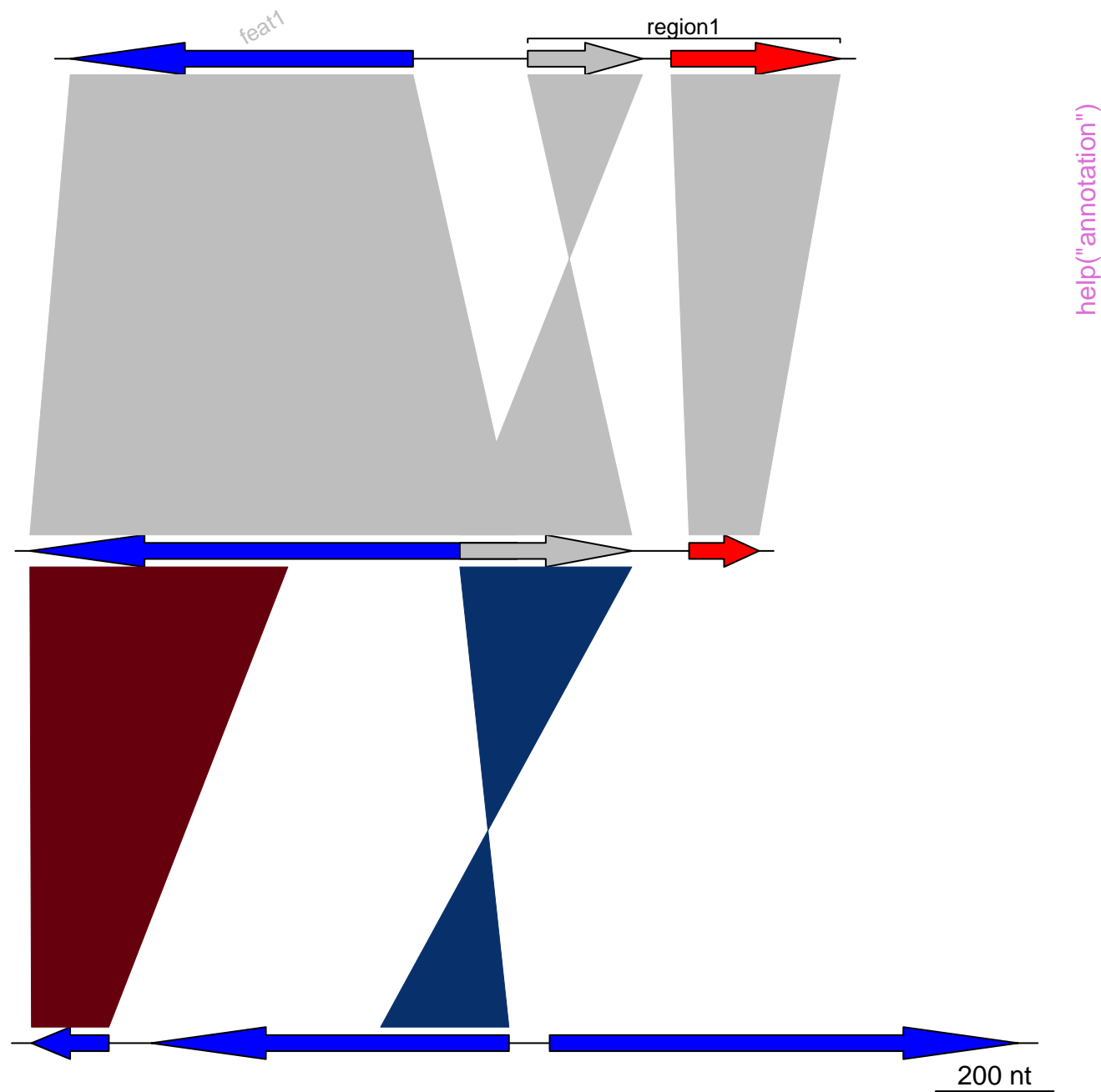
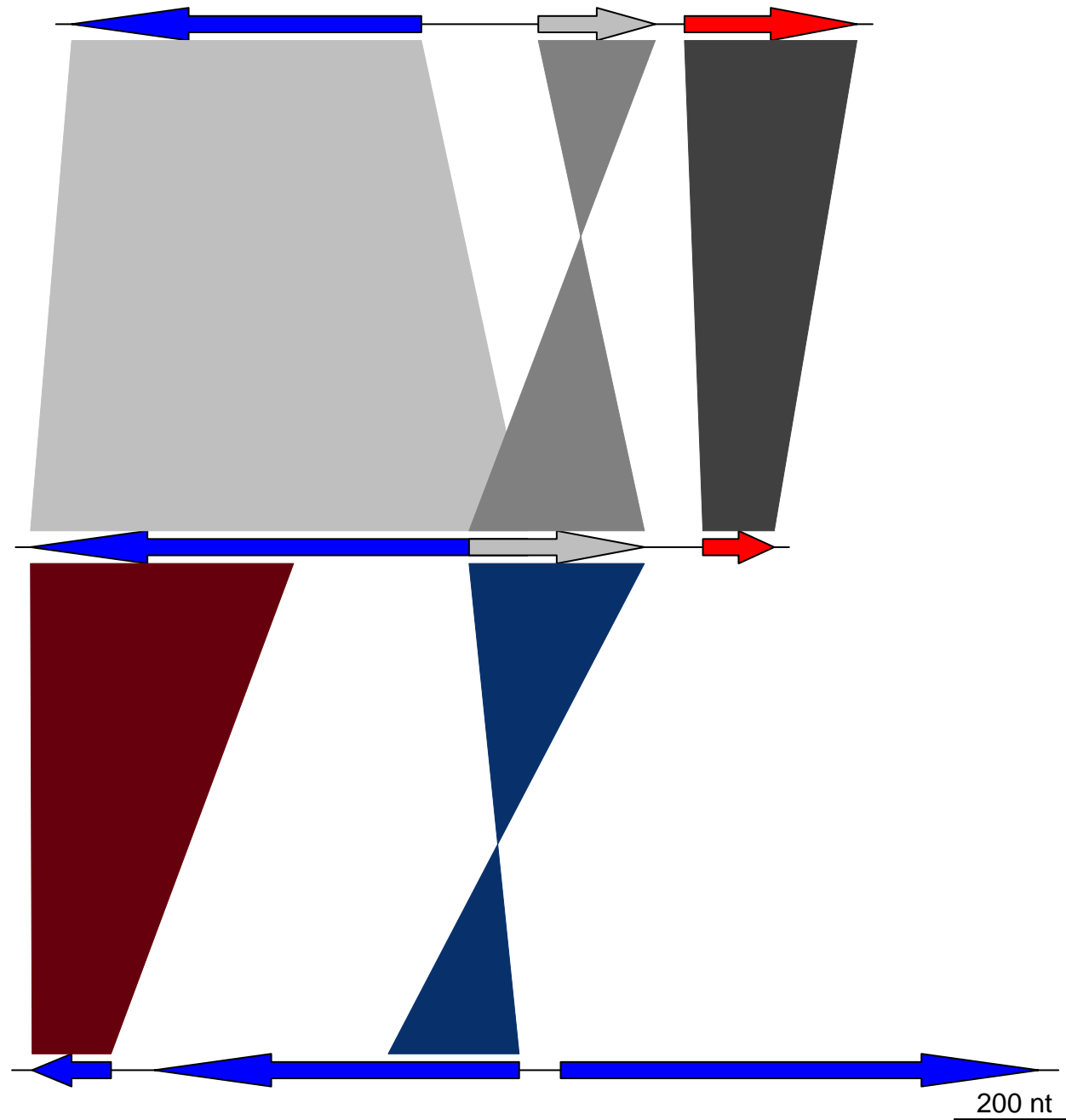
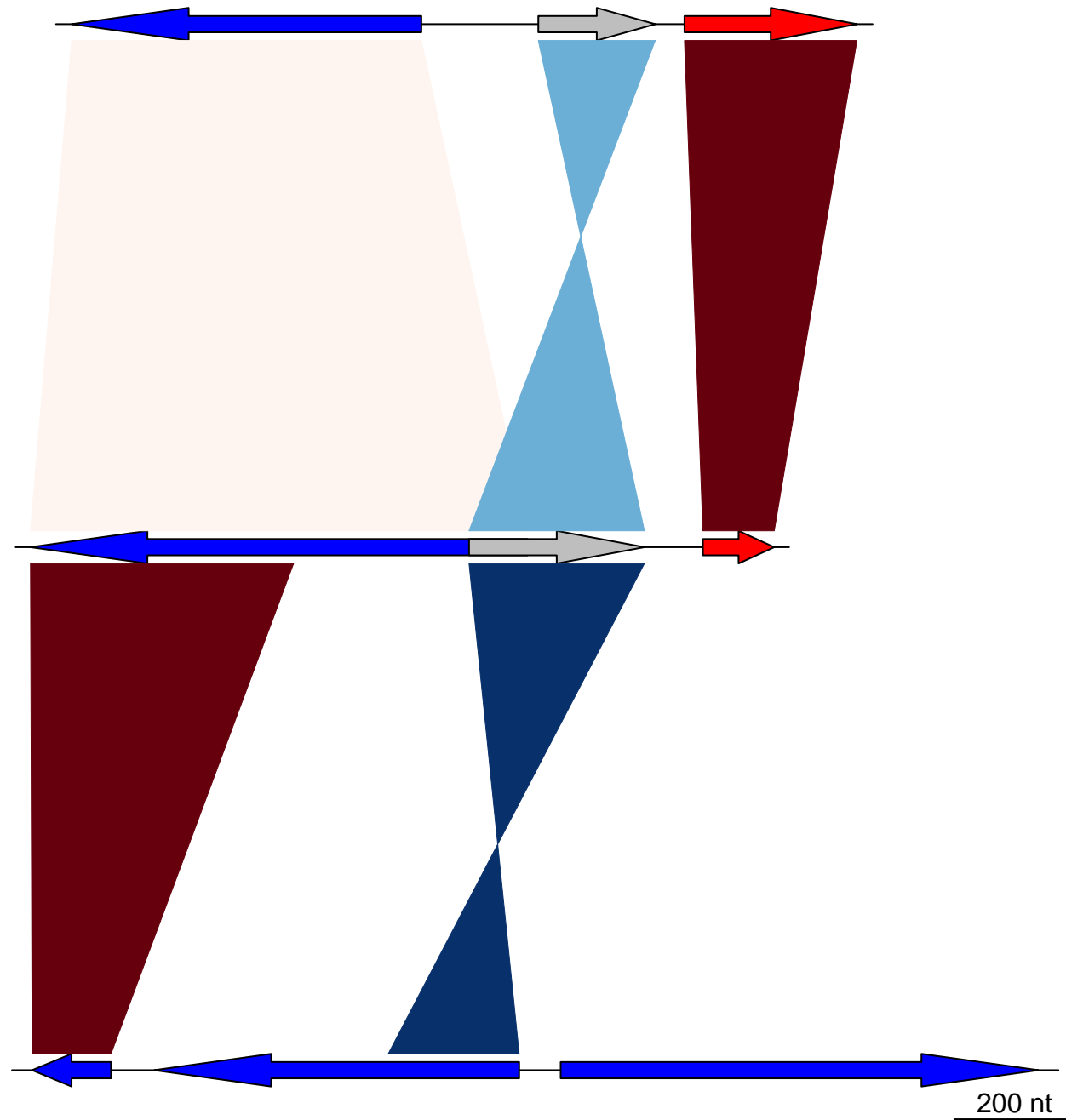


help("annotation")

