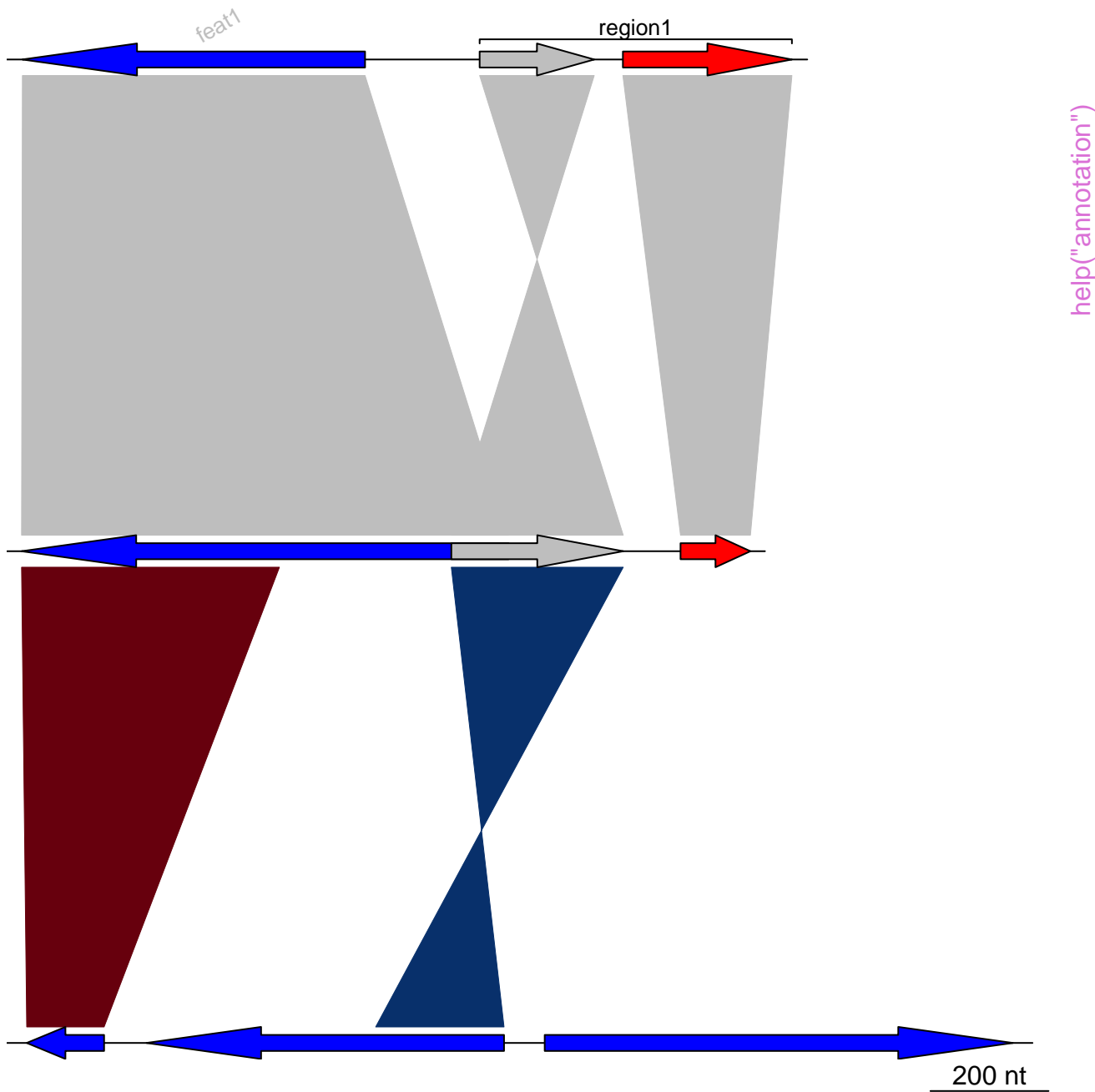
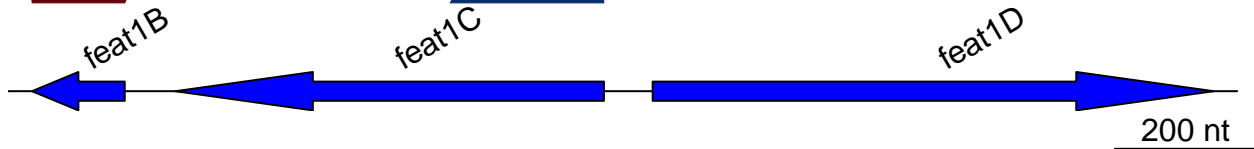
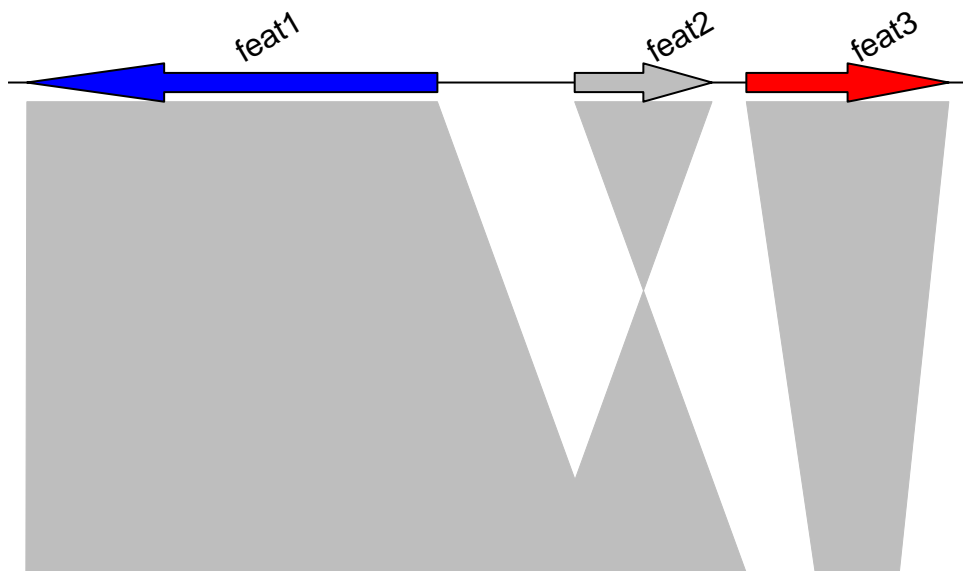


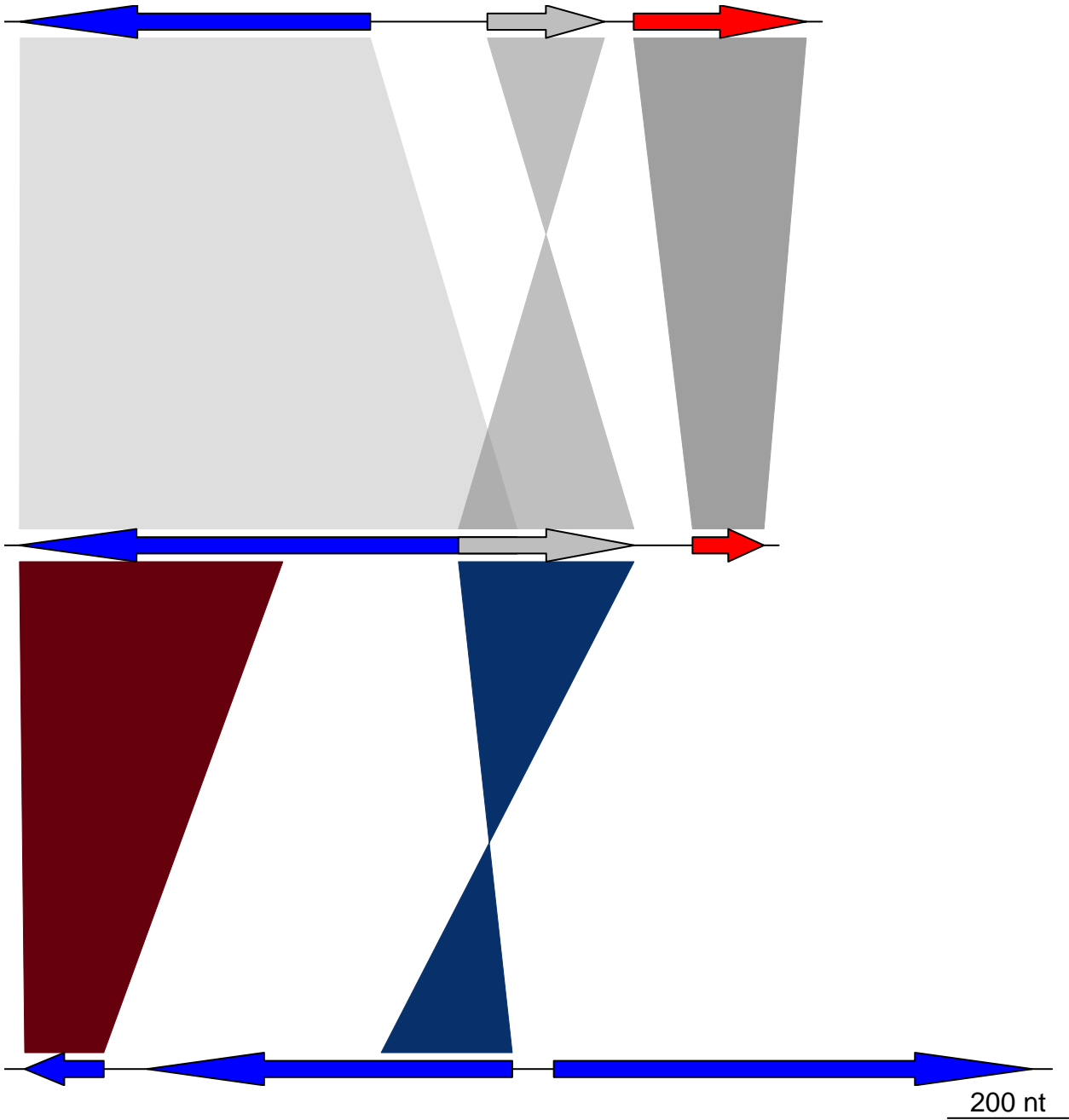
help("annotation")

200 nt

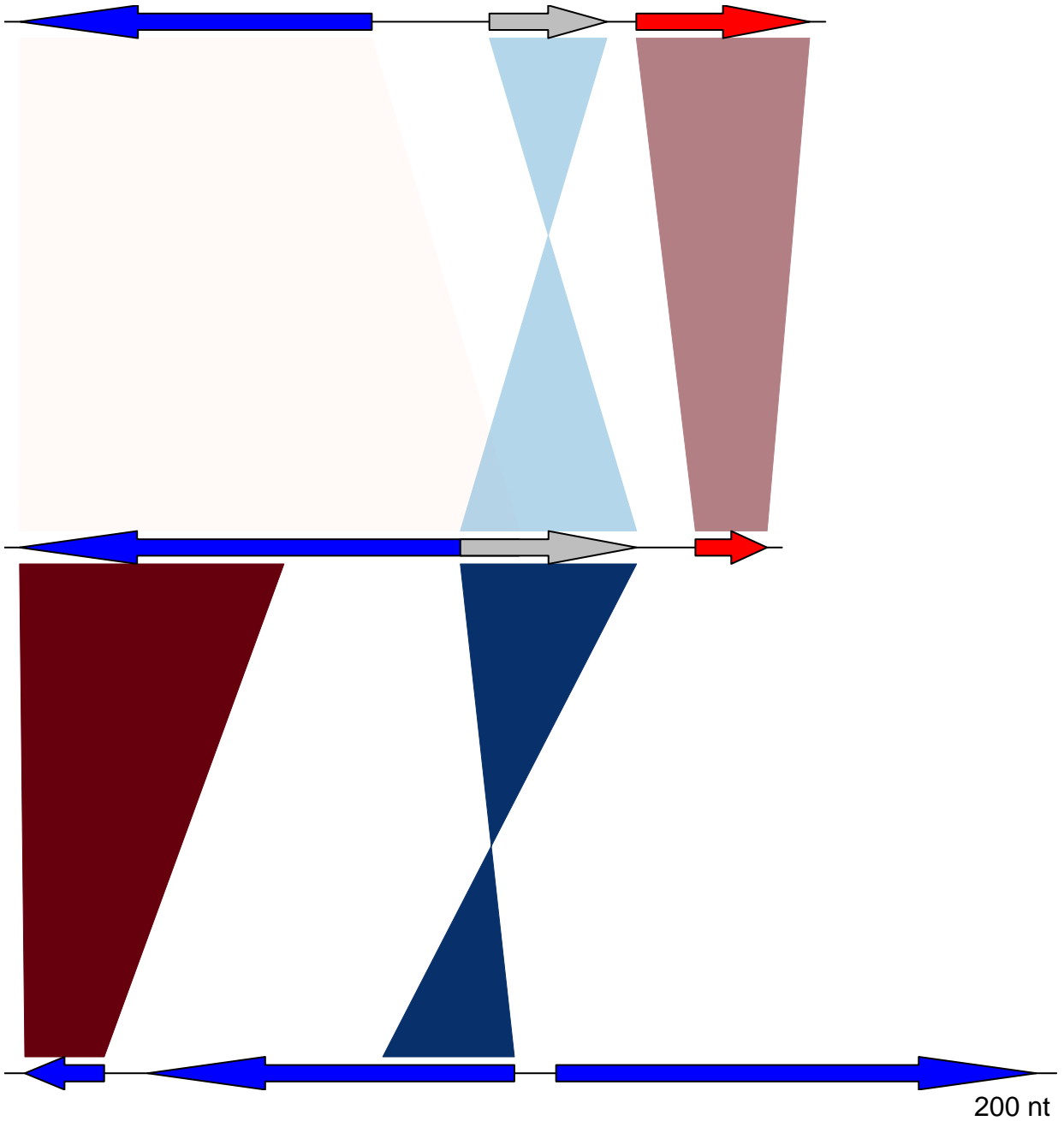




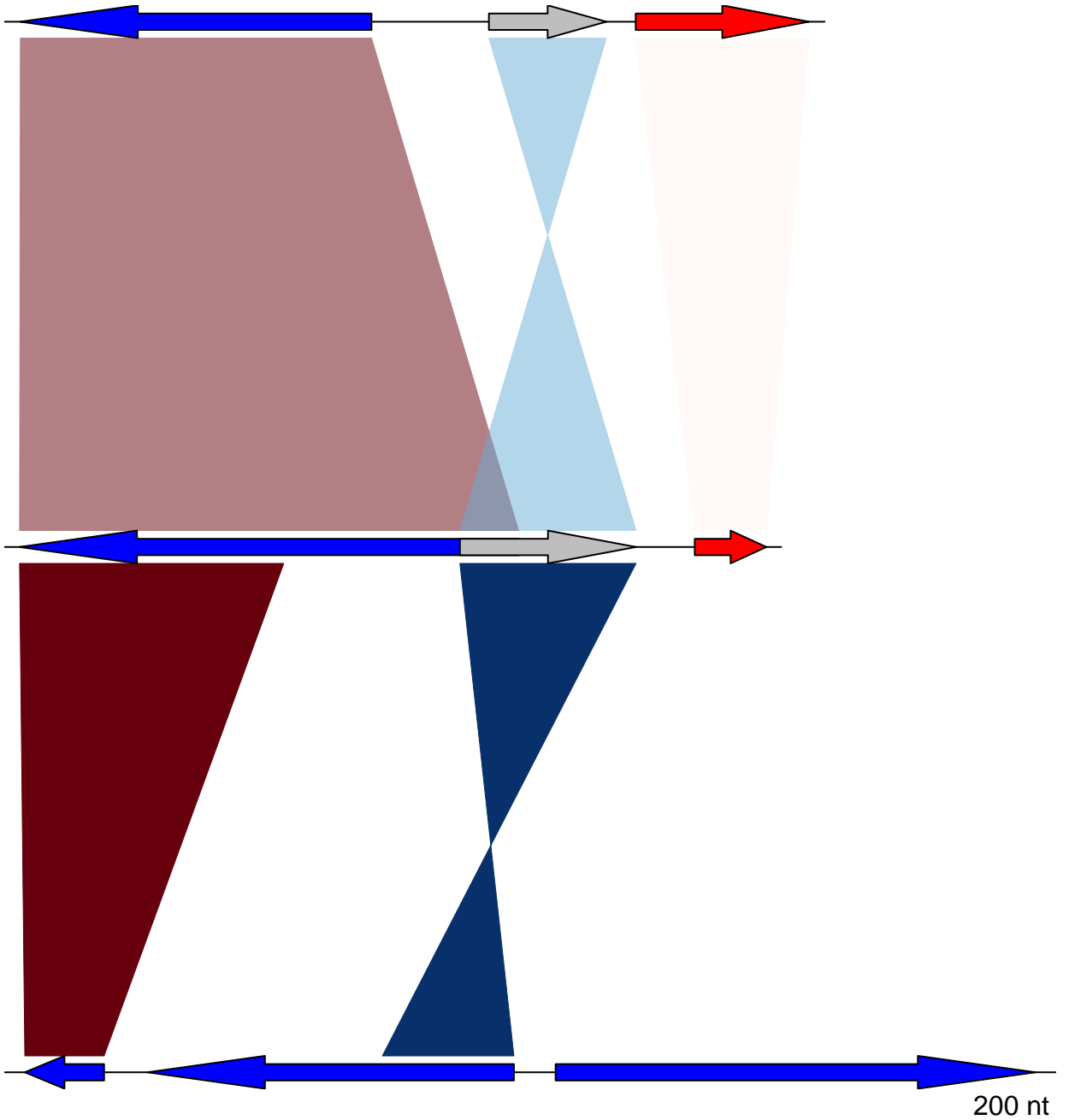
help("annotation")



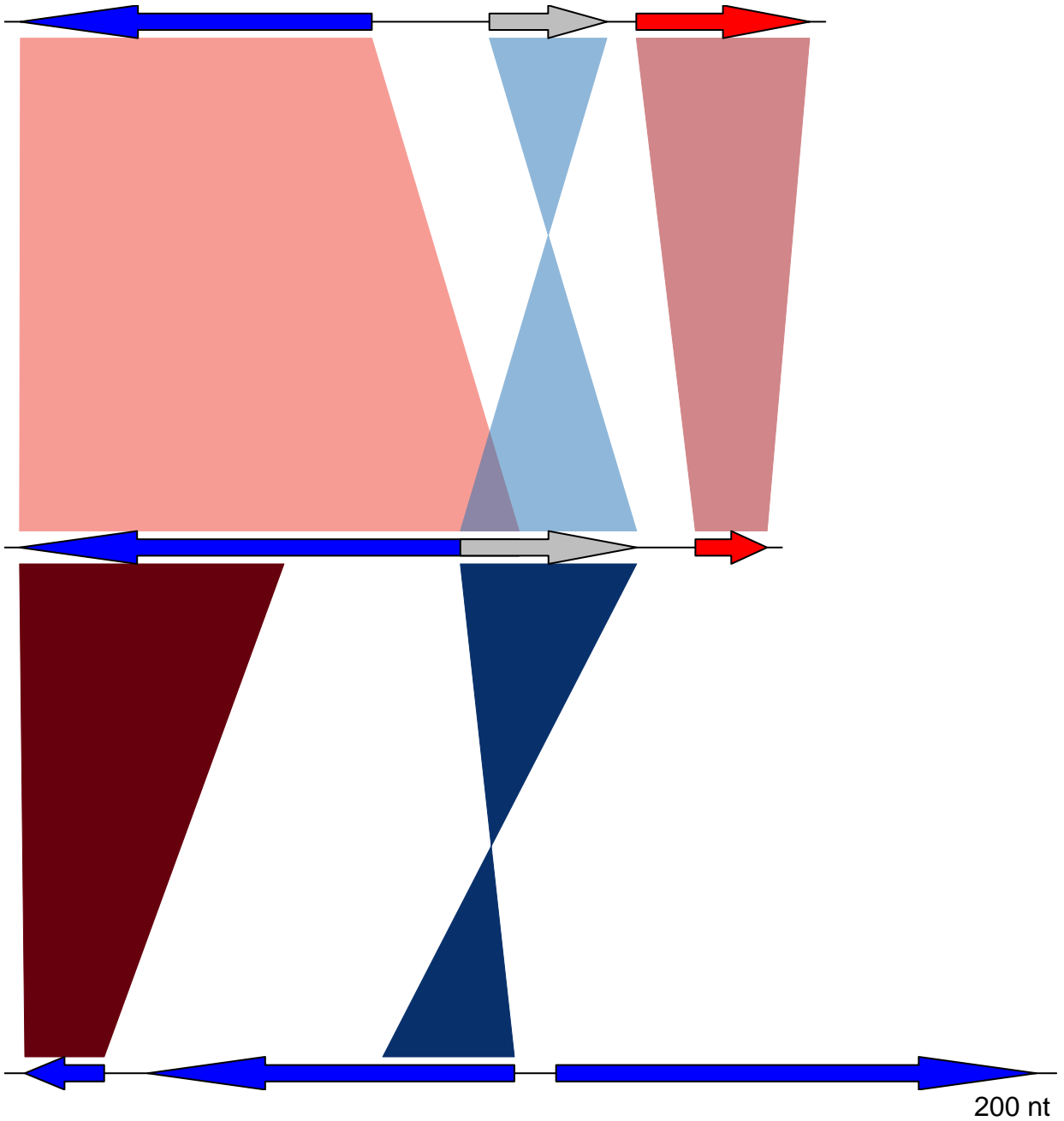
help("apply_color_scheme")



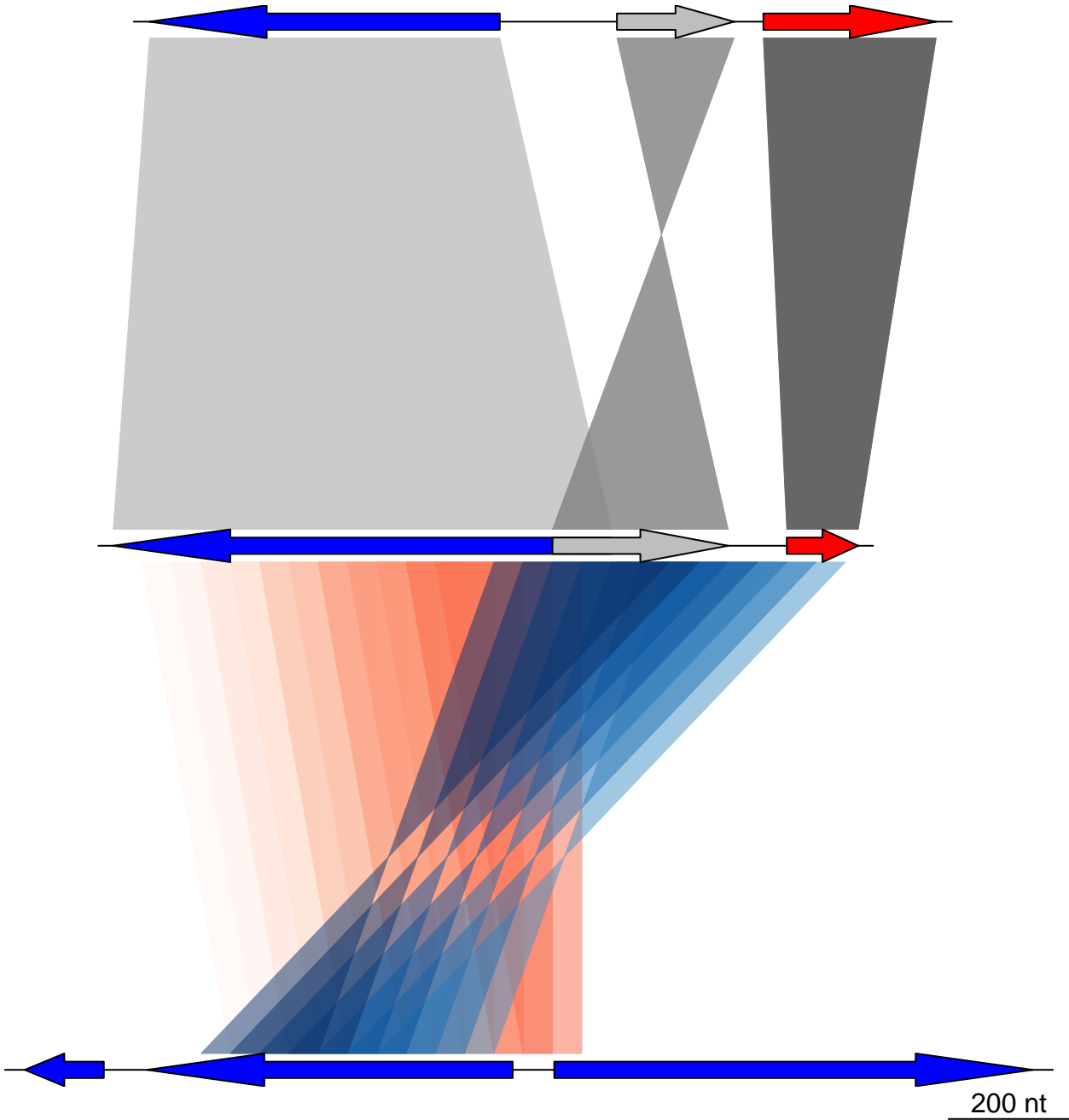
help("apply_color_scheme")



