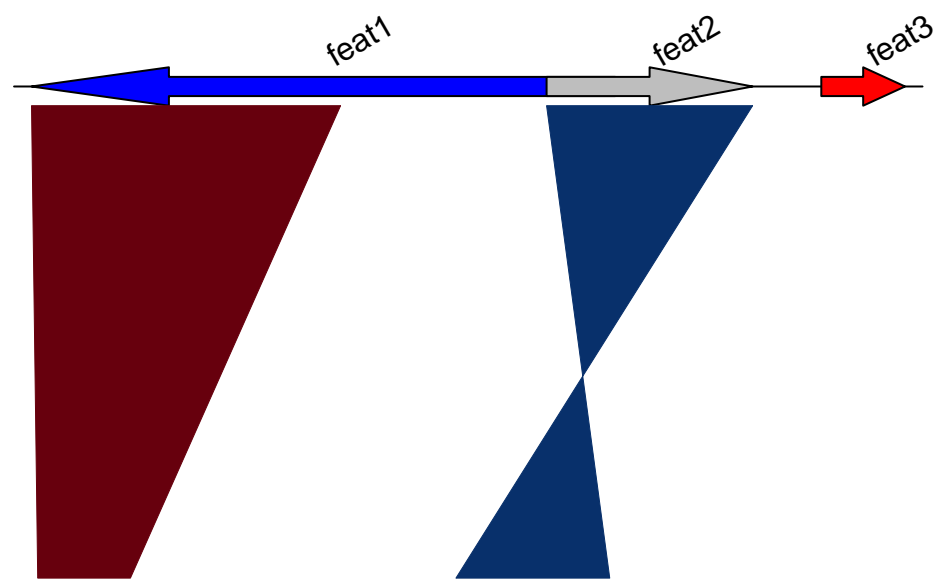
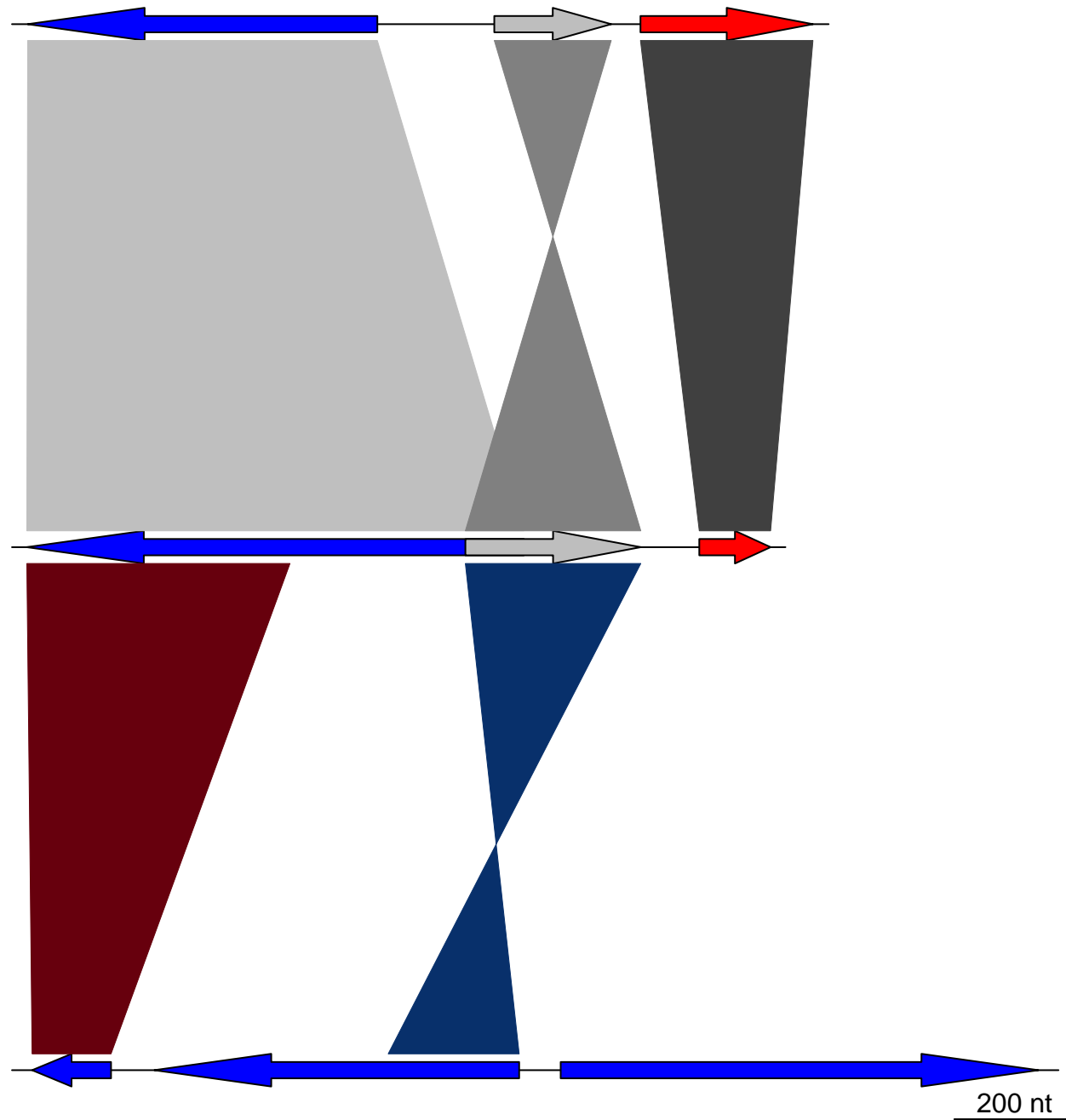
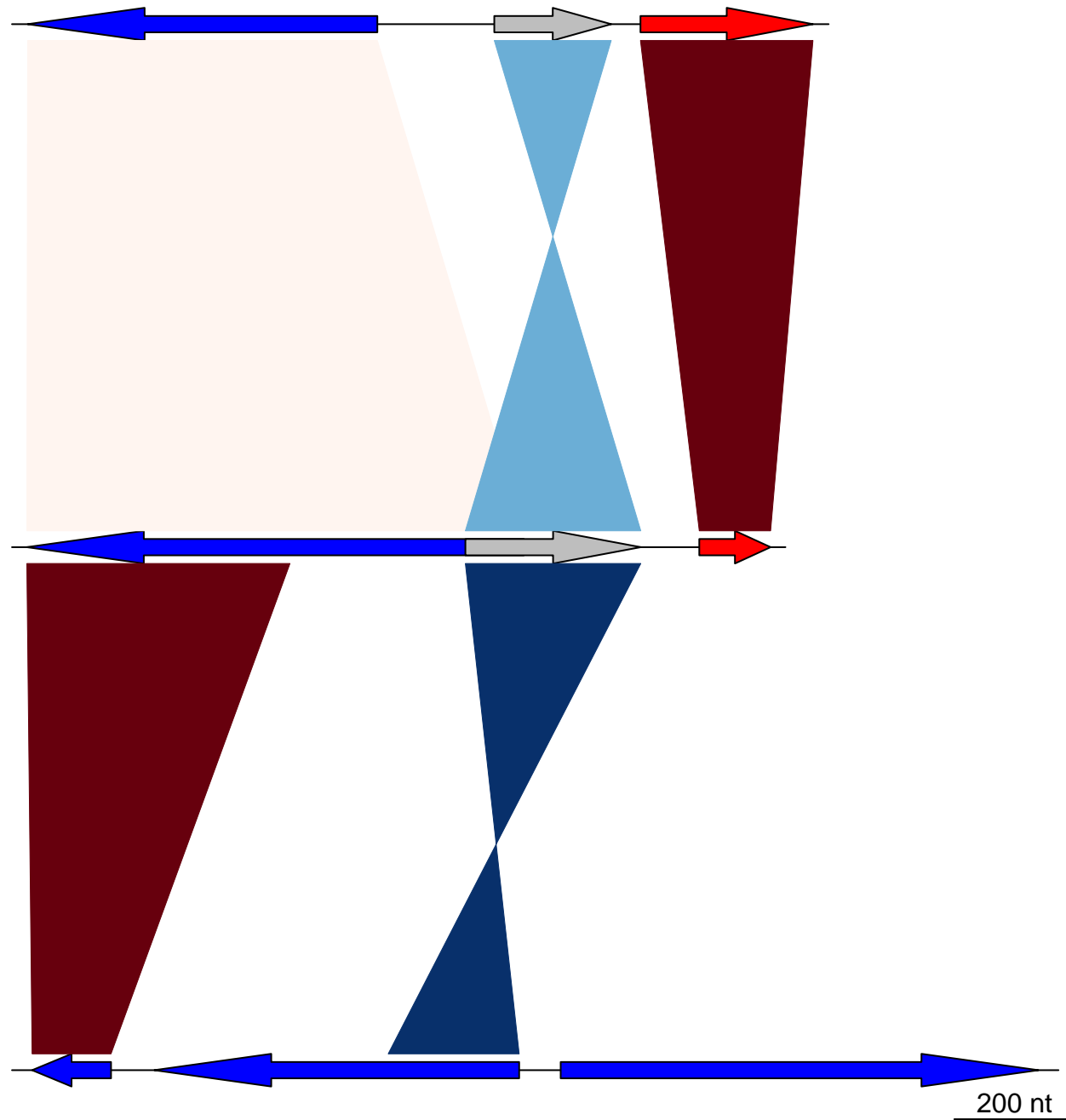


help("annotation")

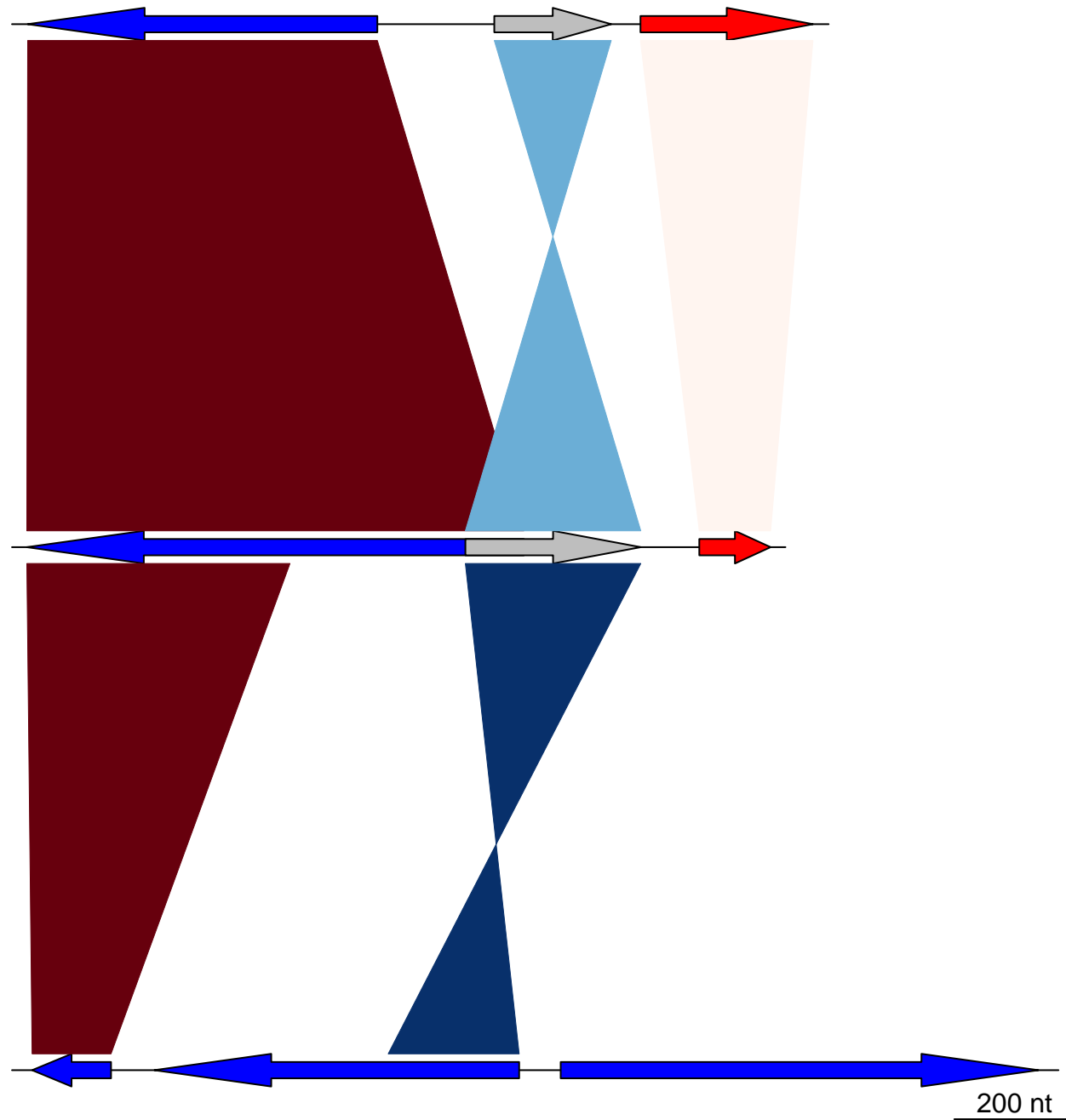




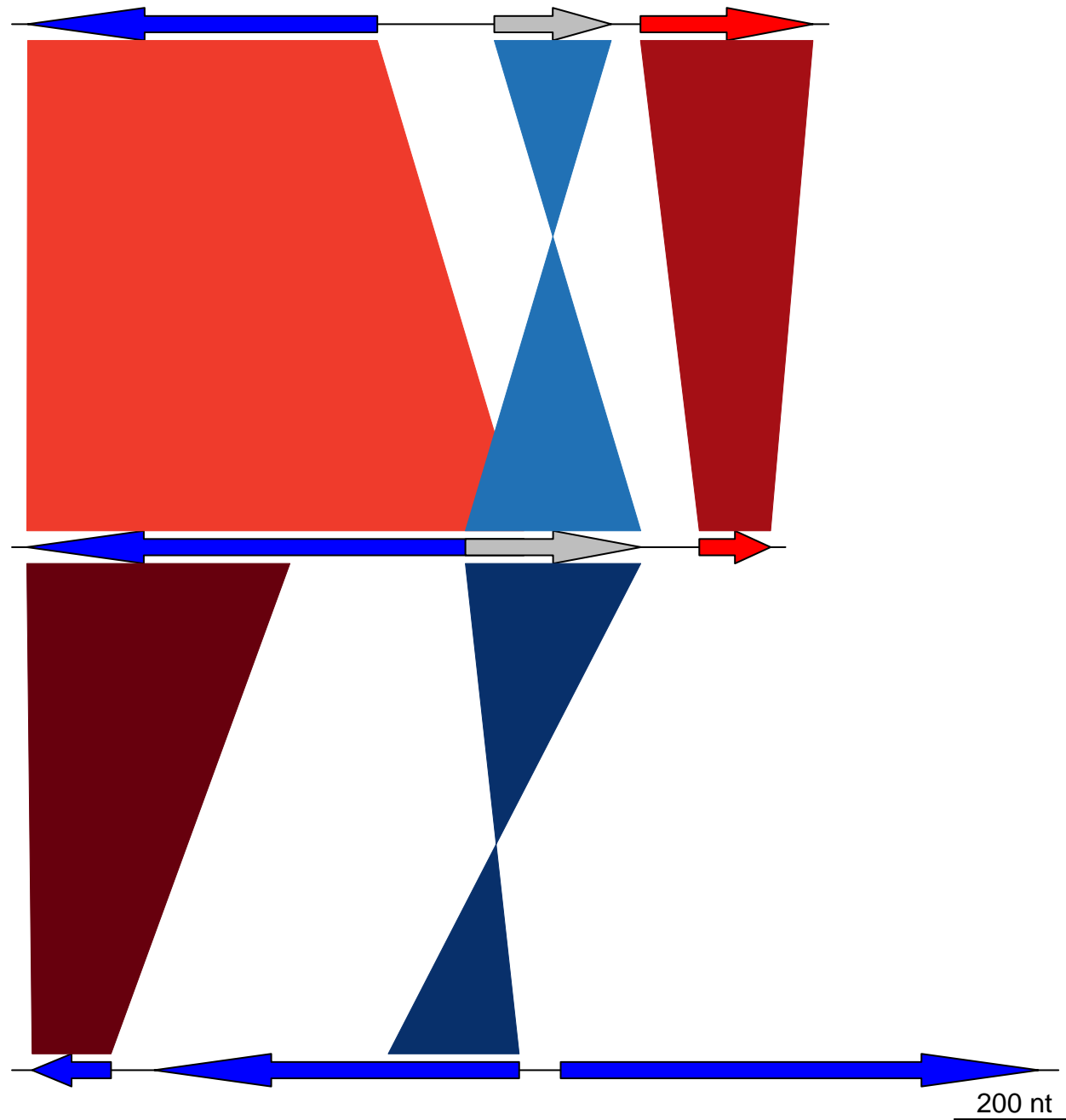
help("apply\_color\_scheme")