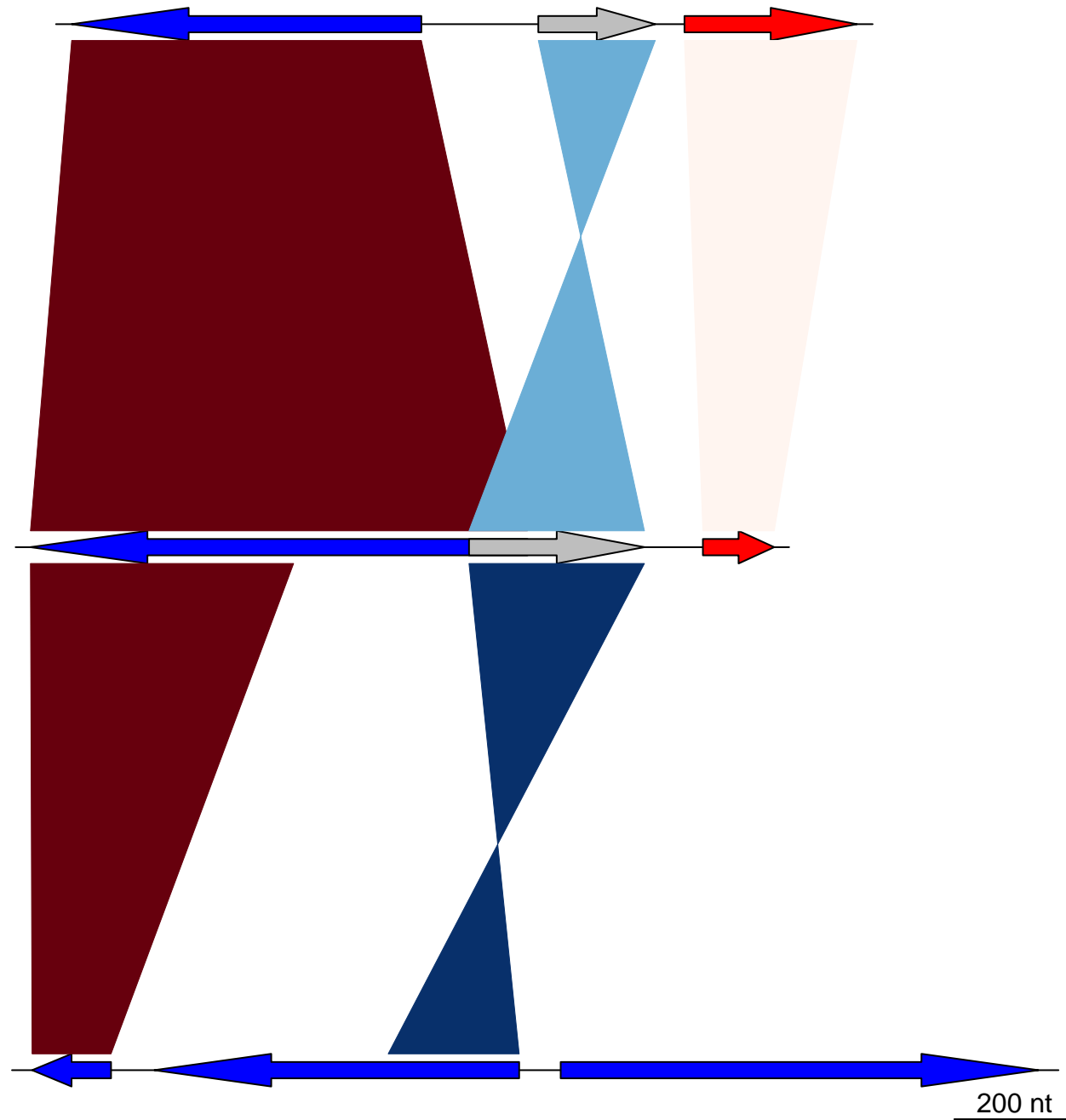




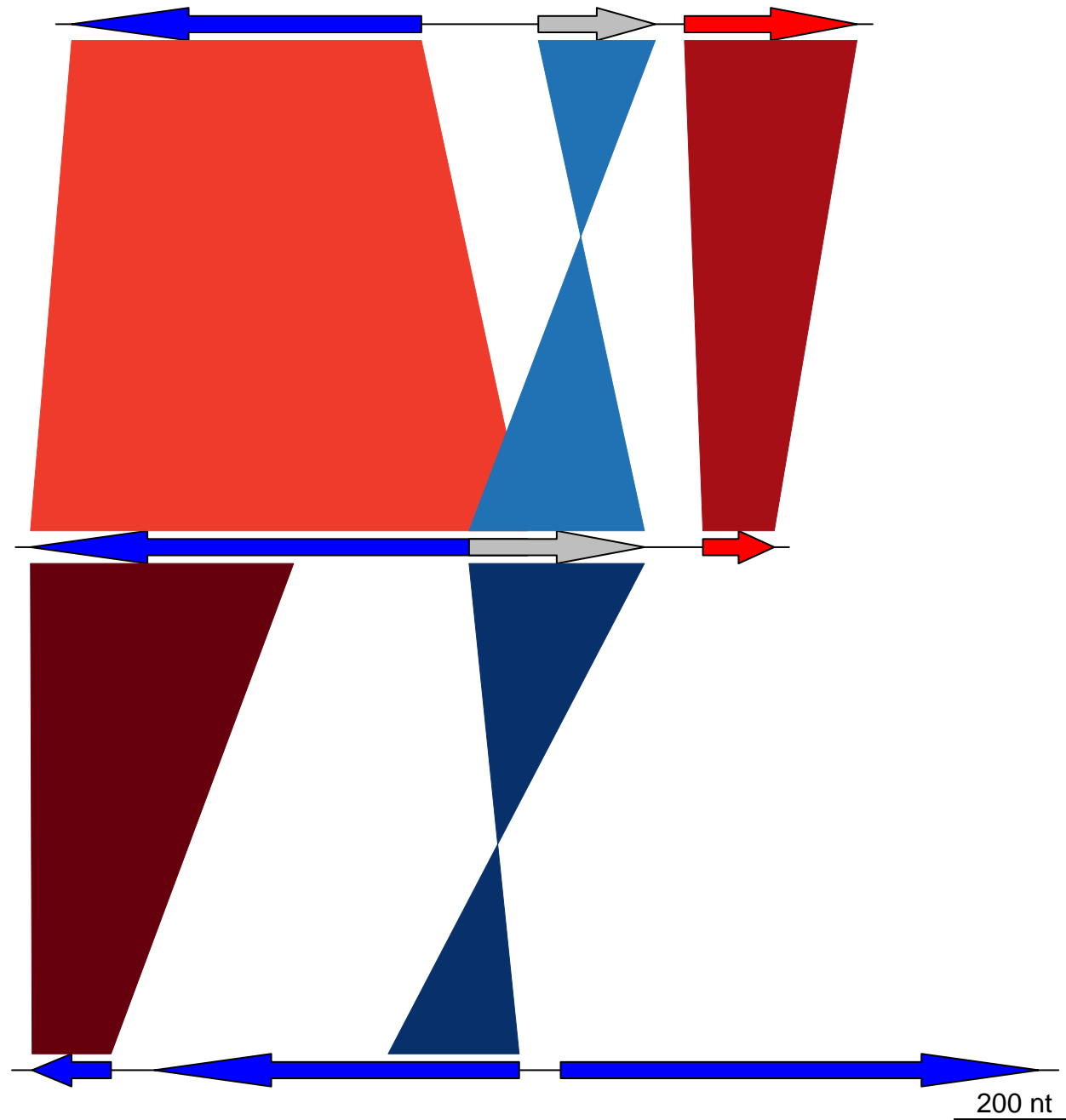
help("apply_color_scheme")



help("apply_color_scheme")



help("apply_color_scheme")



help("apply_color_scheme")

BB

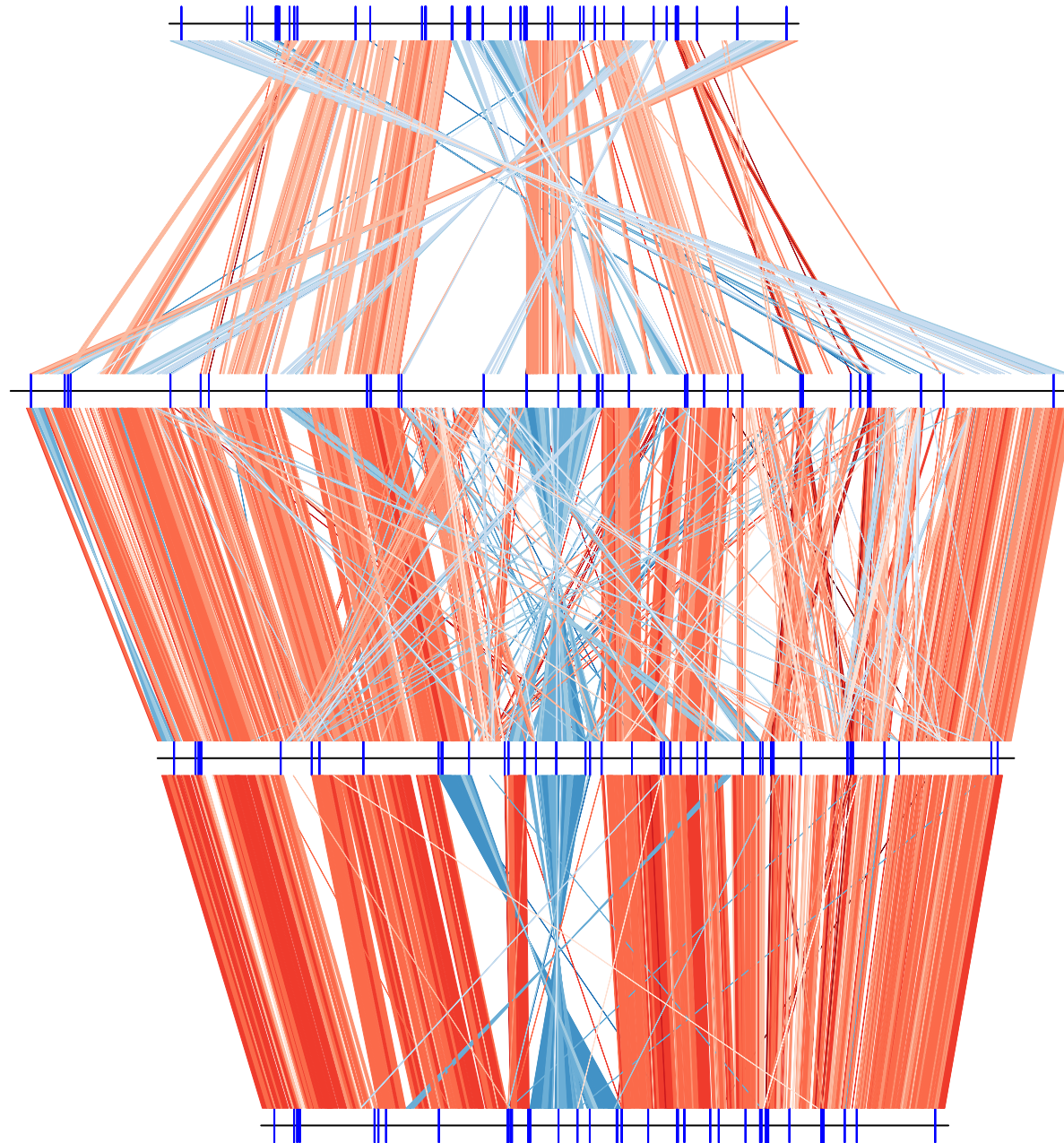
BG

BH

BQ

help("barto")

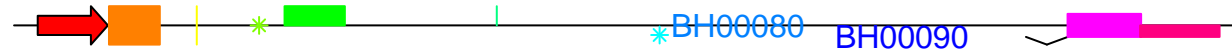
500 kb



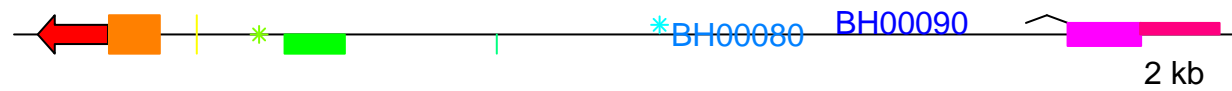


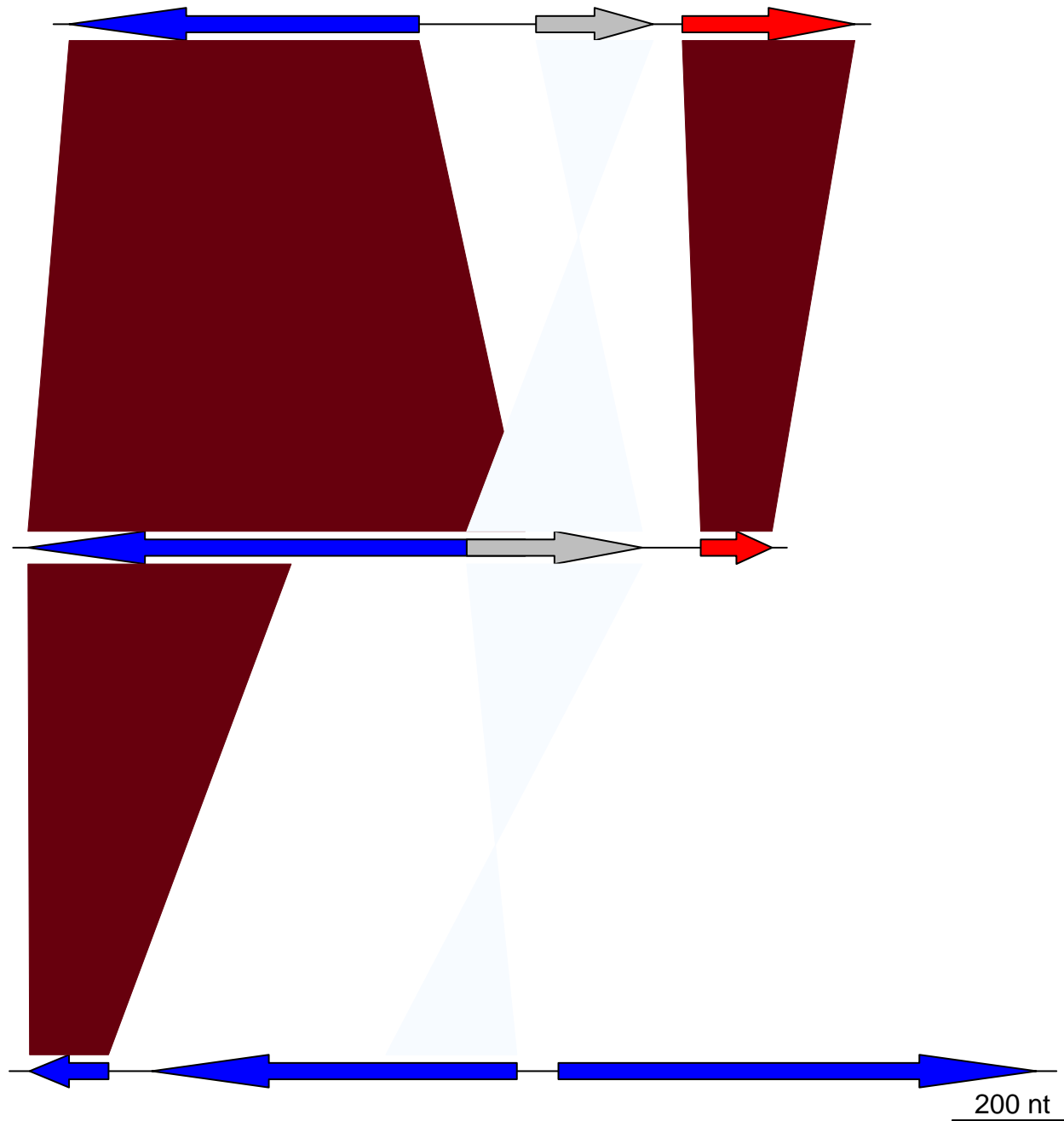
help("gene_types")