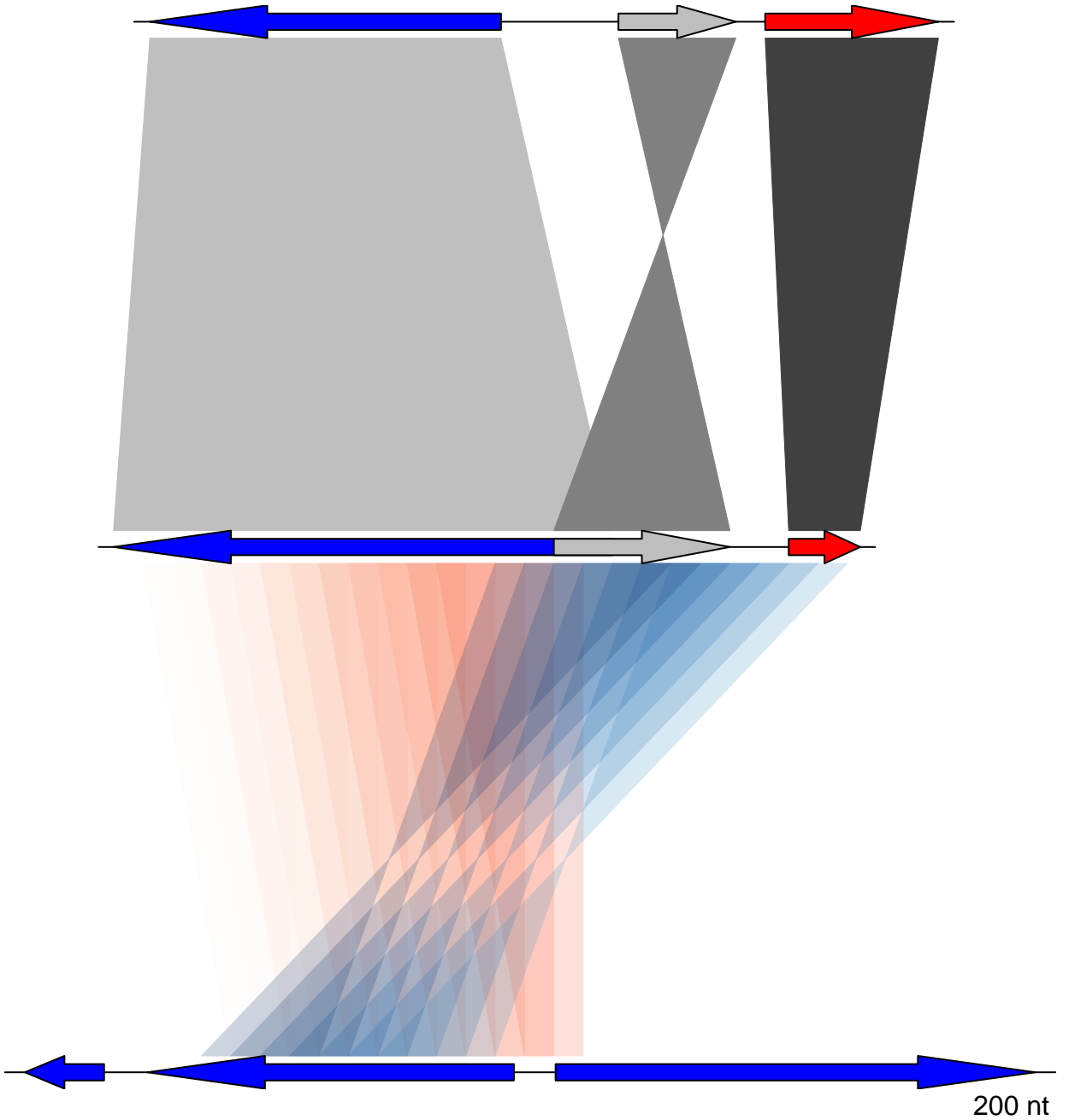
help("apply_color_scheme")



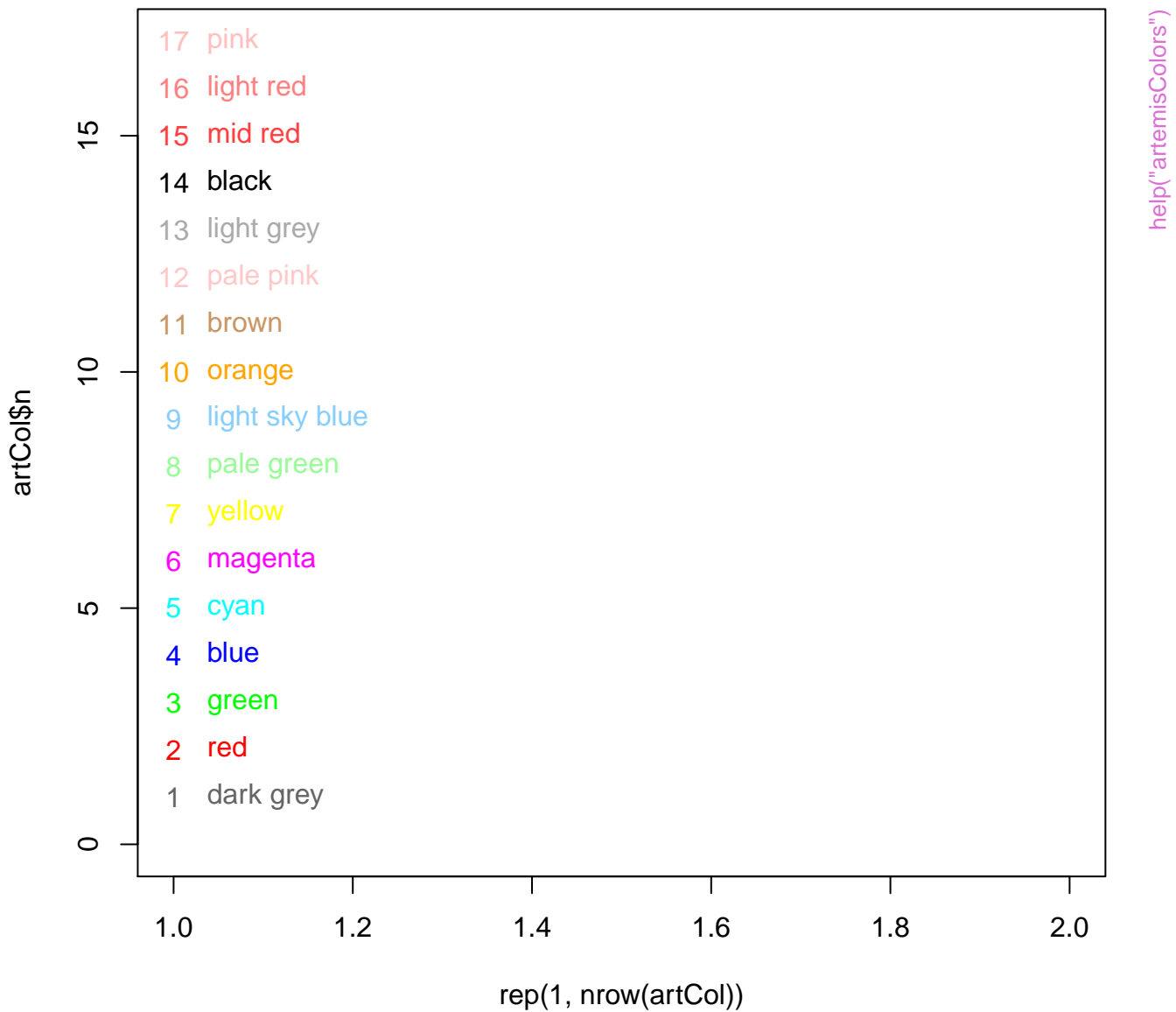
help("apply_color_scheme")



help("apply_color_scheme")



help("apply_color_scheme")



BB

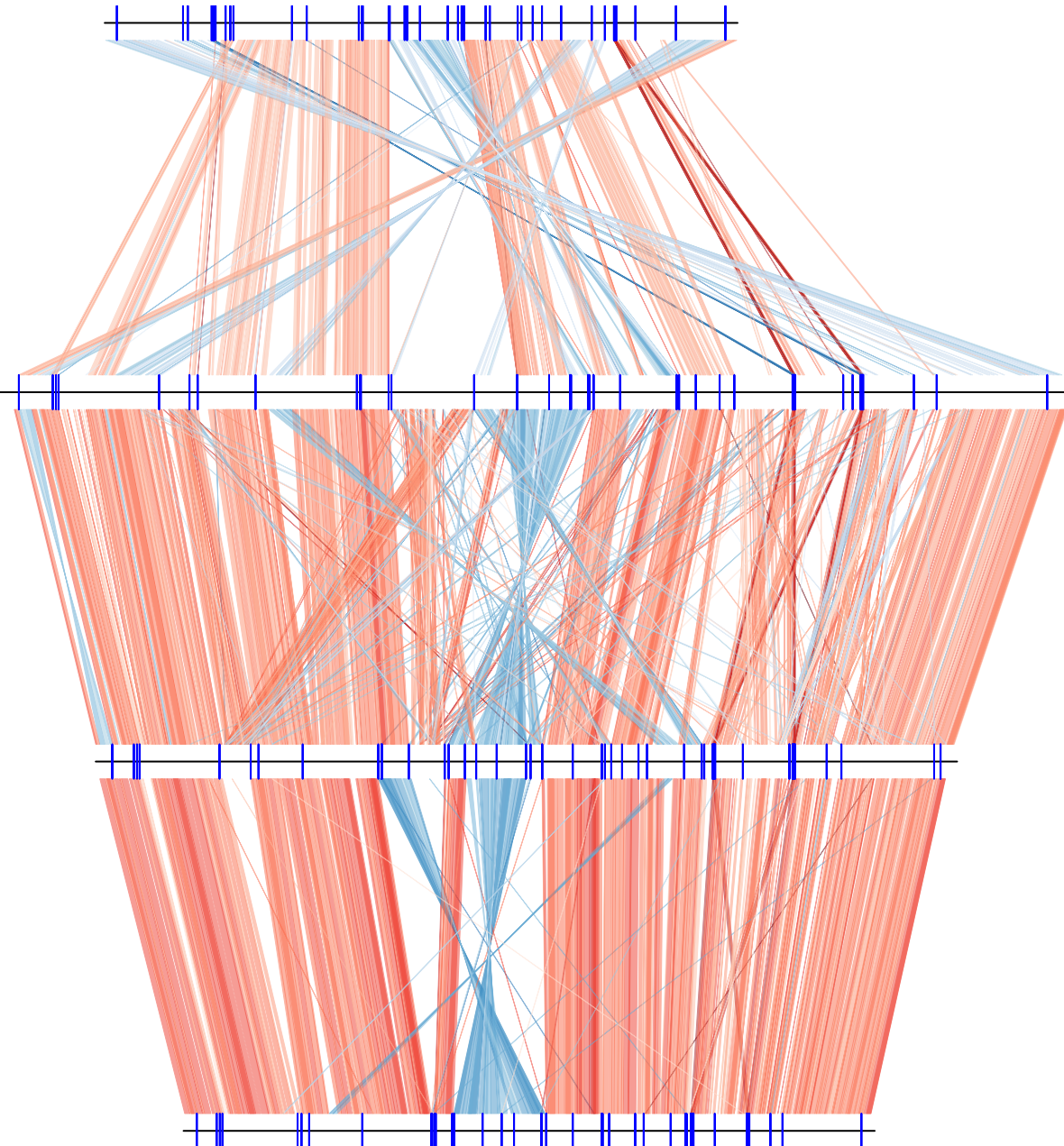
BG

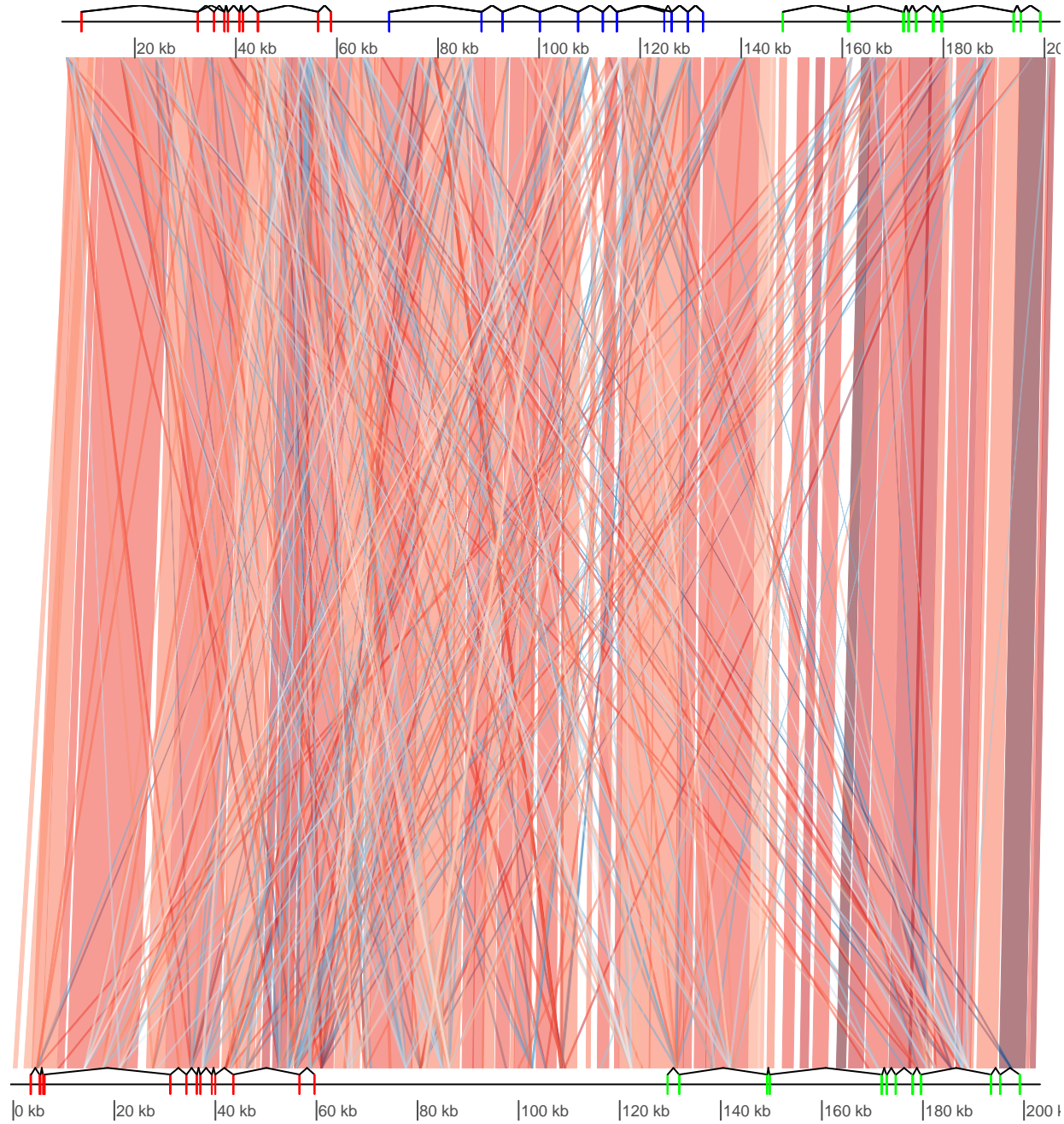
BH

BQ

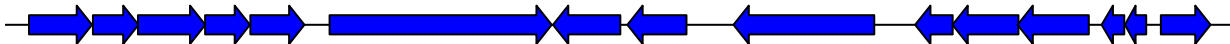
help("barto")

500 kb



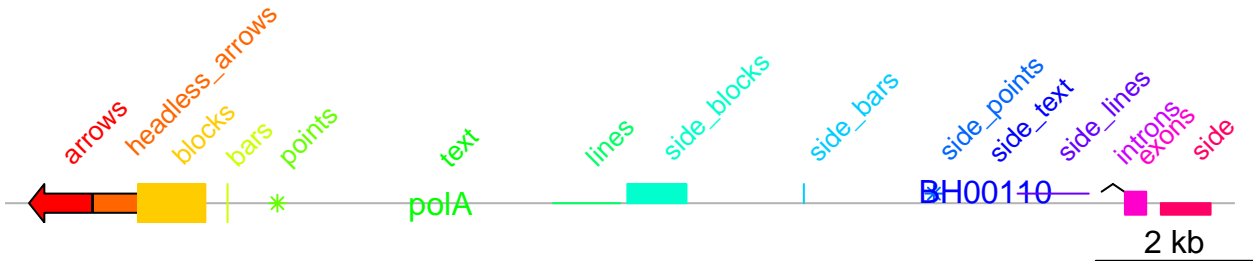
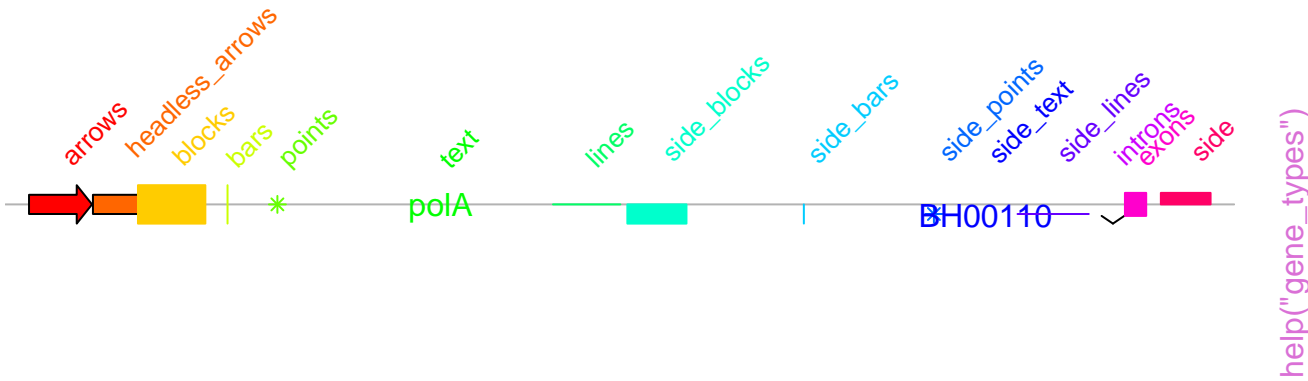


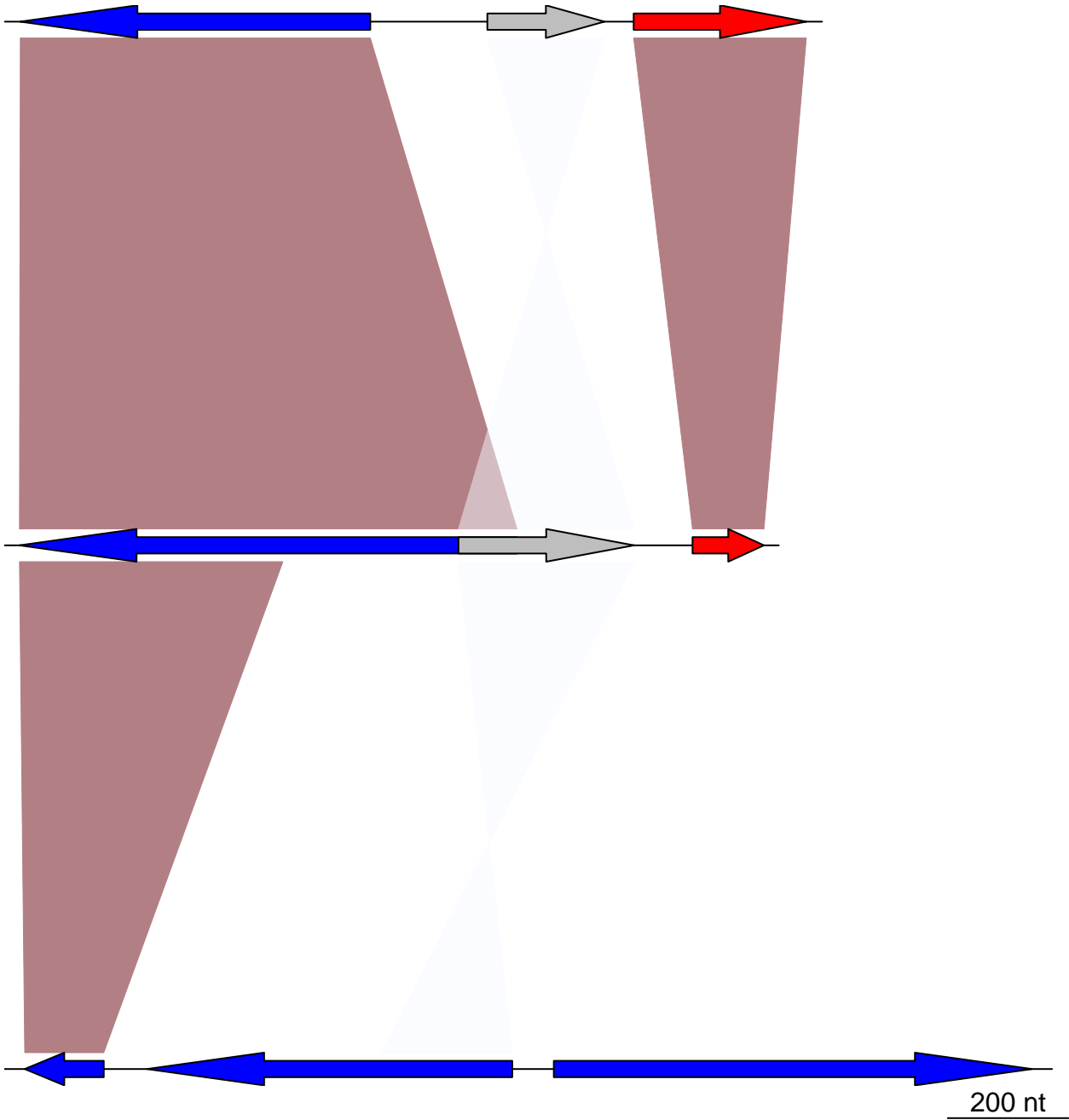
help("chrY_subseg")



2 kb

help("gene_types")