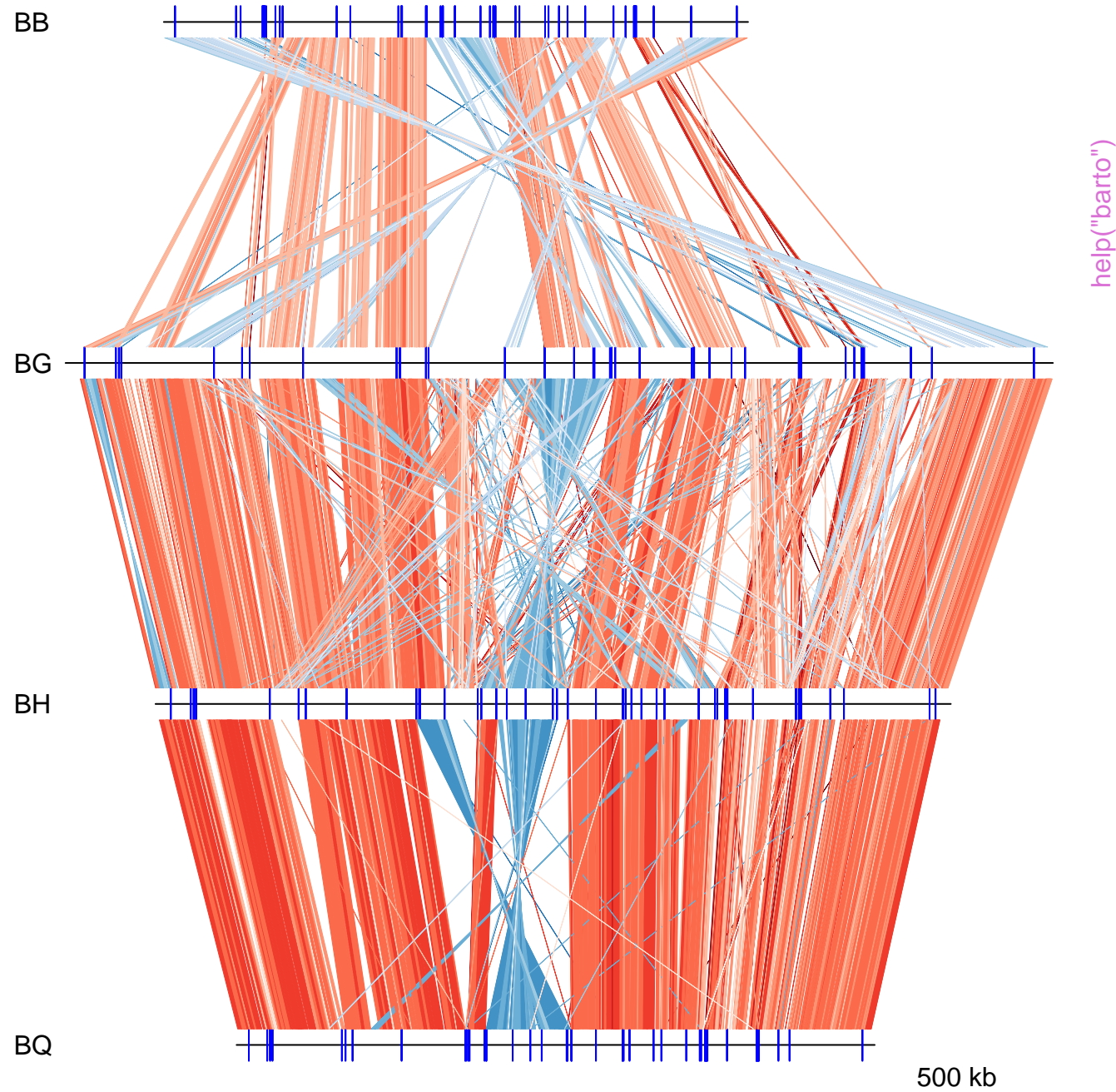
help("apply\_color\_scheme")



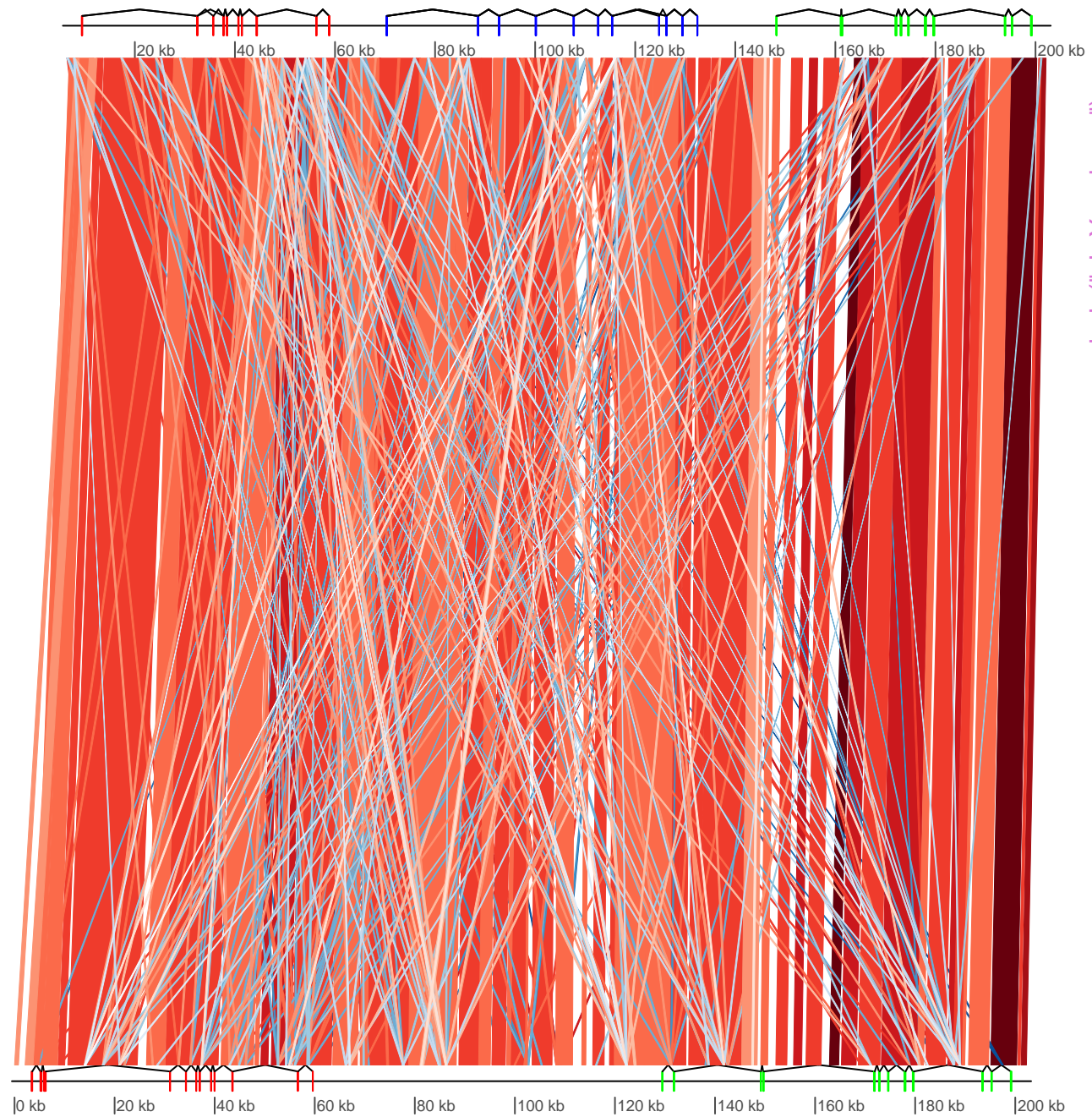
help("apply\_color\_scheme")



help("apply\_color\_scheme")





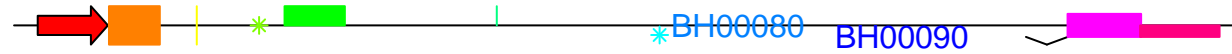


help("chrY\_subseg")

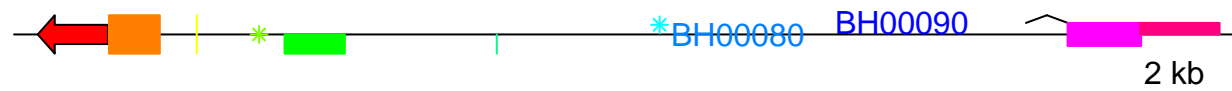


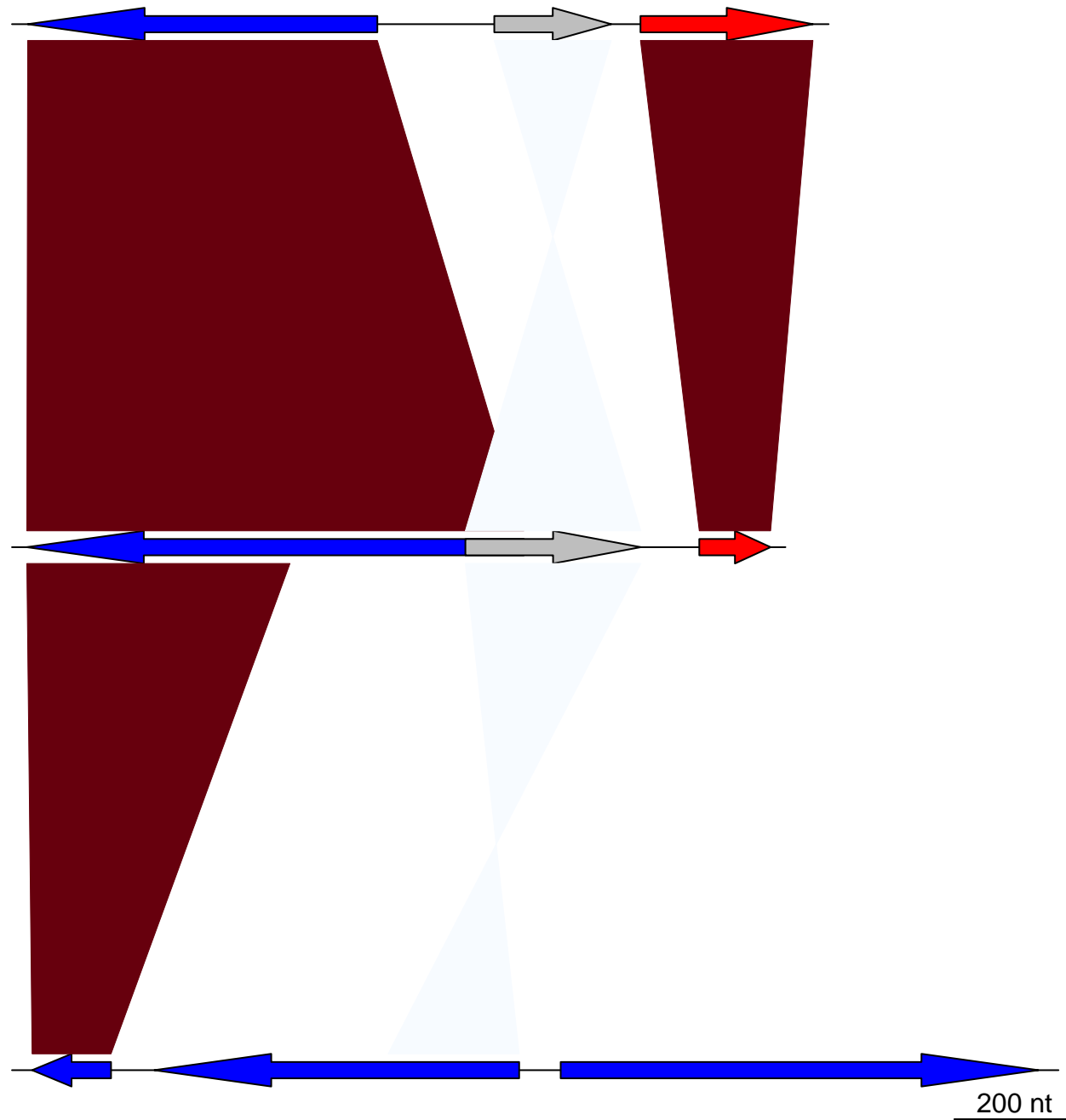
help("gene\_types")

2 kb



help("gene\_types")





help("genoPlotR-package")

B\_bacilliformis

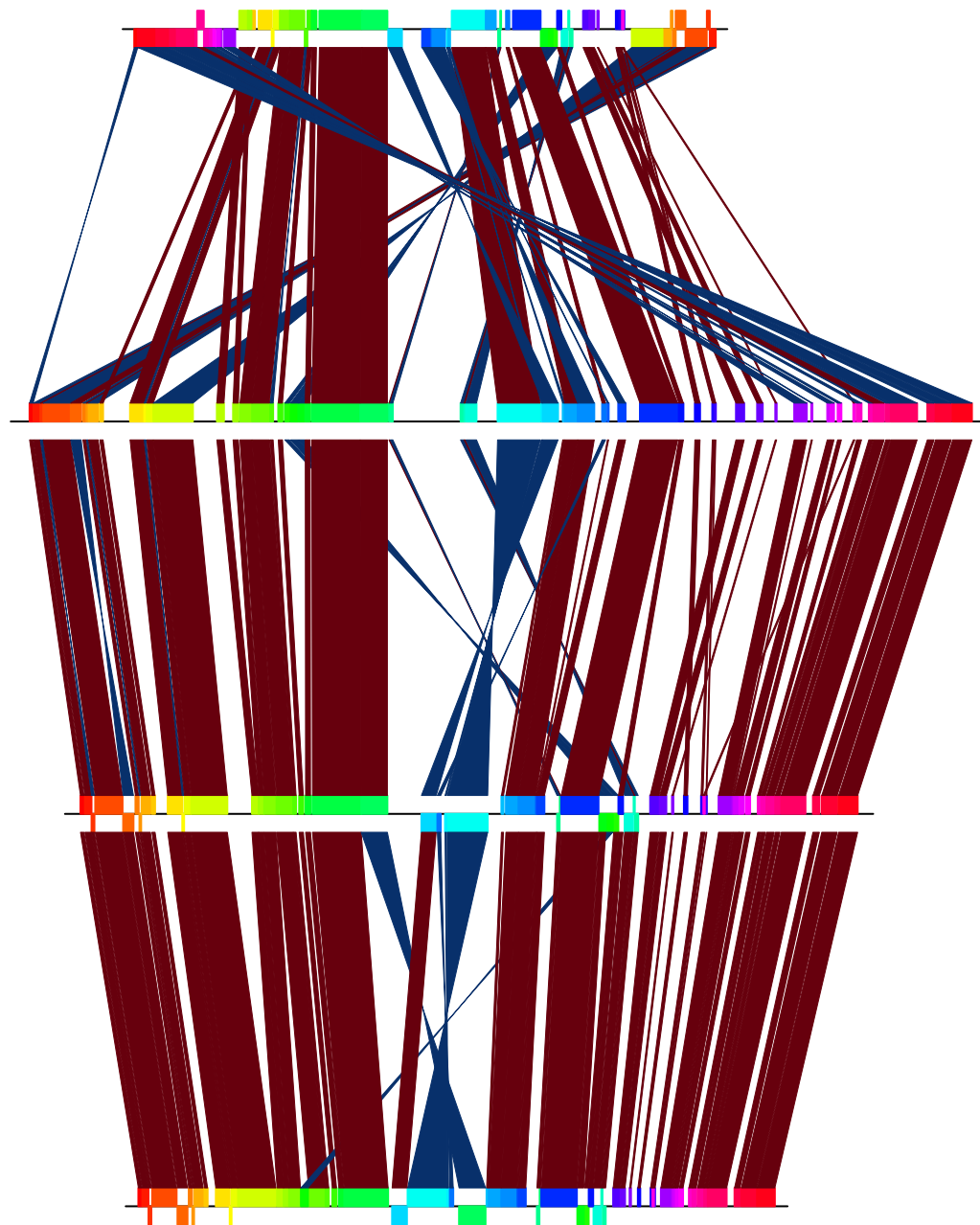
B\_grahamii

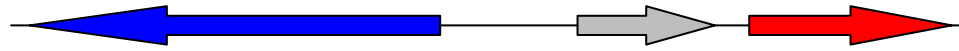
B\_henselae

B\_quintana

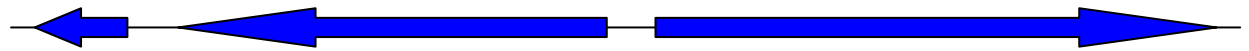
help("mauve\_bbone")