2 kb



help("gene_types")





help("genoPlotR-package")

B_bacilliformis

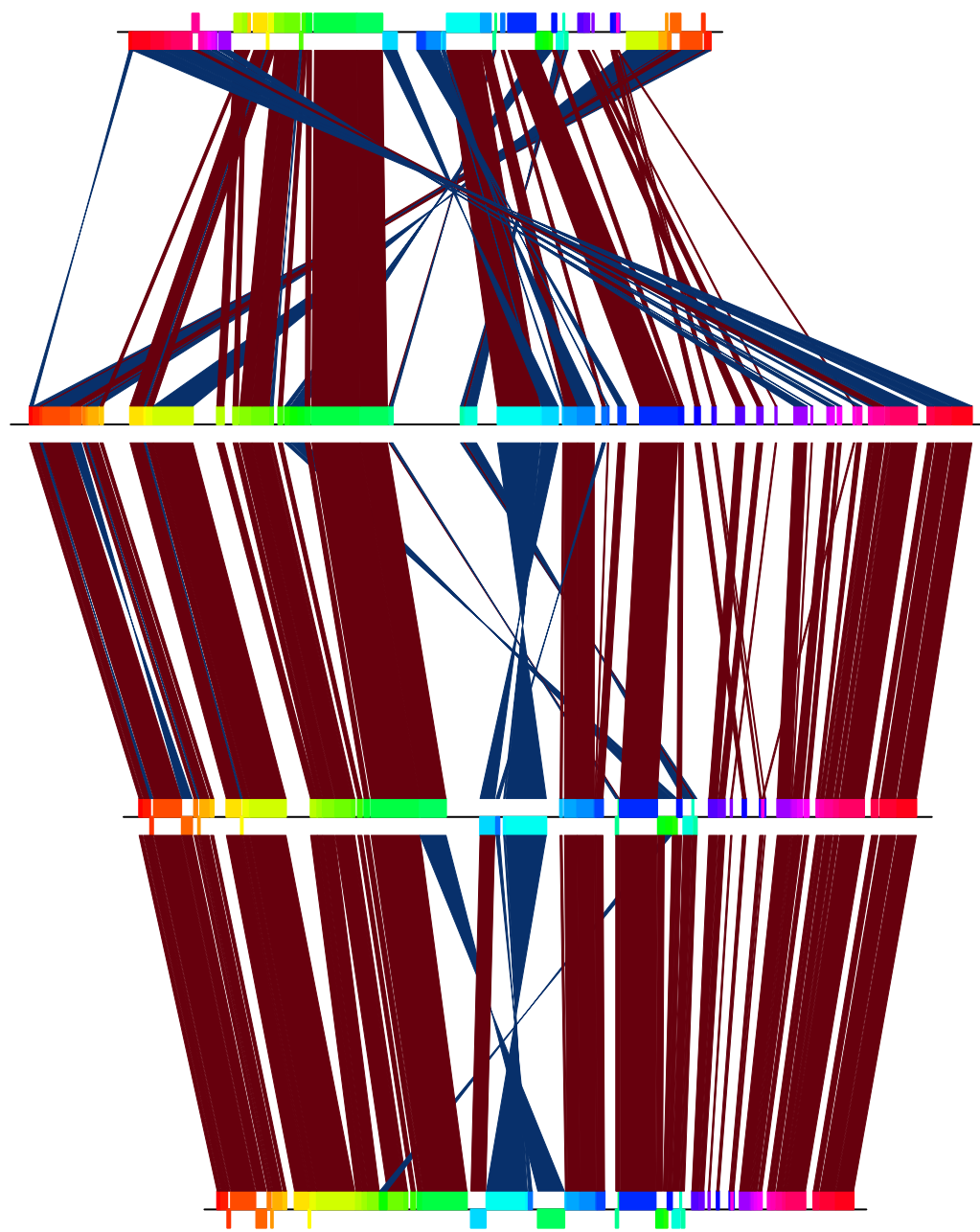
B_grahamii

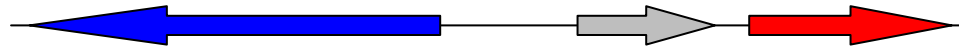
B_henselae

B_quintana

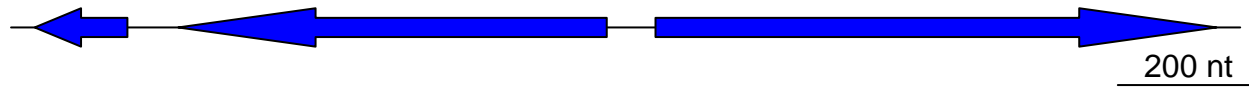
help("mauve_bbone")

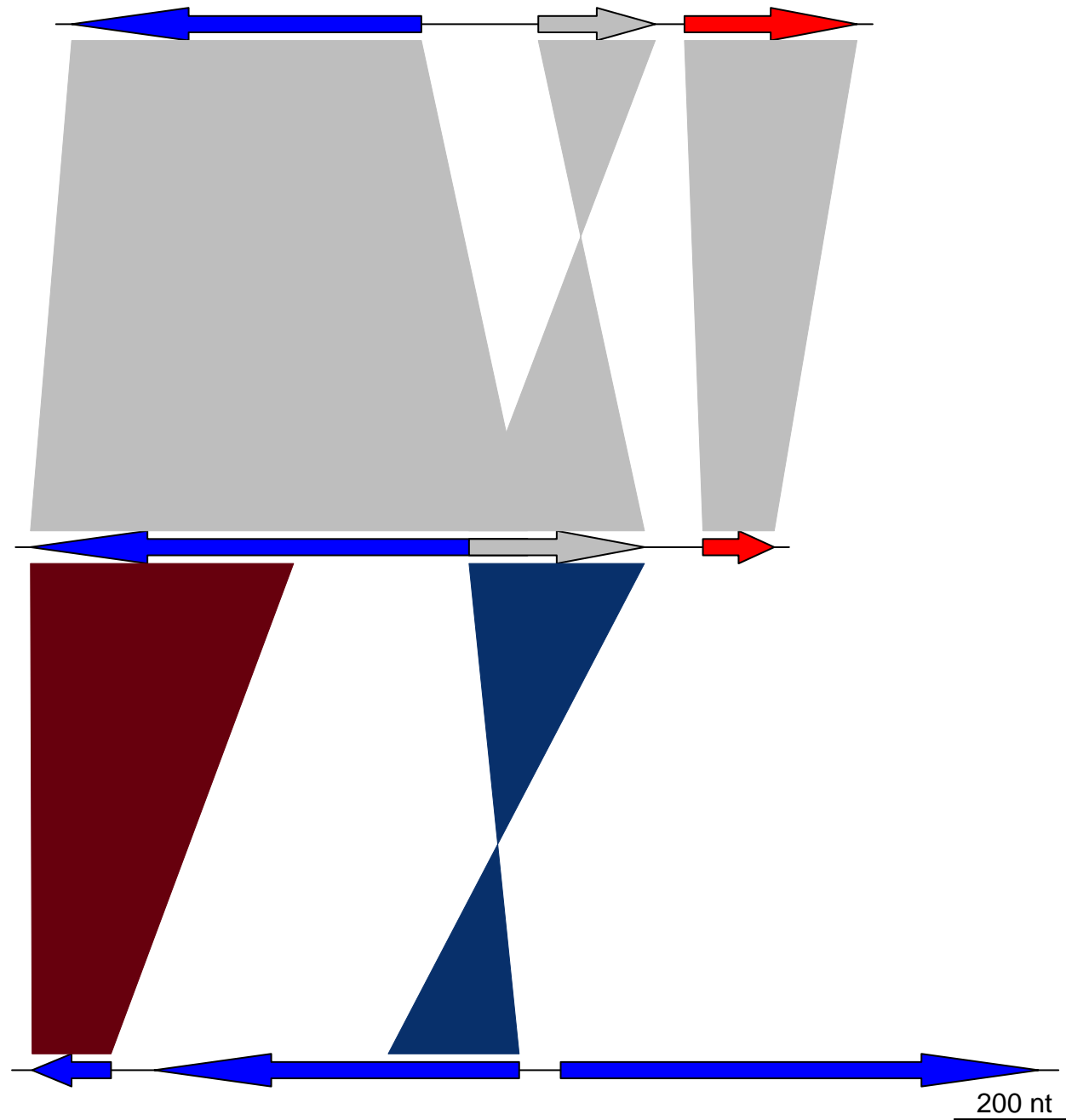
500 kb



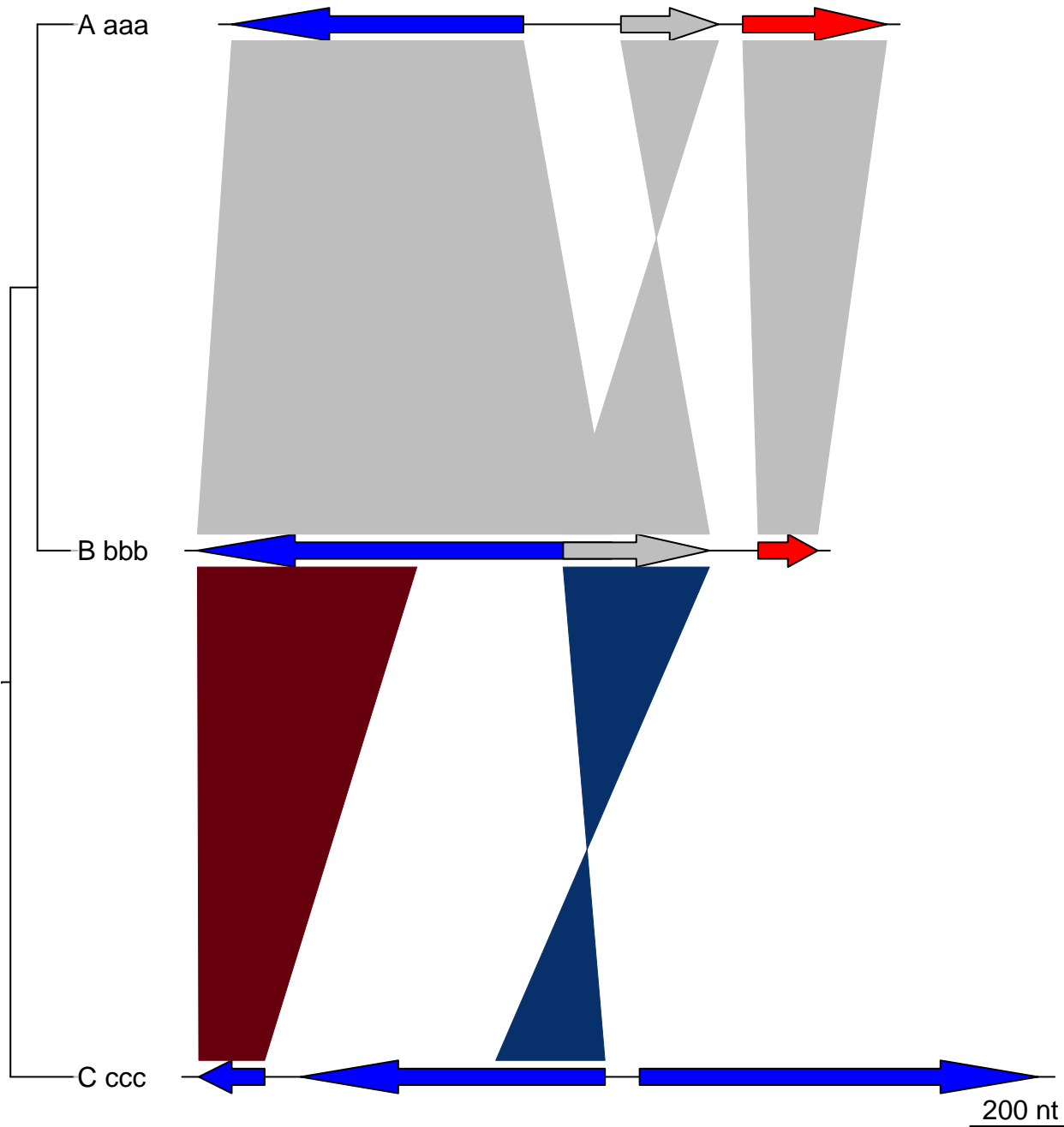


help("plot_gene_map")