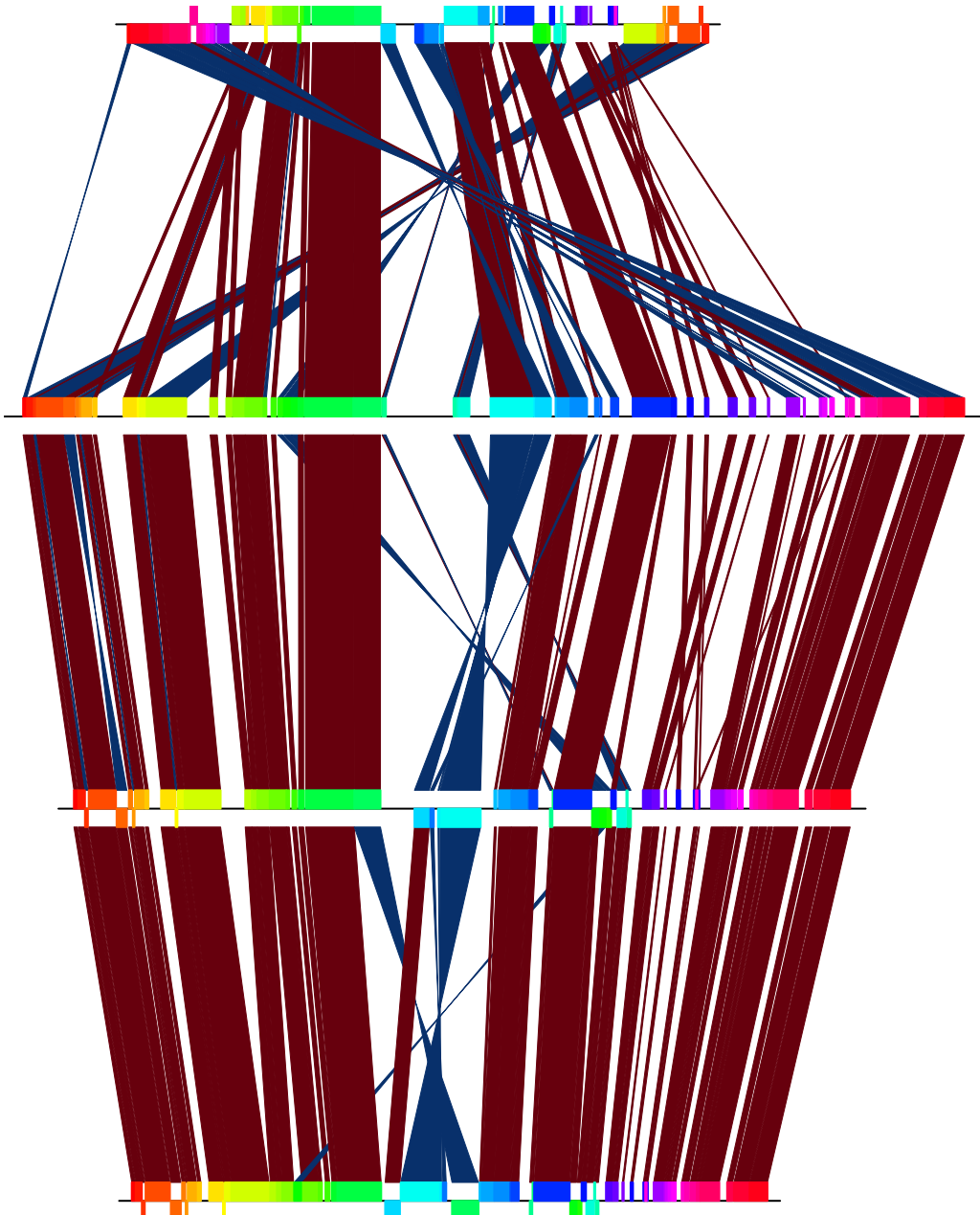
help("genoPlotR-package")

B_bacilliformis

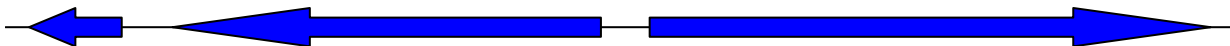
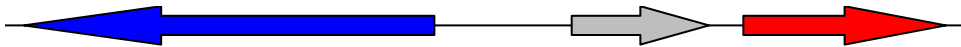
B_grahamii

B_henselae

B_quintana

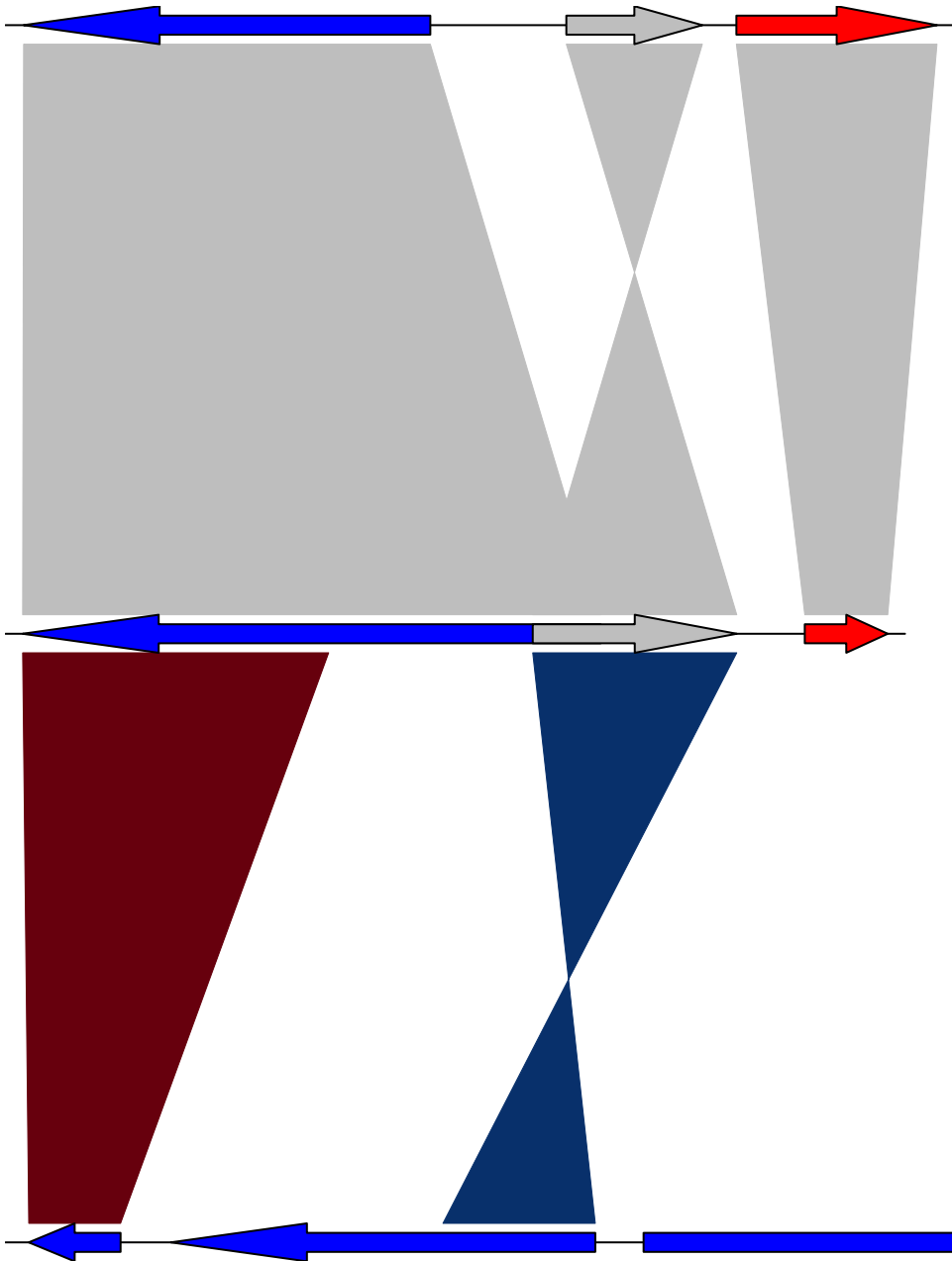


help("mauve.bbone")



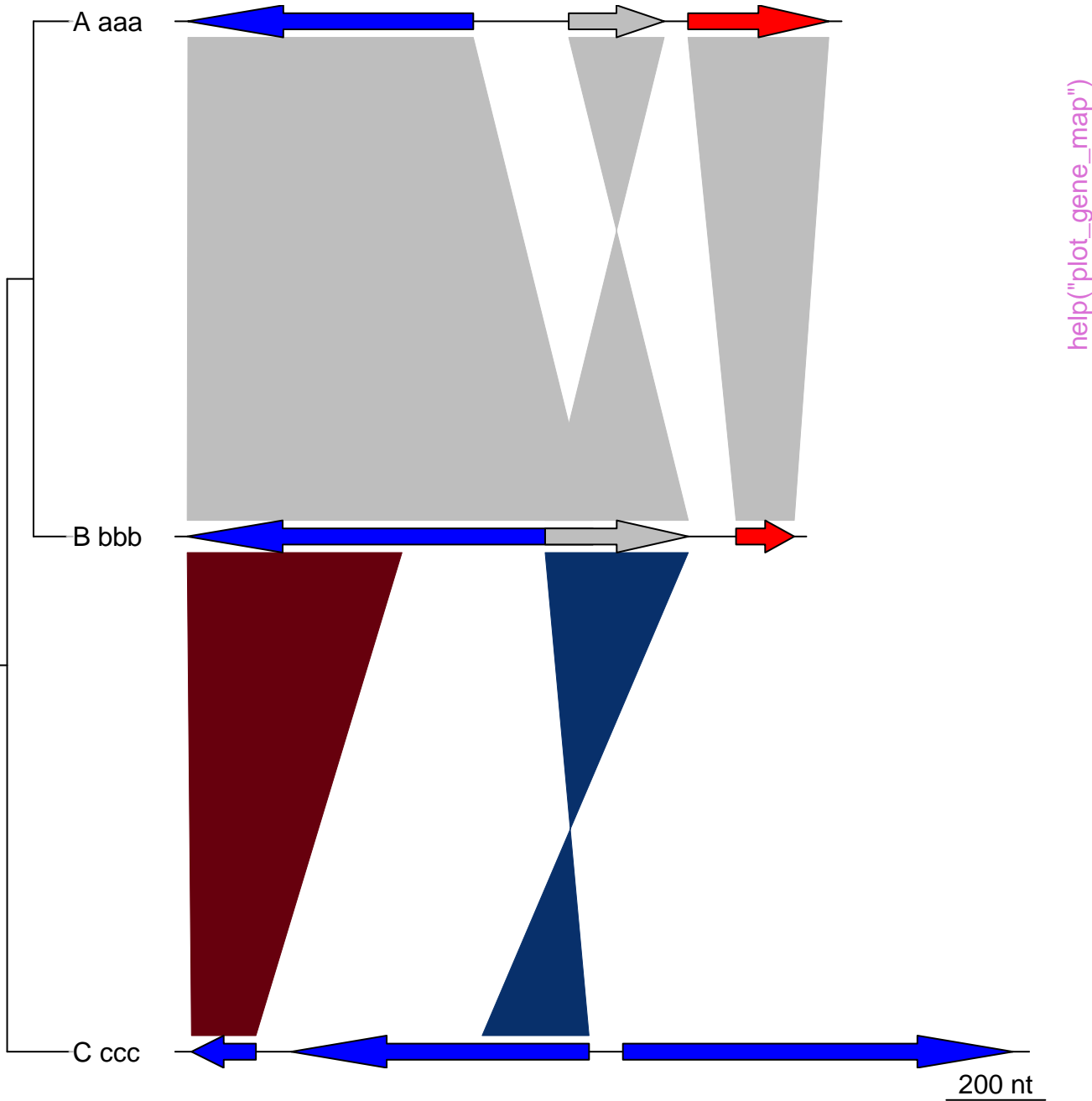
200 nt

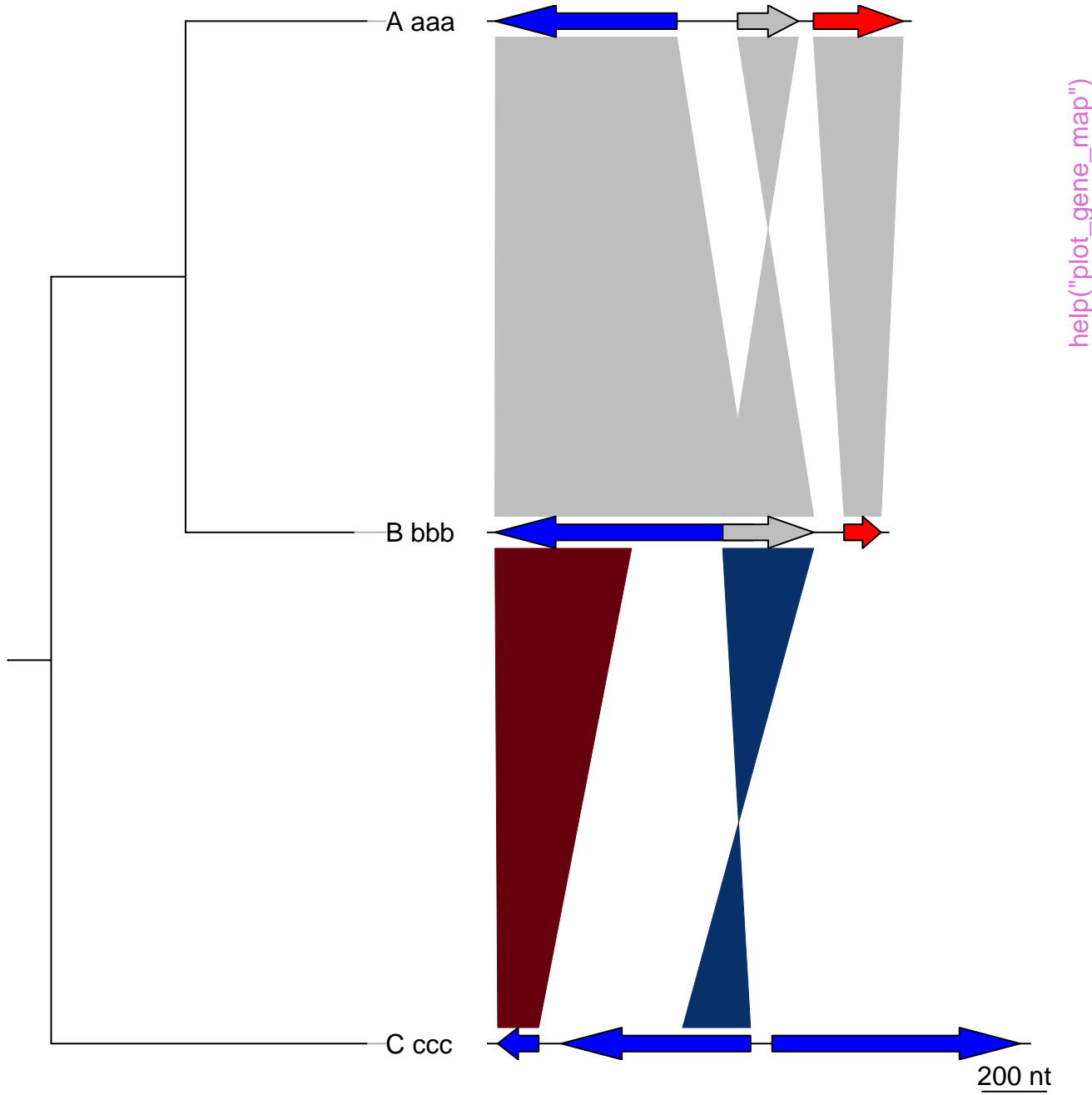
help("plot_gene_map")

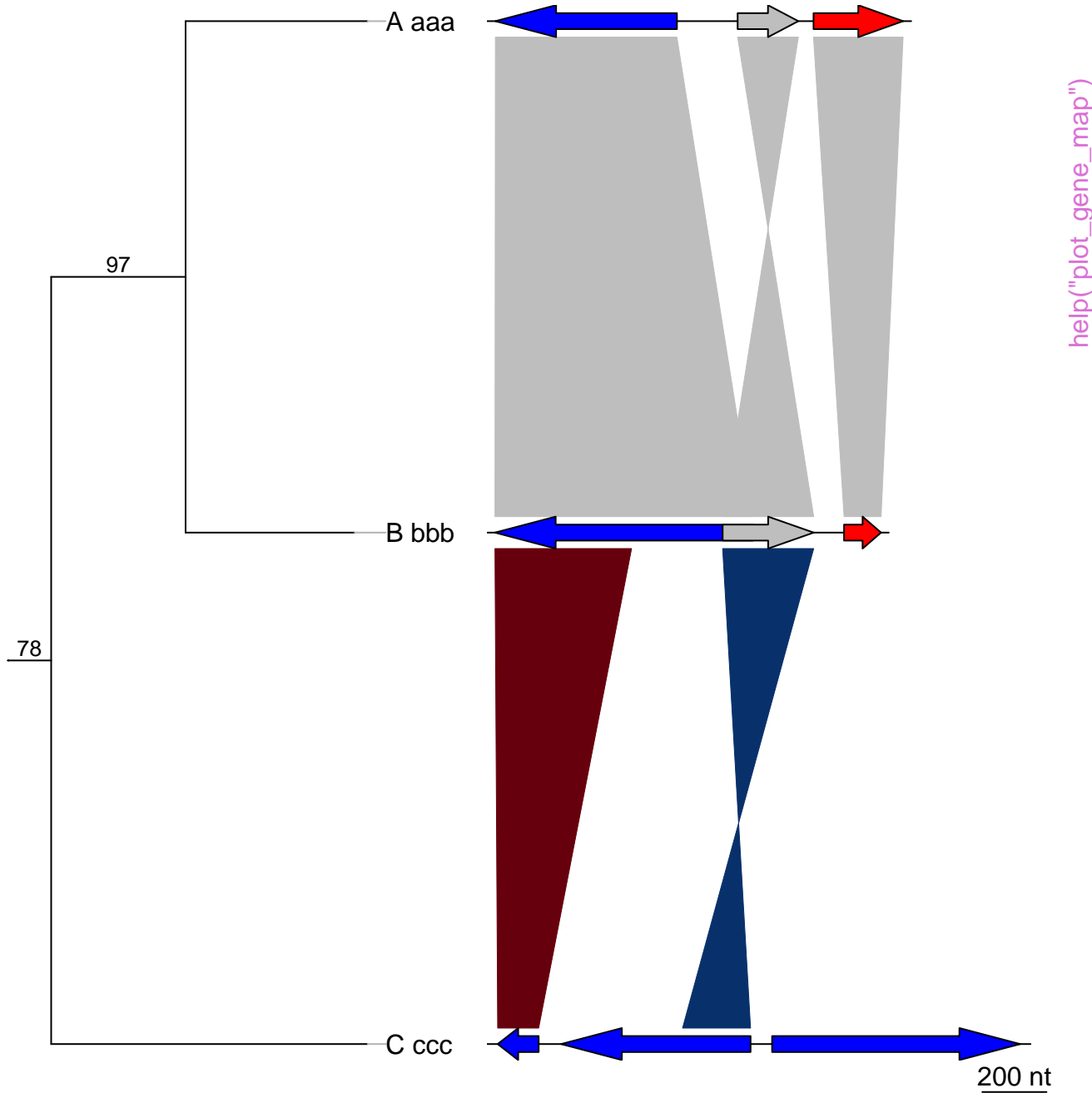


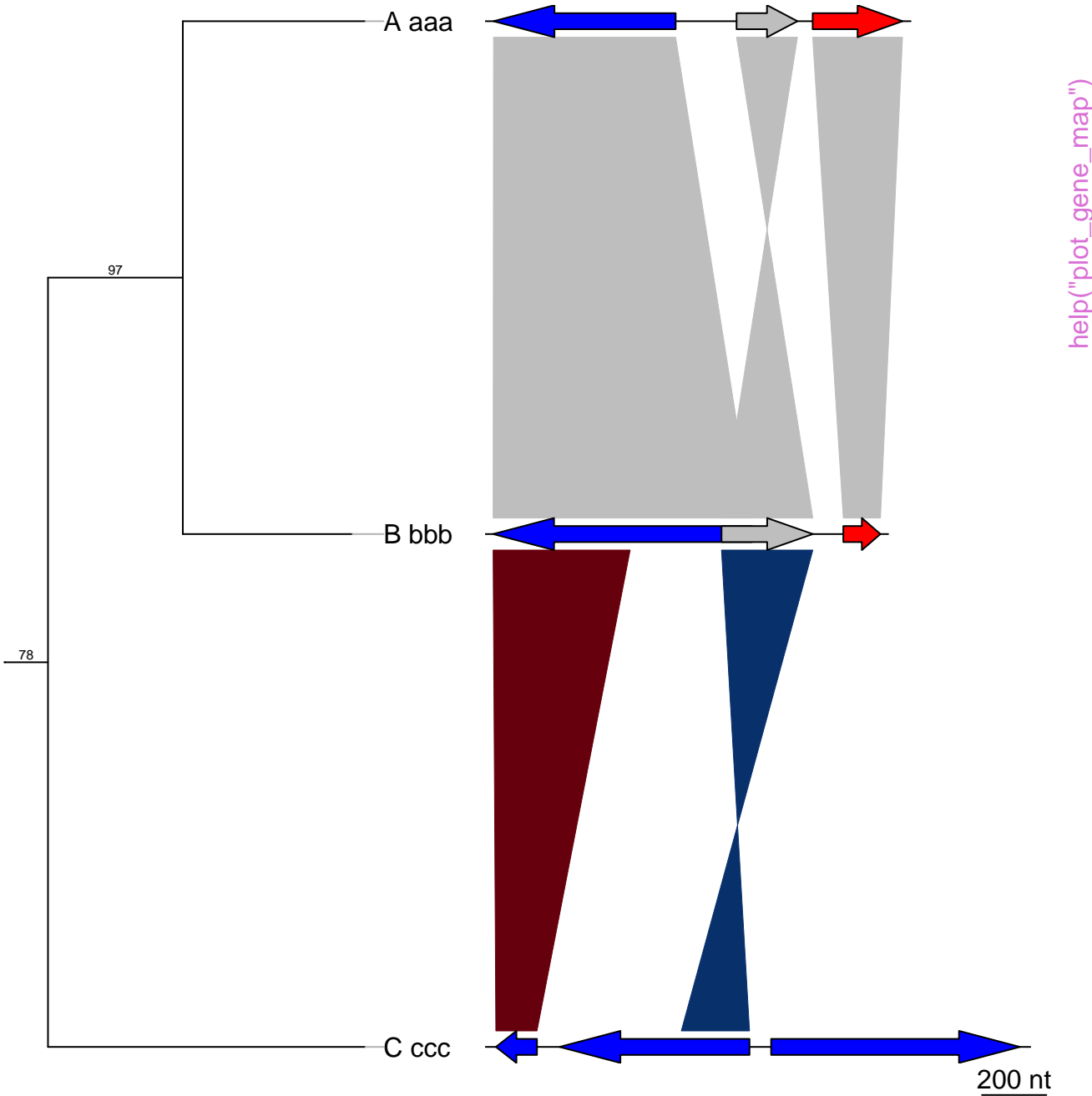
help("plot_gene_map")