500 kb

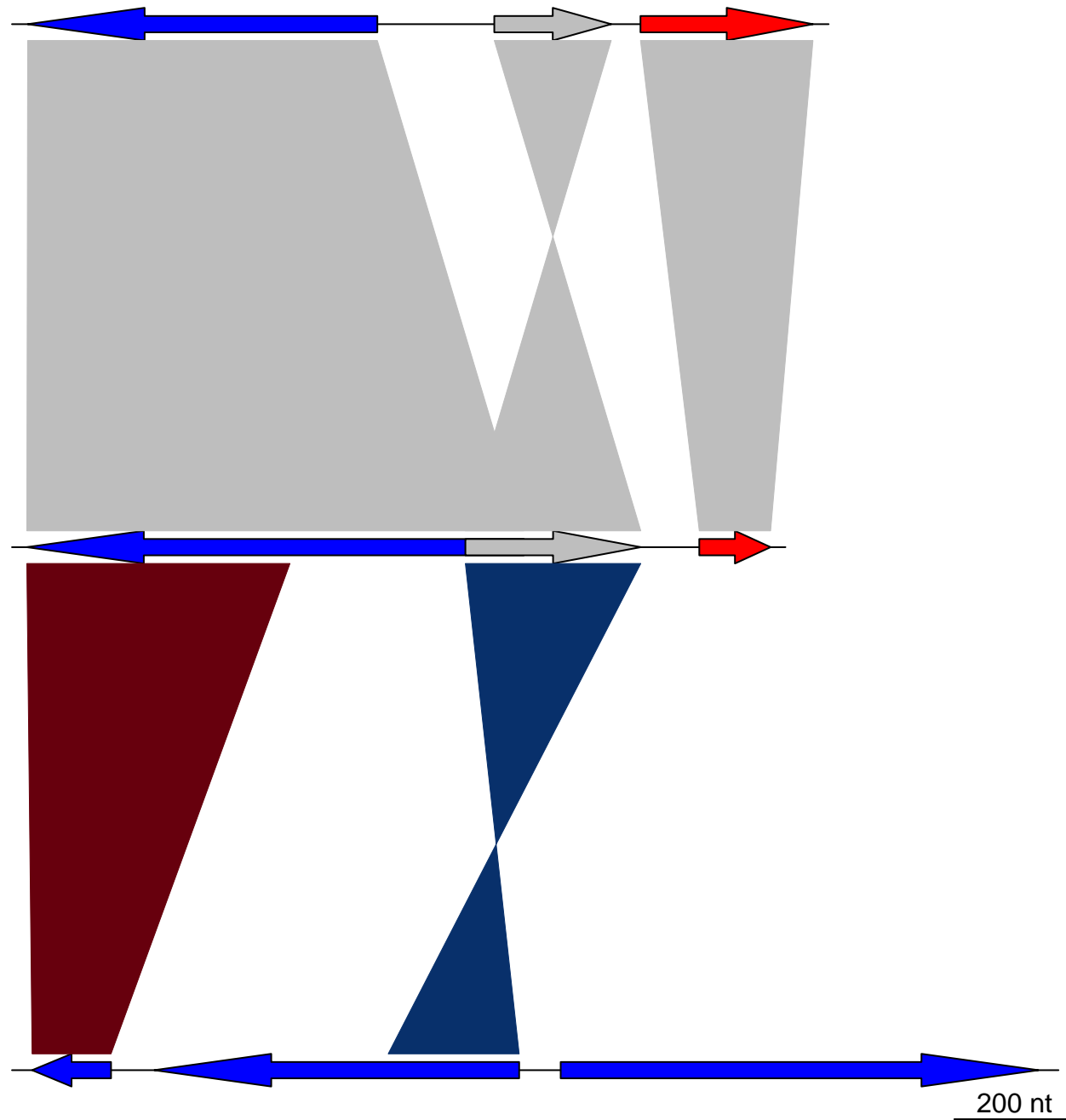




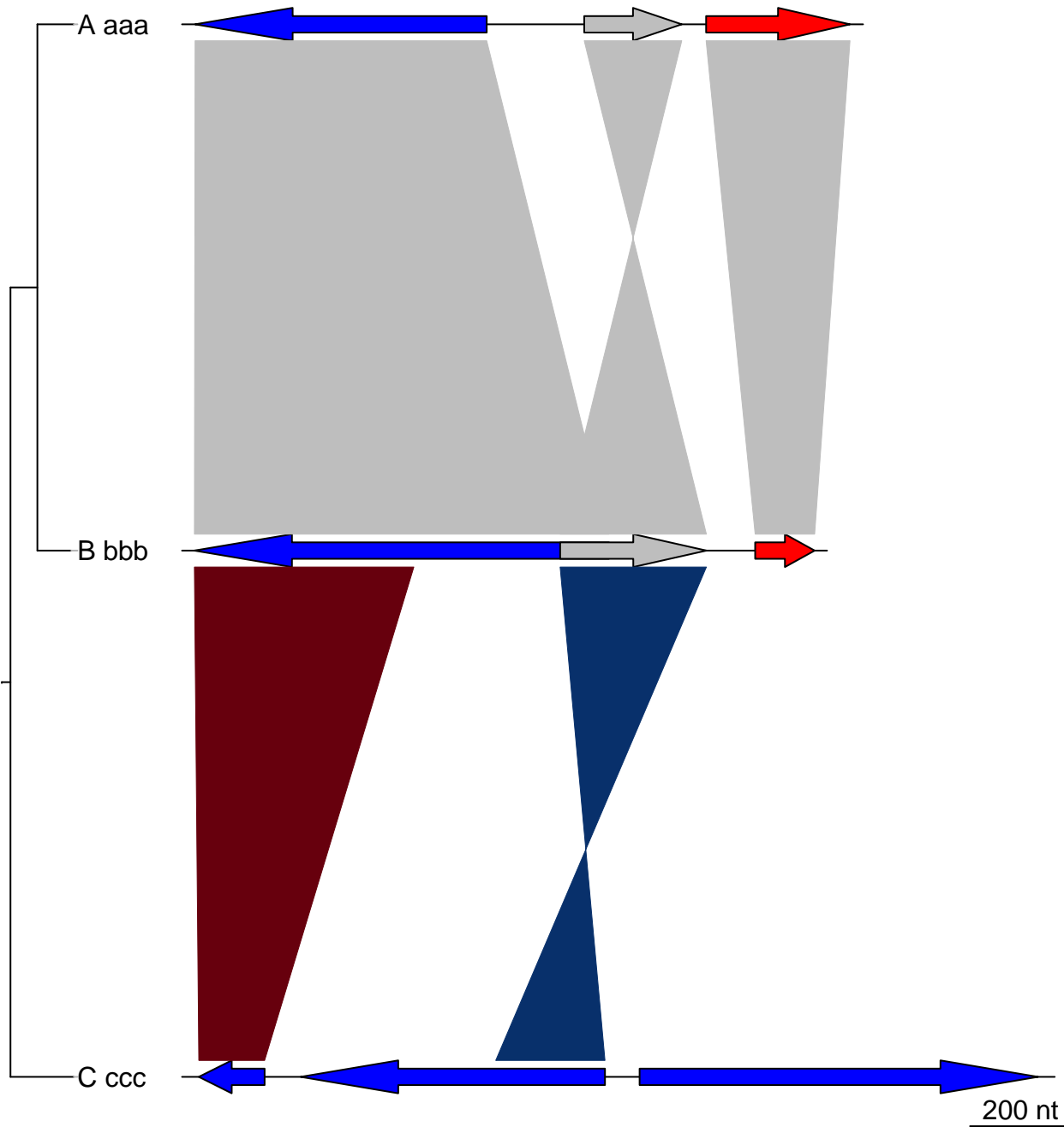
help("plot\_gene\_map")



200 nt

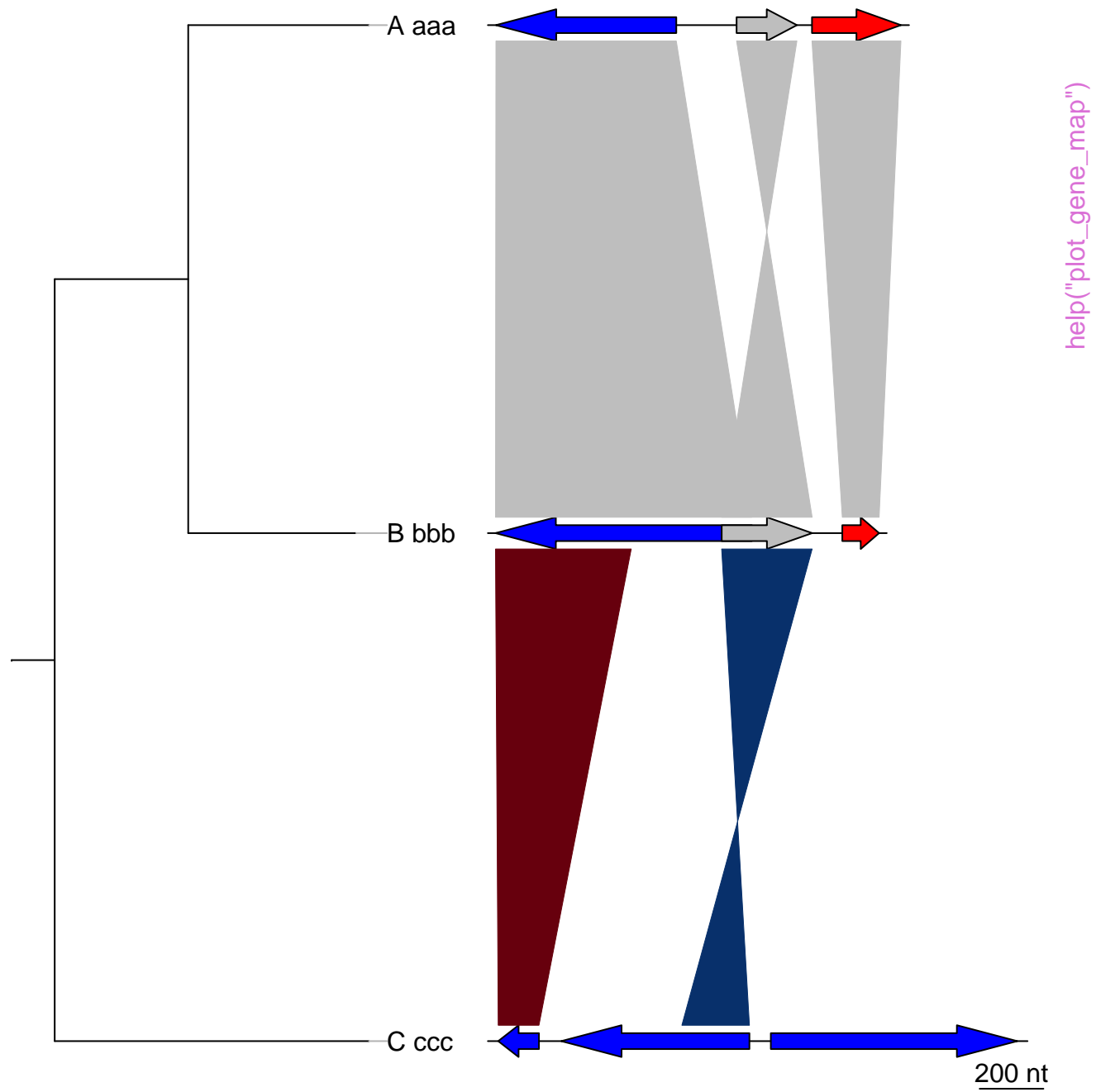


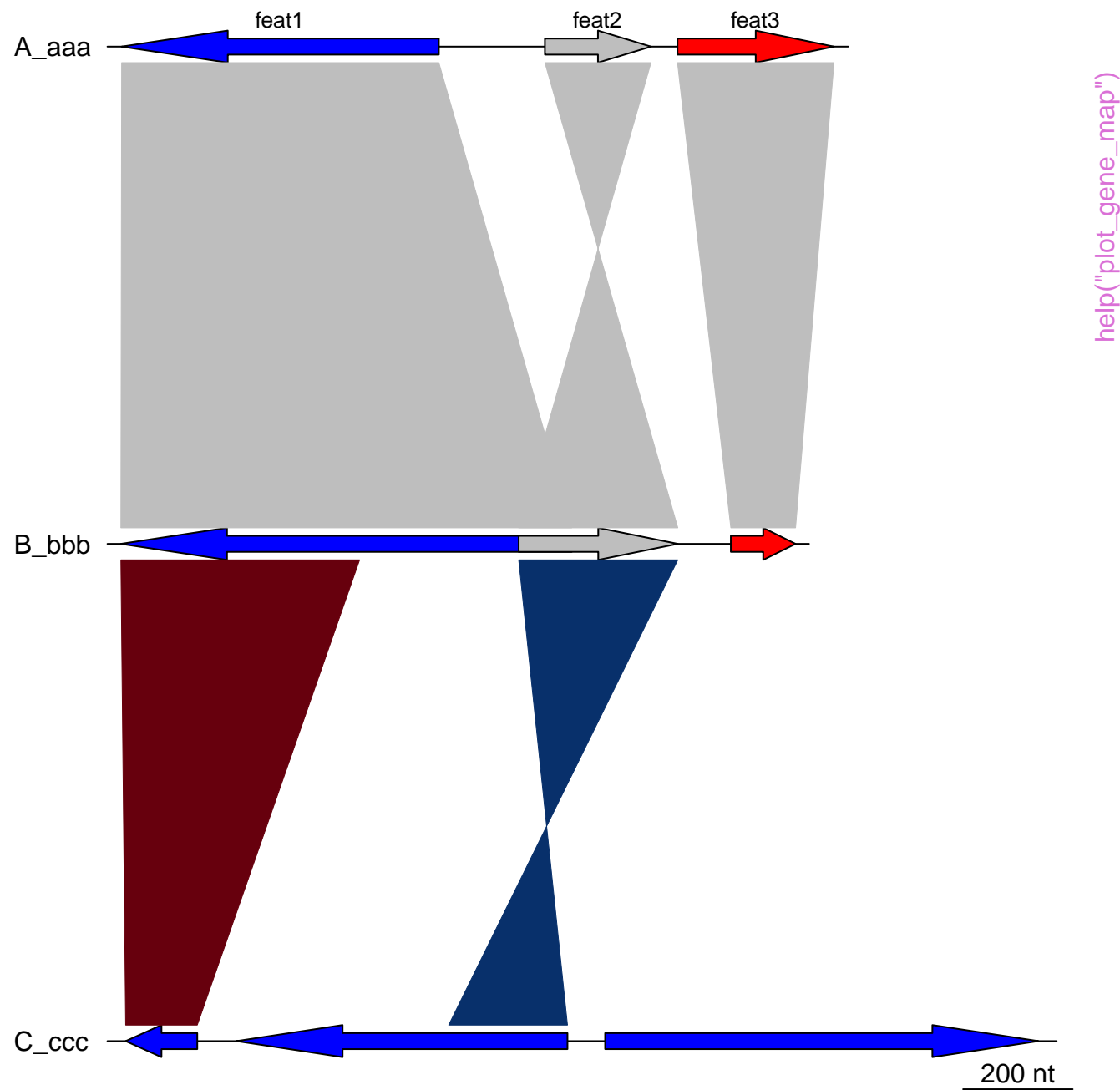
help("plot\_gene\_map")

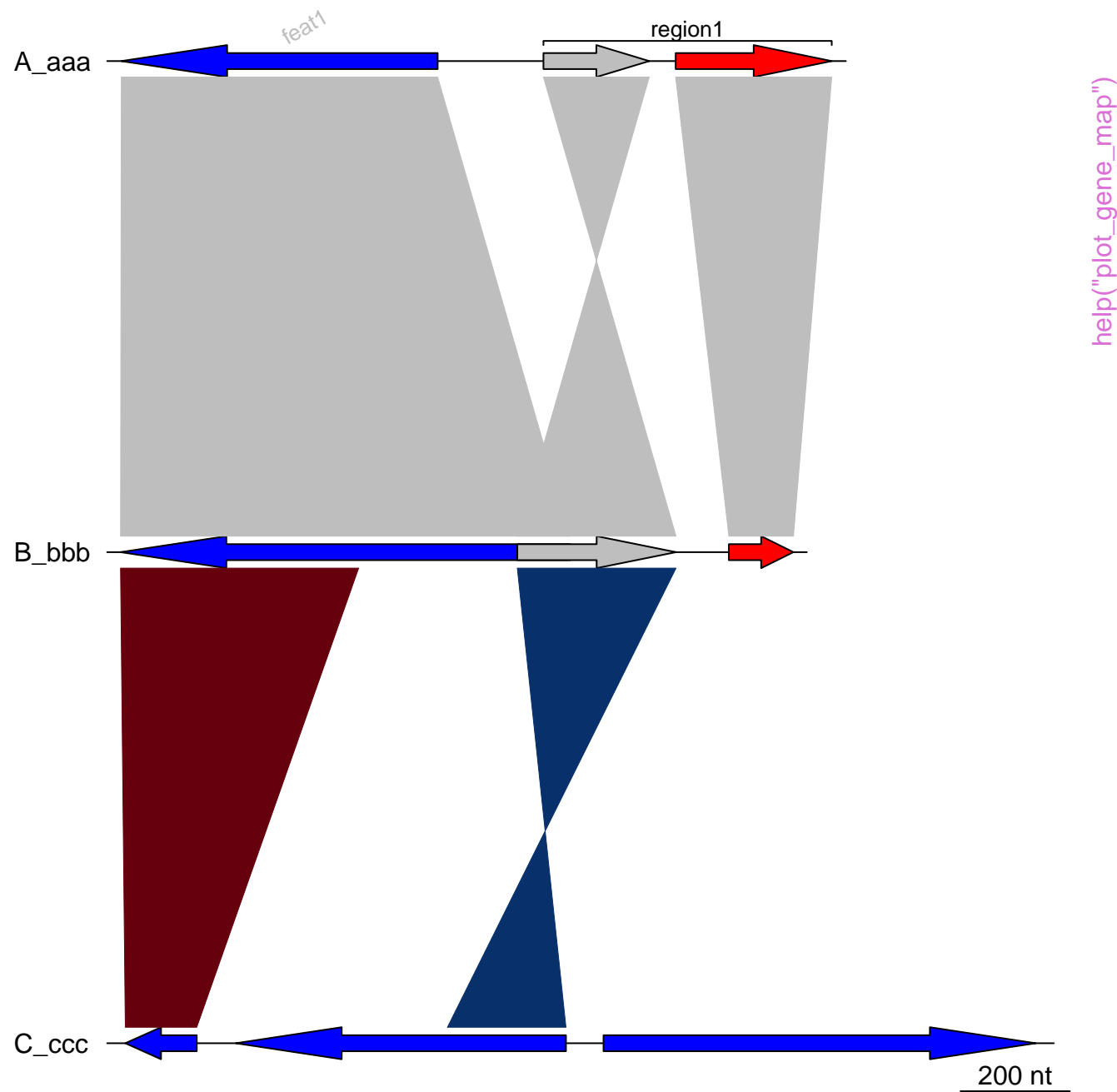


help("plot\_gene\_map")







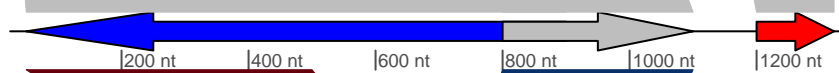


A\_aaa

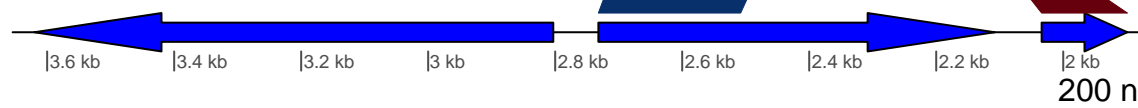


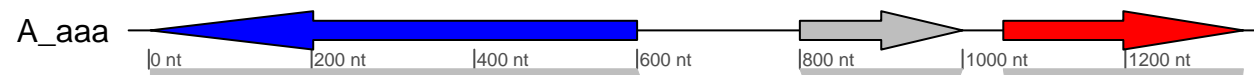
help("plot\_gene\_map")

B\_bbb

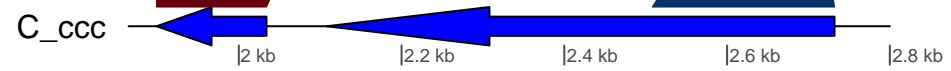
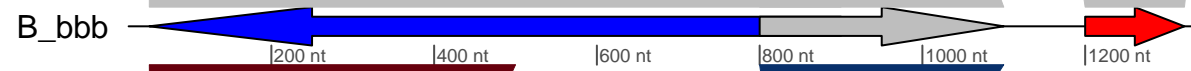


C\_ccc

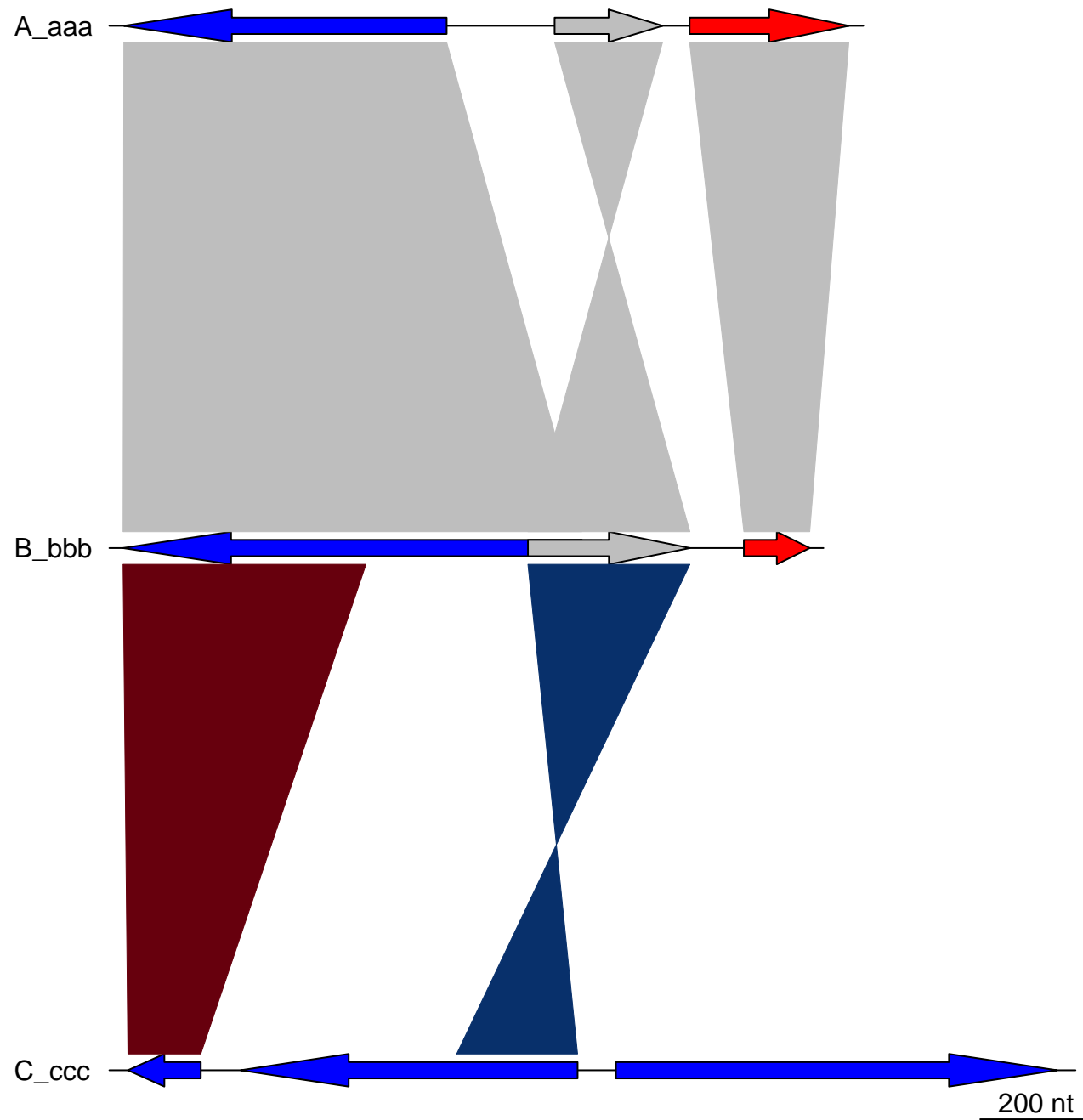




help("plot\_gene\_map")