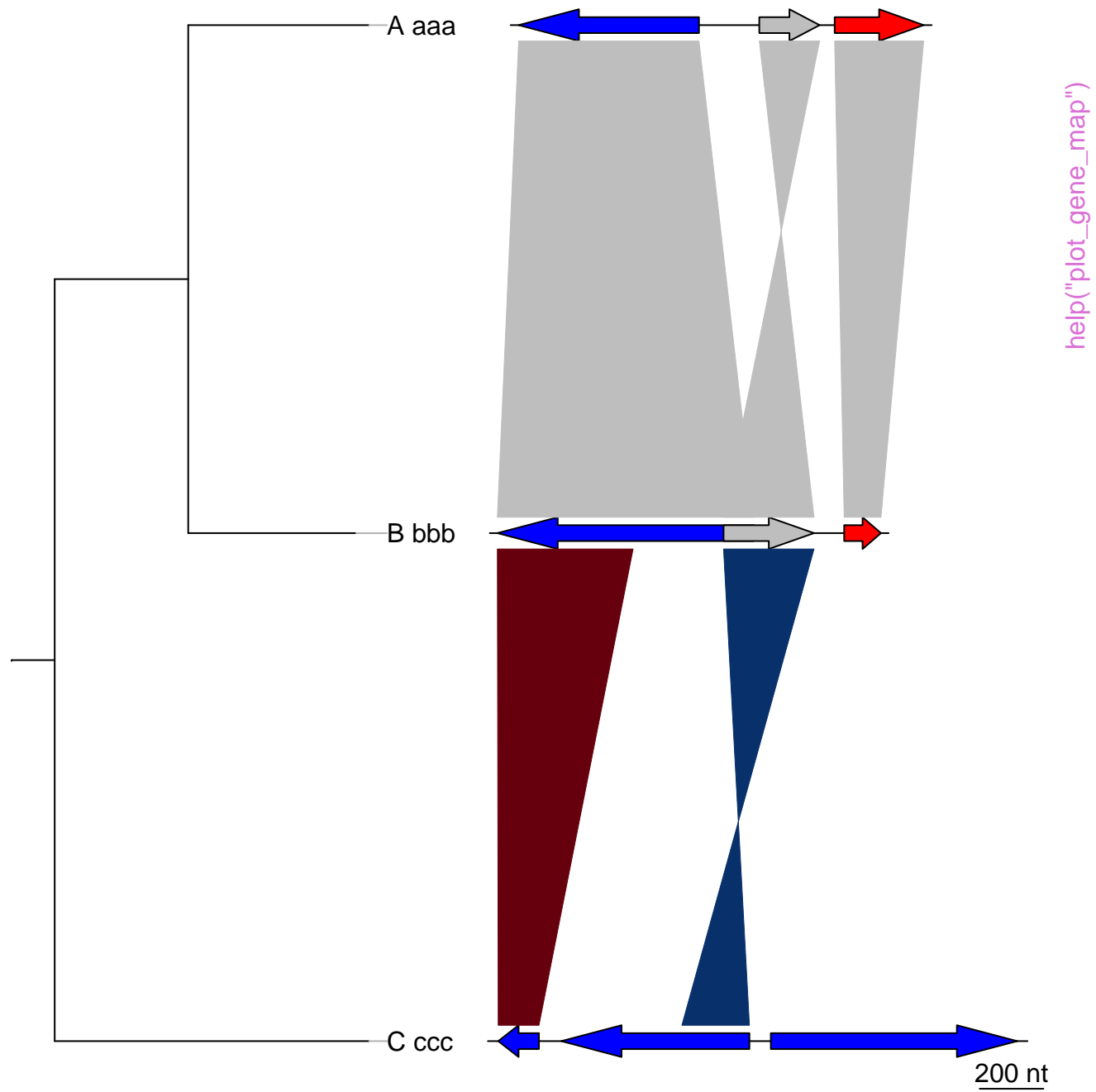


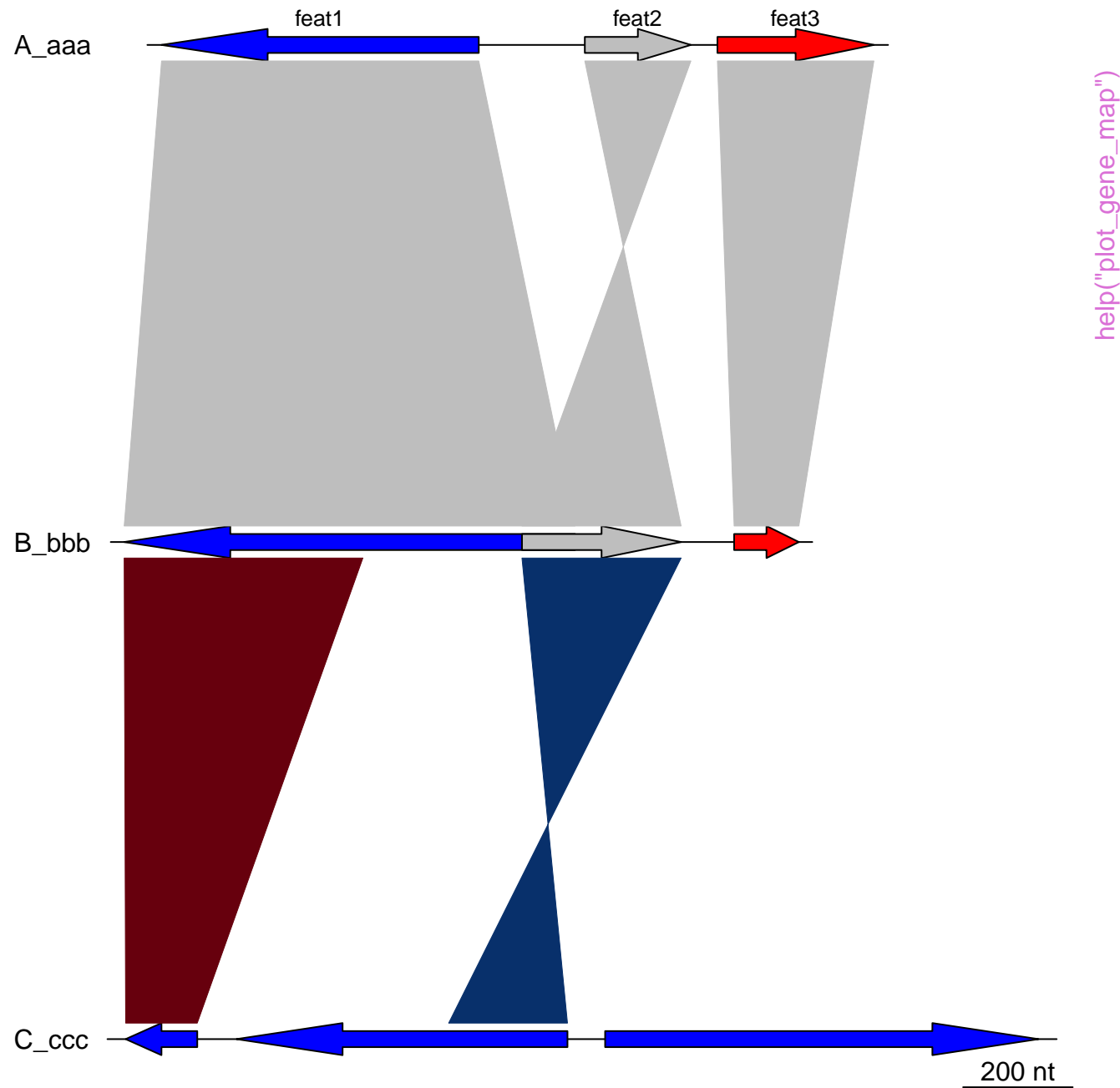


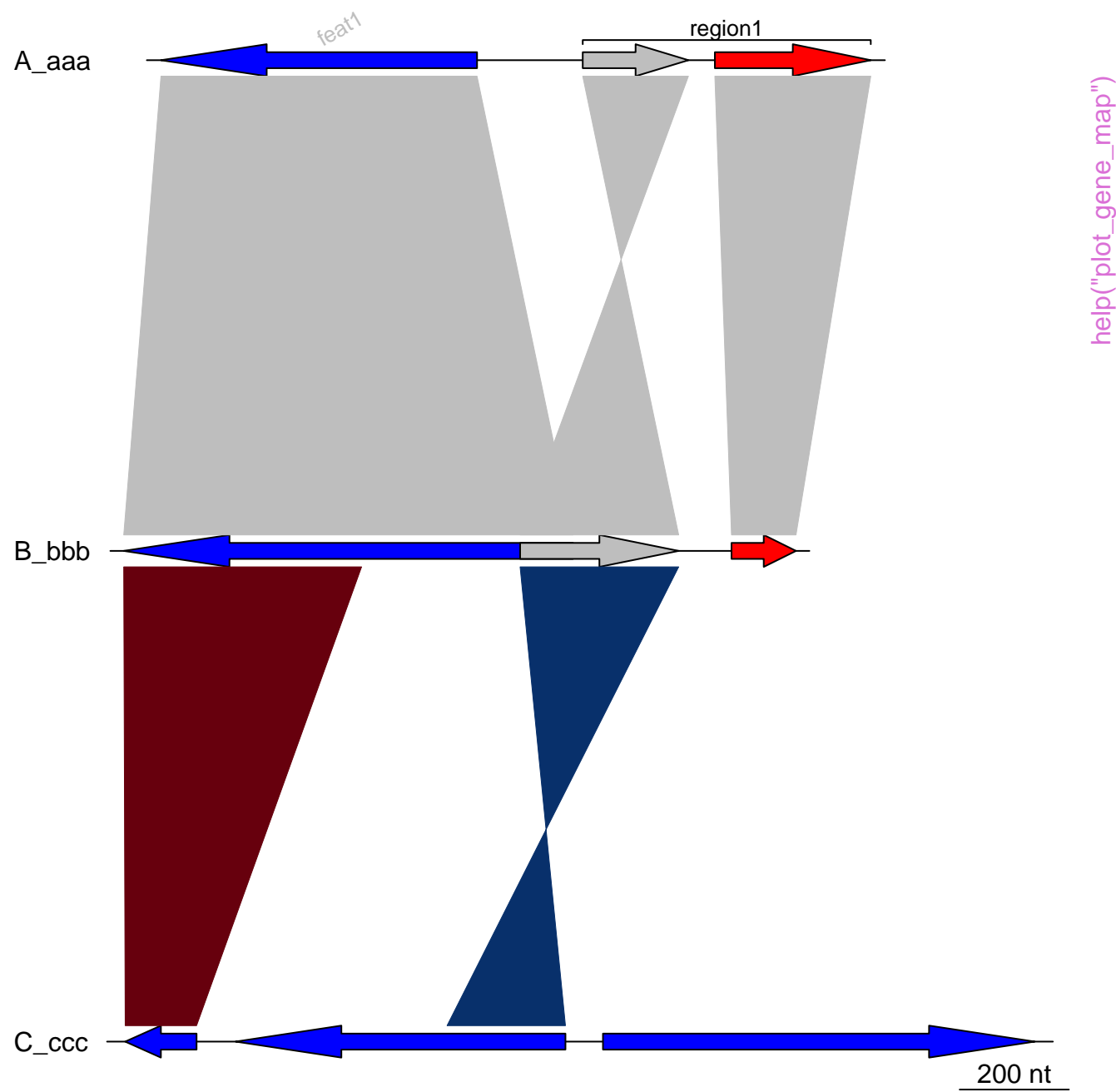
help("plot_gene_map")