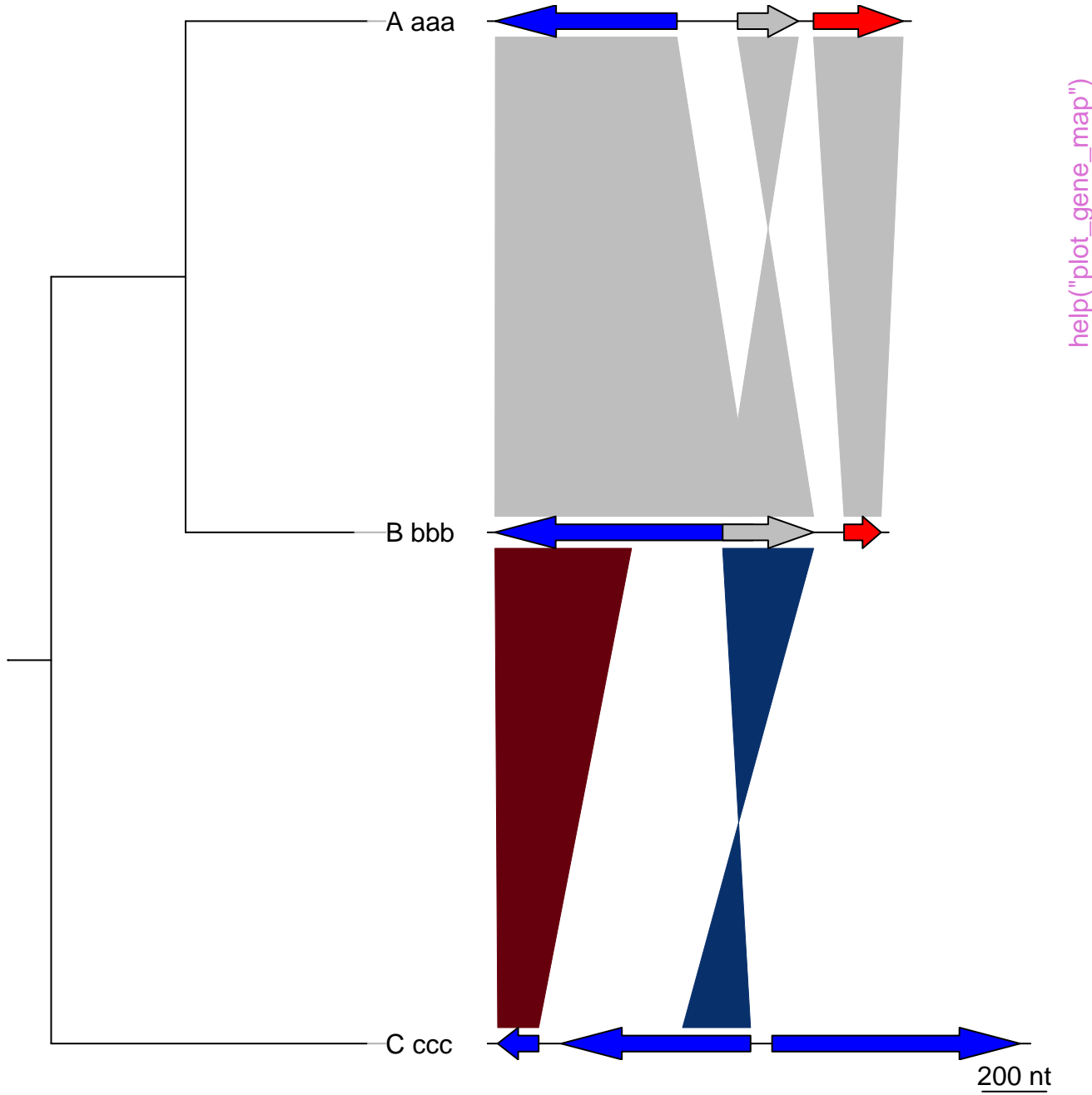
200 nt

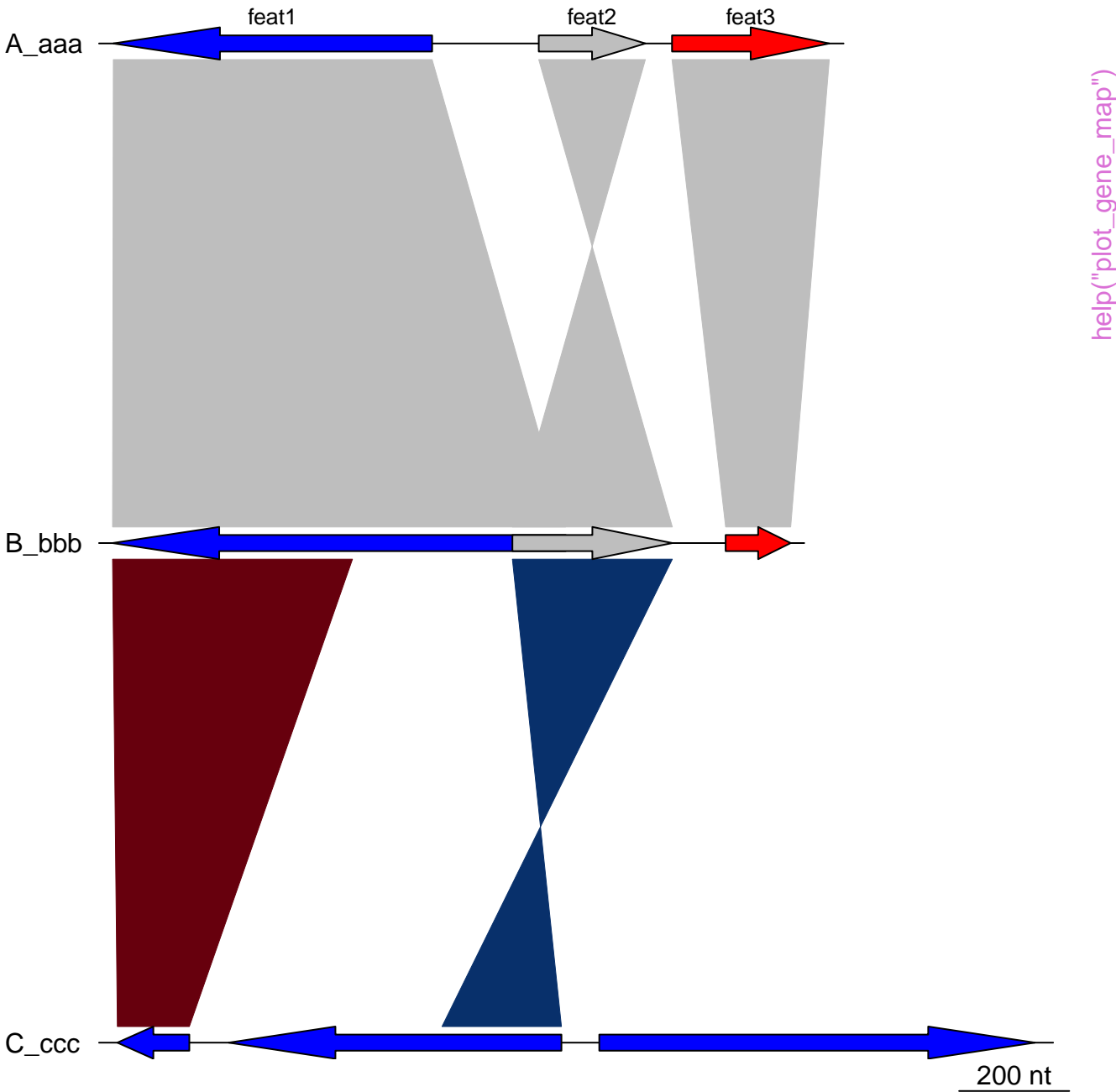


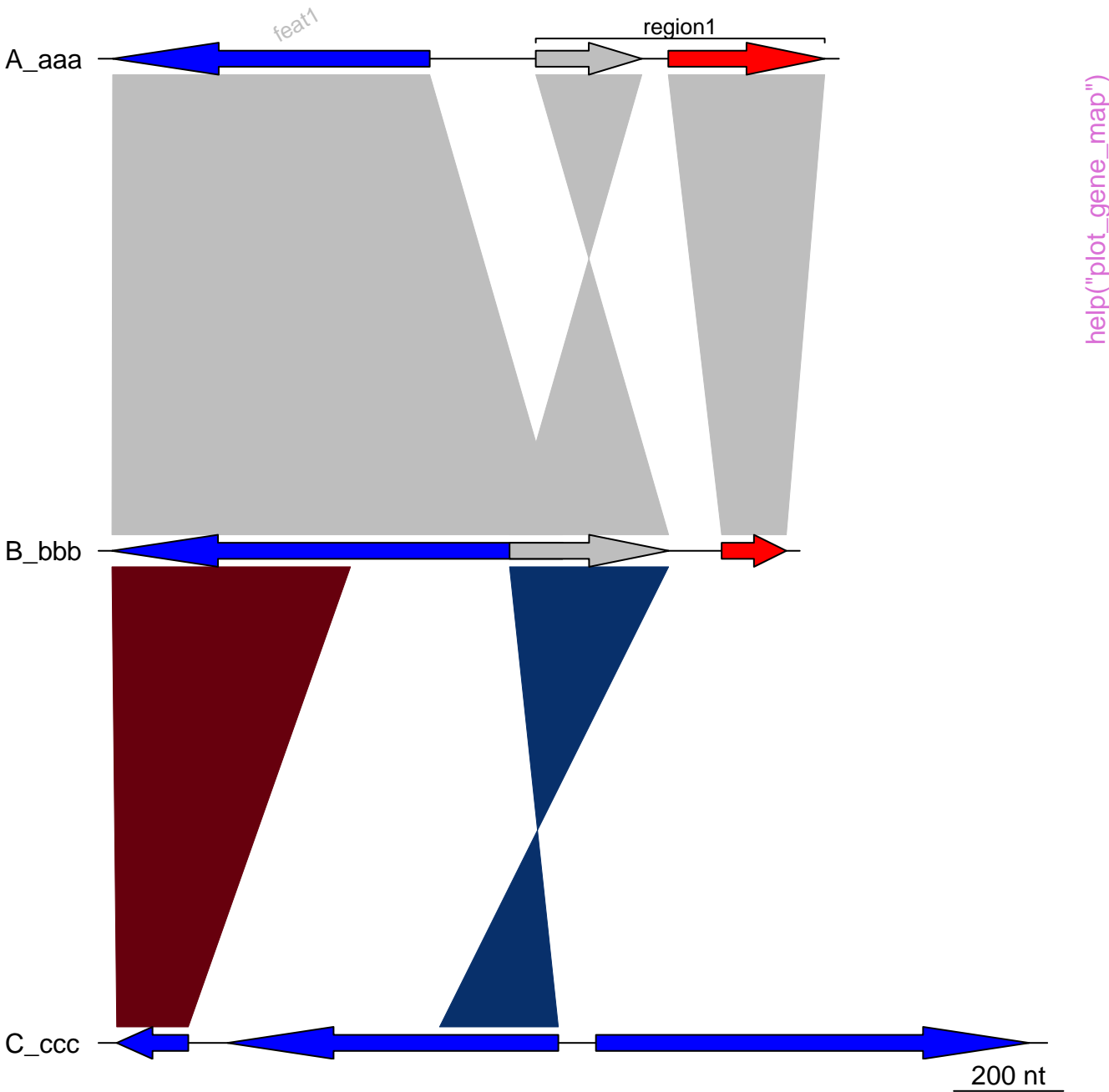


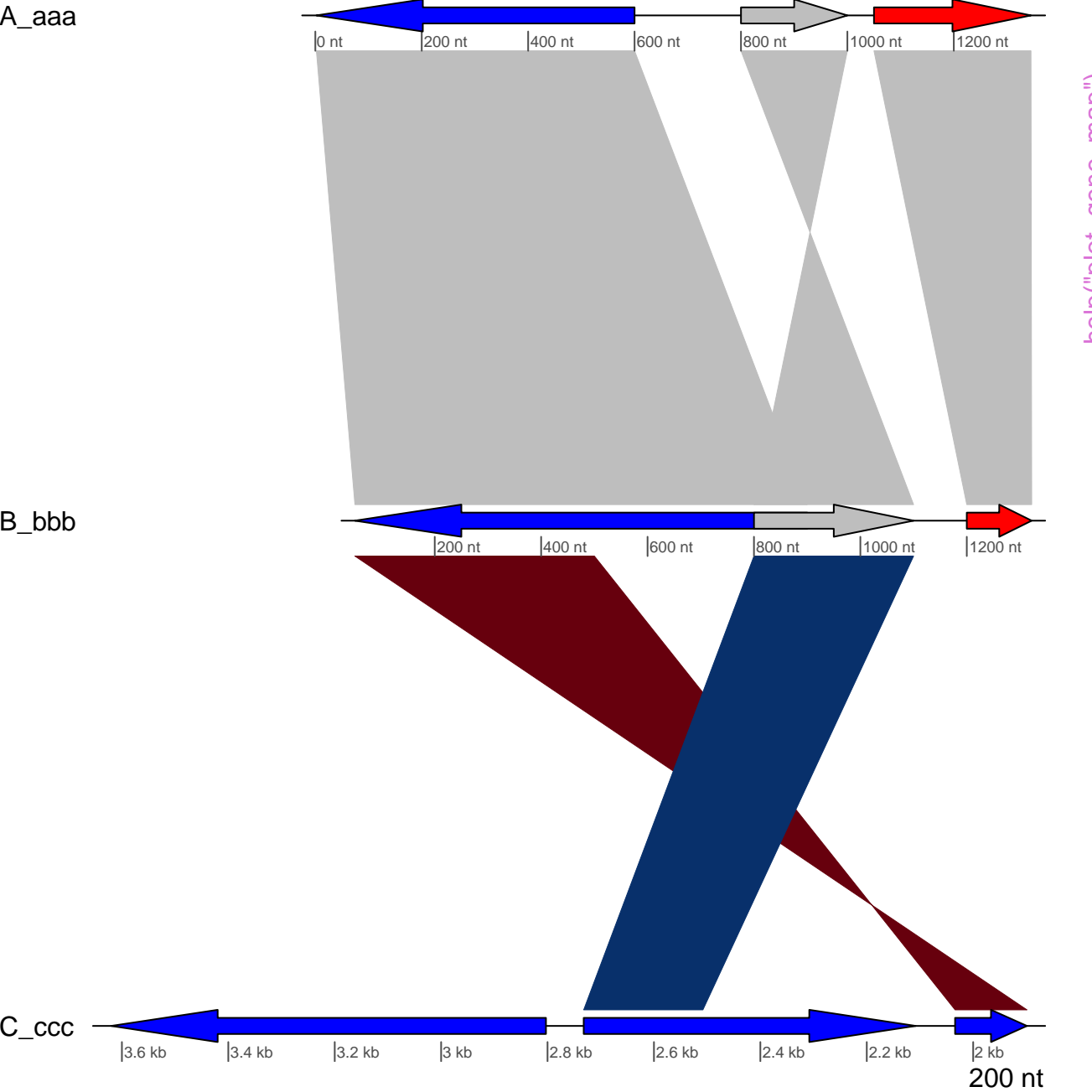


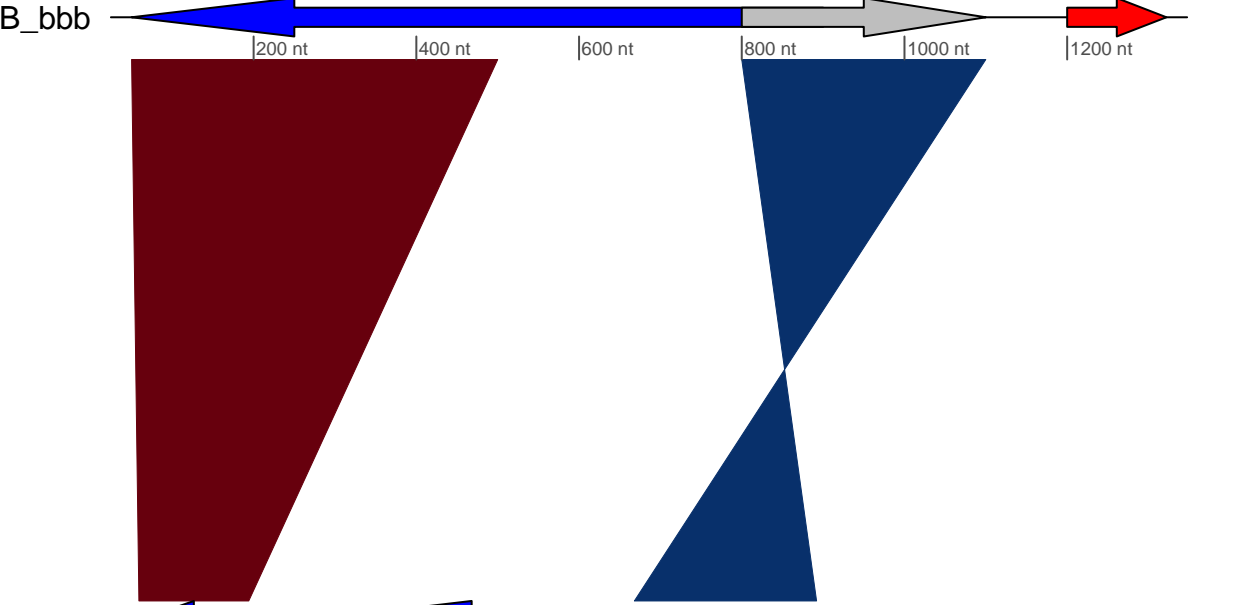
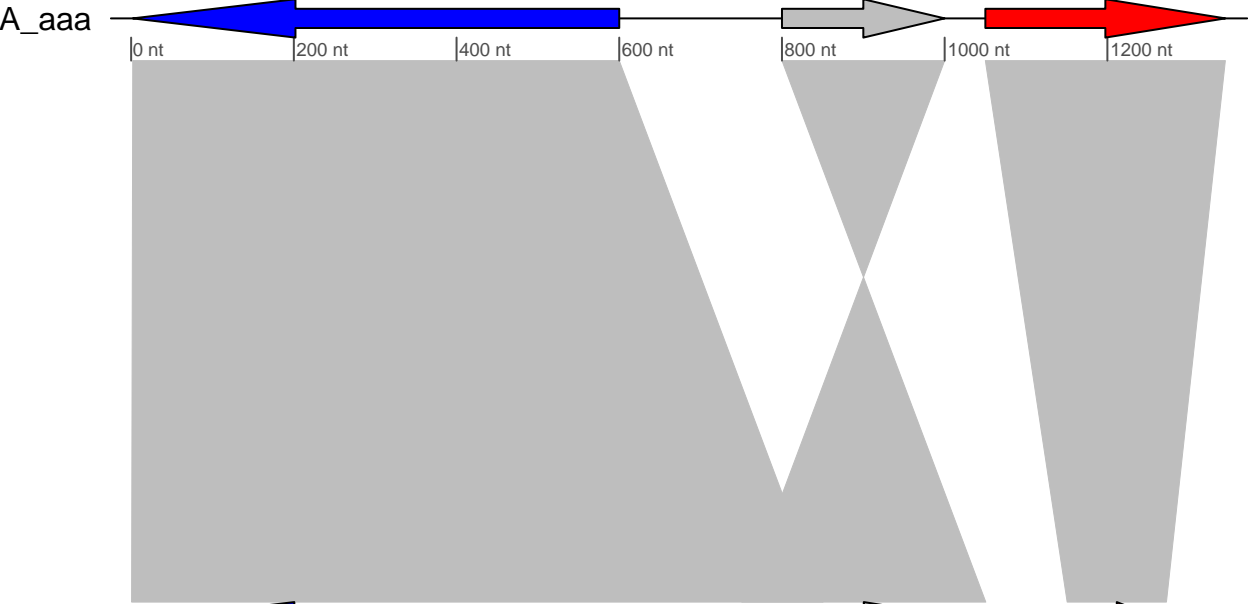








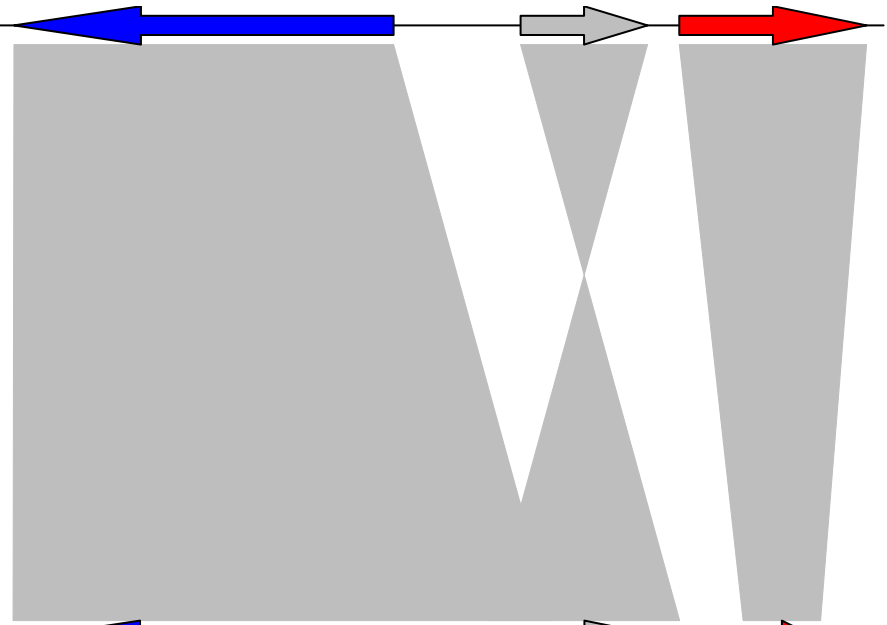




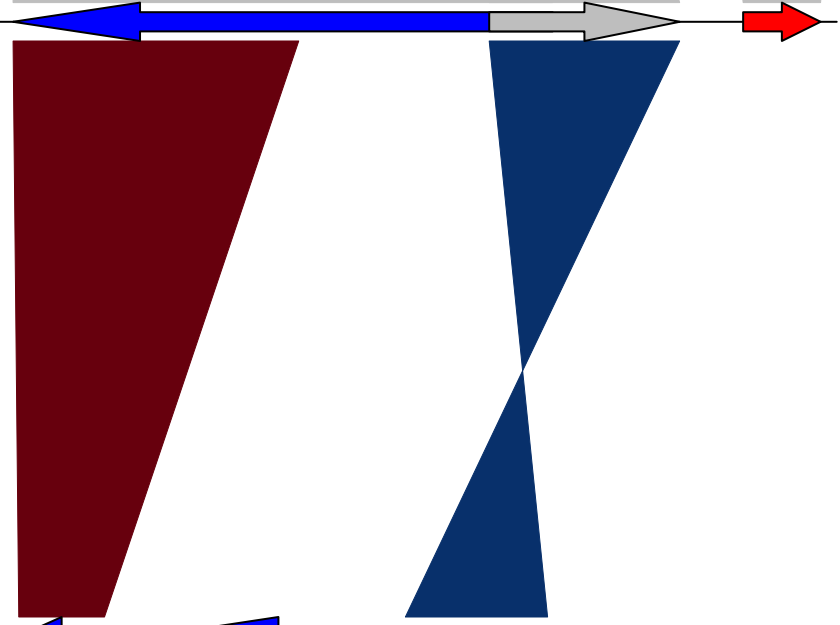
200 nt

help("plot_gene_map")

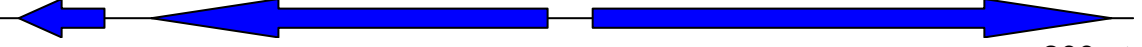
A_aaa



B_bbb



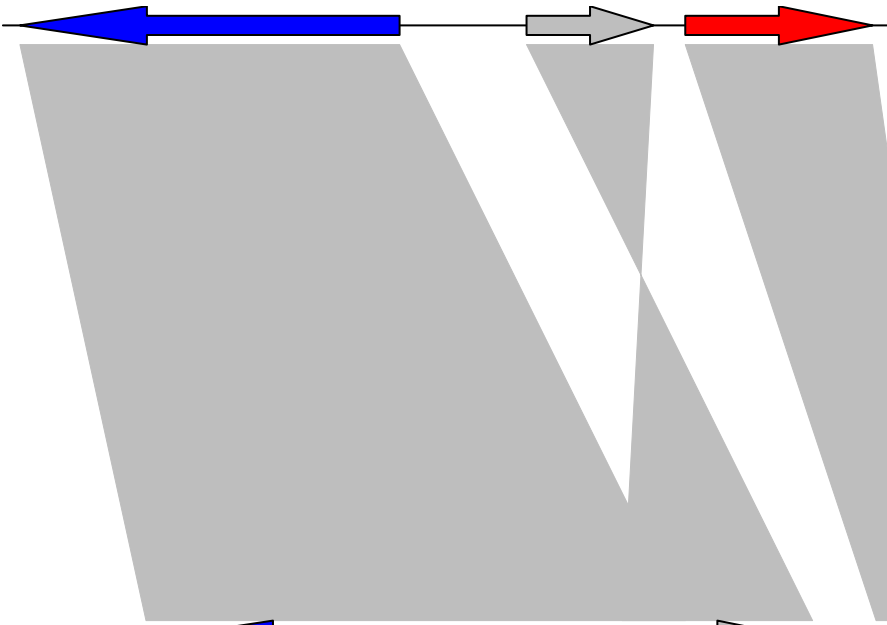
C_ccc



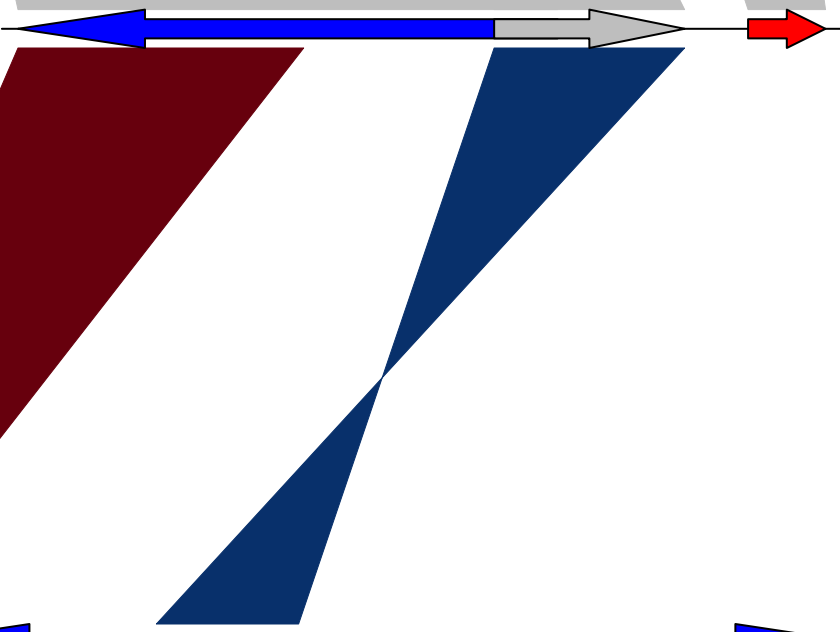
200 nt

help("plot_gene_map")