200 nt



help("plot\_gene\_map")

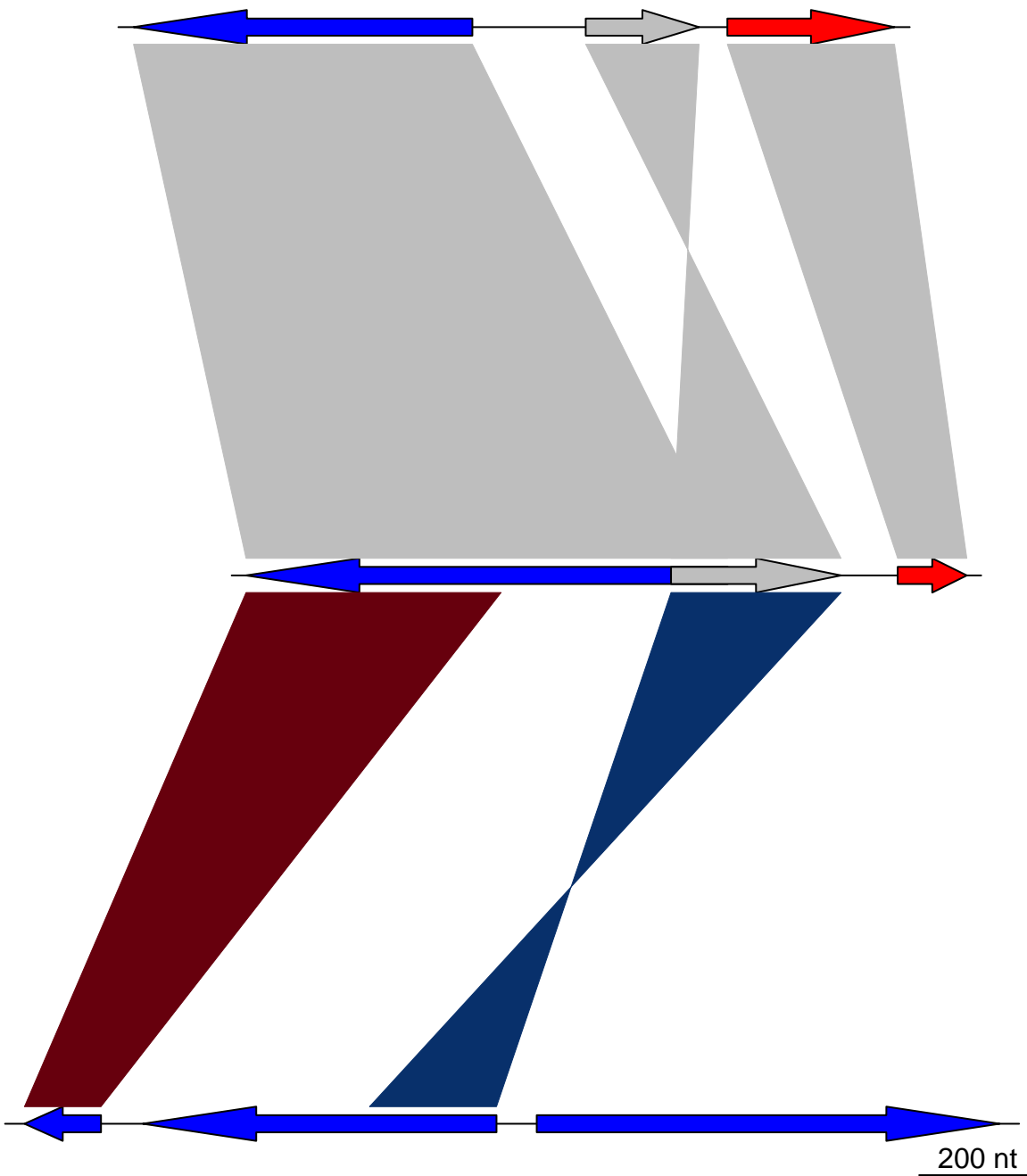
A\_aaa

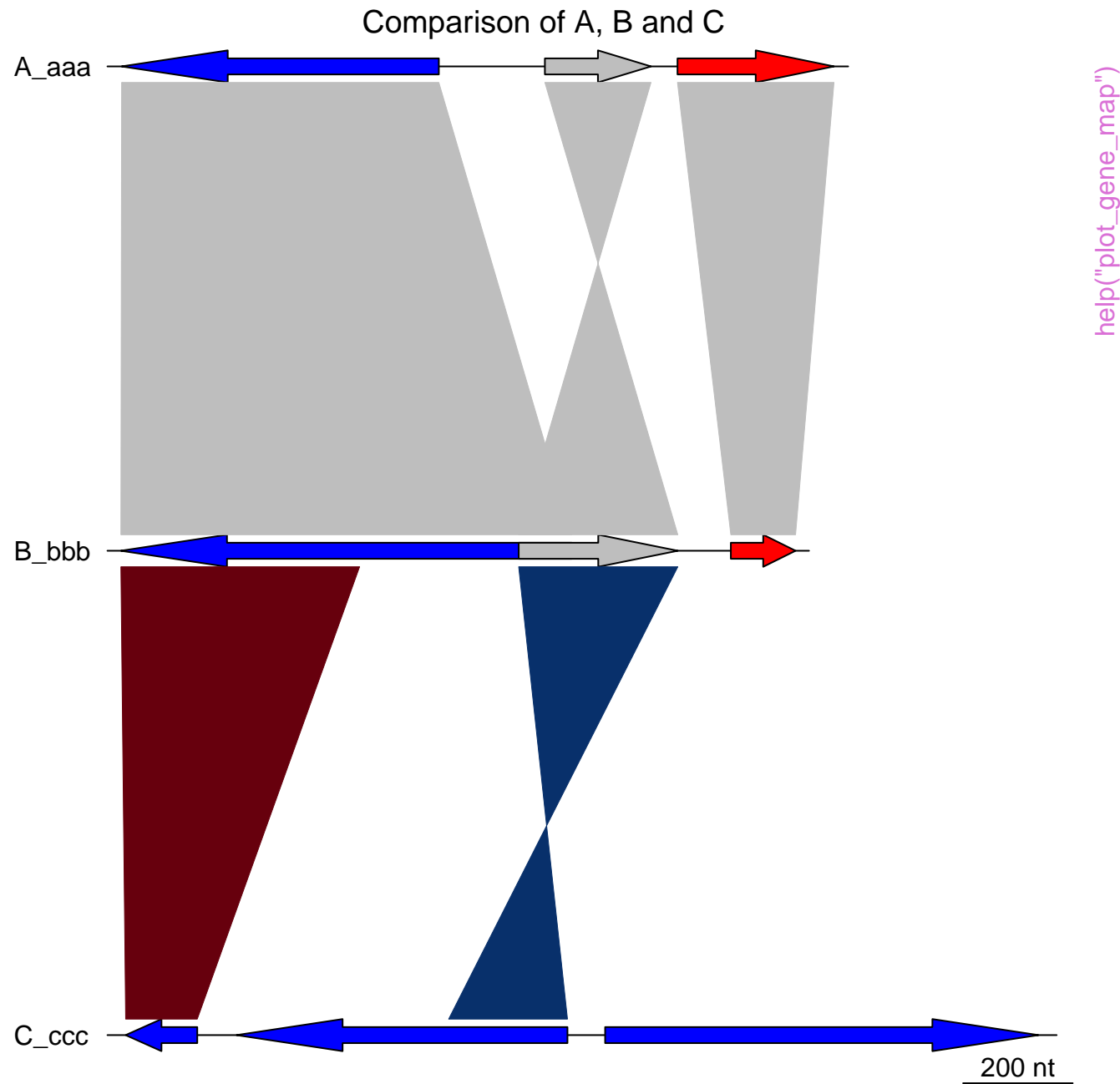
B\_bbb

C\_ccc

help("plot\_gene\_map")