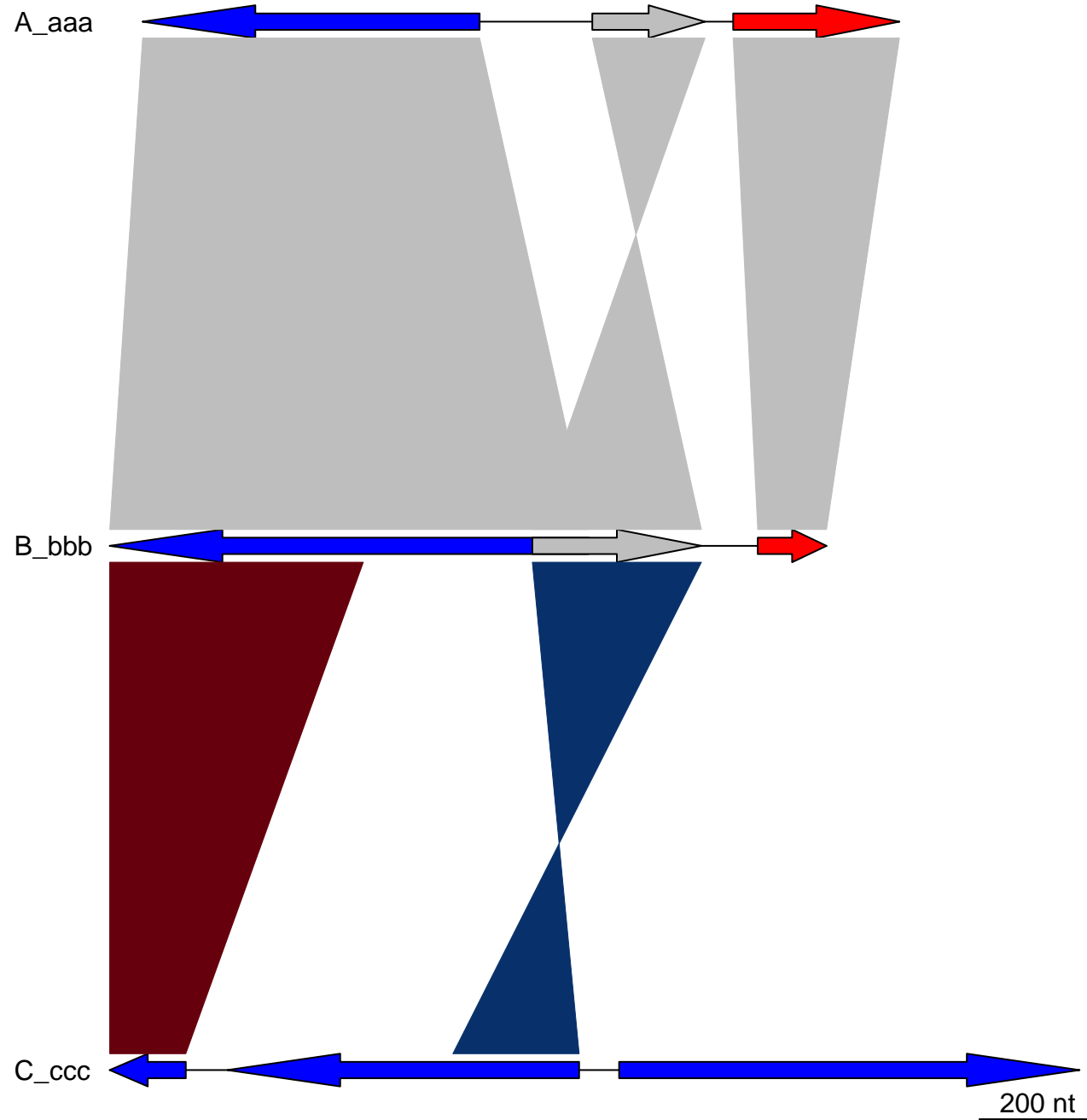


help("plot_gene_map")

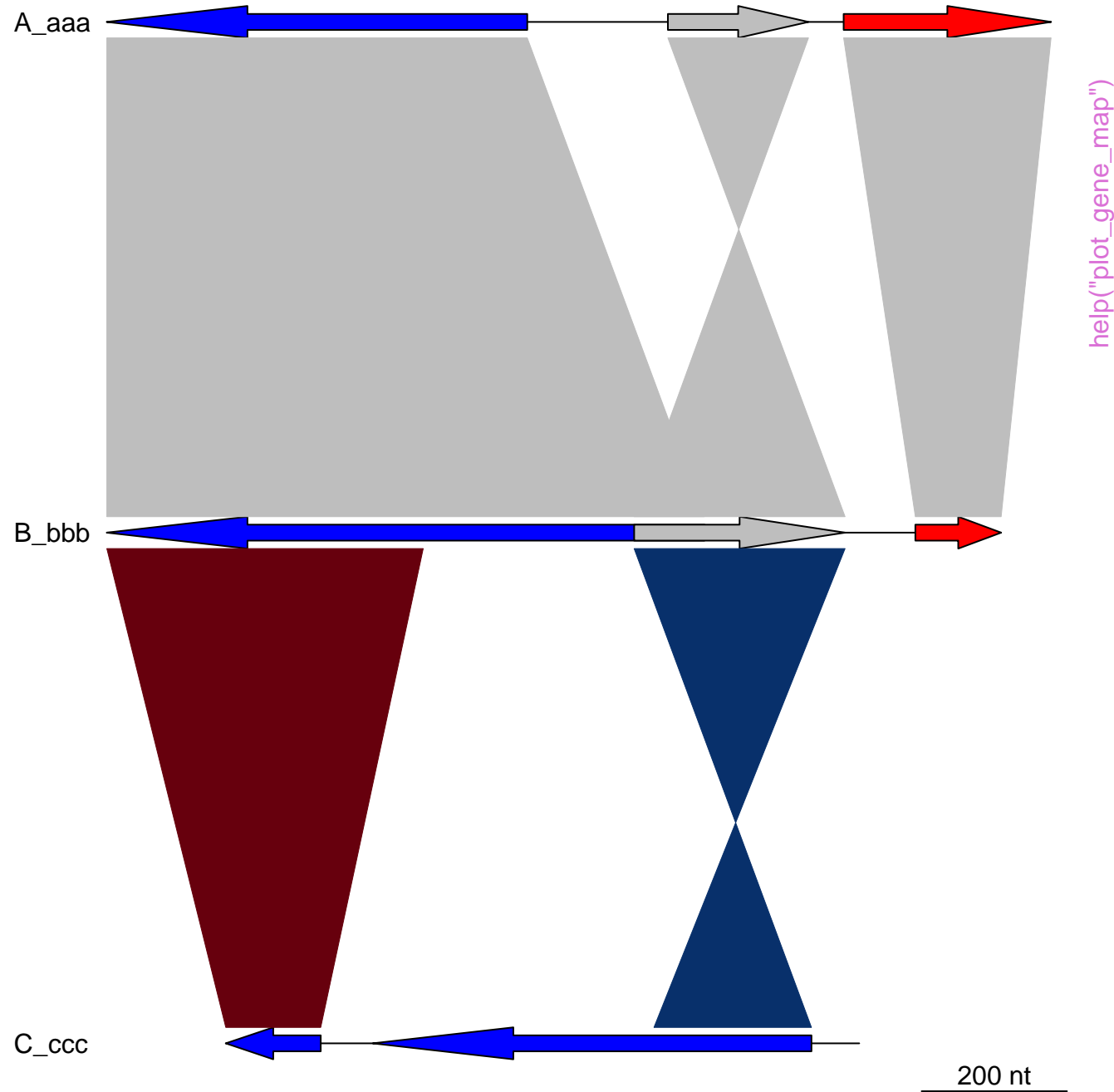








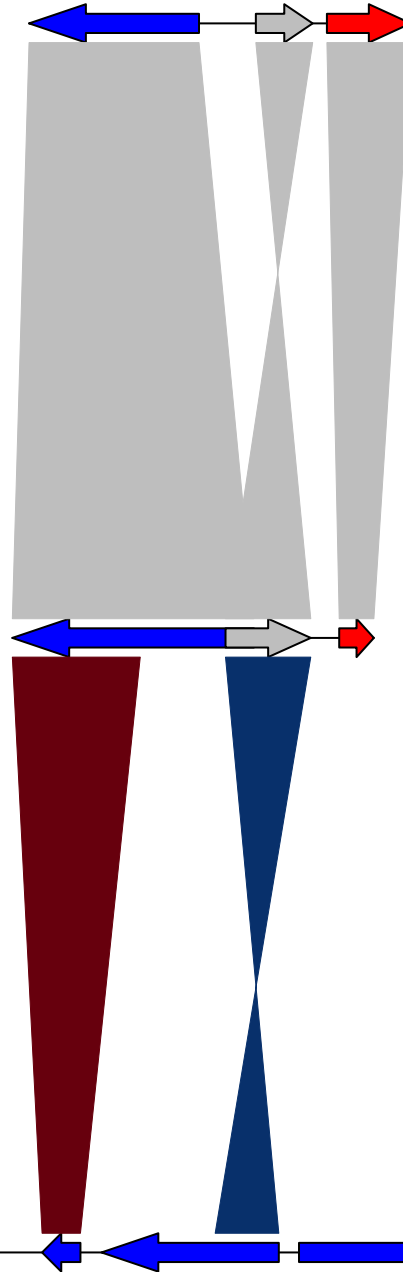
help("plot_gene_map")



A_aaa

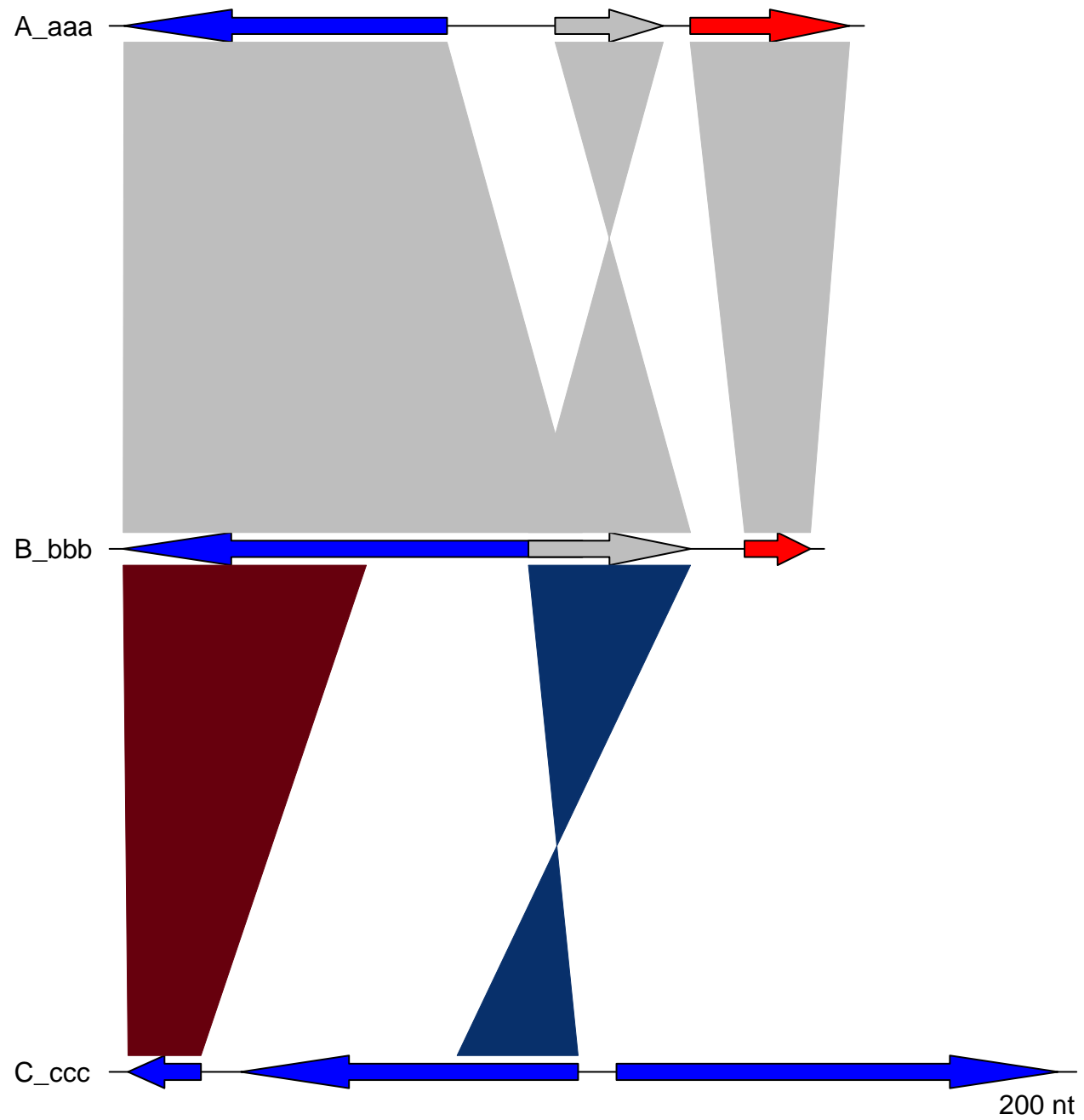
B_bbb

C_ccc



500 nt

help("plot_gene_map")



help("plot_gene_map")