A_aaa



B_bbb



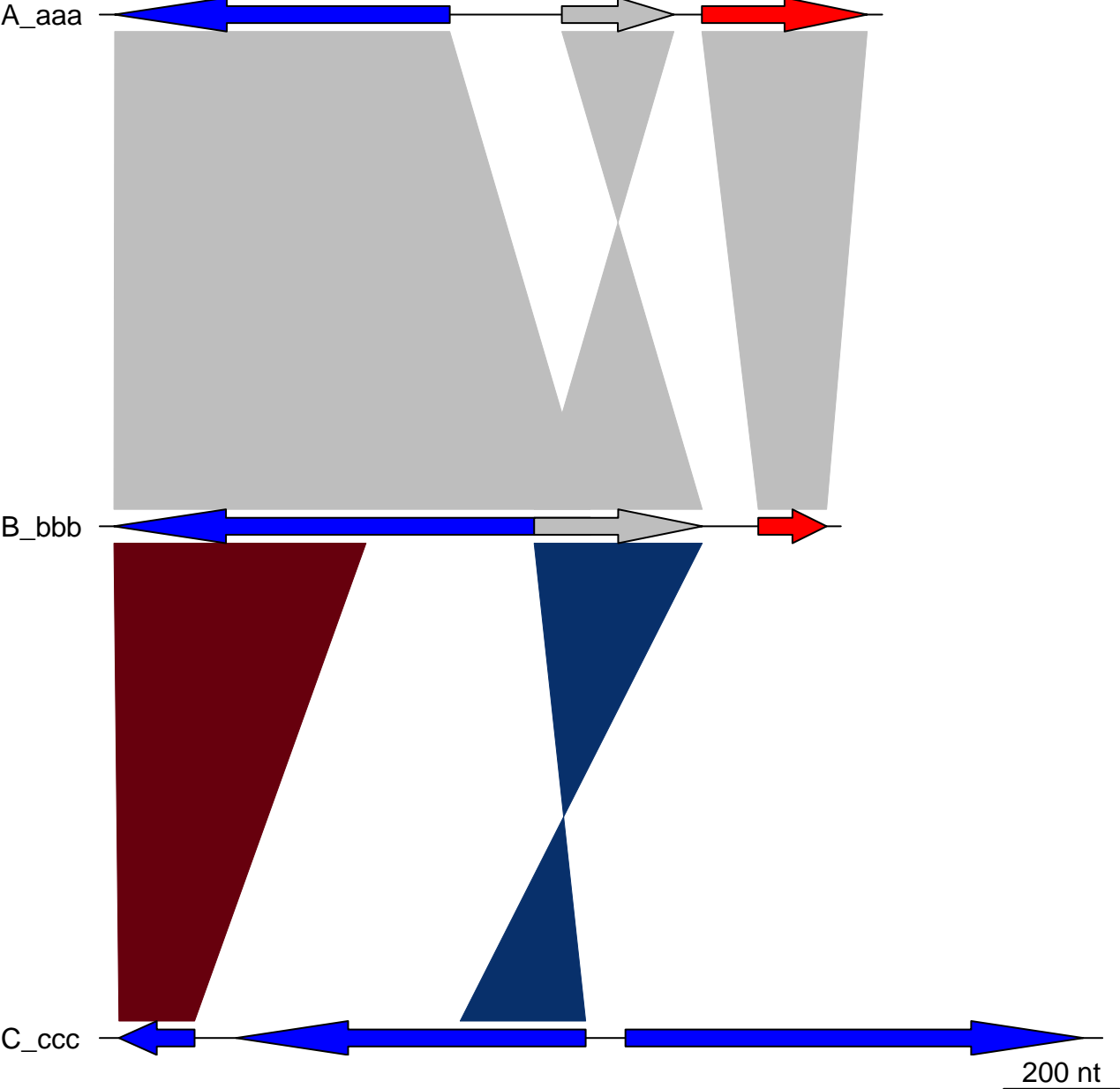
C_ccc



200 nt

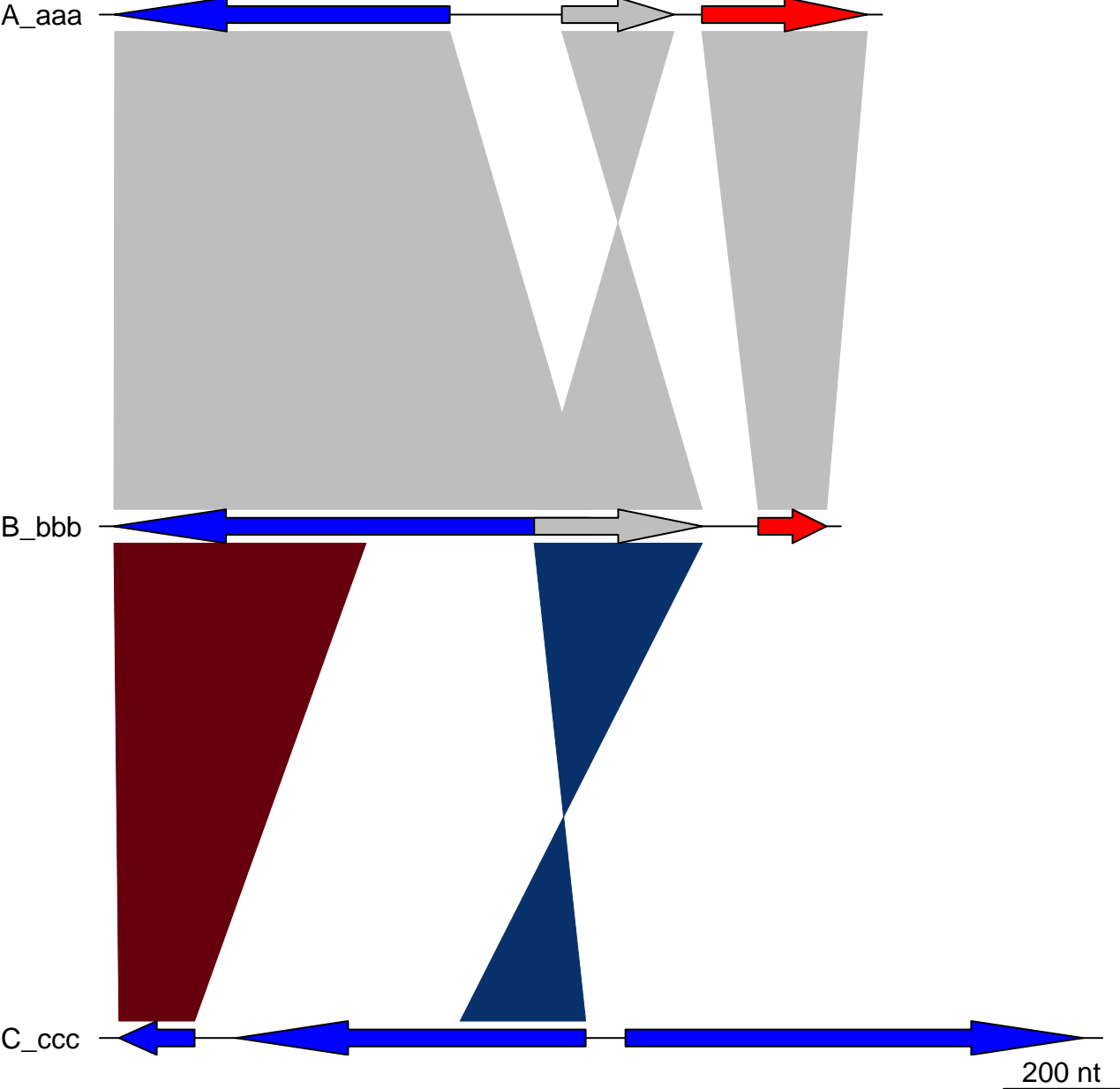
help("plot_gene_map")

Comparison of A, B and C



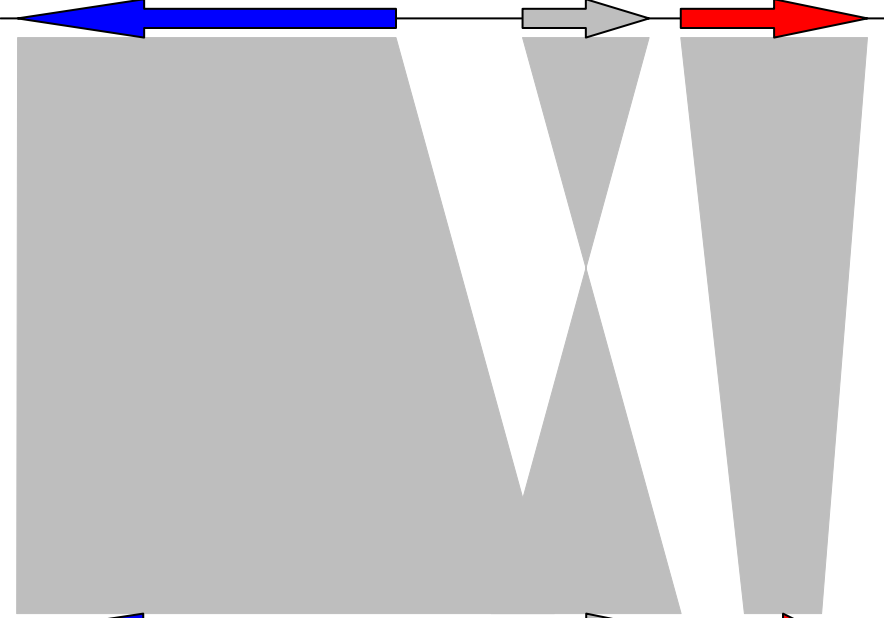
help("plot_gene_map")

Comparison of A, B and C

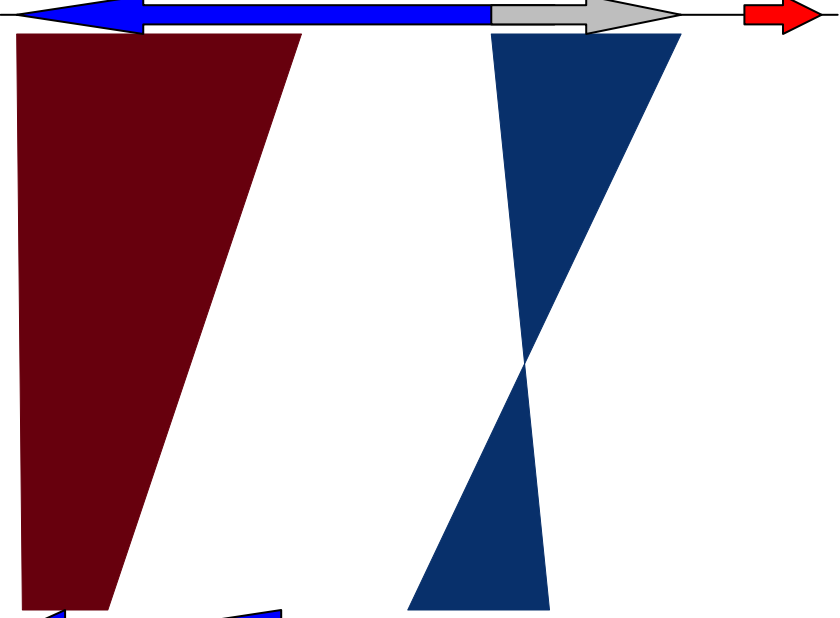


help("plot_gene_map")

Huey



Dewey



Louie



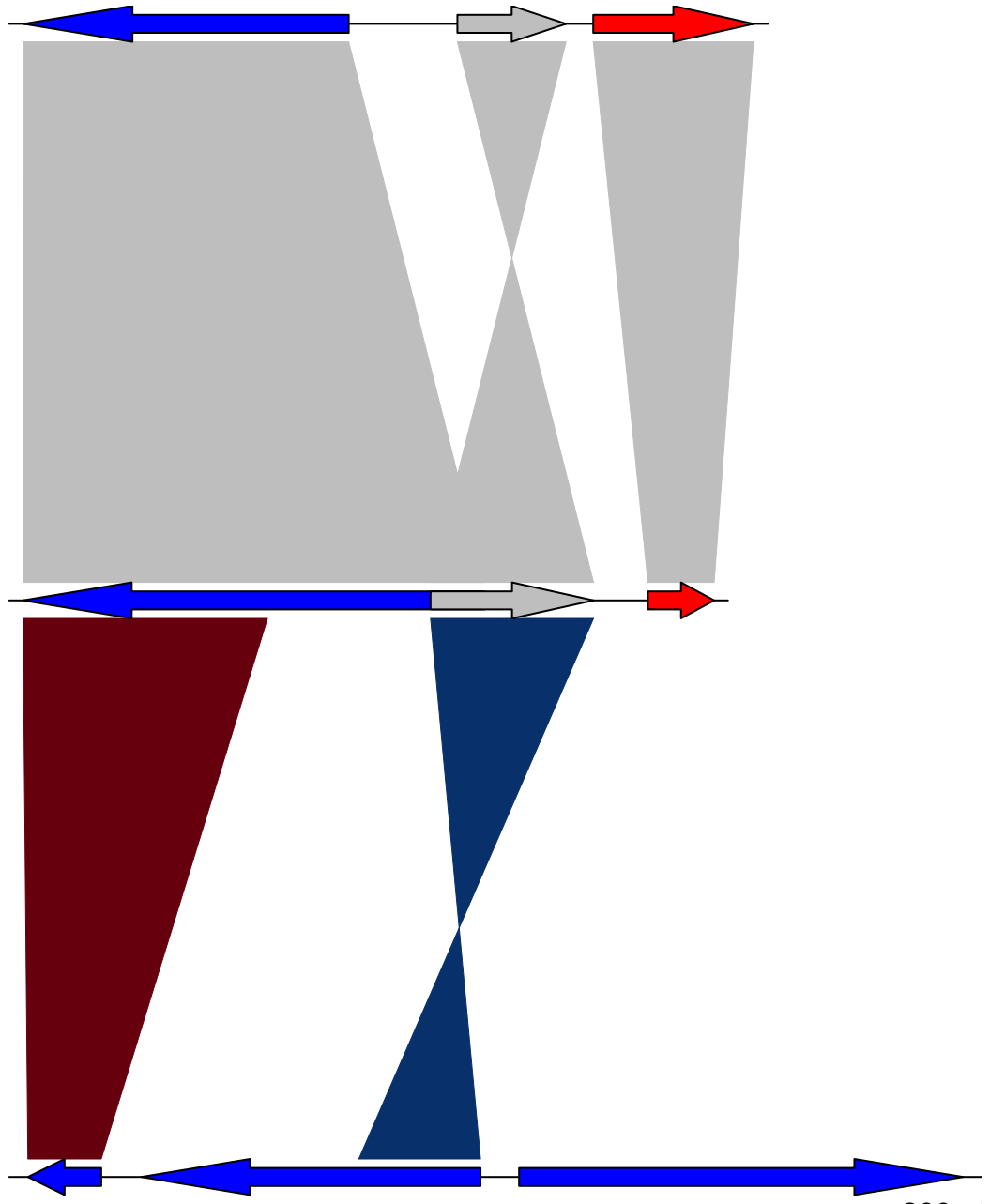
200 nt

help("plot_gene_map")