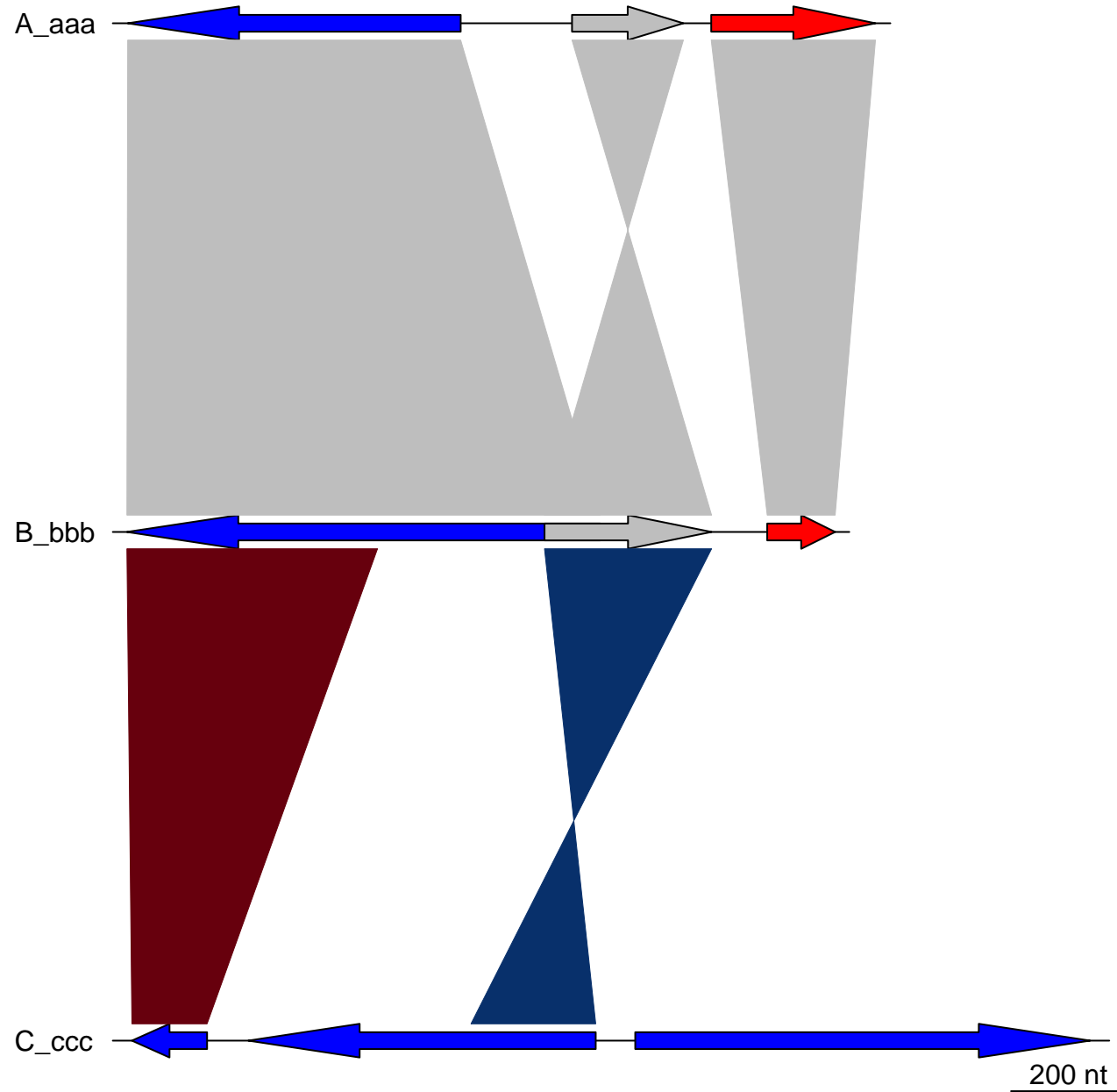
200 nt



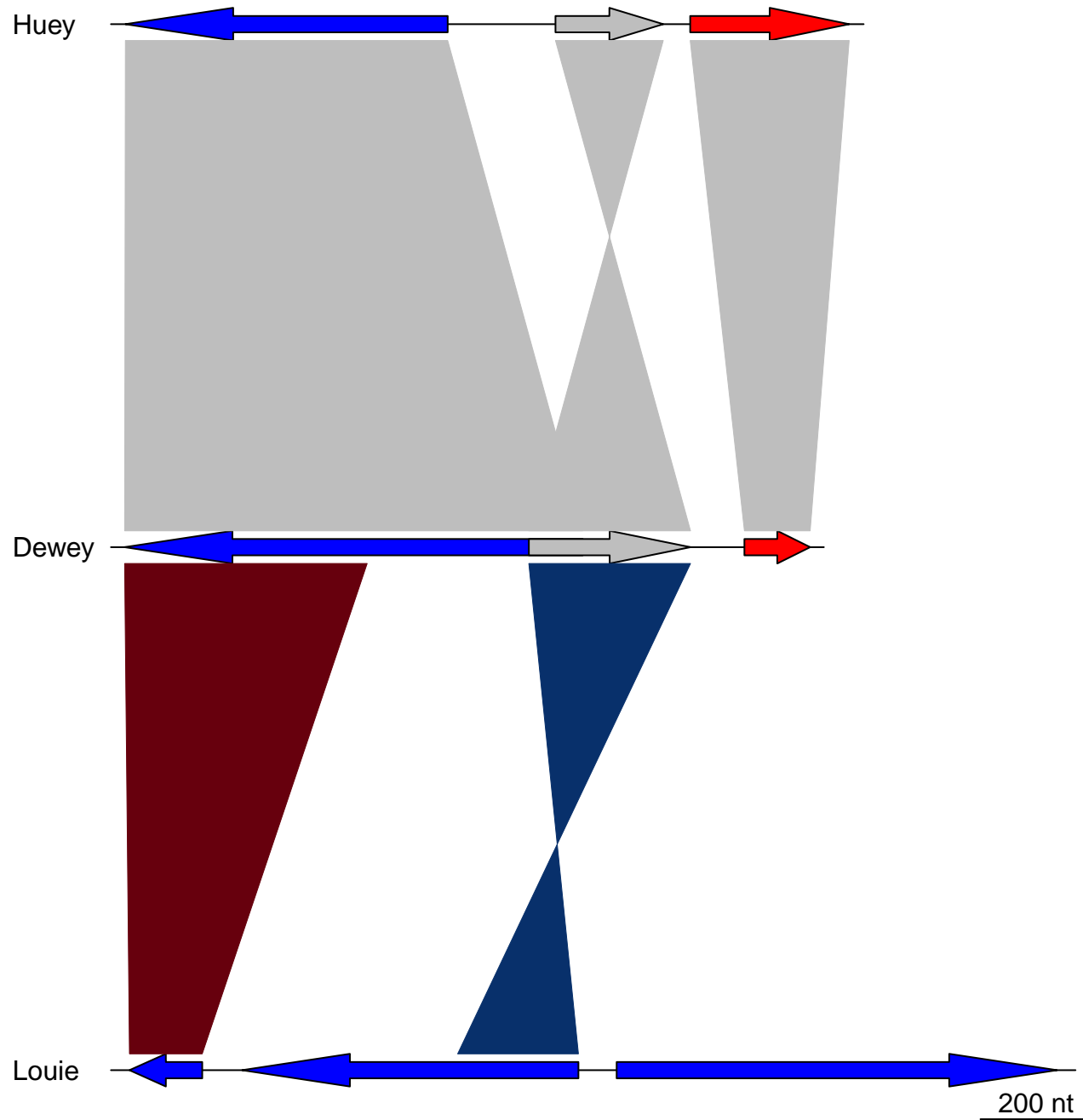




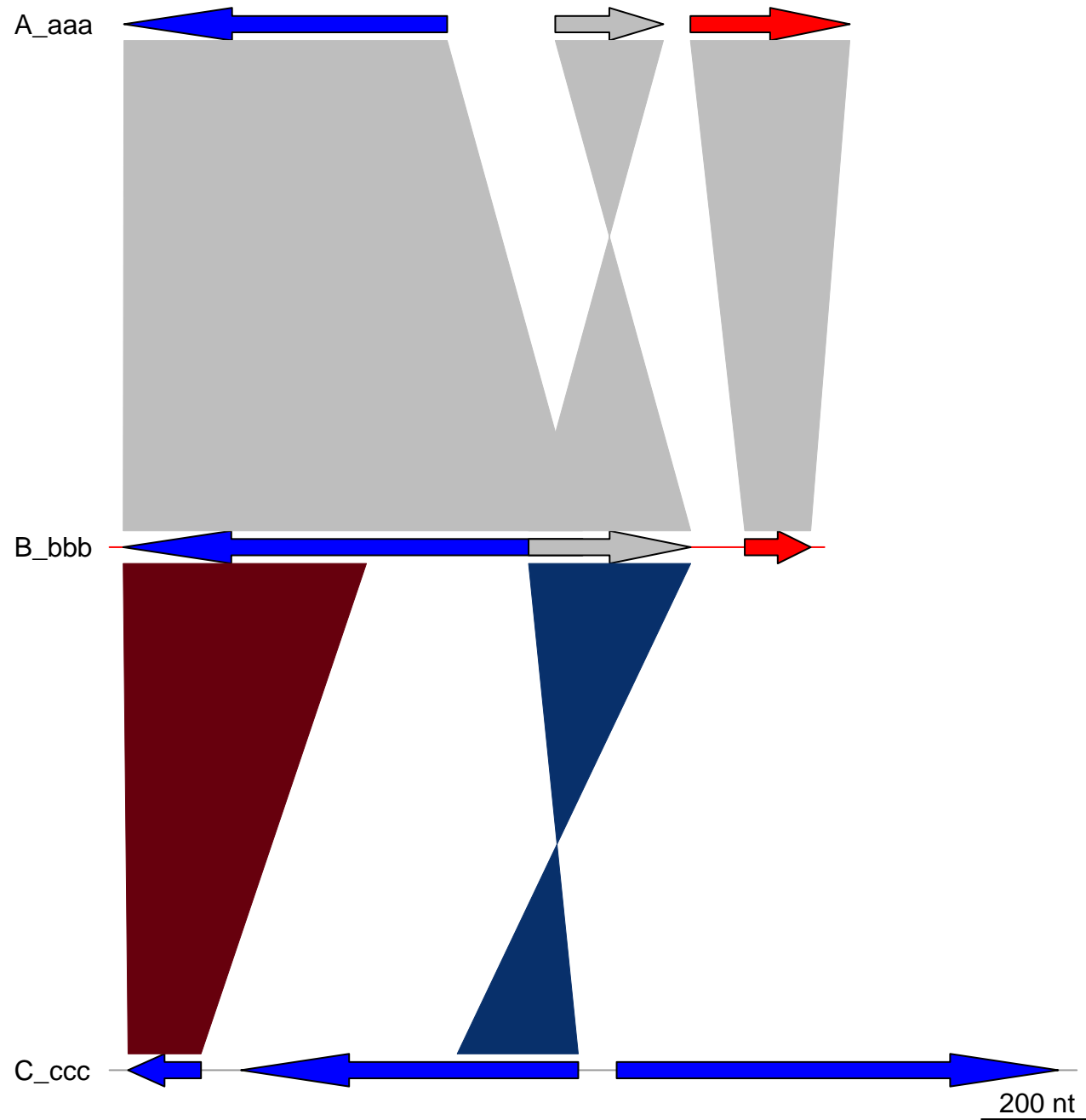
# Comparison of A, B and C



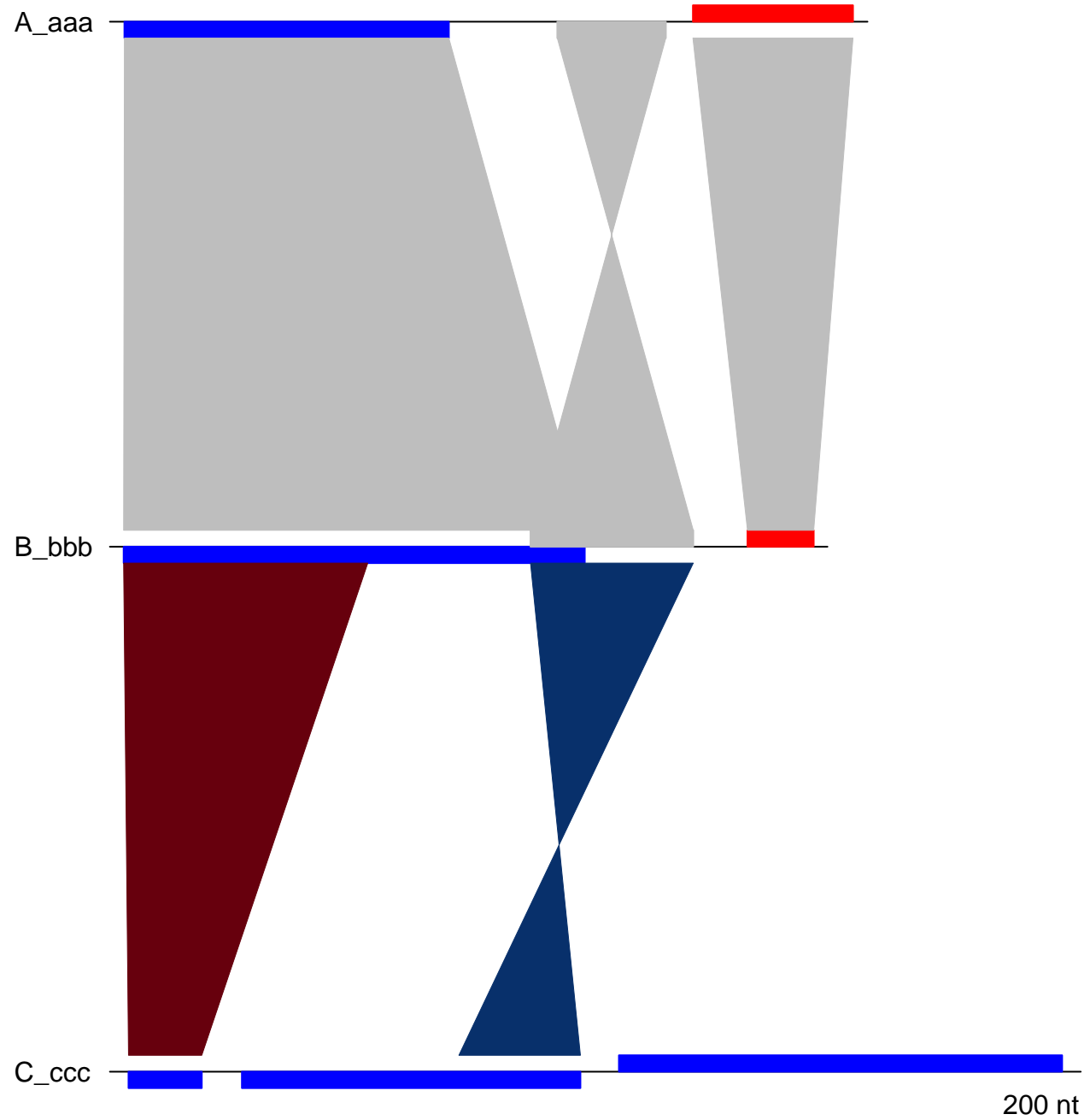
help("plot\_gene\_map")



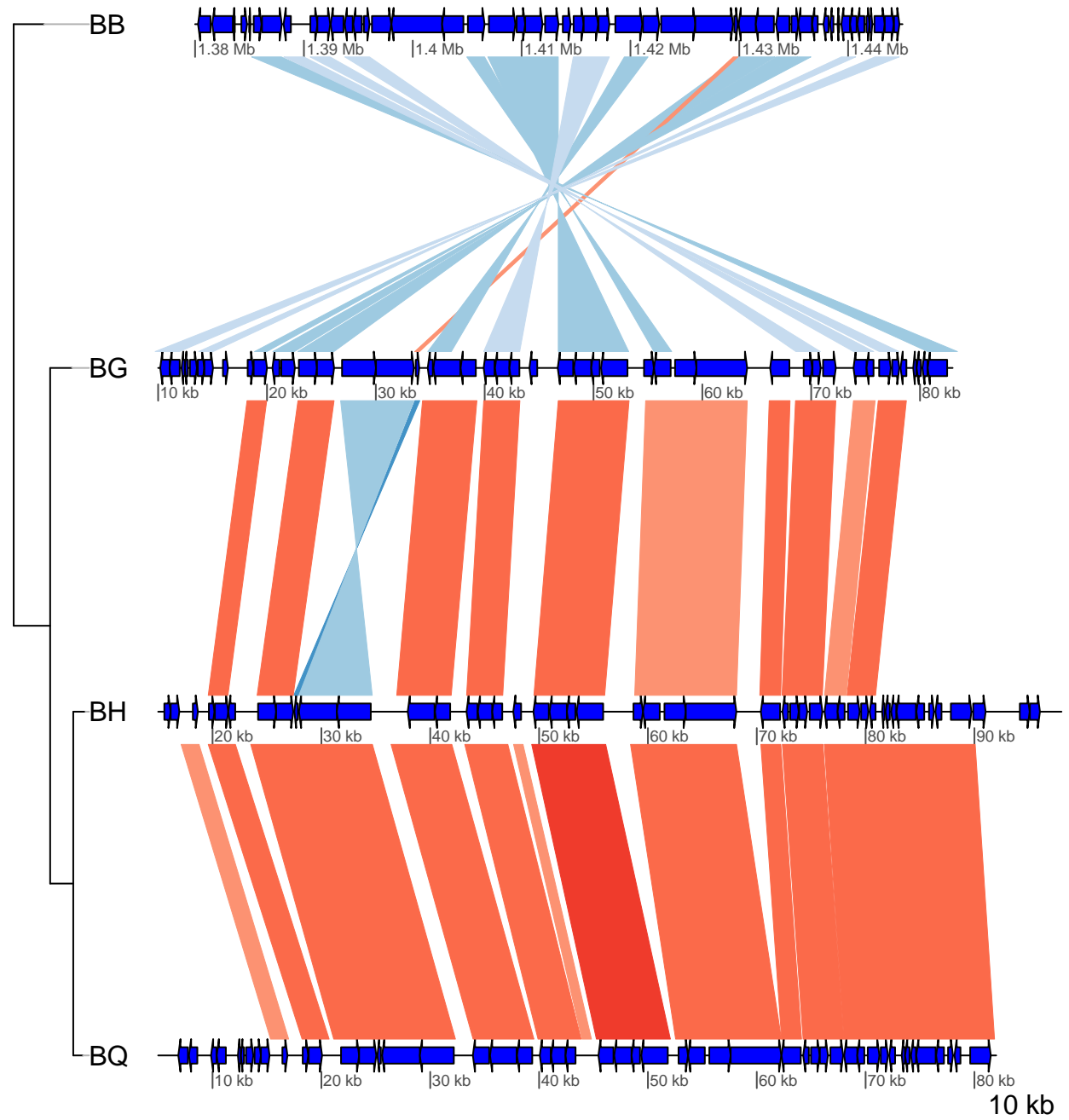
help("plot\_gene\_map")



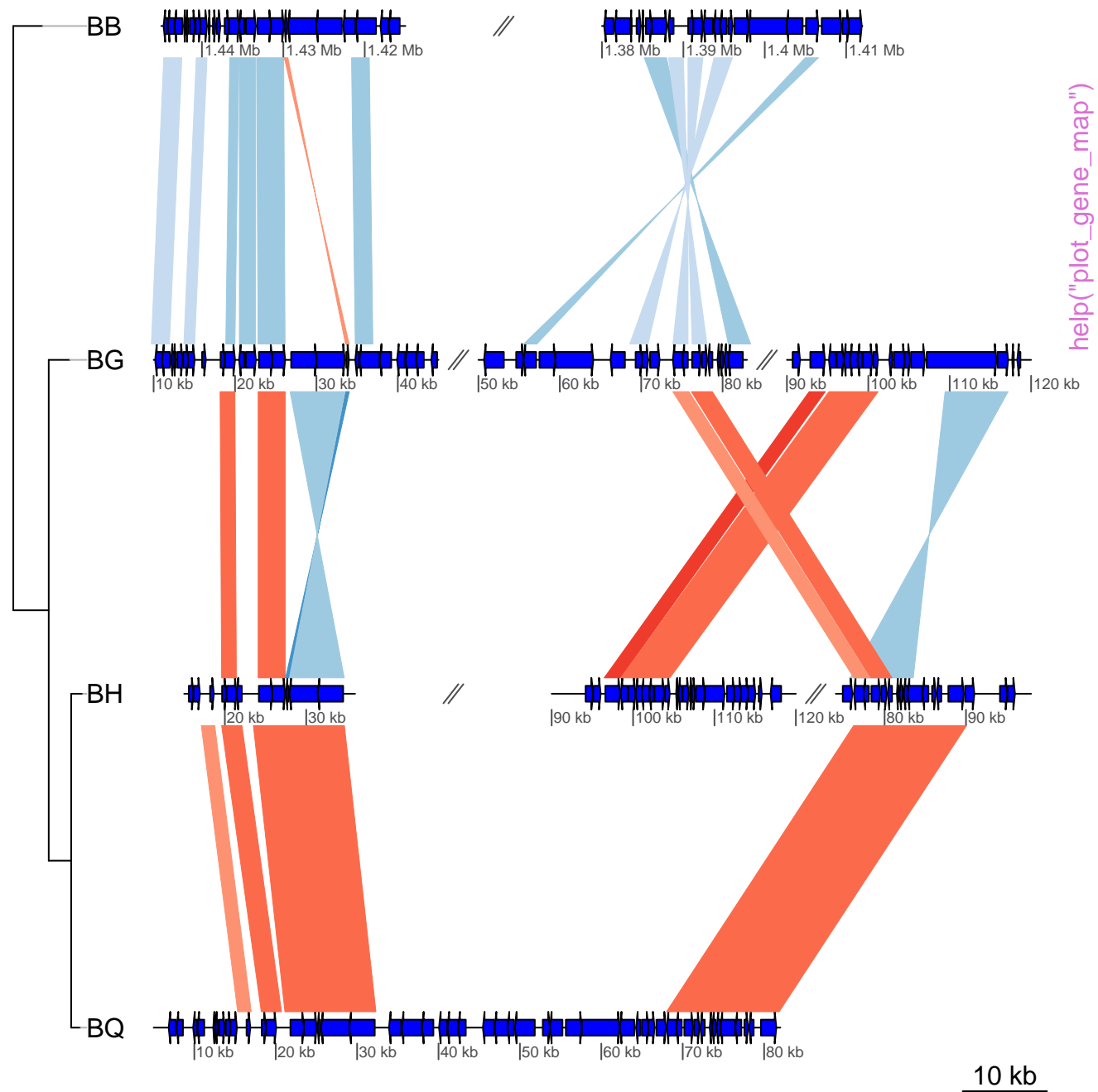
help("plot\_gene\_map")

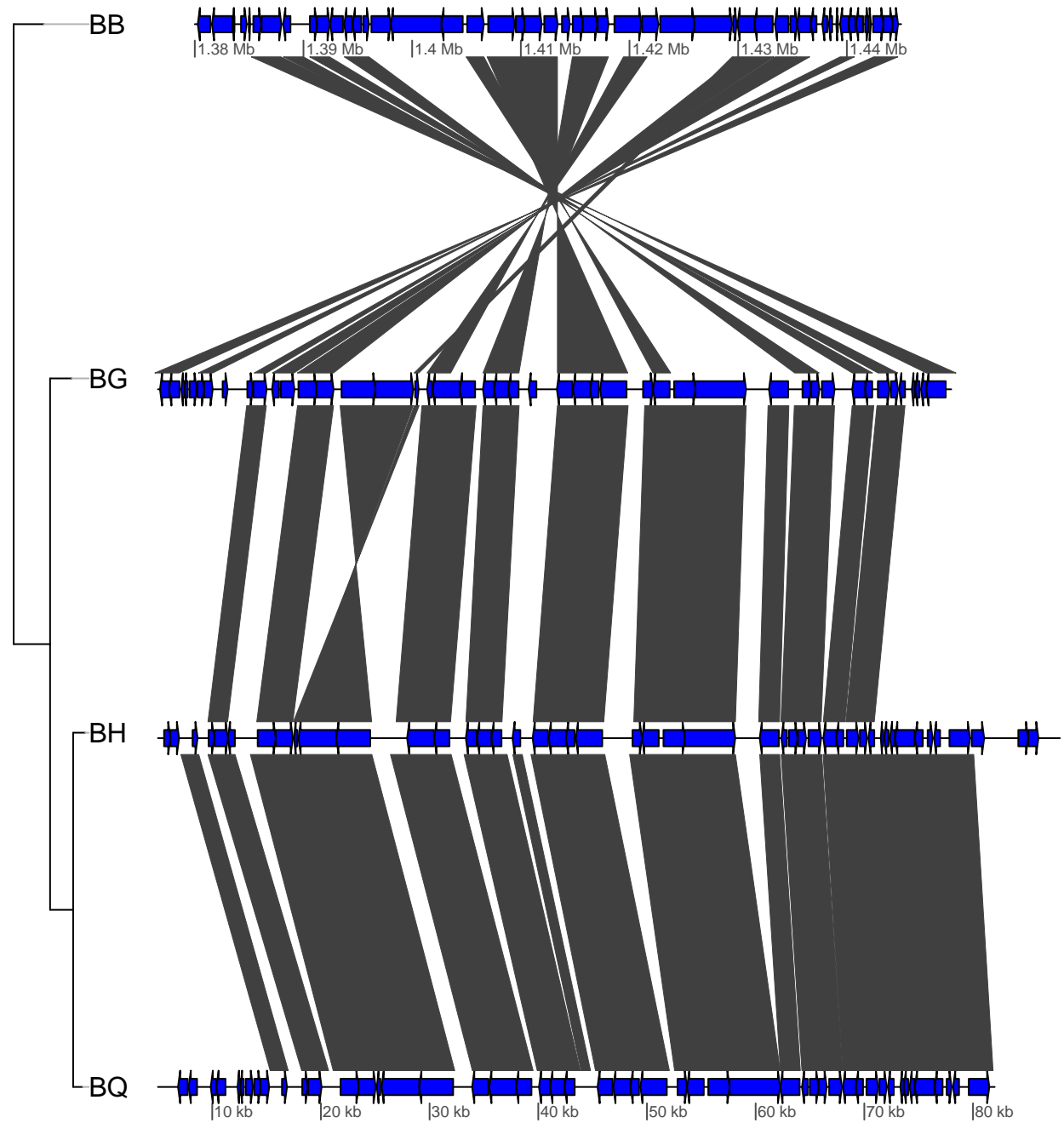


help("plot\_gene\_map")