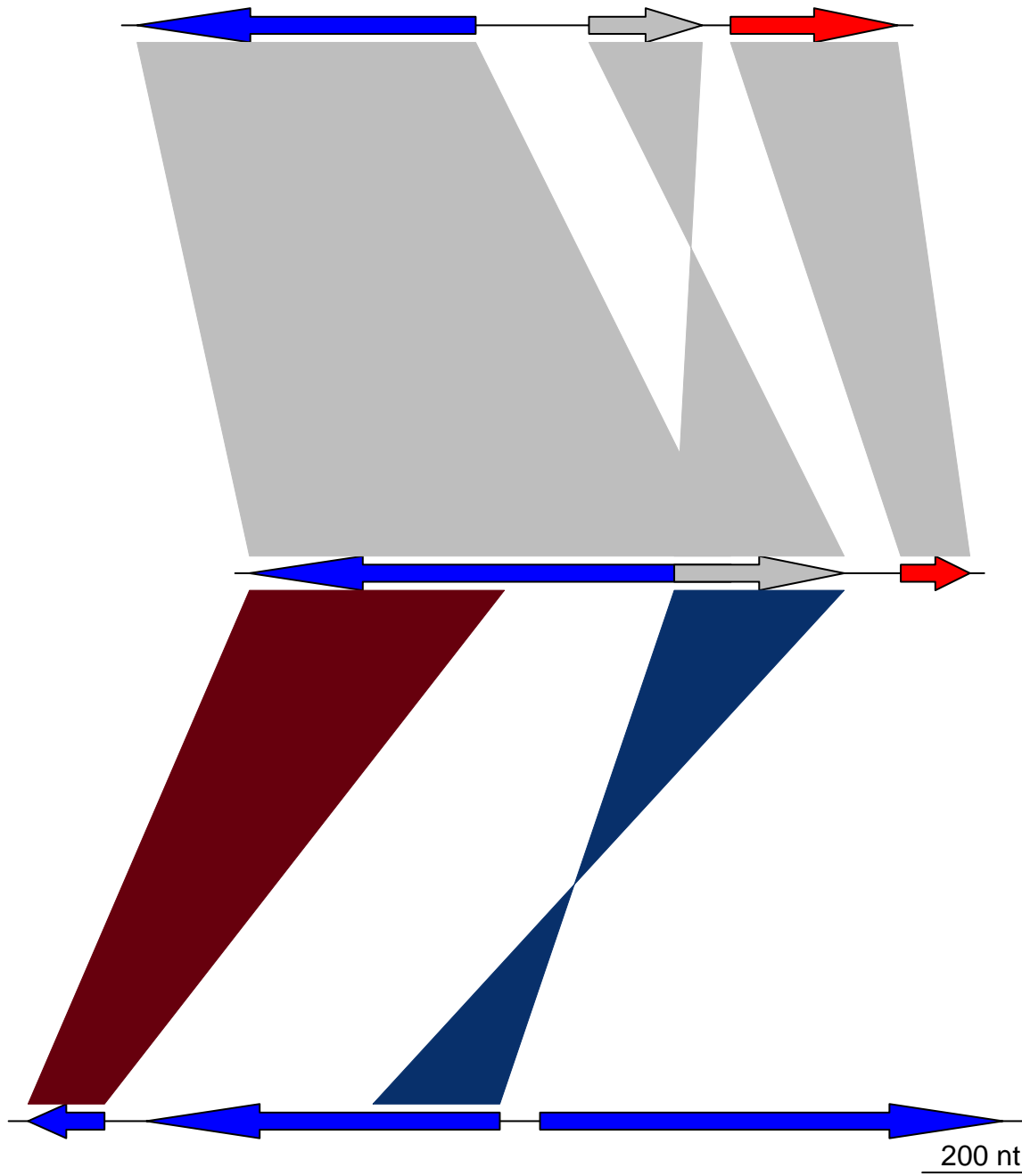
A_aaa

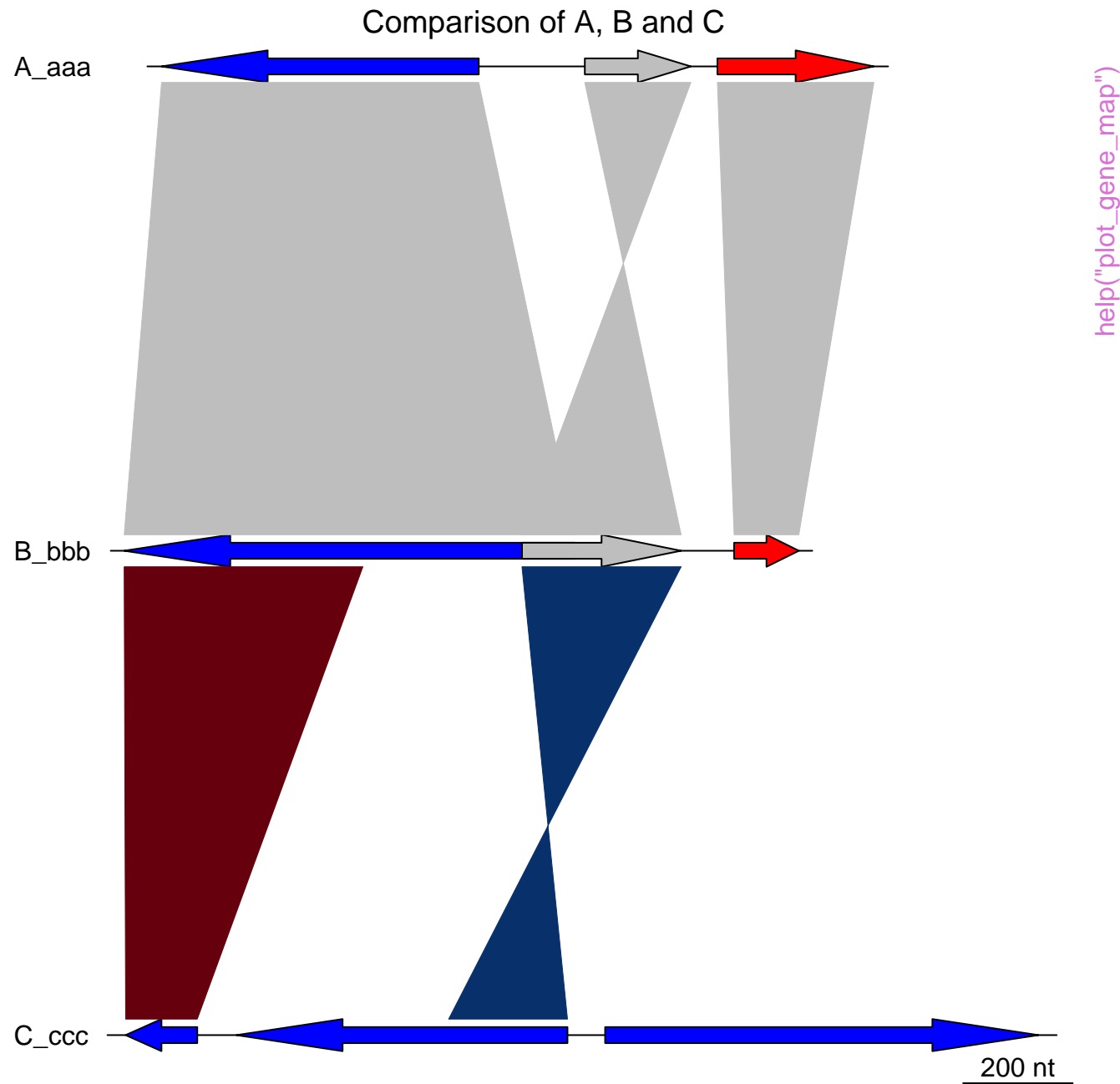
B_bbb

C_ccc

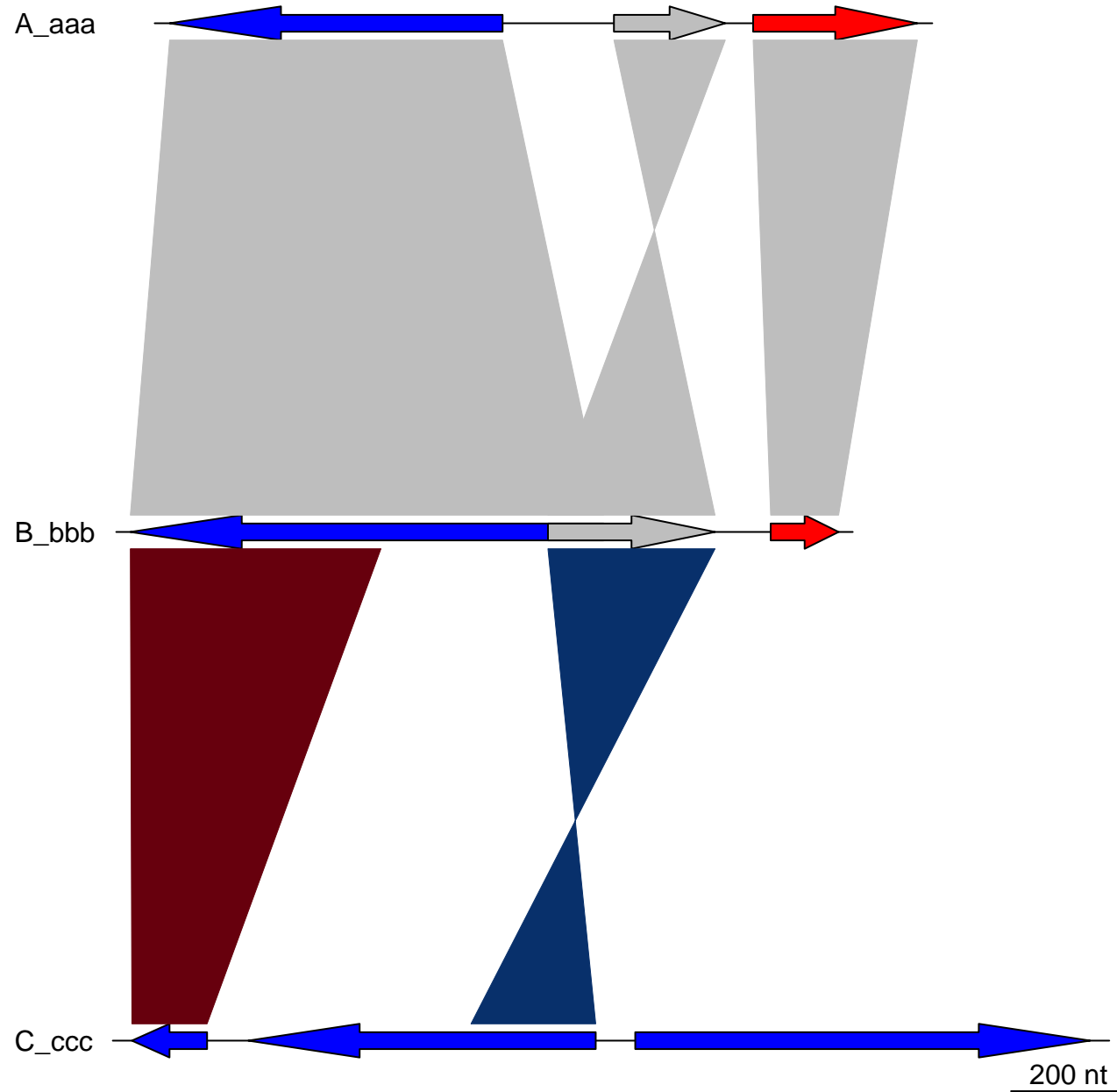
help("plot_gene_map")