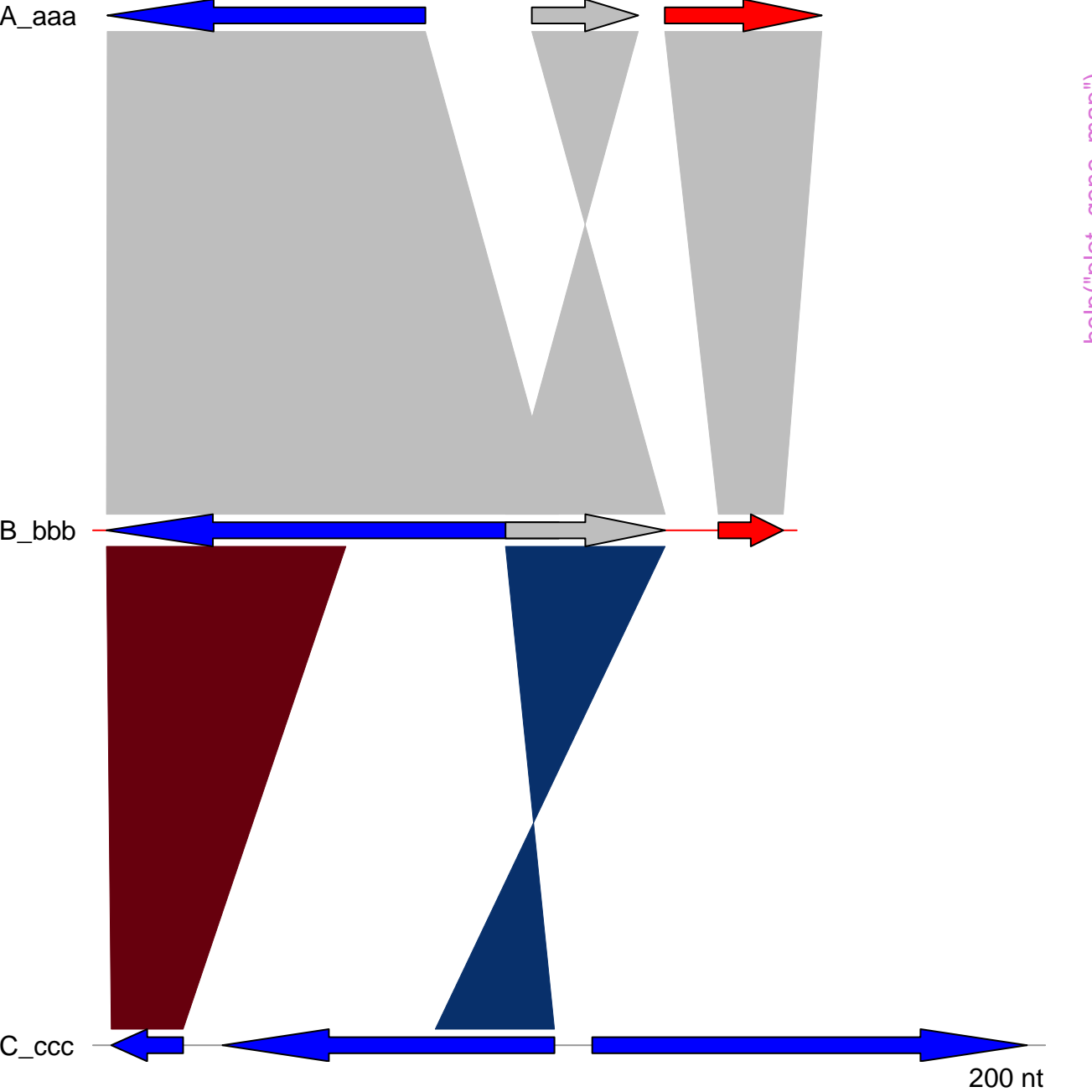
Huey

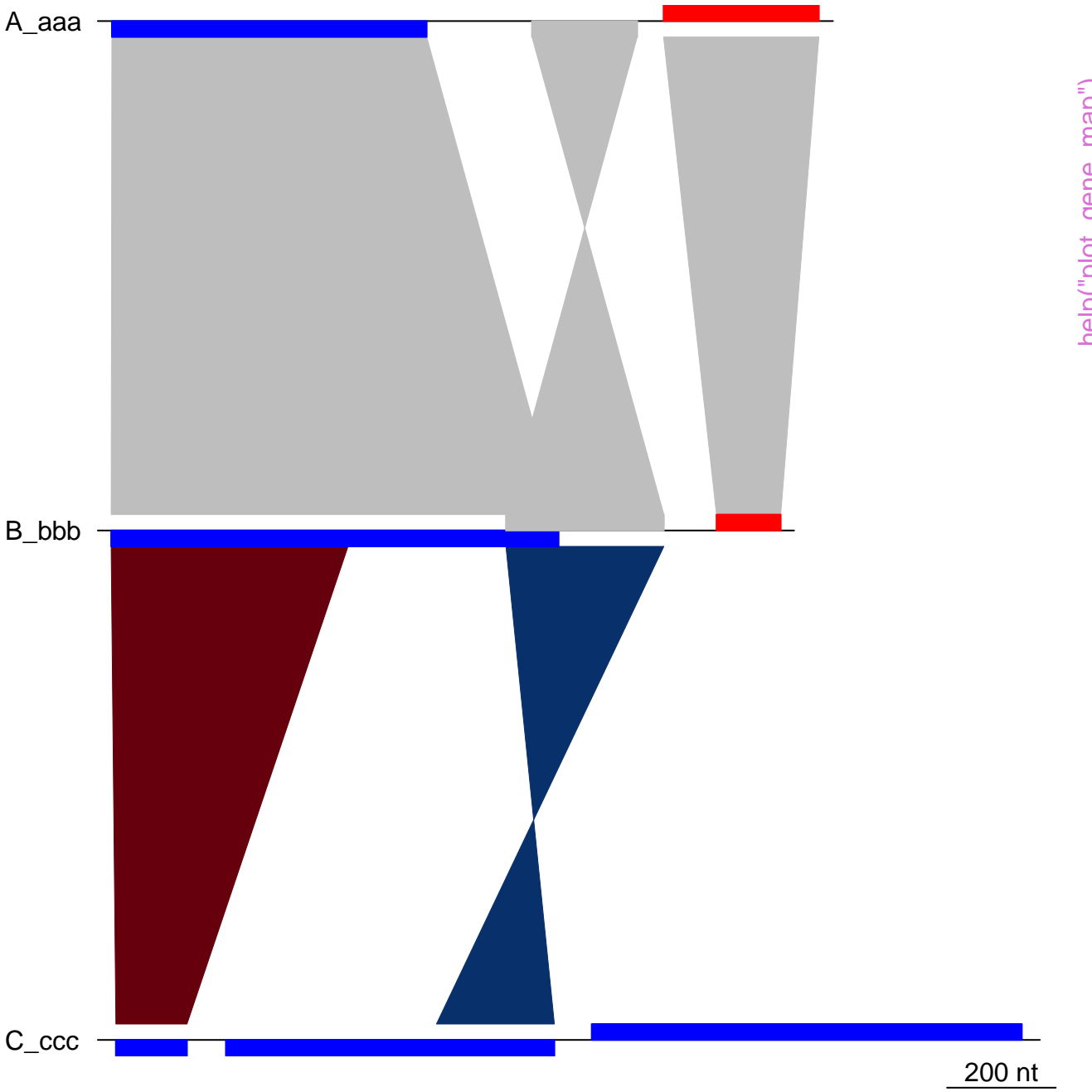
Dewey

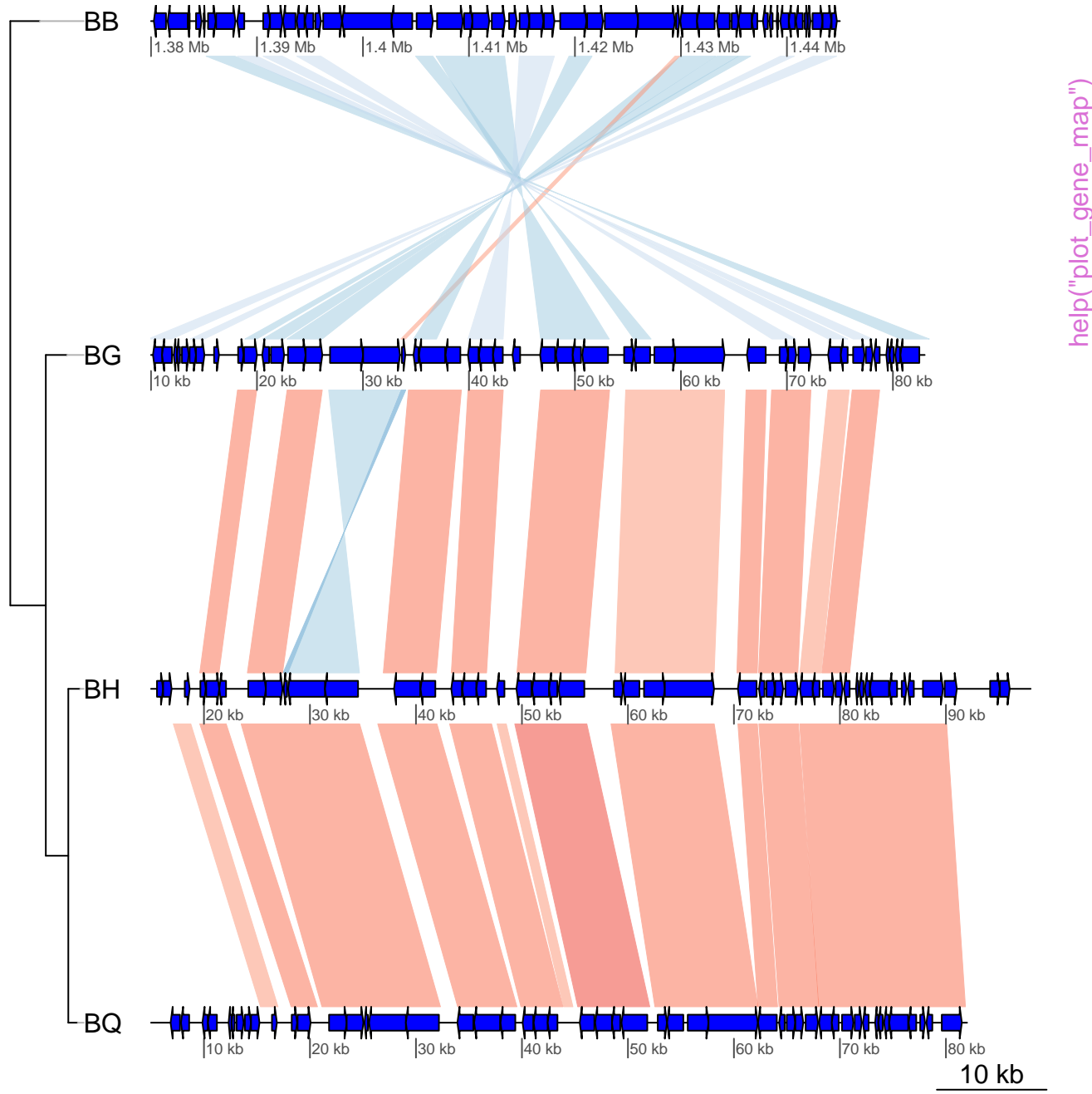
Louie

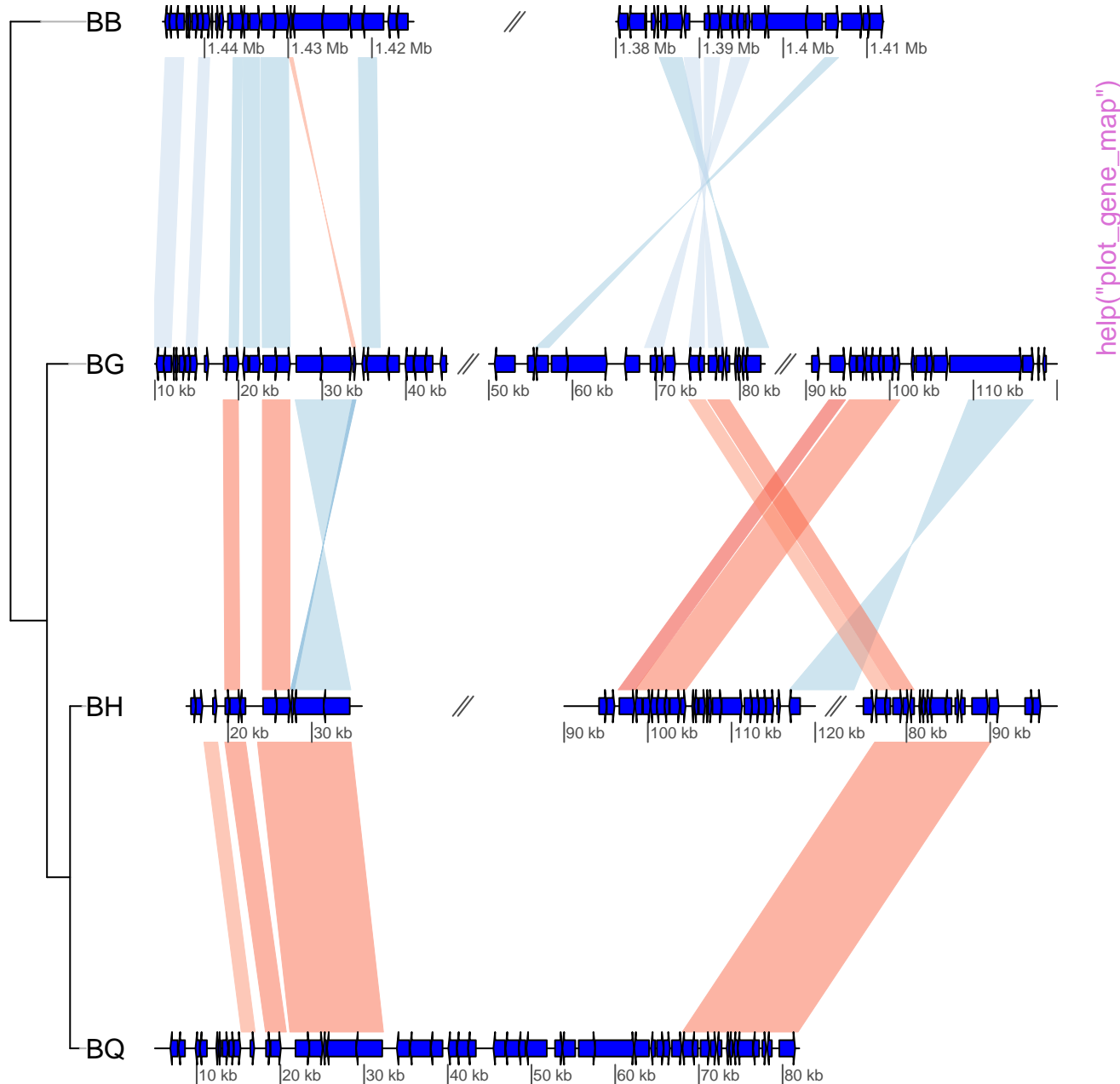


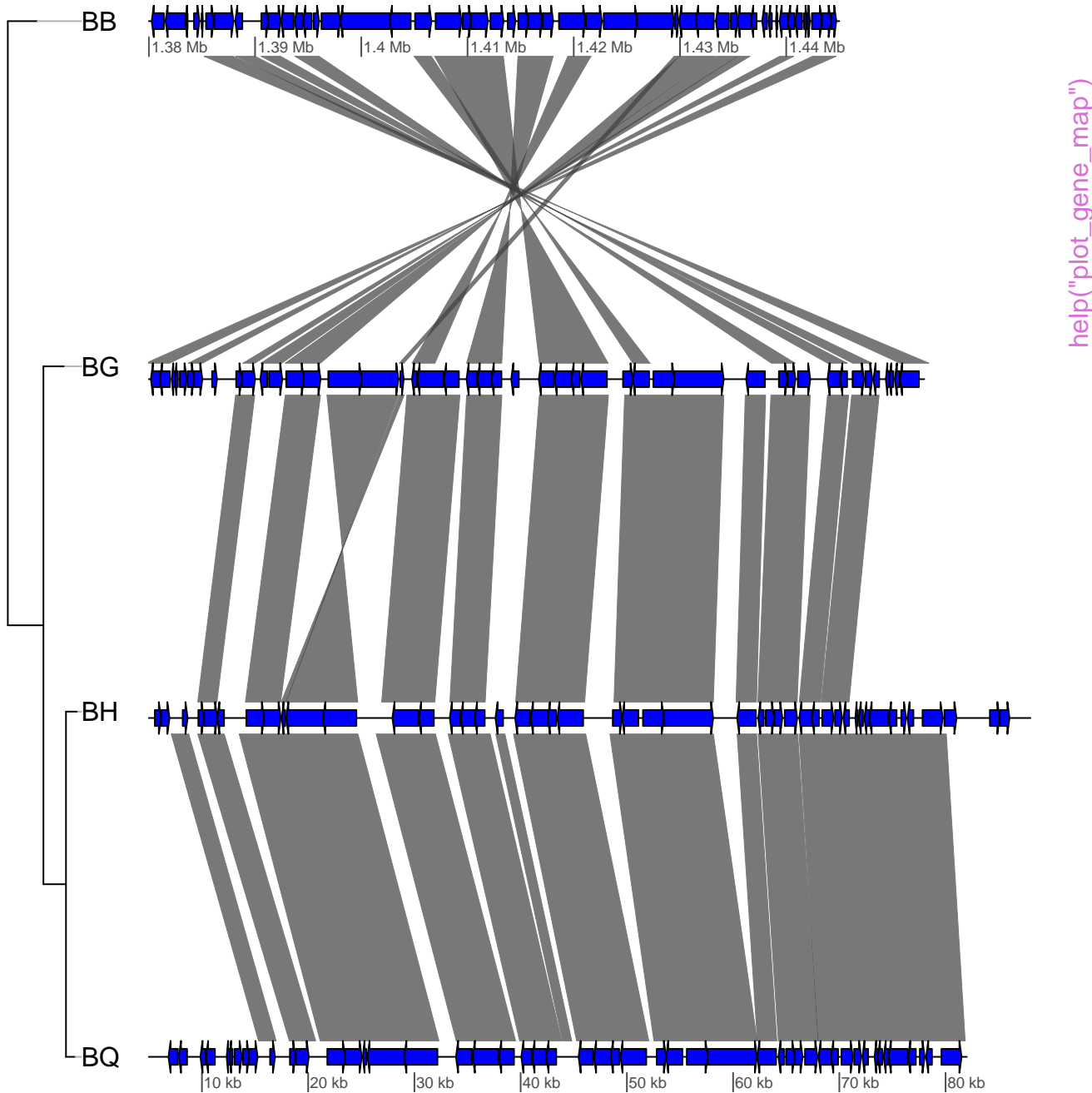
help("plot_gene_map")

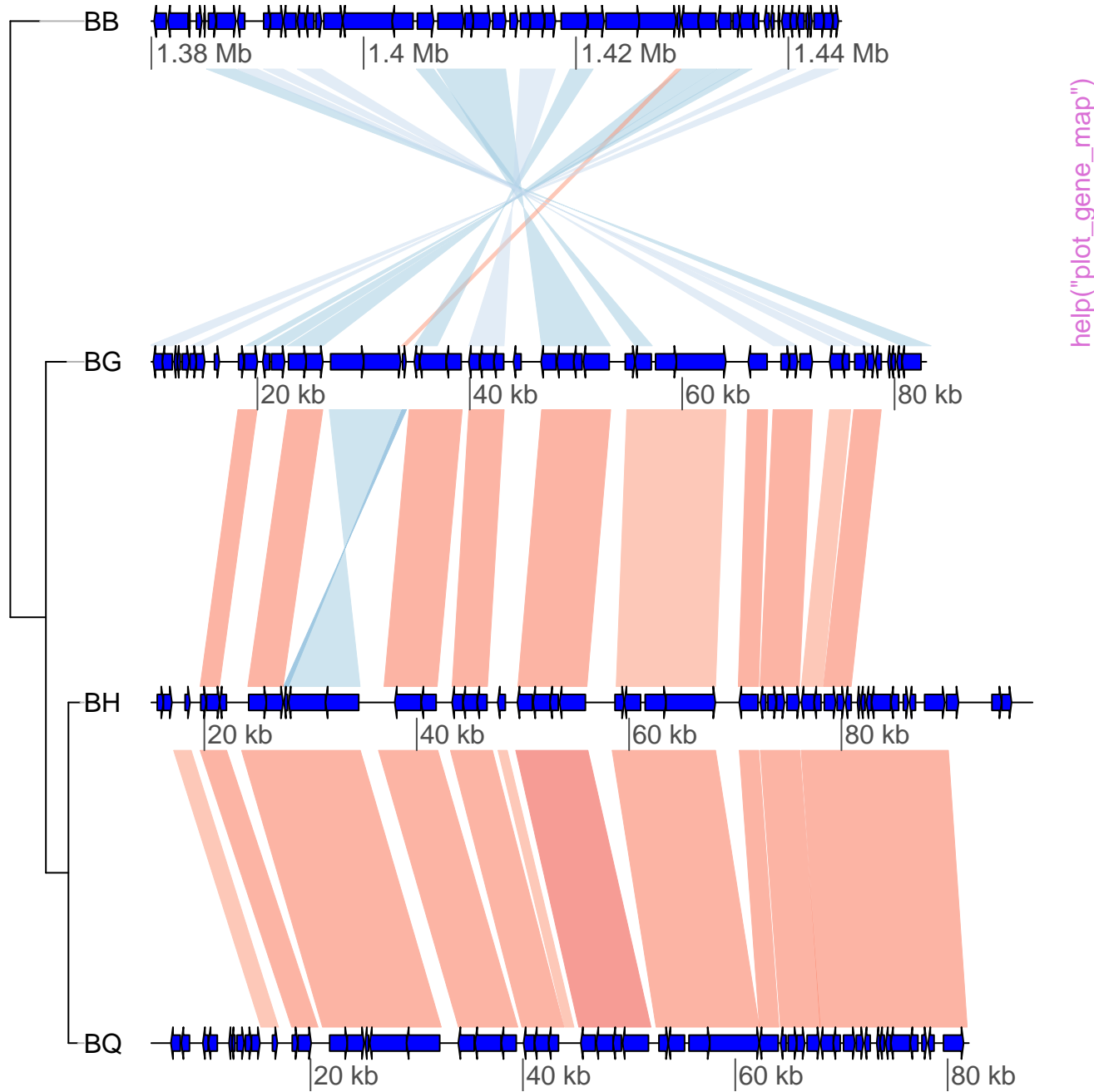




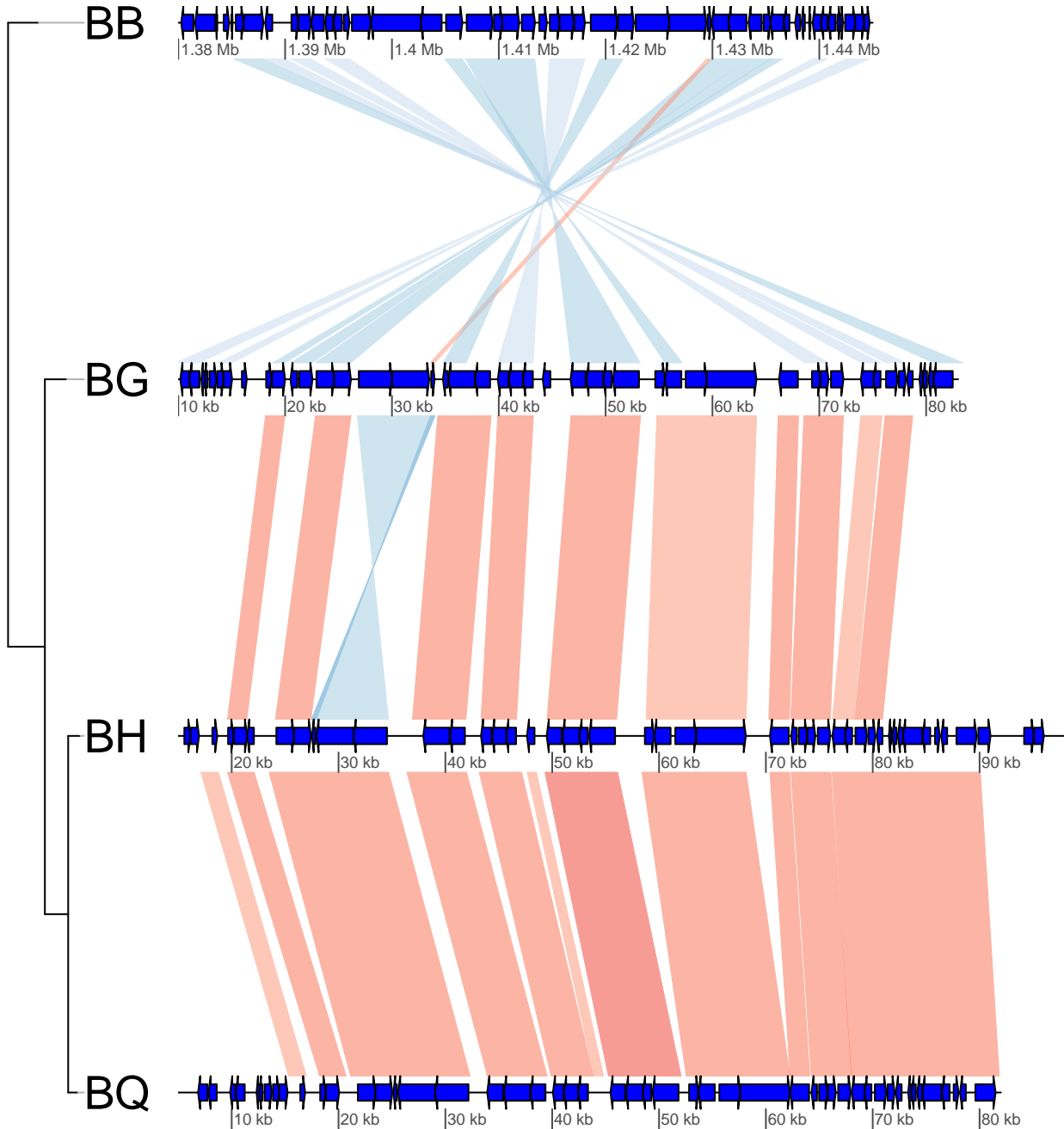


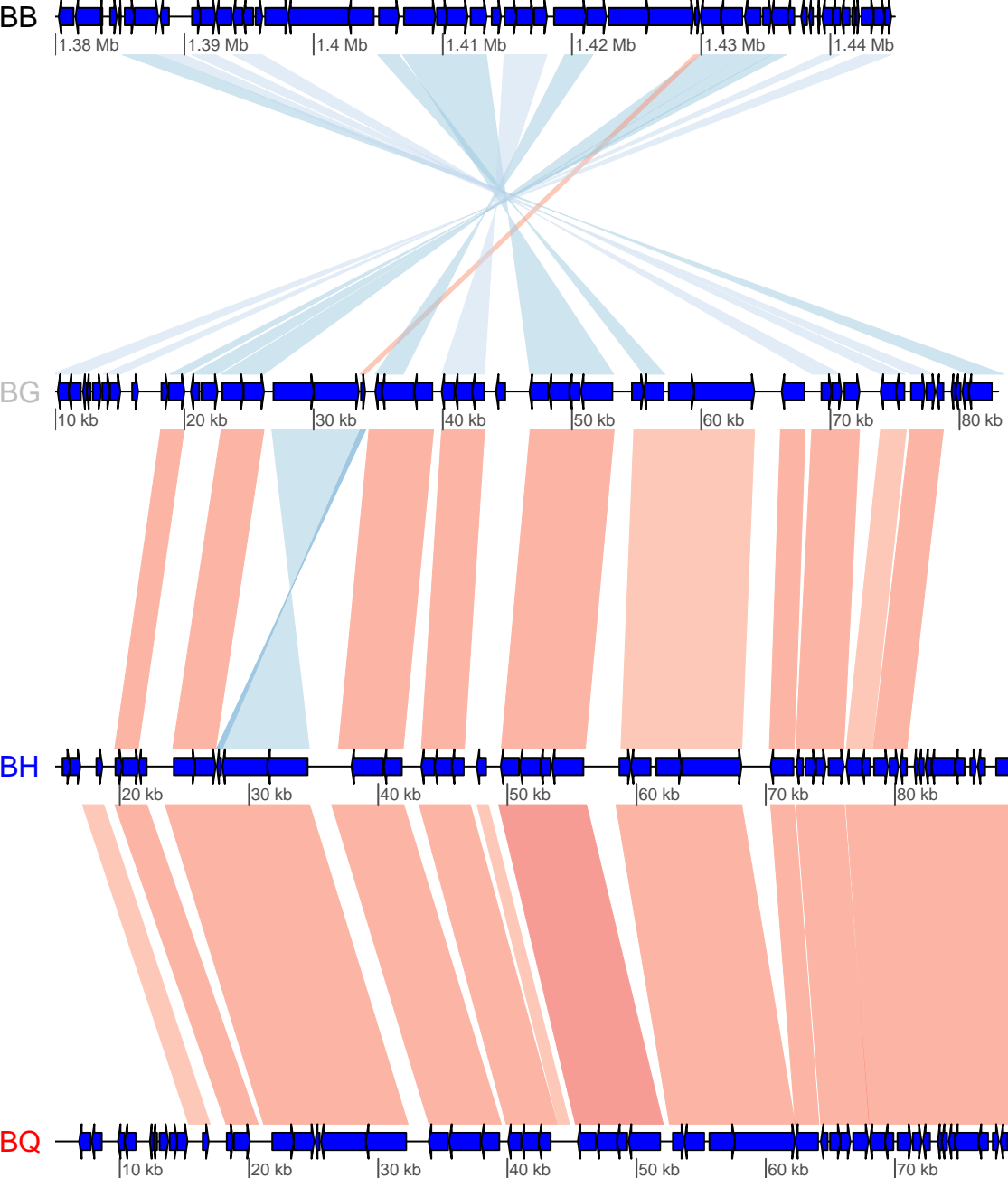




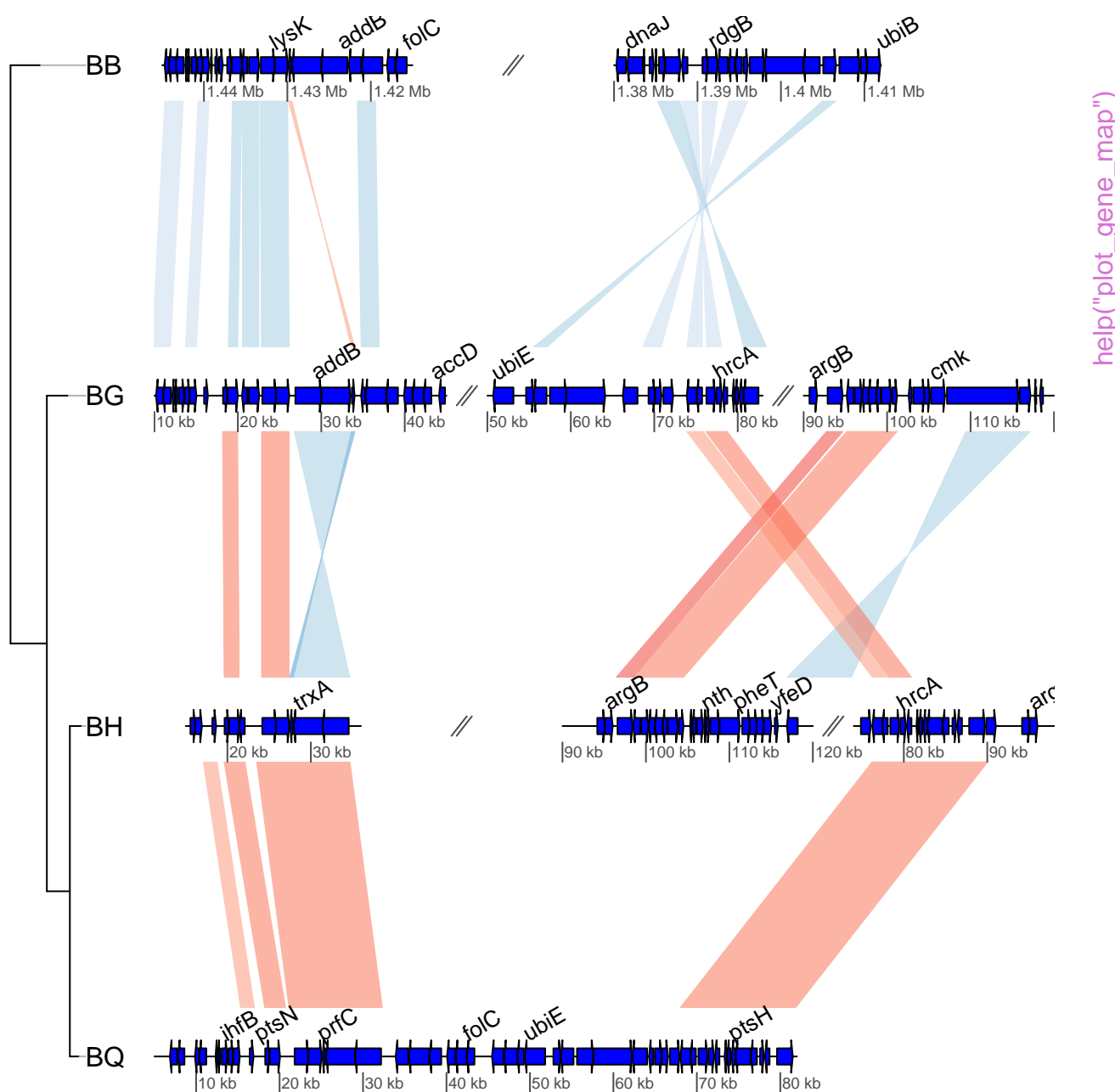


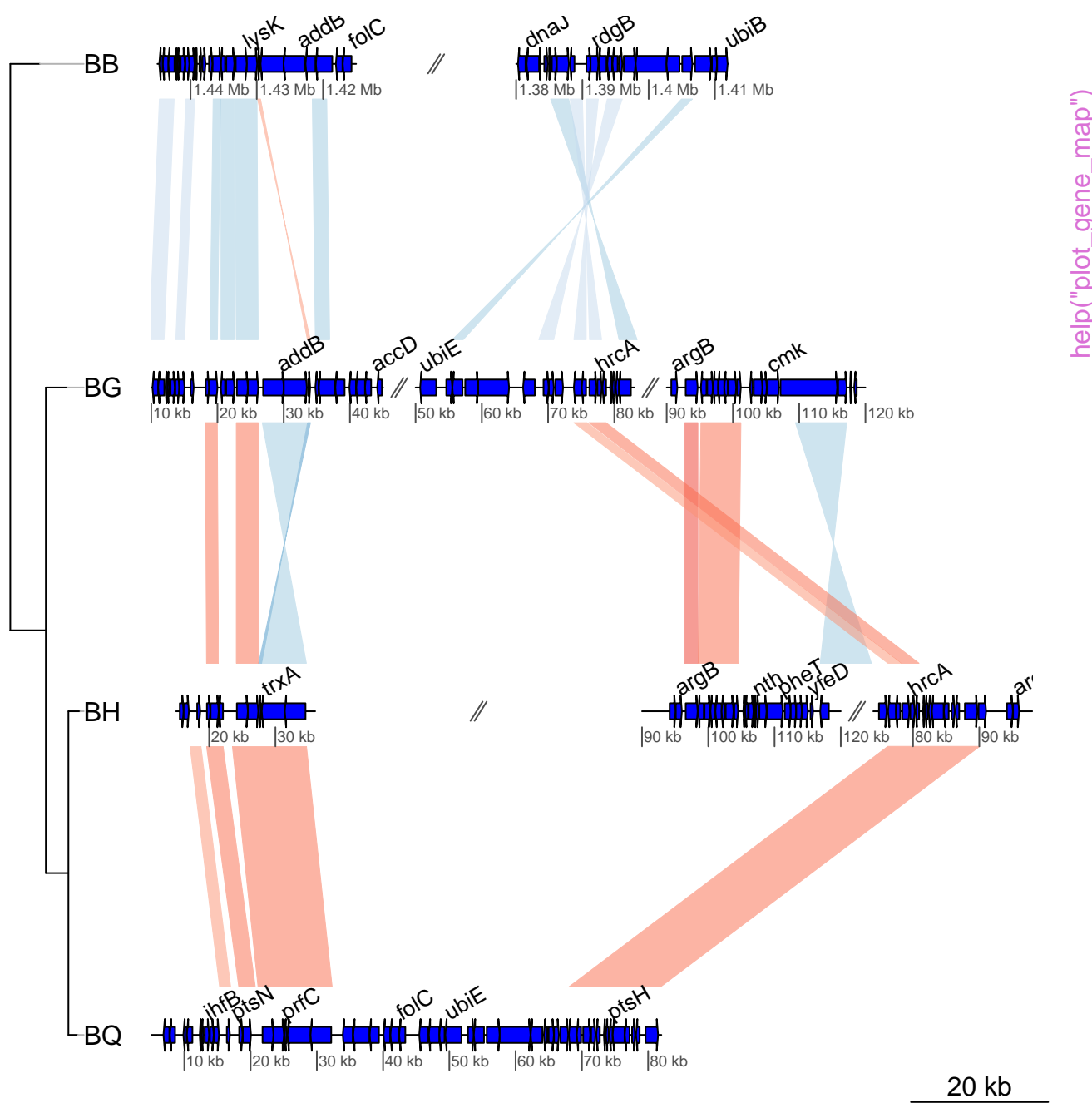
help("plot_gene_map")