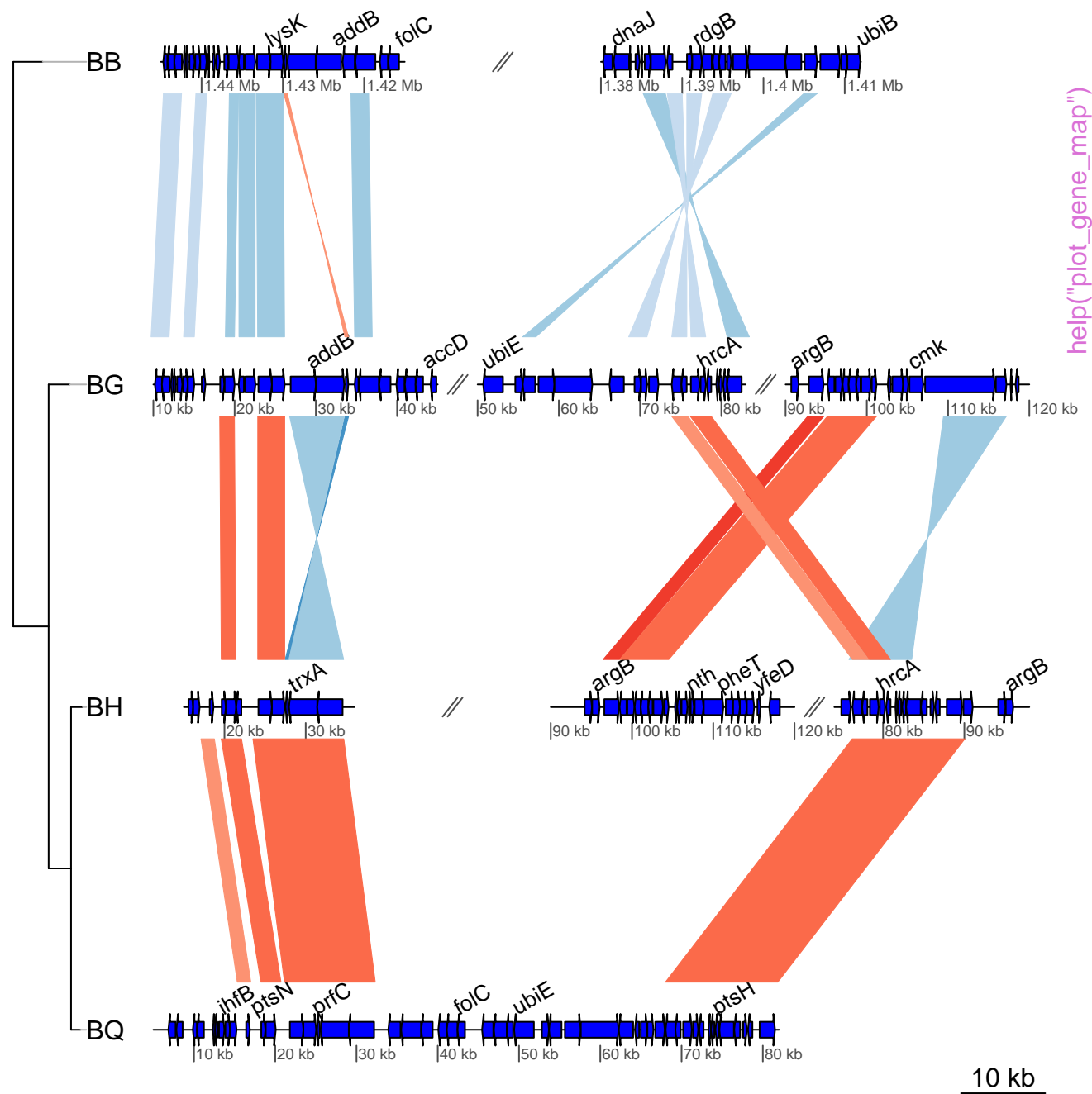


help("plot\_gene\_map")

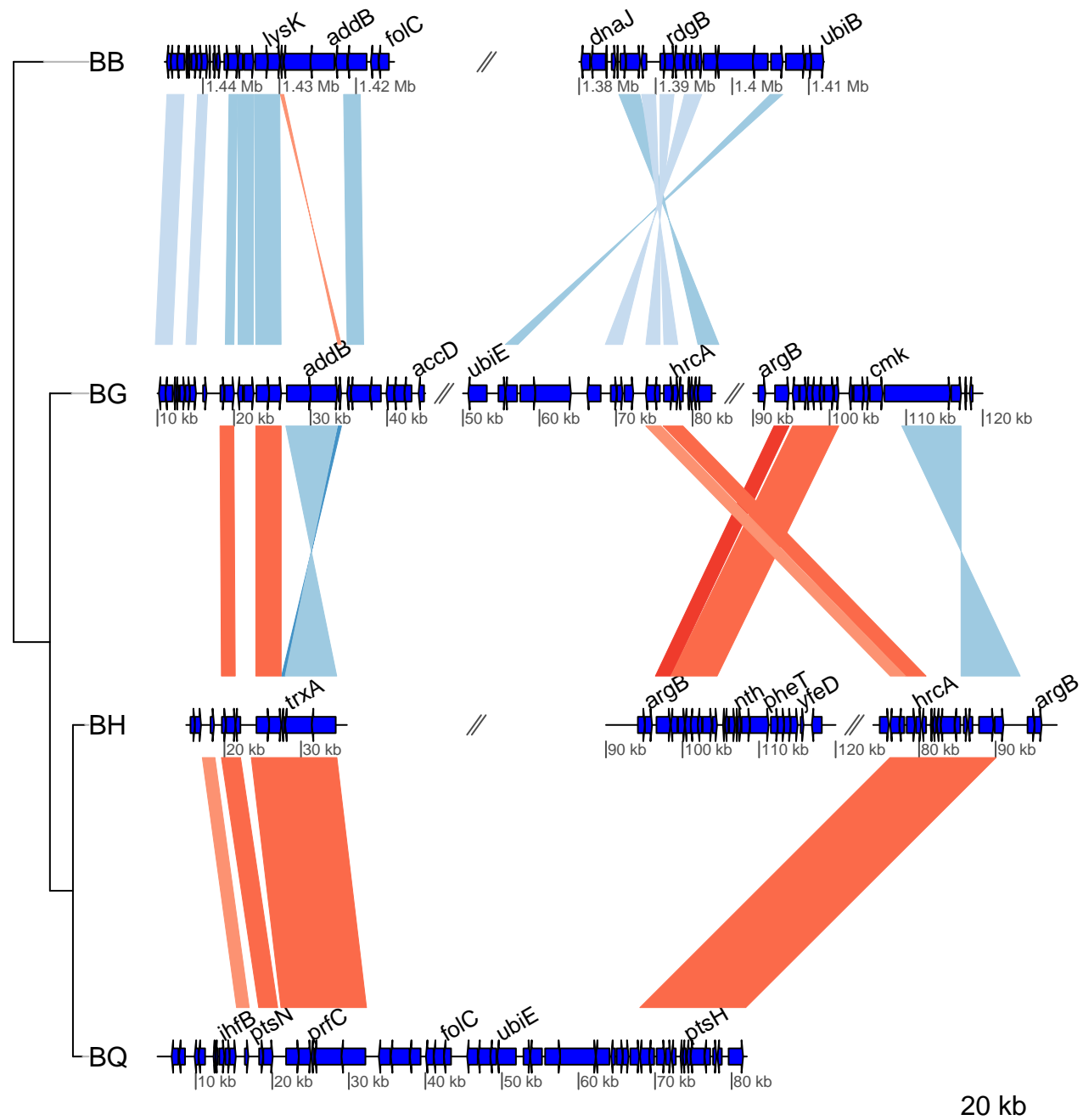




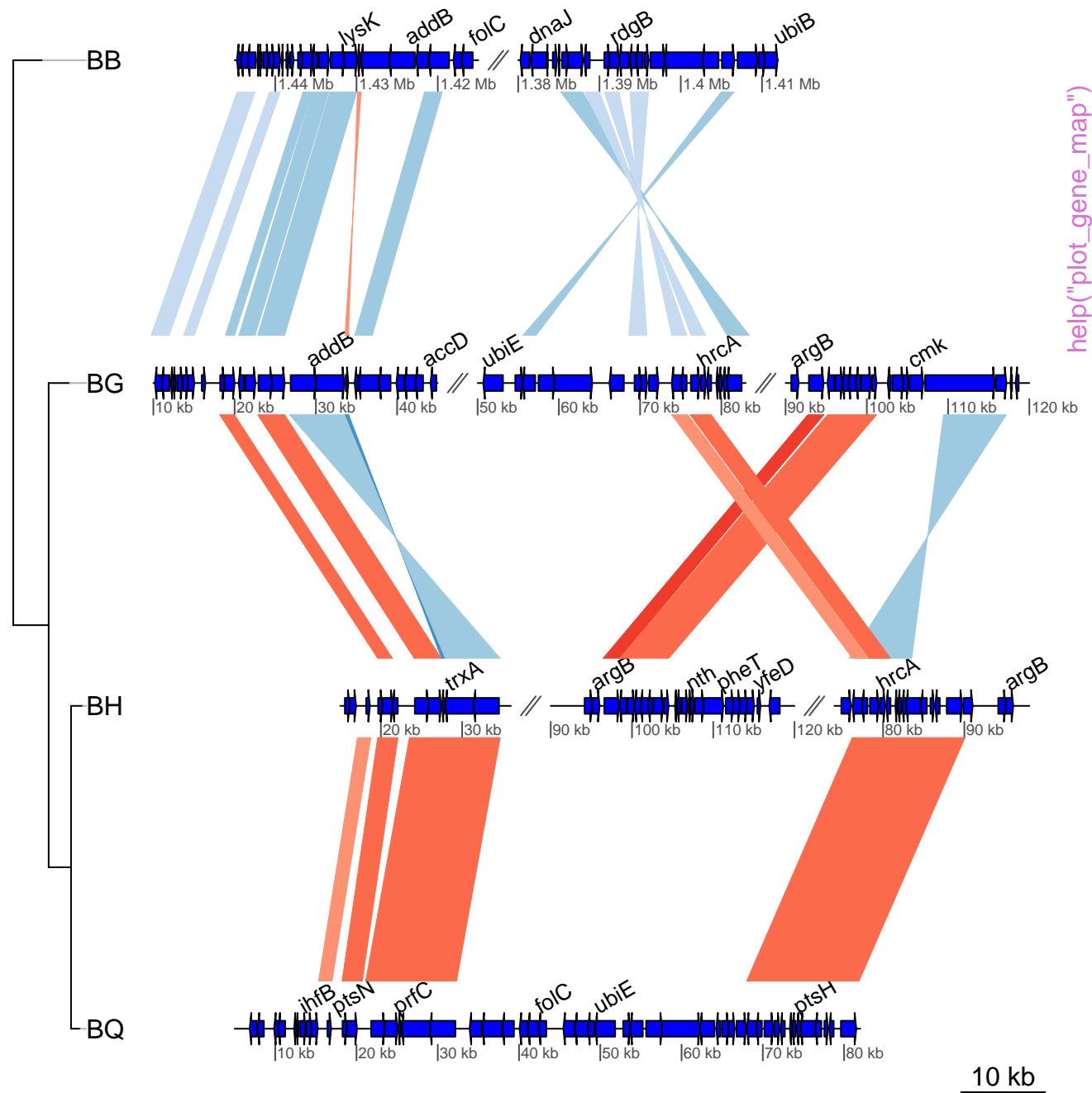
help("plot\_gene\_map")