200 nt

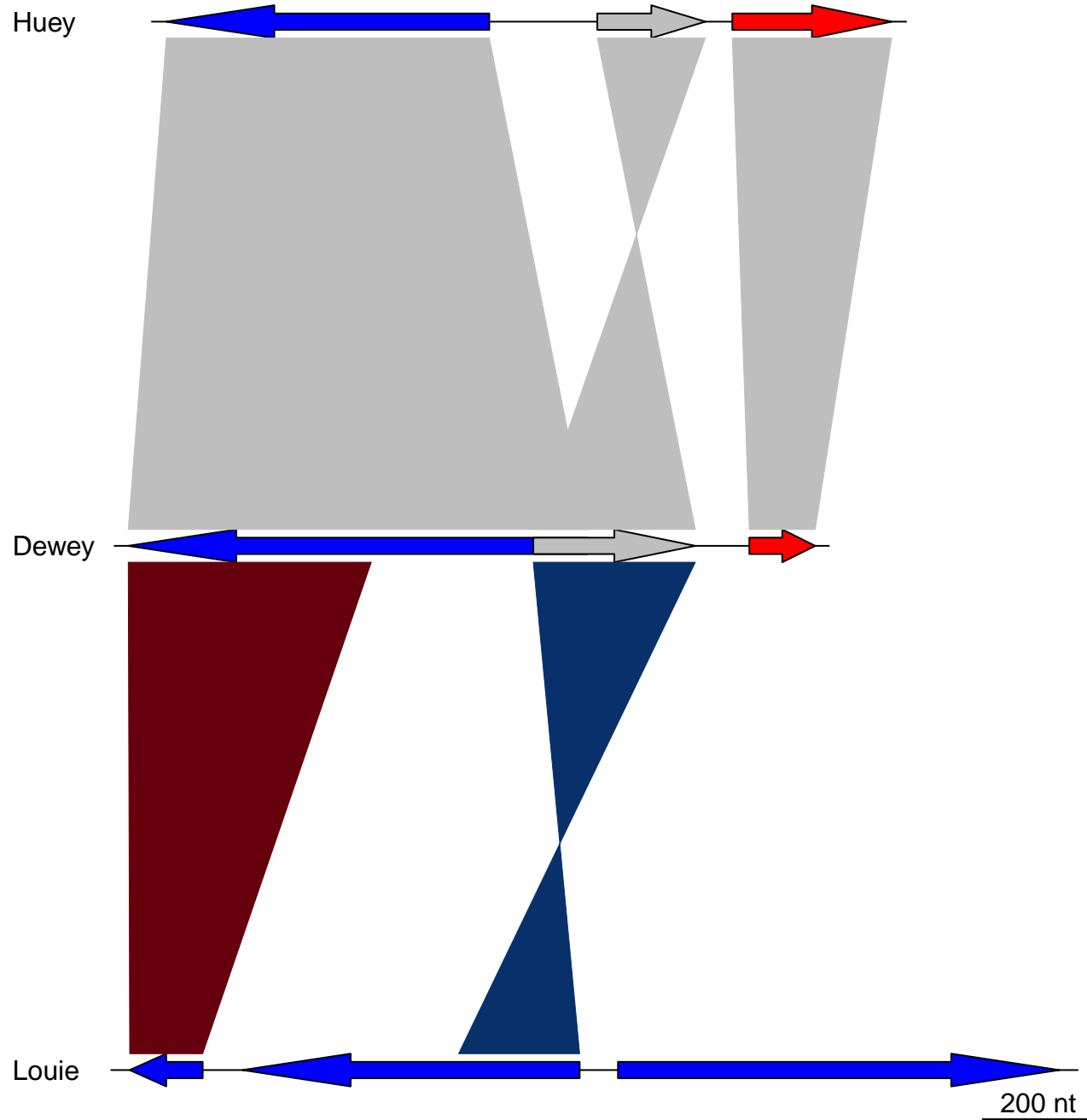




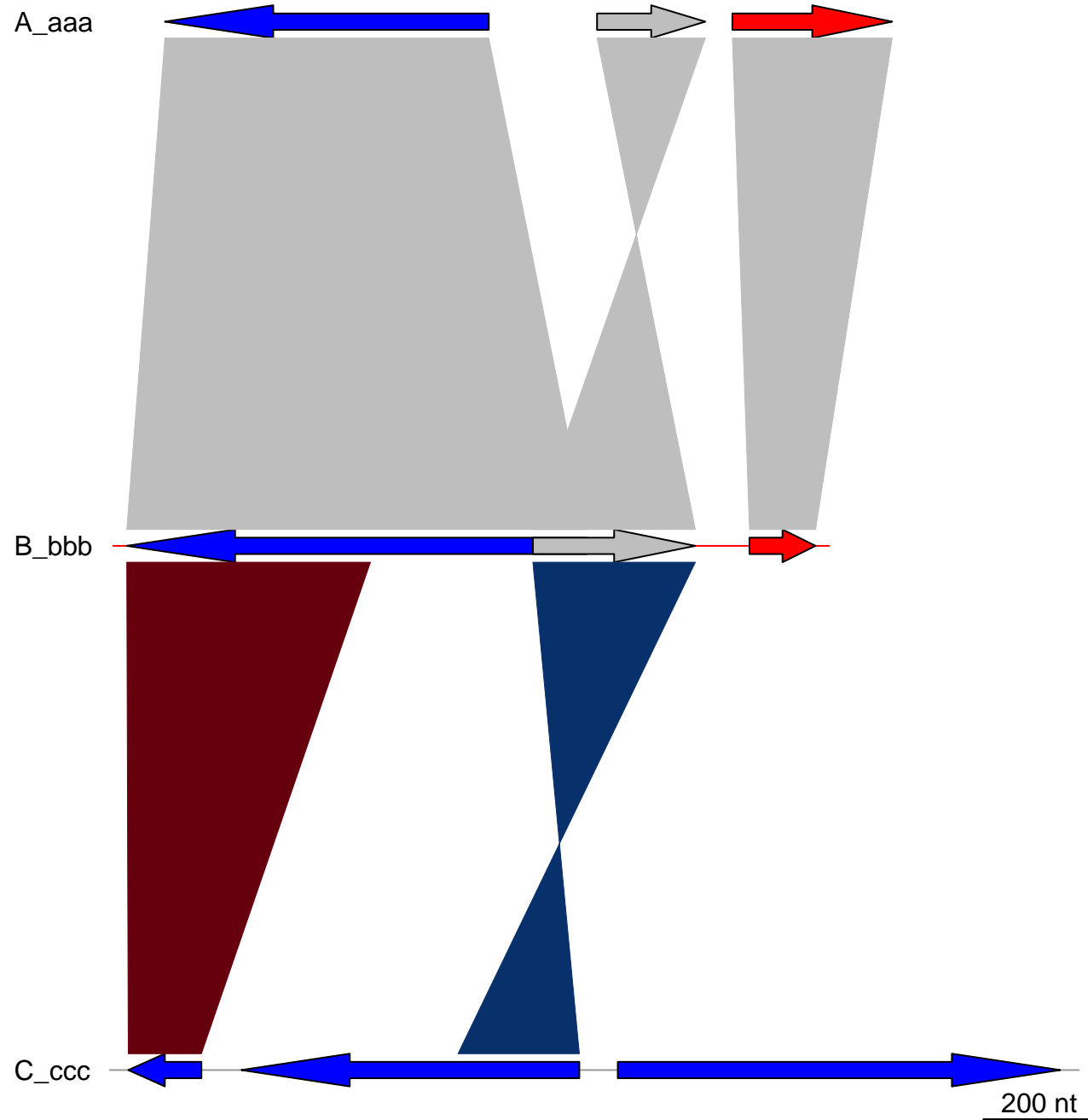
Comparison of A, B and C



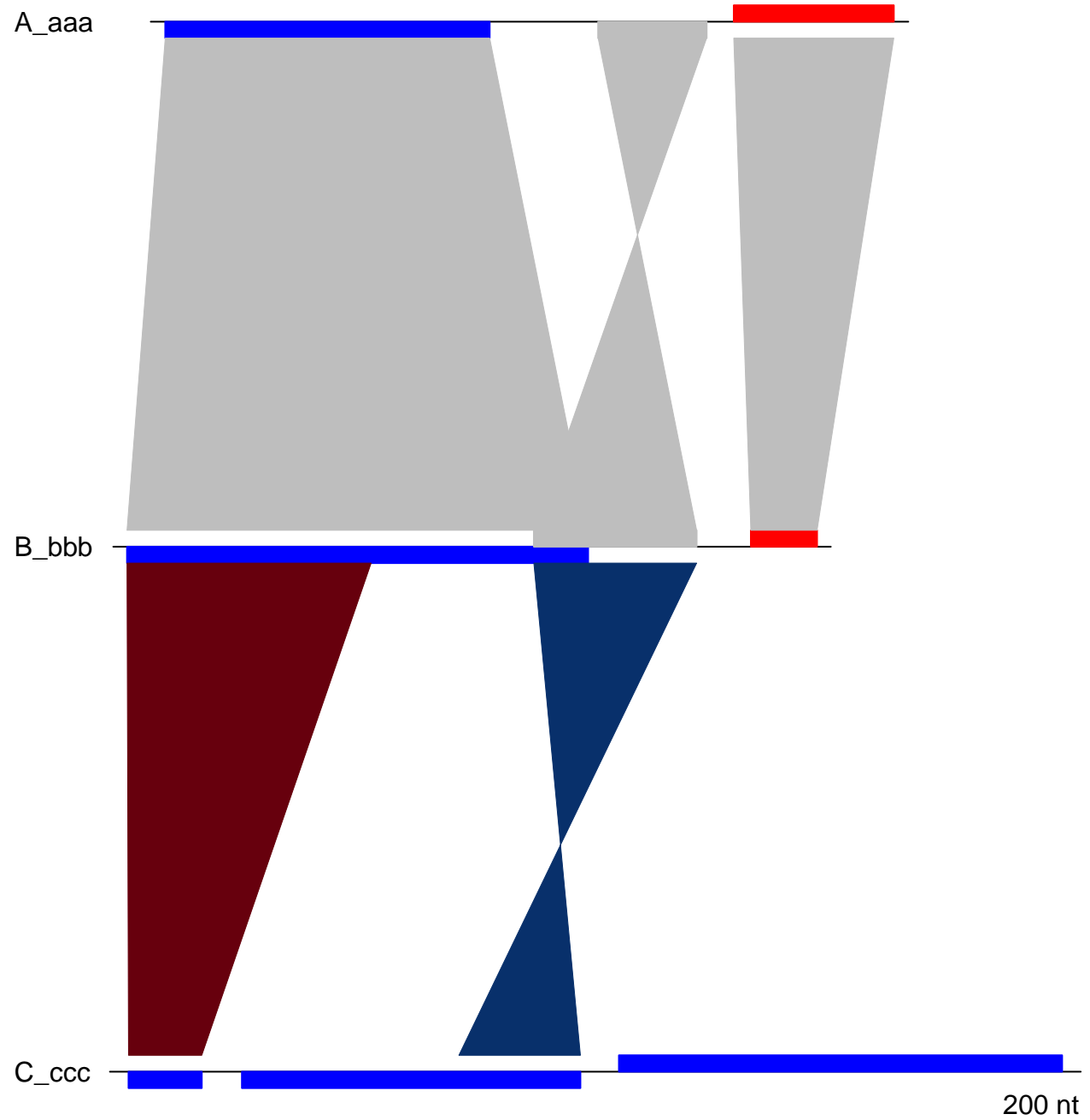
help("plot_gene_map")



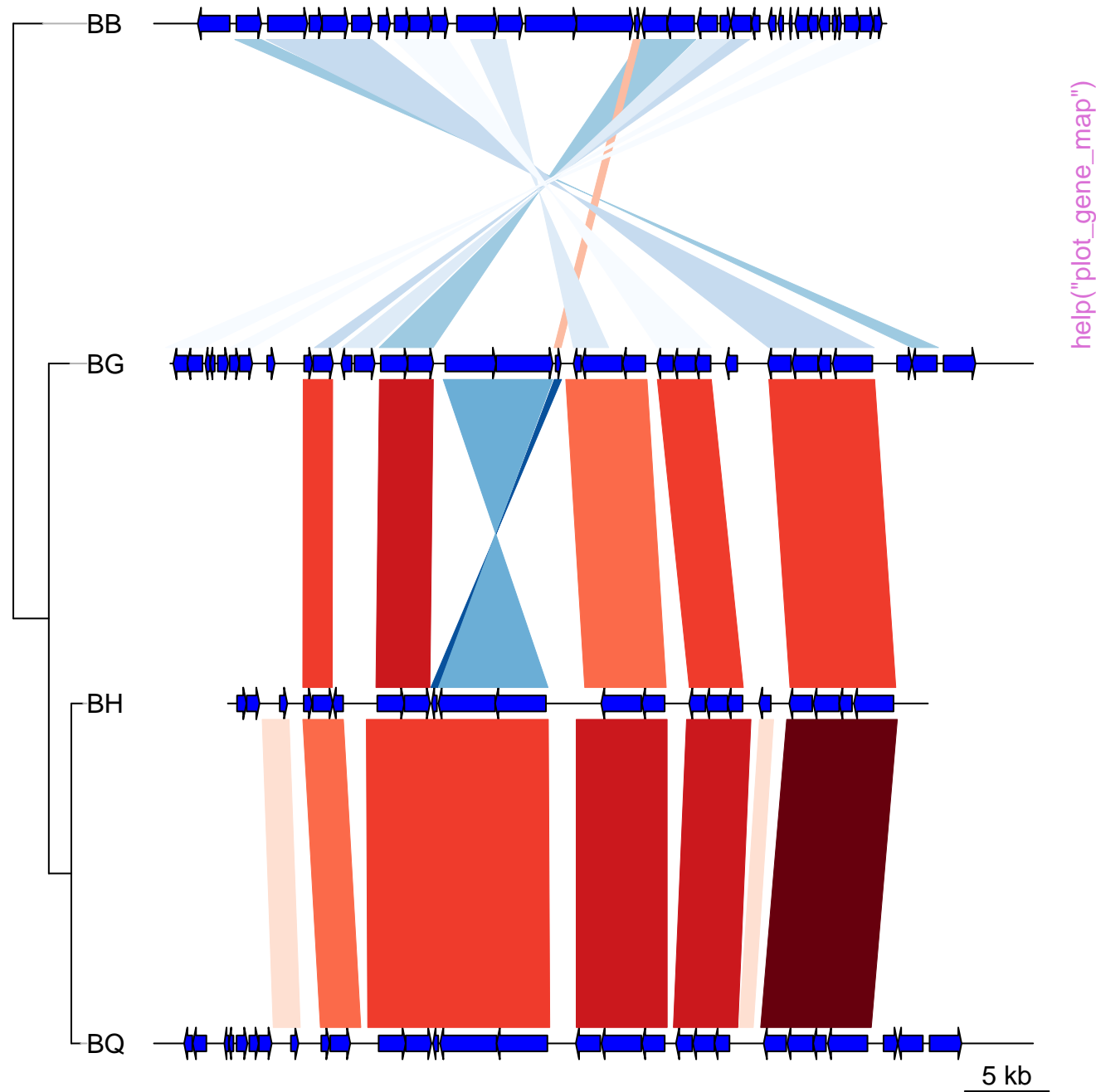
help("plot_gene_map")