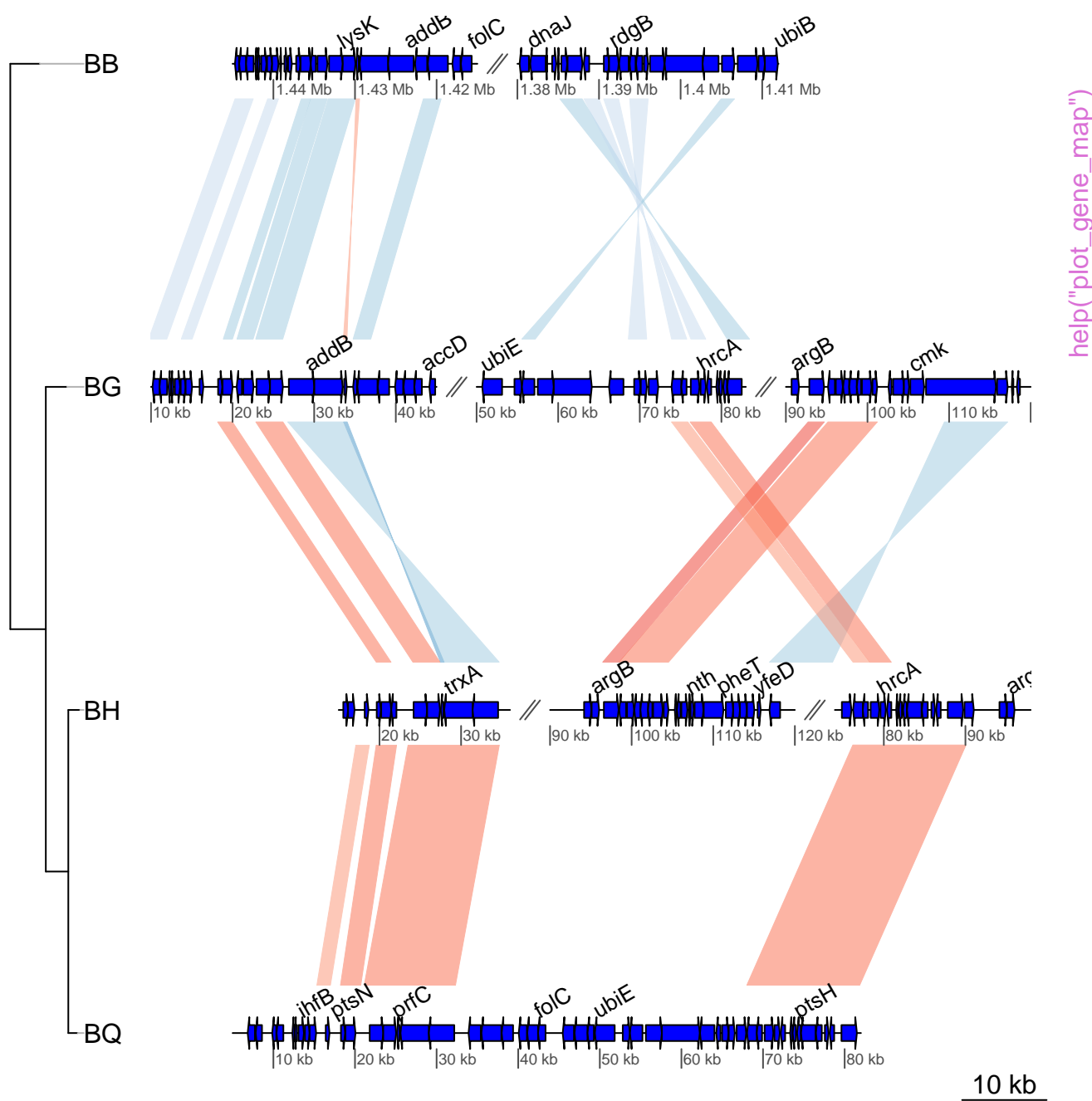


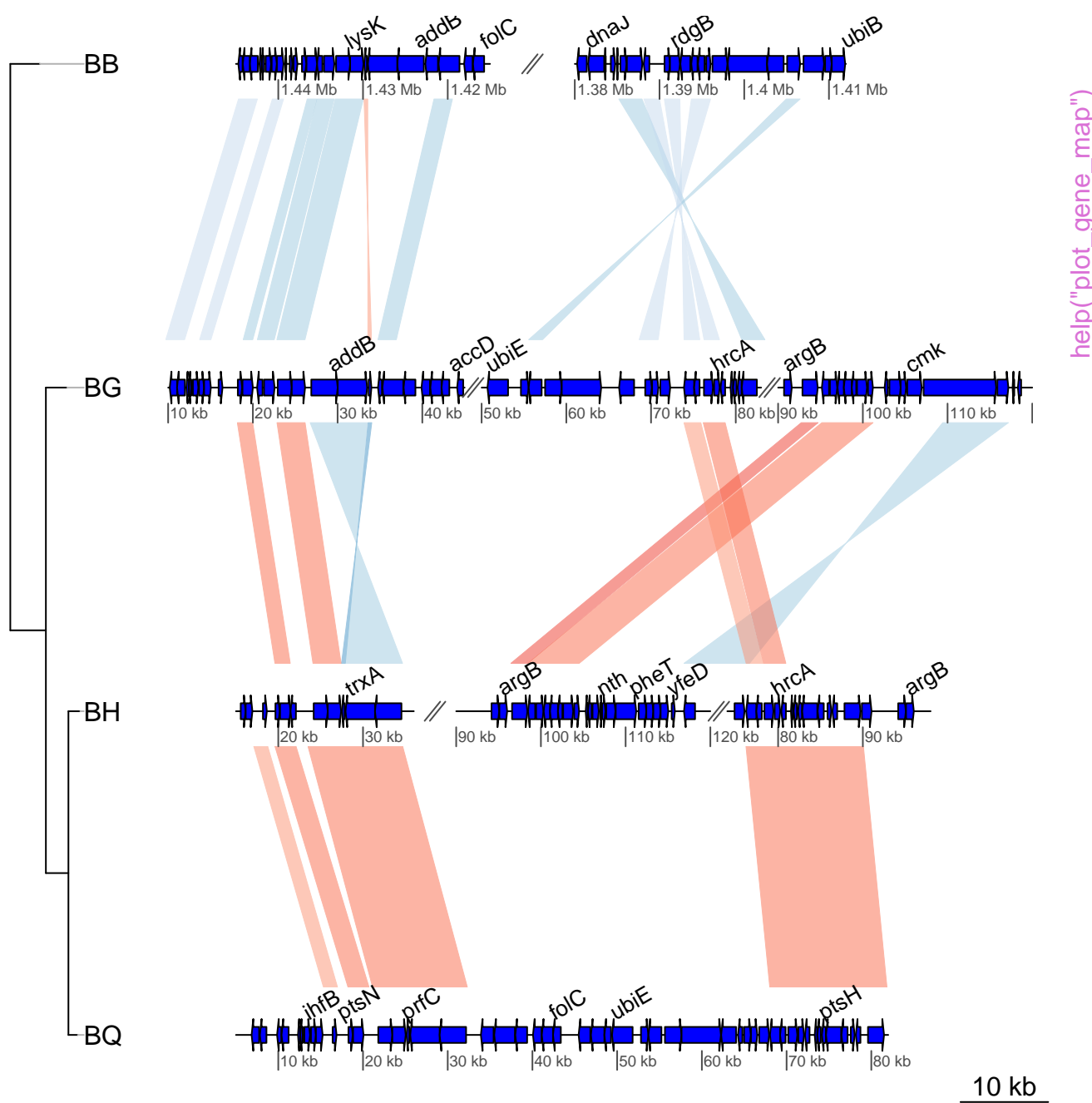


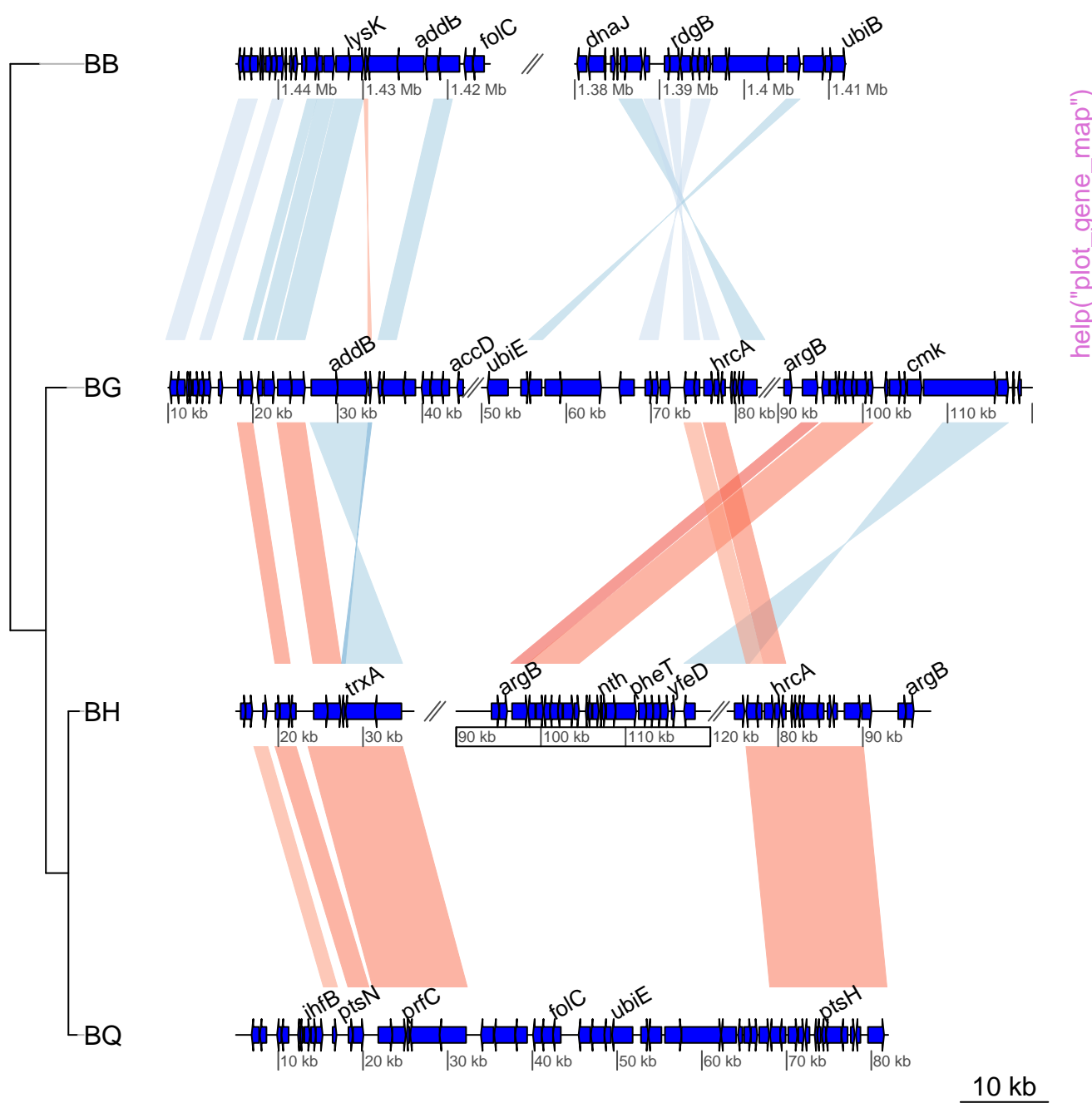
help("plot_gene_map")

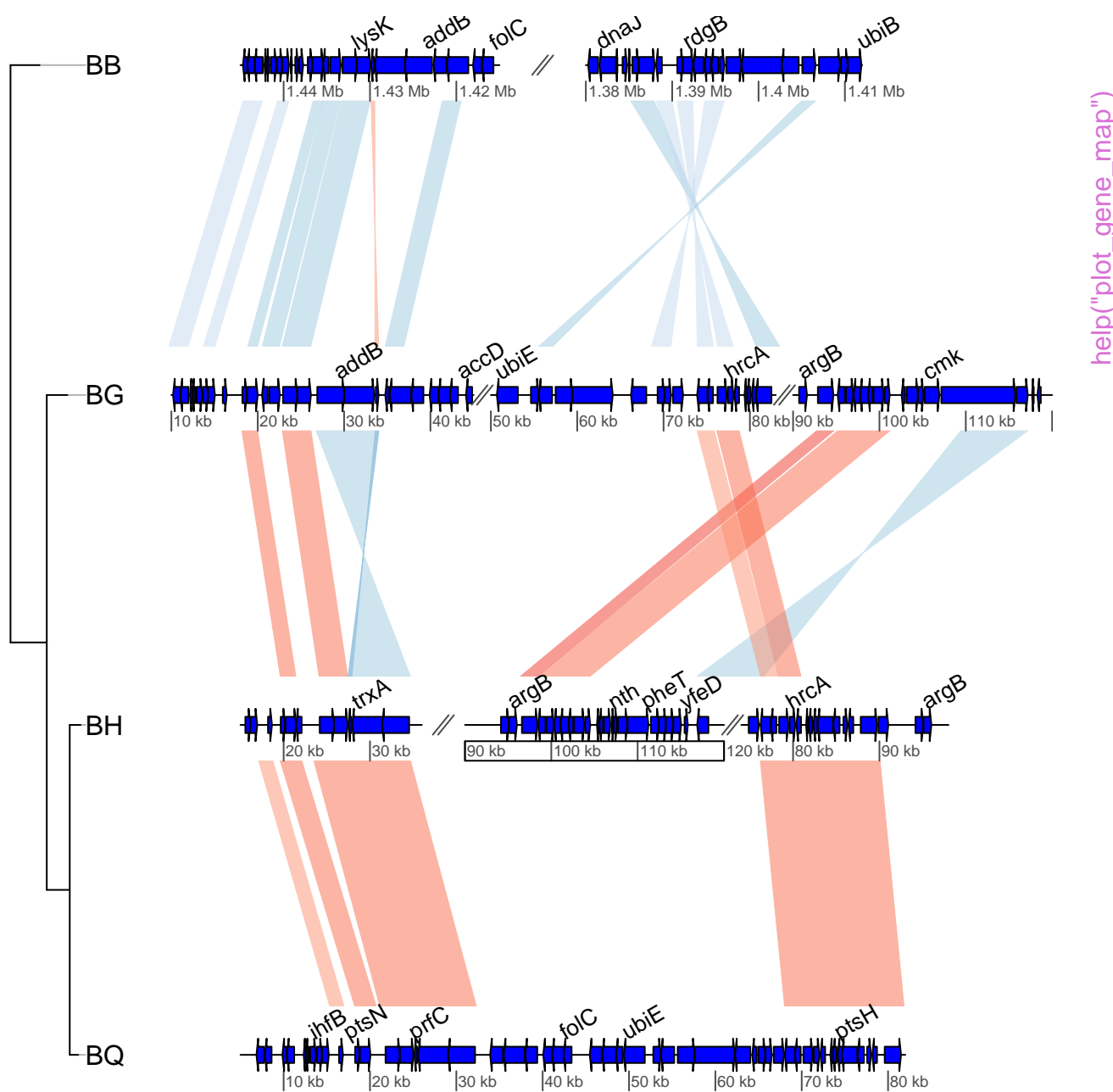


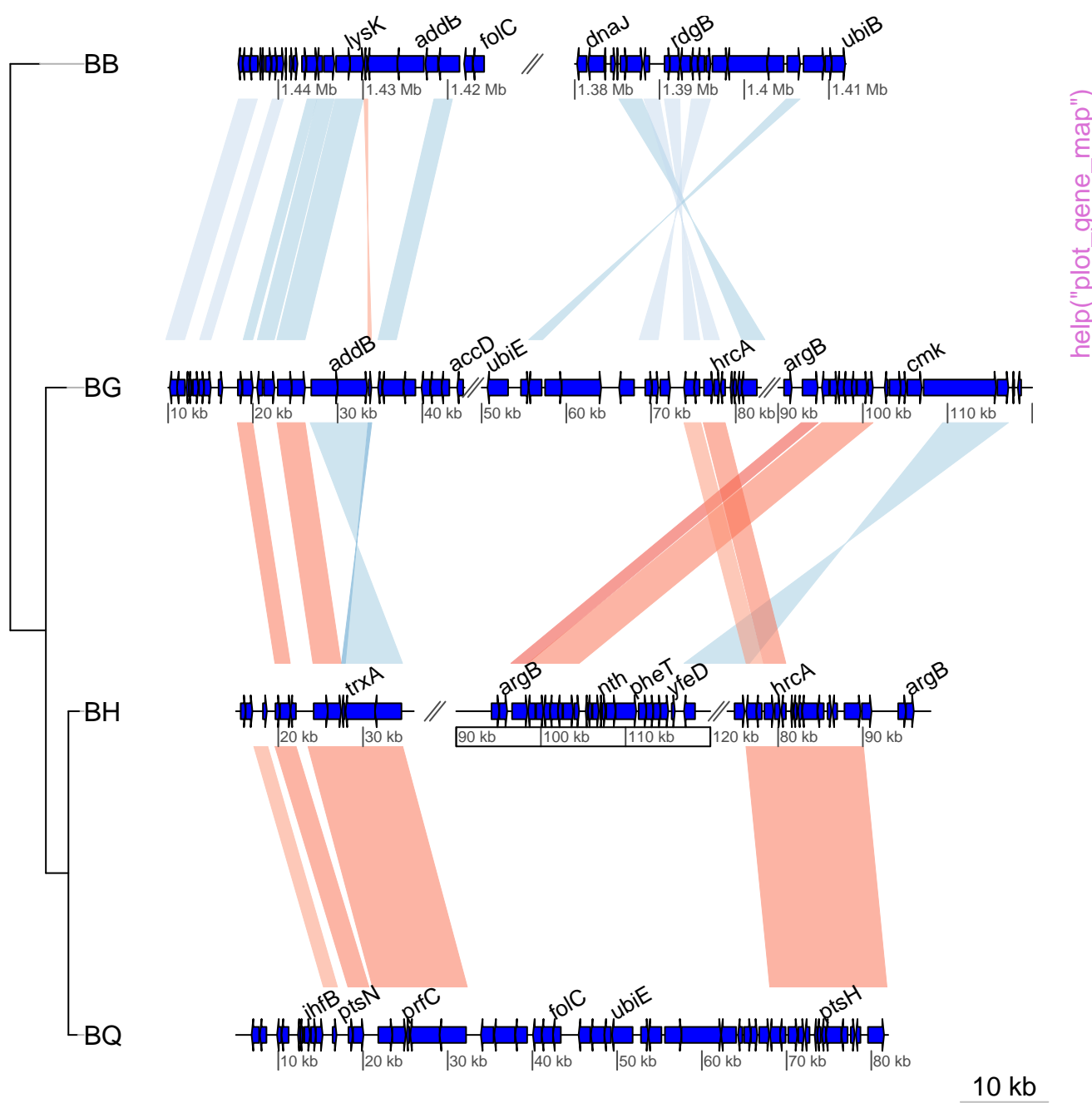












geno

R

Plot

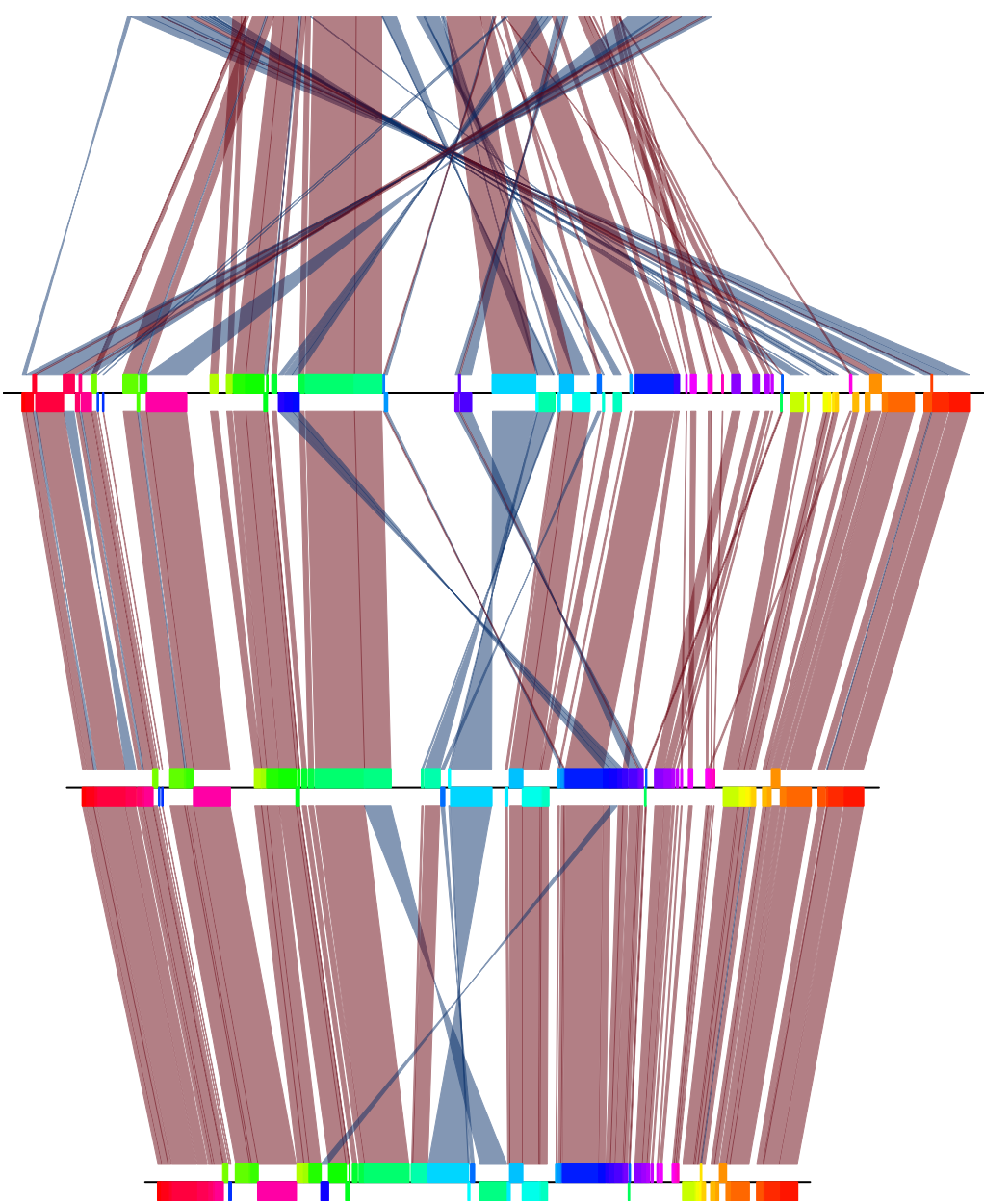
help("plot_gene_map")