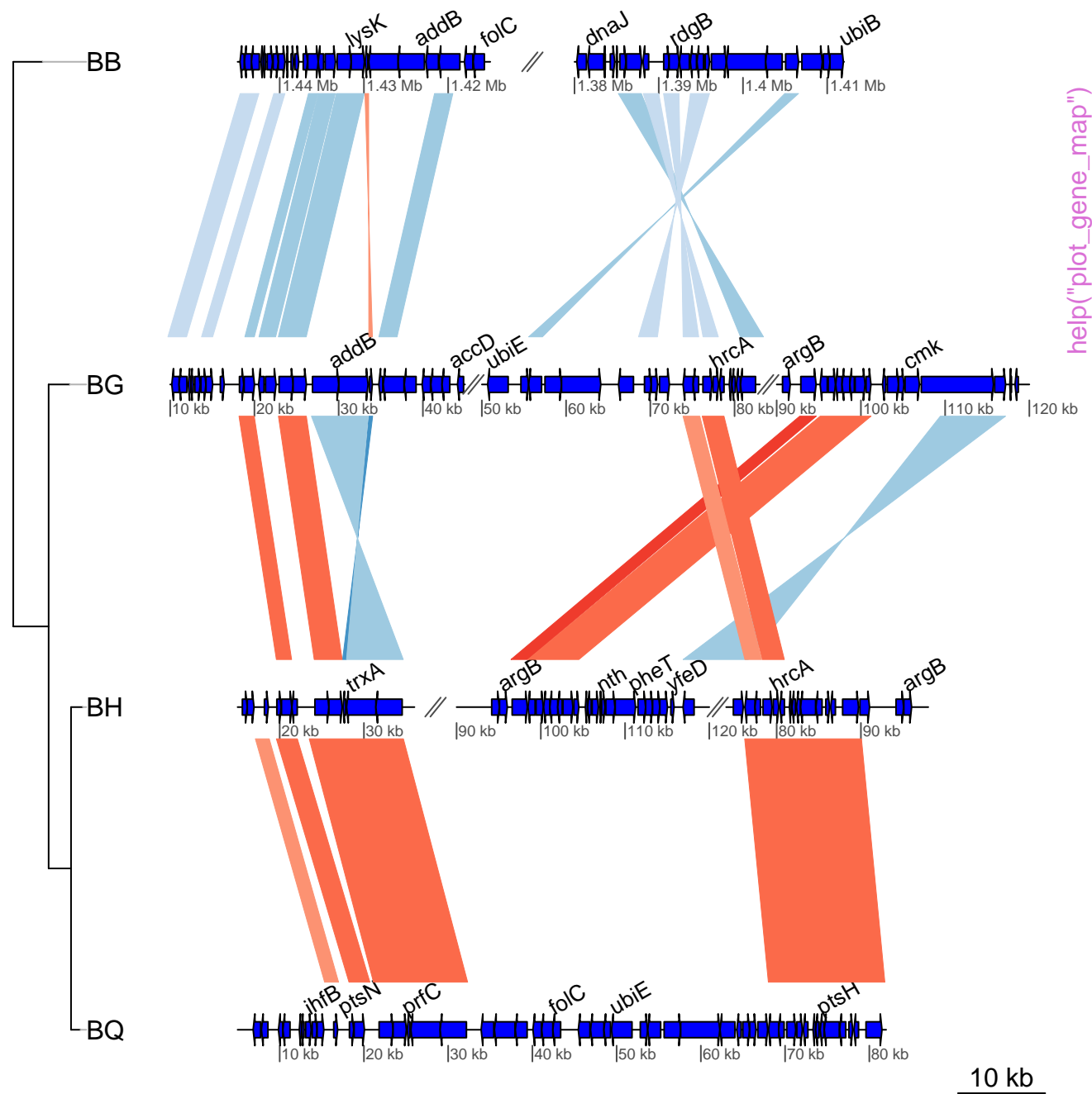


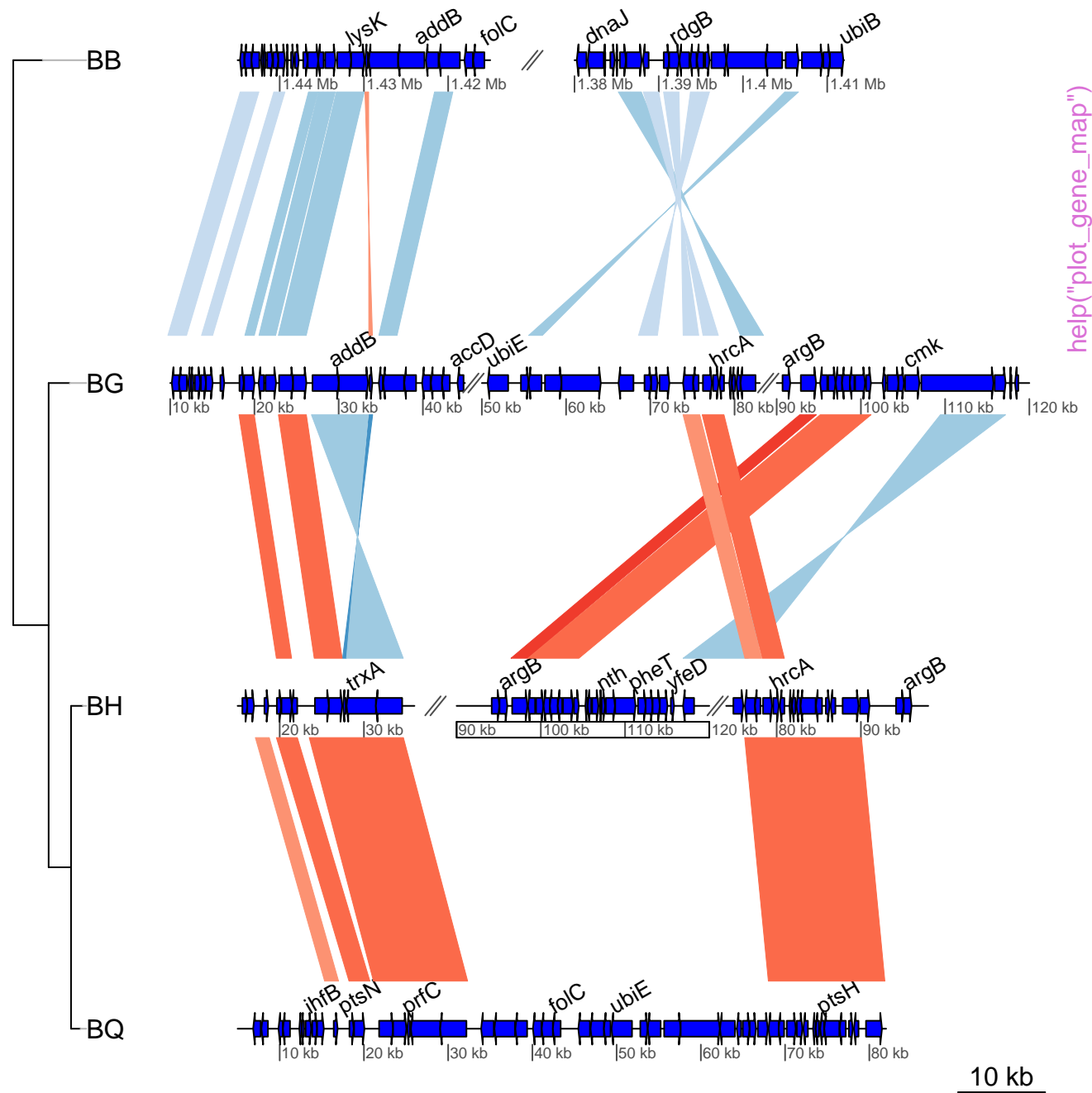


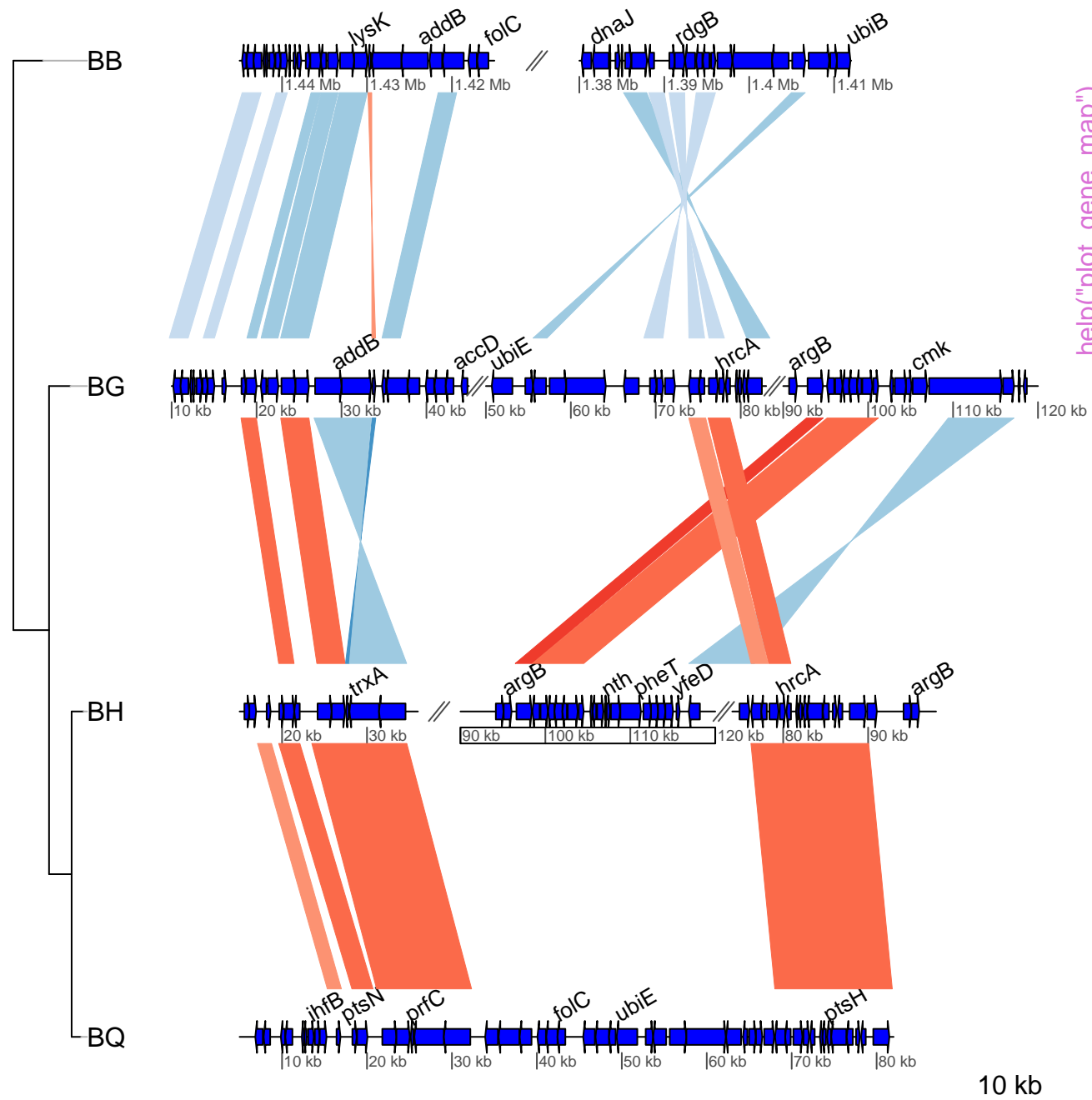


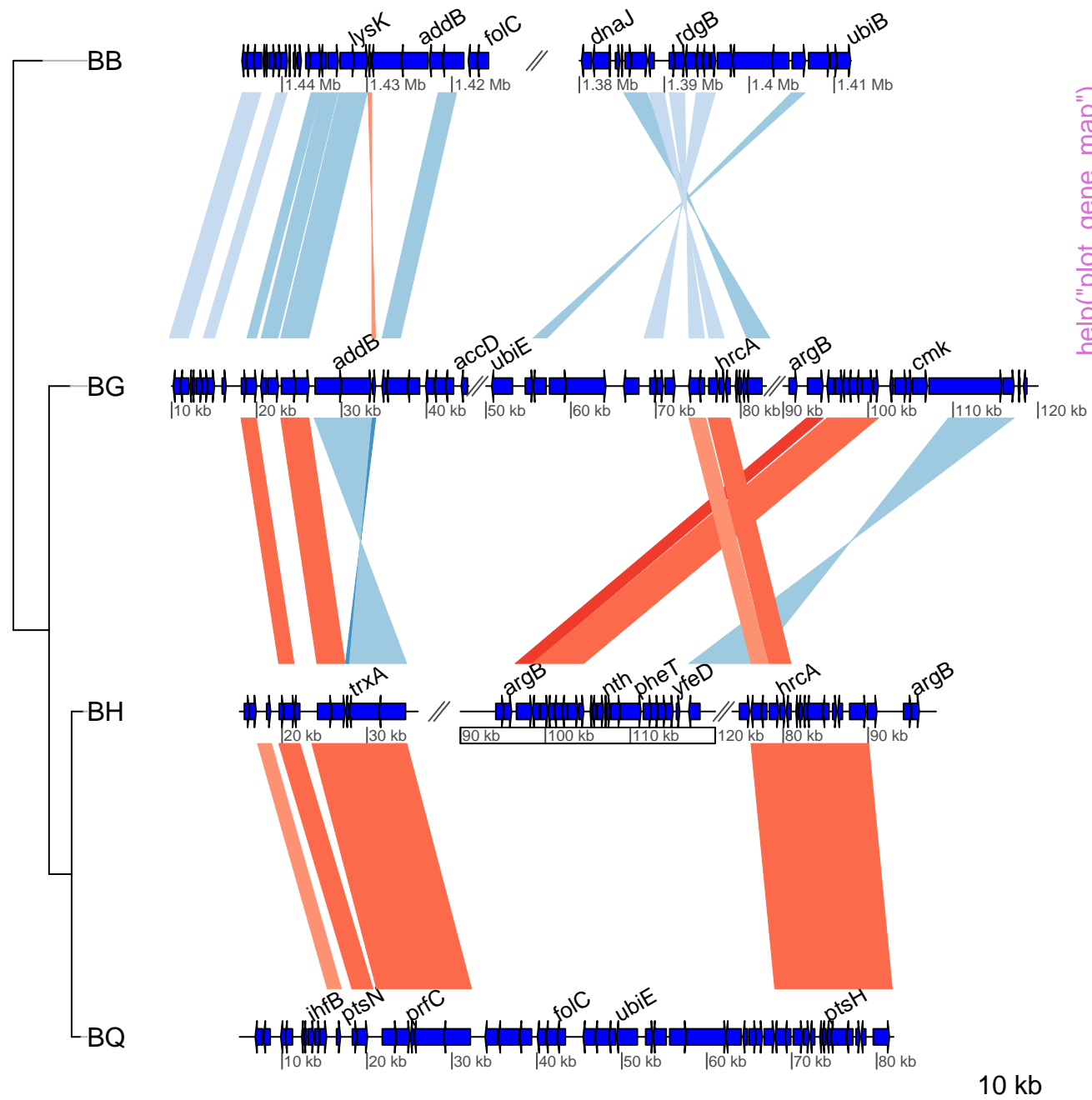
help("plot\_gene\_map")



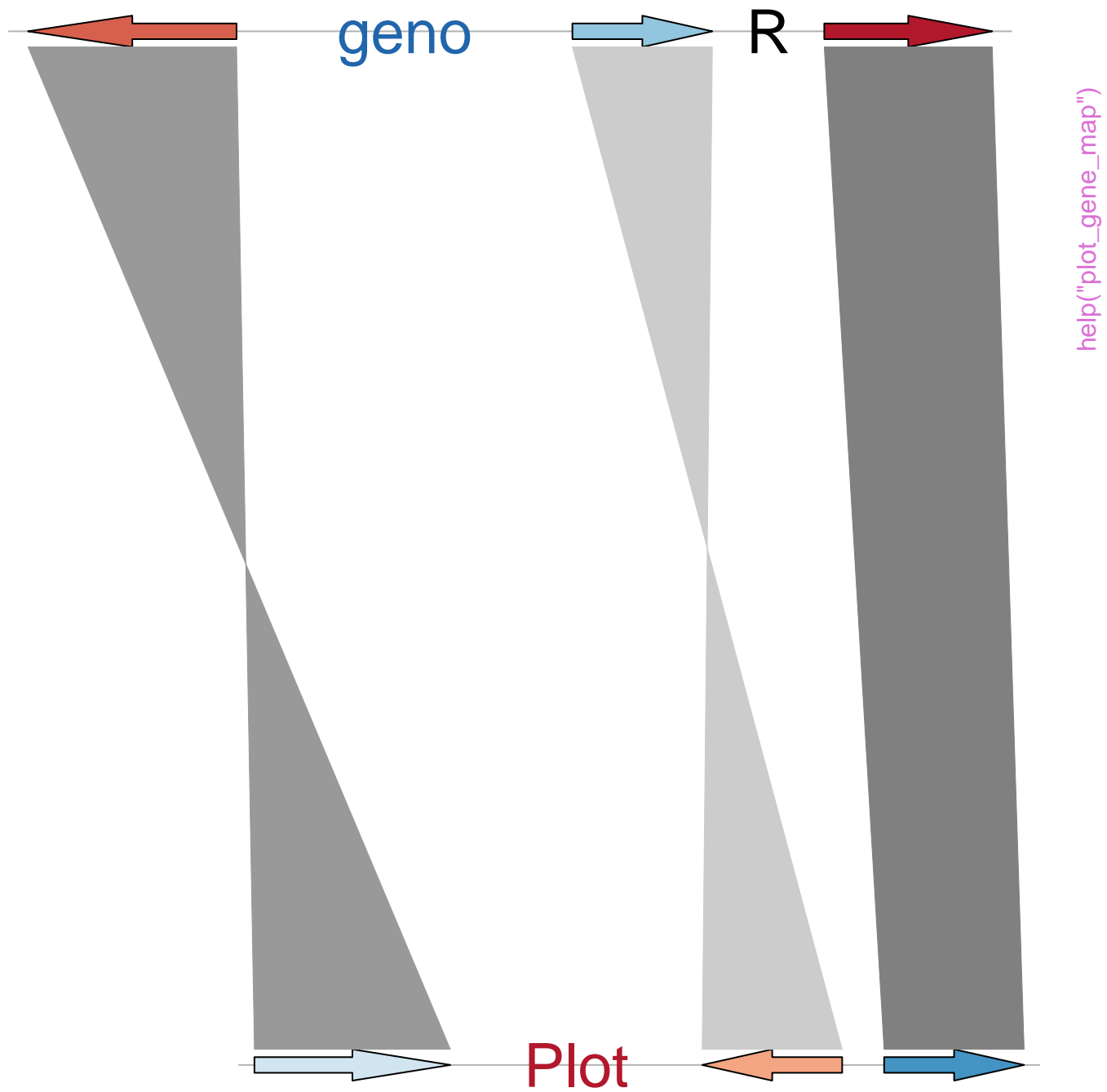








help("plot\_gene\_map")



B\_bacilliformis

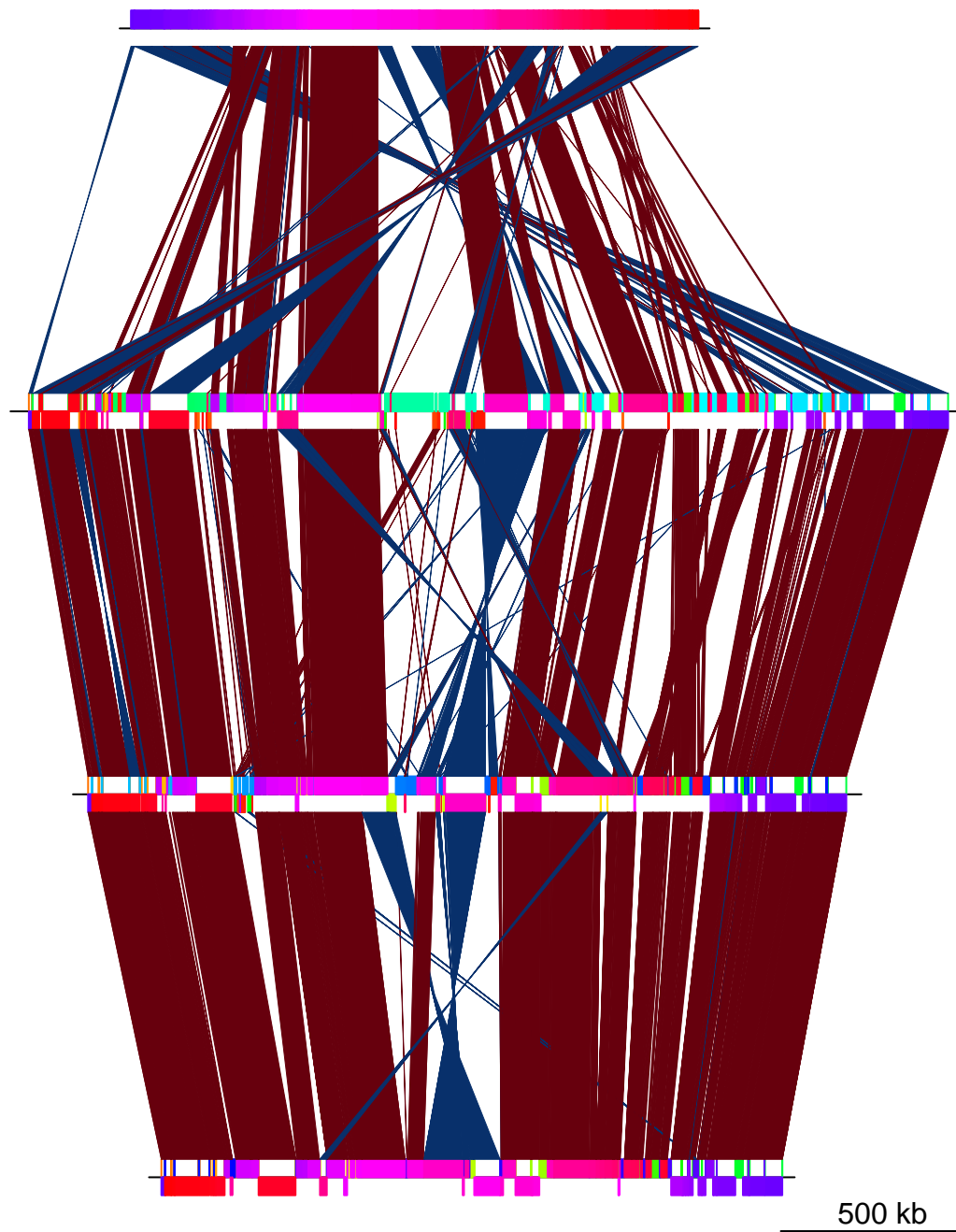
B\_grahamii

B\_henselae

B\_quintana

help("read\_functions")