B_bacilliformis

B_grahamii

B_henselae

B_quintana



help("read_functions")

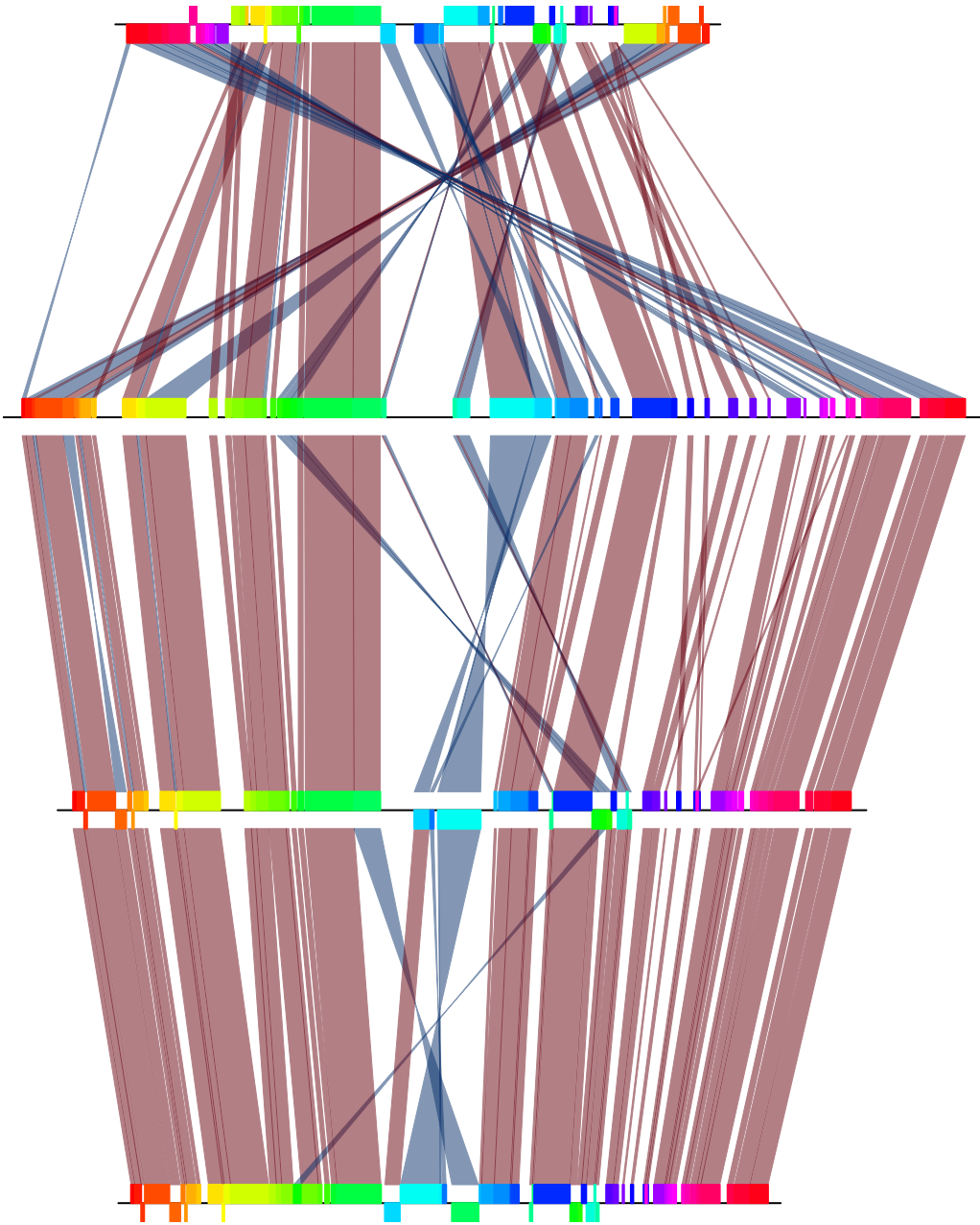
500 kb

B_bacilliformis

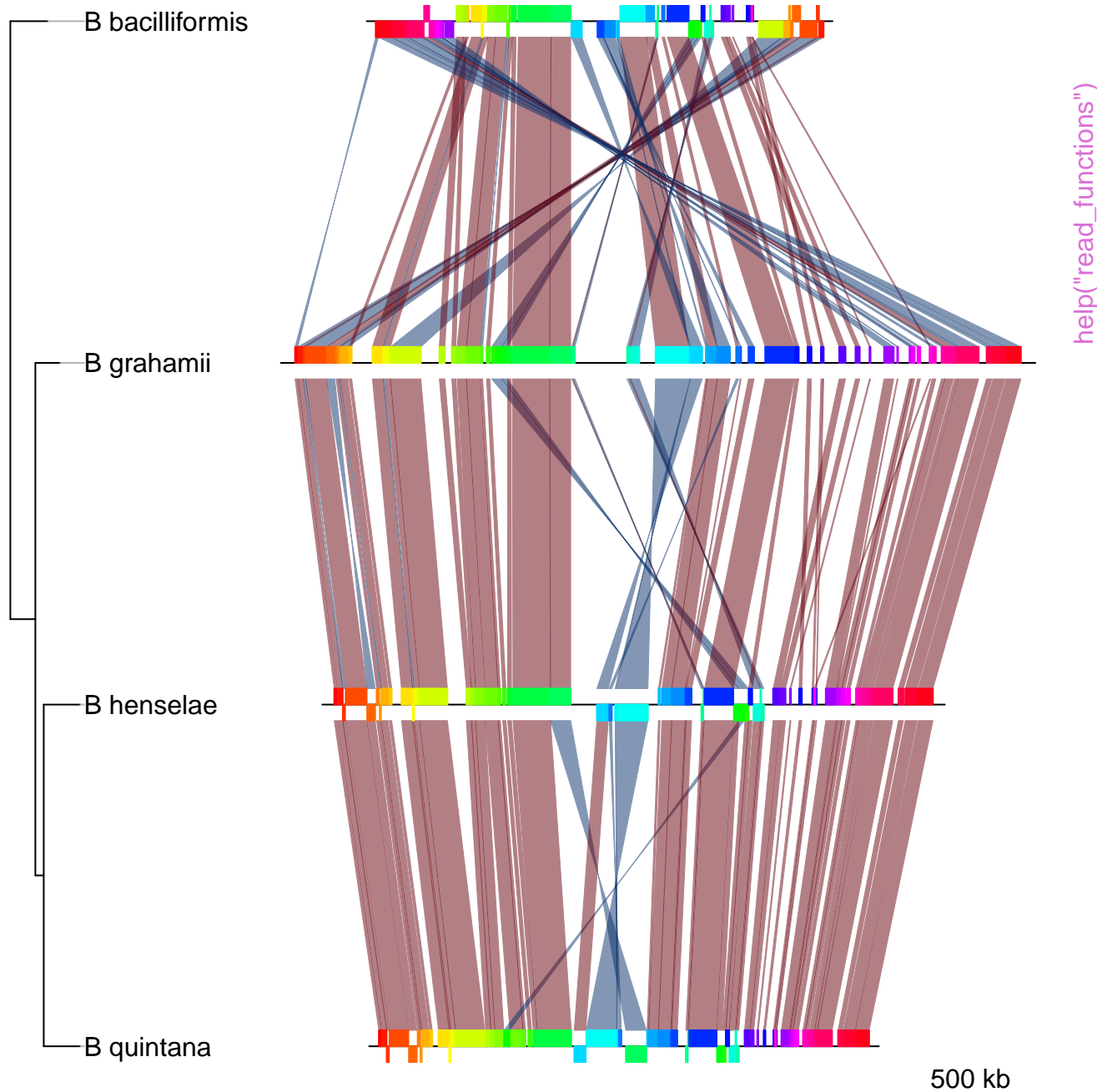
B_grahamii

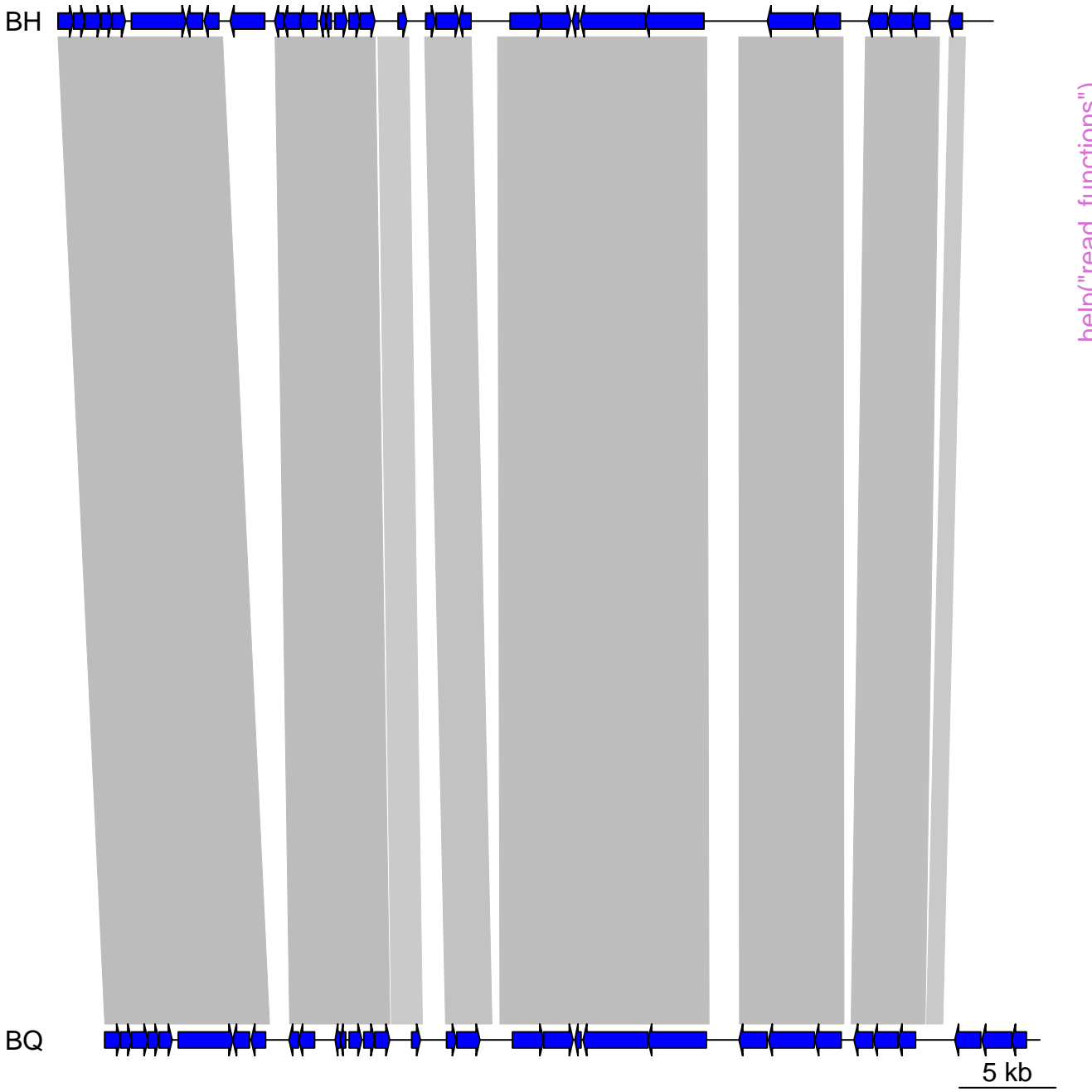
B_henselae

B_quintana



help("read_functions")



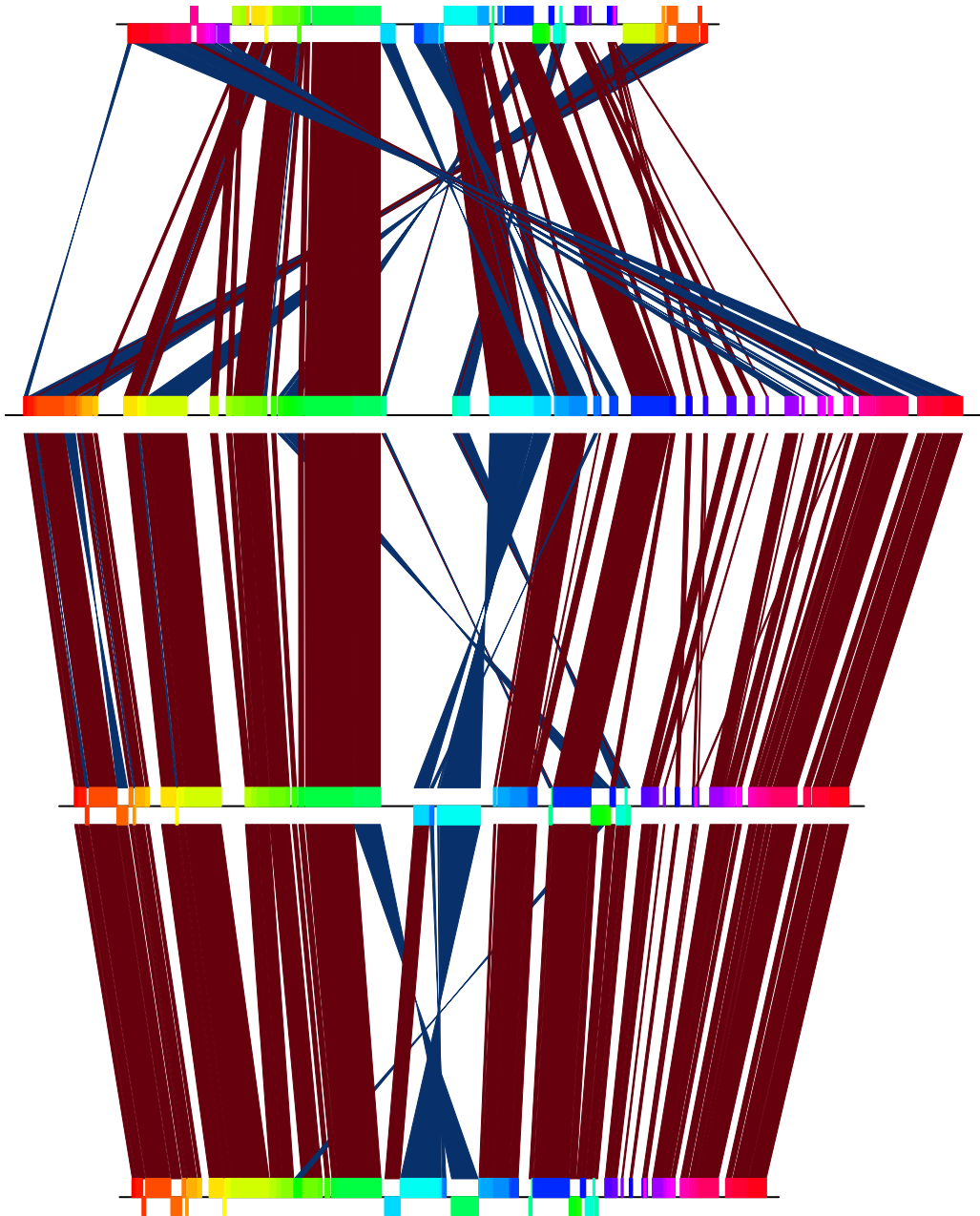


B_bacilliformis

B_grahamii

B_henselae

B_quintana



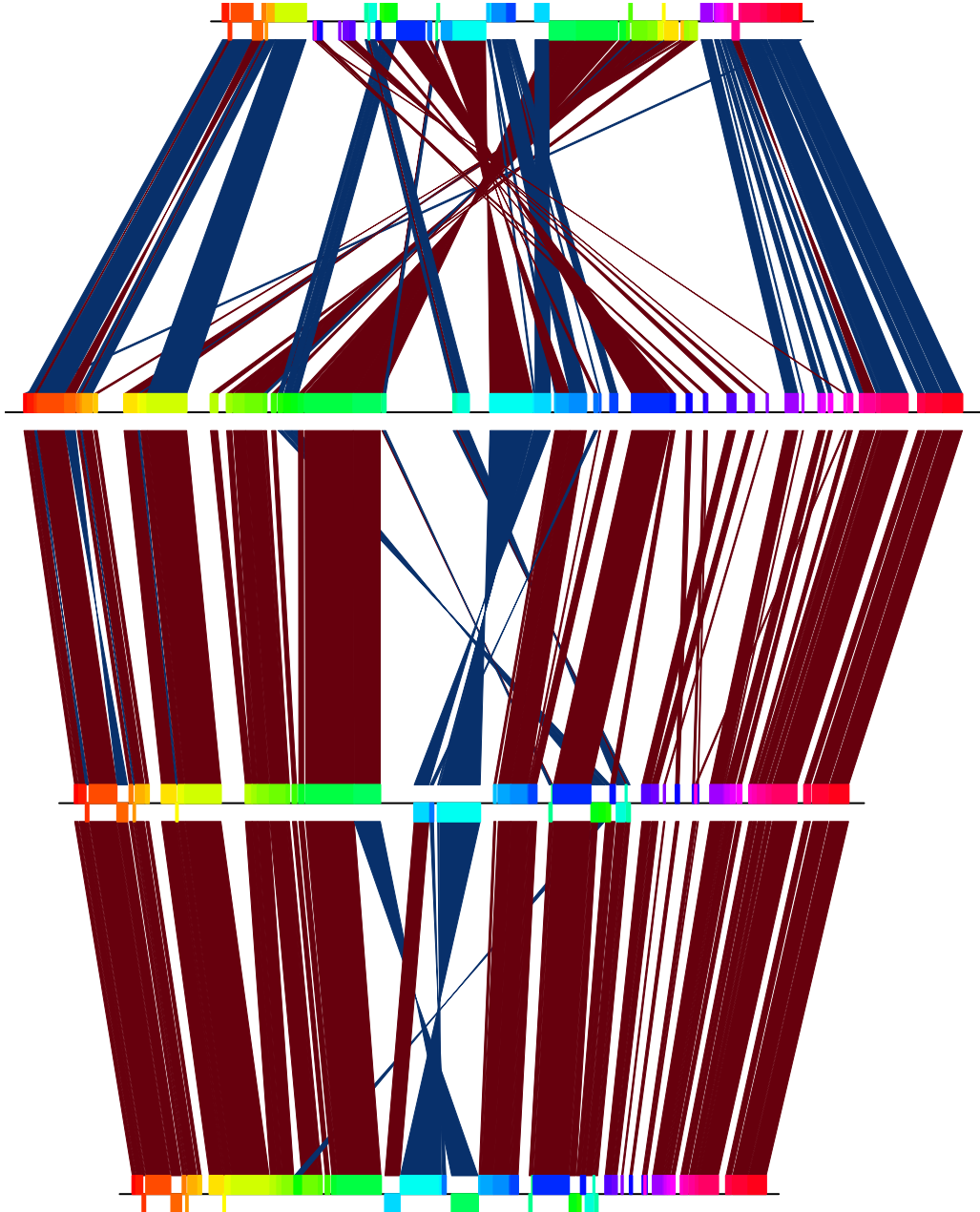
help("reverse")

B_bacilliformis

B_grahamii

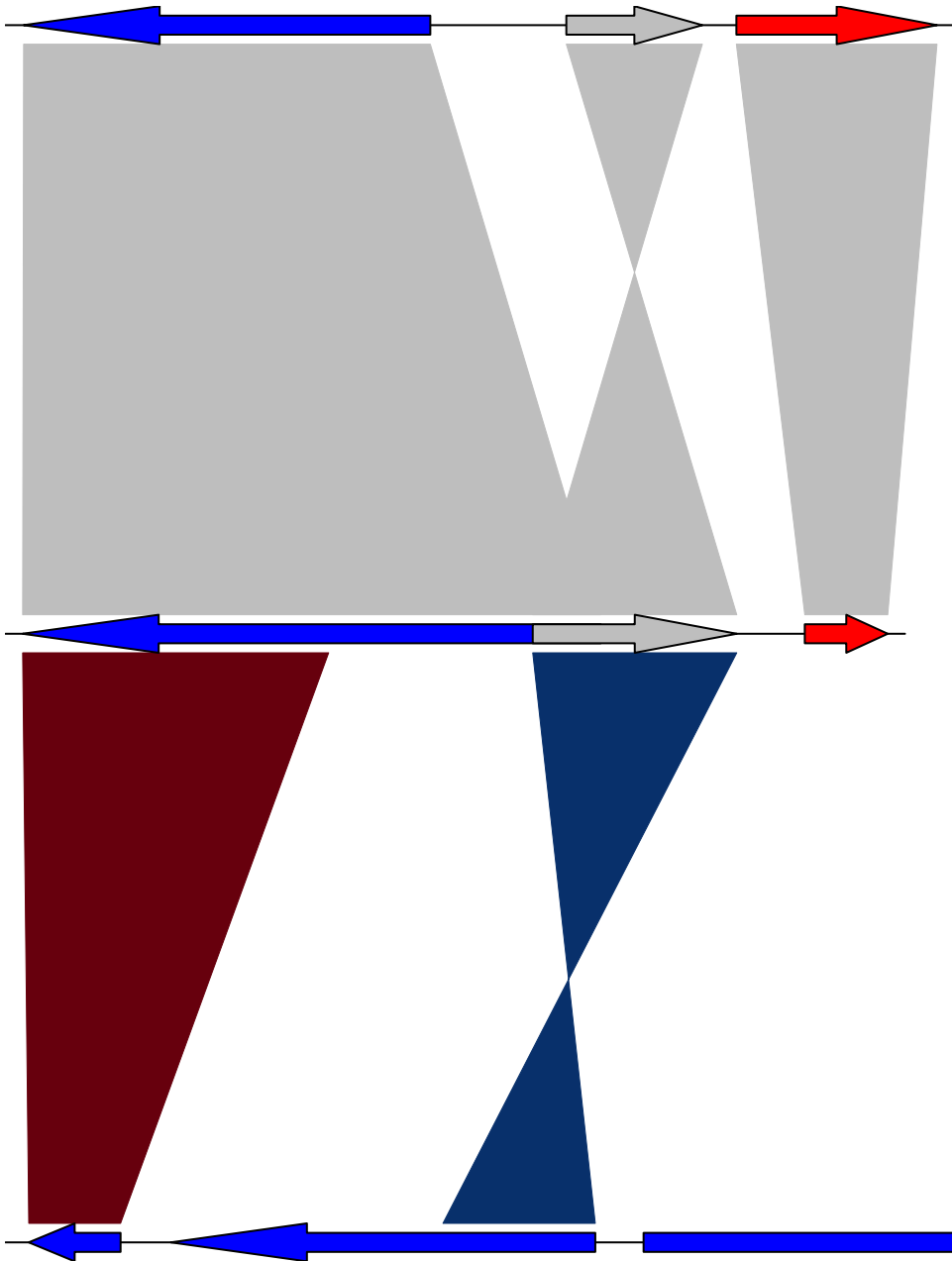
B_henselae

B_quintana



help("reverse")

500 kb



help("three_genes")