500 kb





B\_bacilliformis

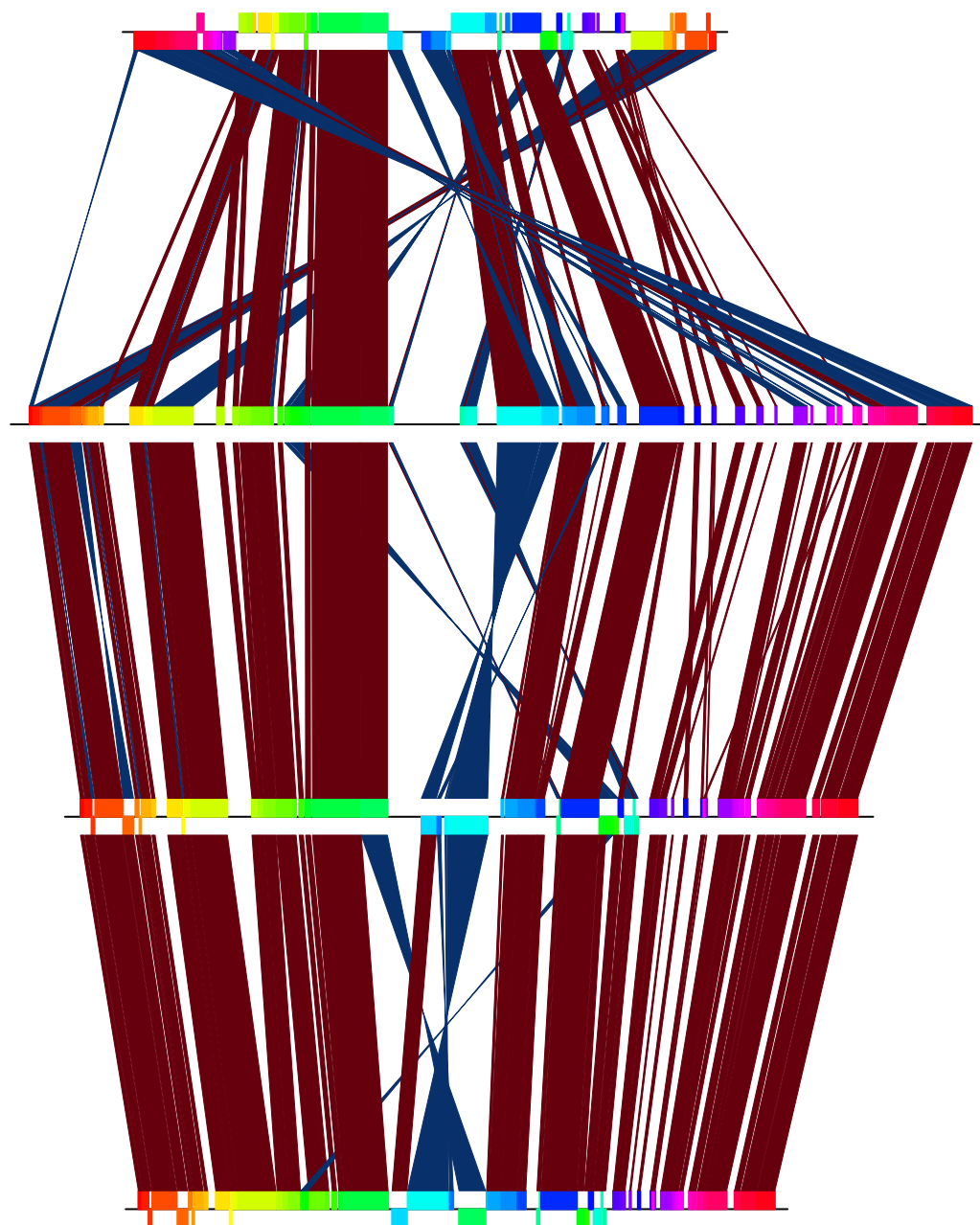
B\_grahamii

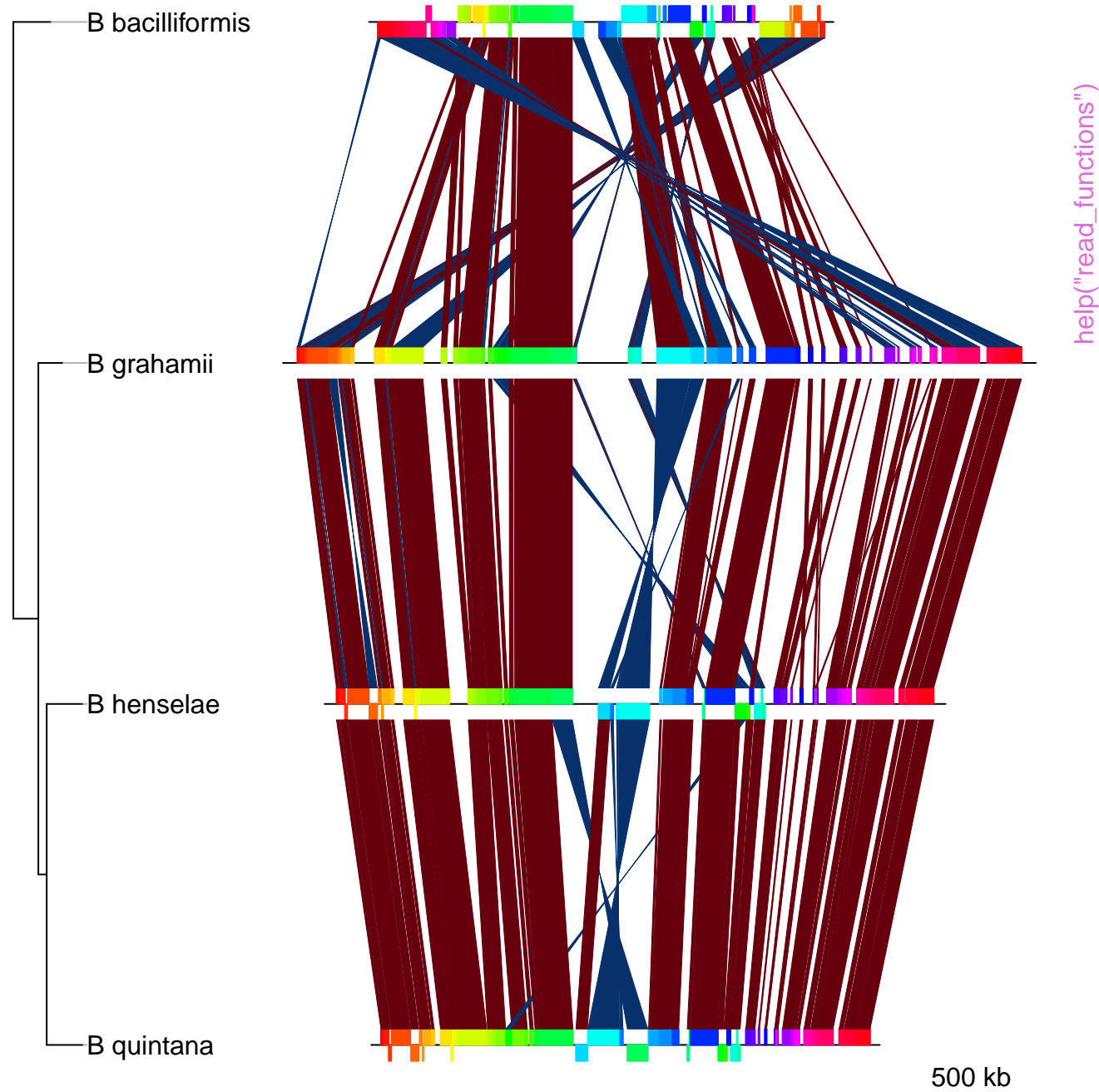
B\_henselae

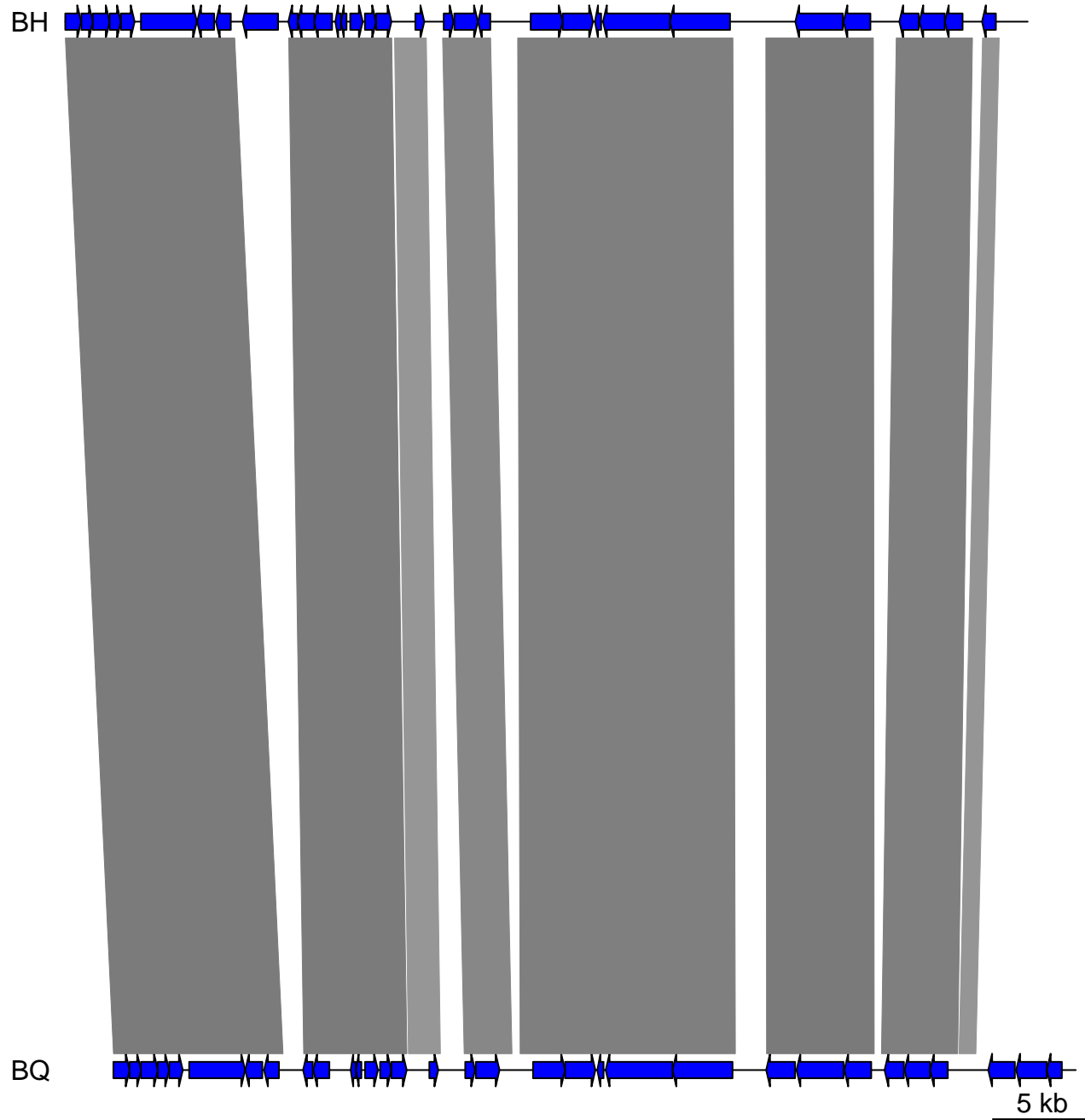
B\_quintana

help("read\_functions")

500 kb







help("read\_functions")

B\_bacilliformis

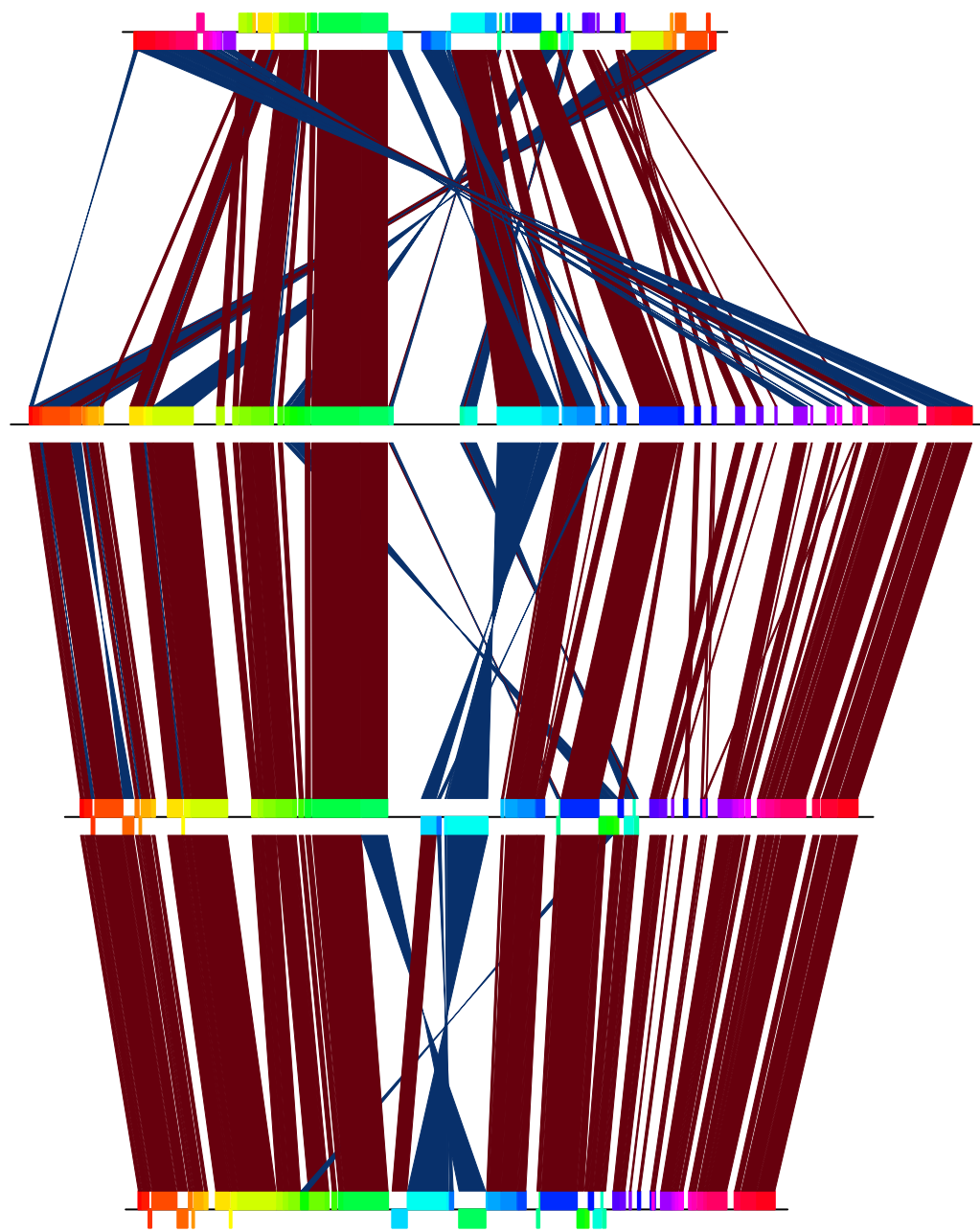
B\_grahamii

B\_henselae

B\_quintana

help("reverse")

500 kb



B\_bacilliformis

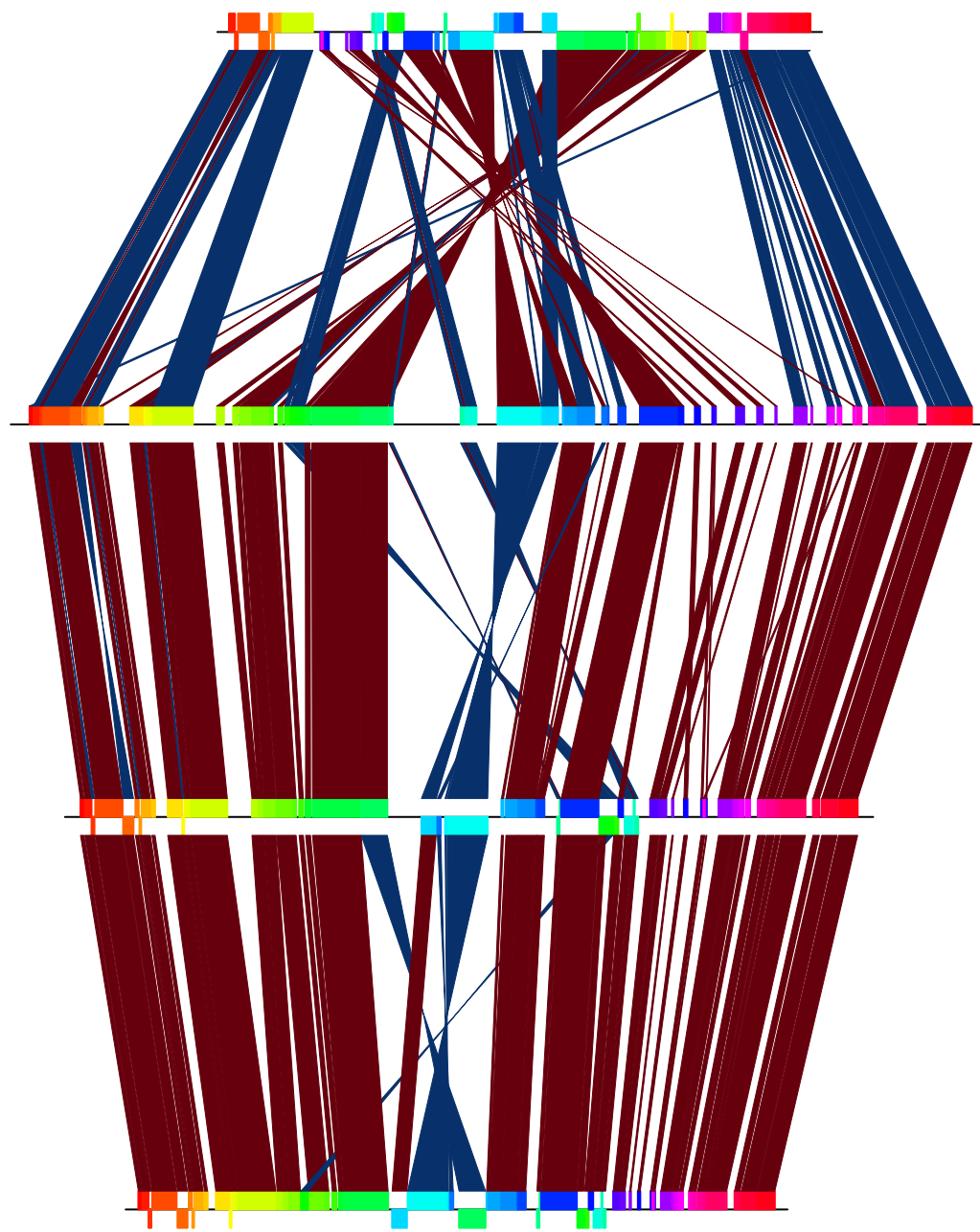
B\_grahamii

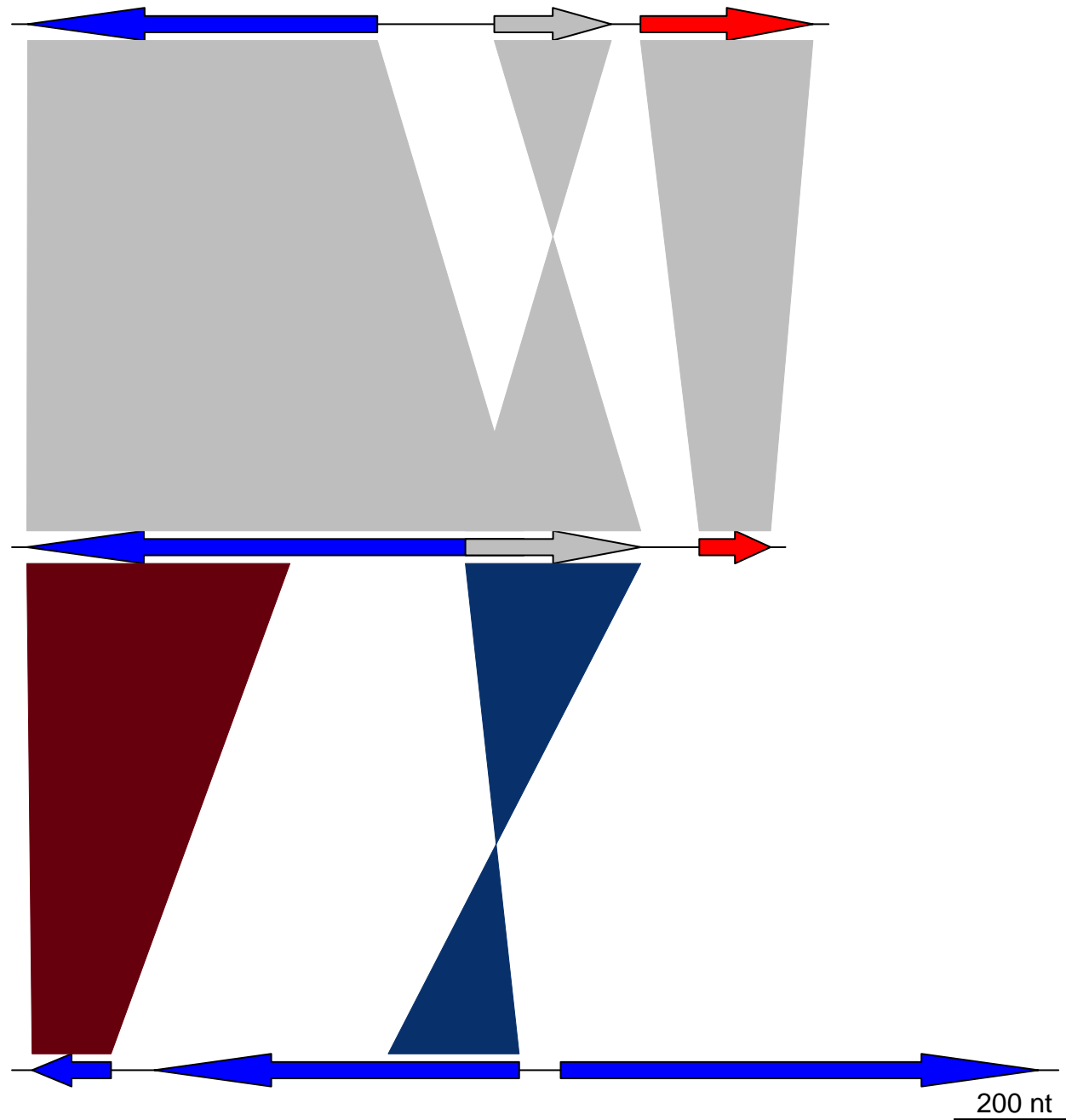
B\_henselae

B\_quintana

help("reverse")

500 kb





help("three\_genes")