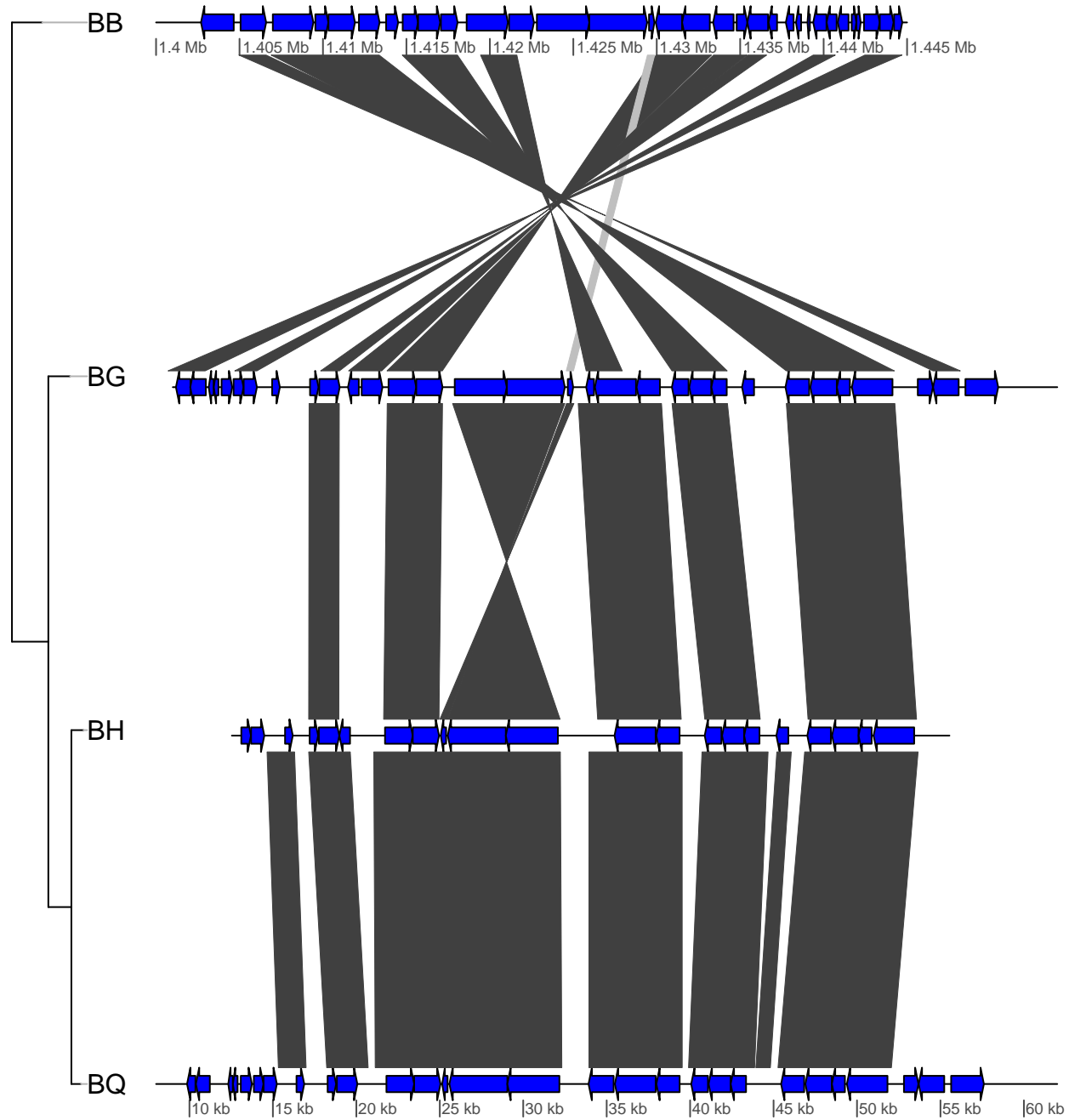


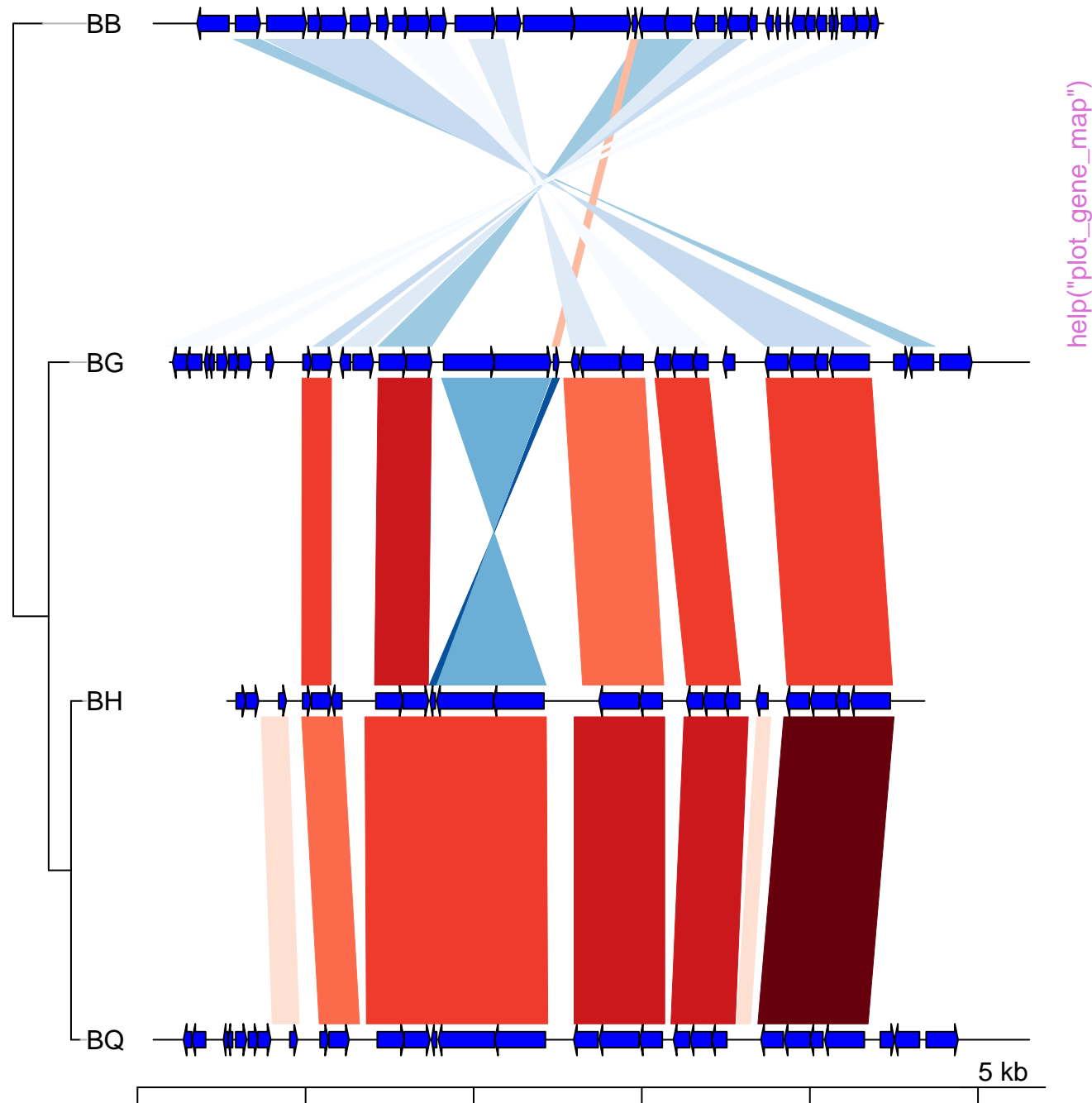
help("plot_gene_map")

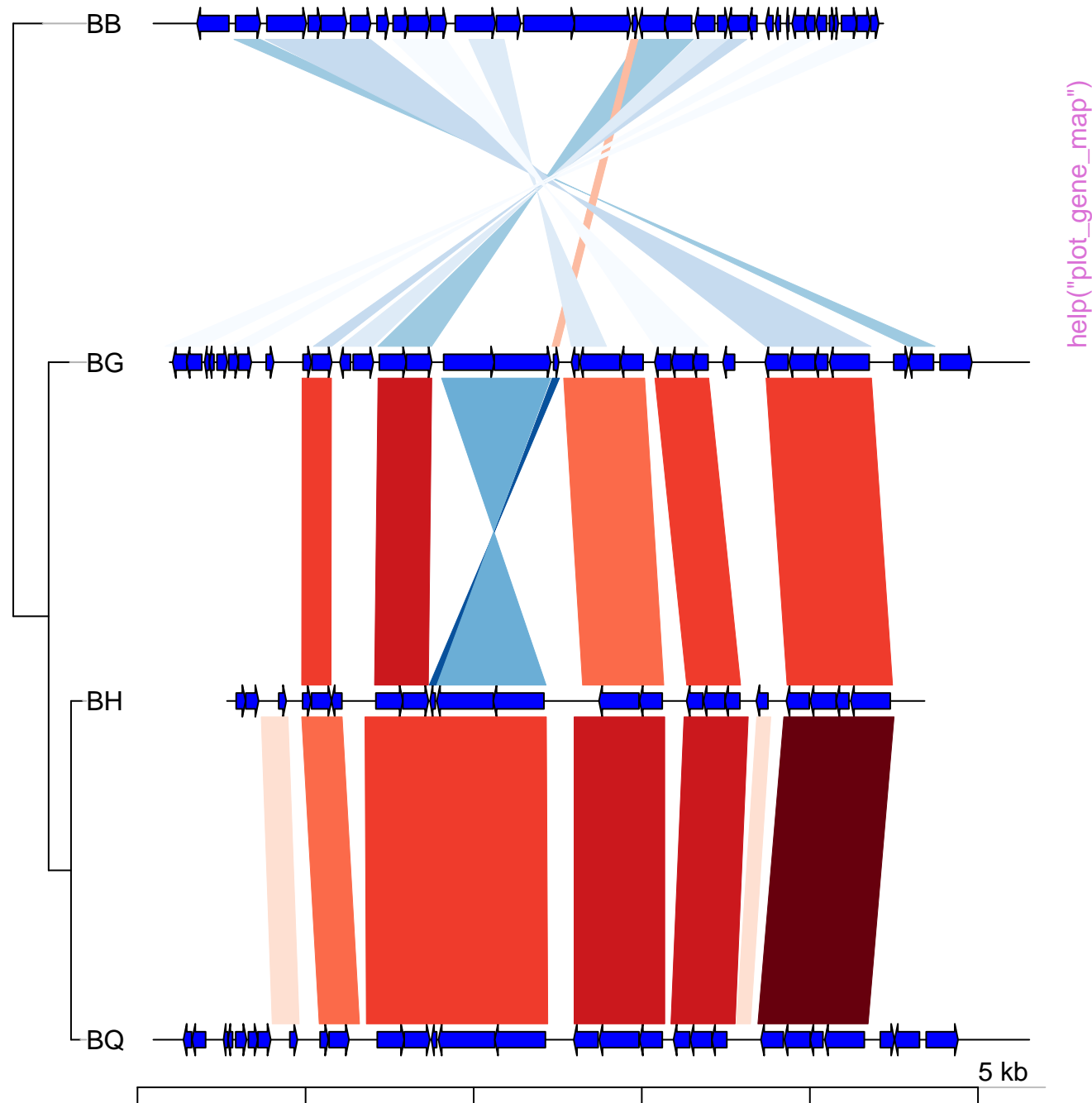


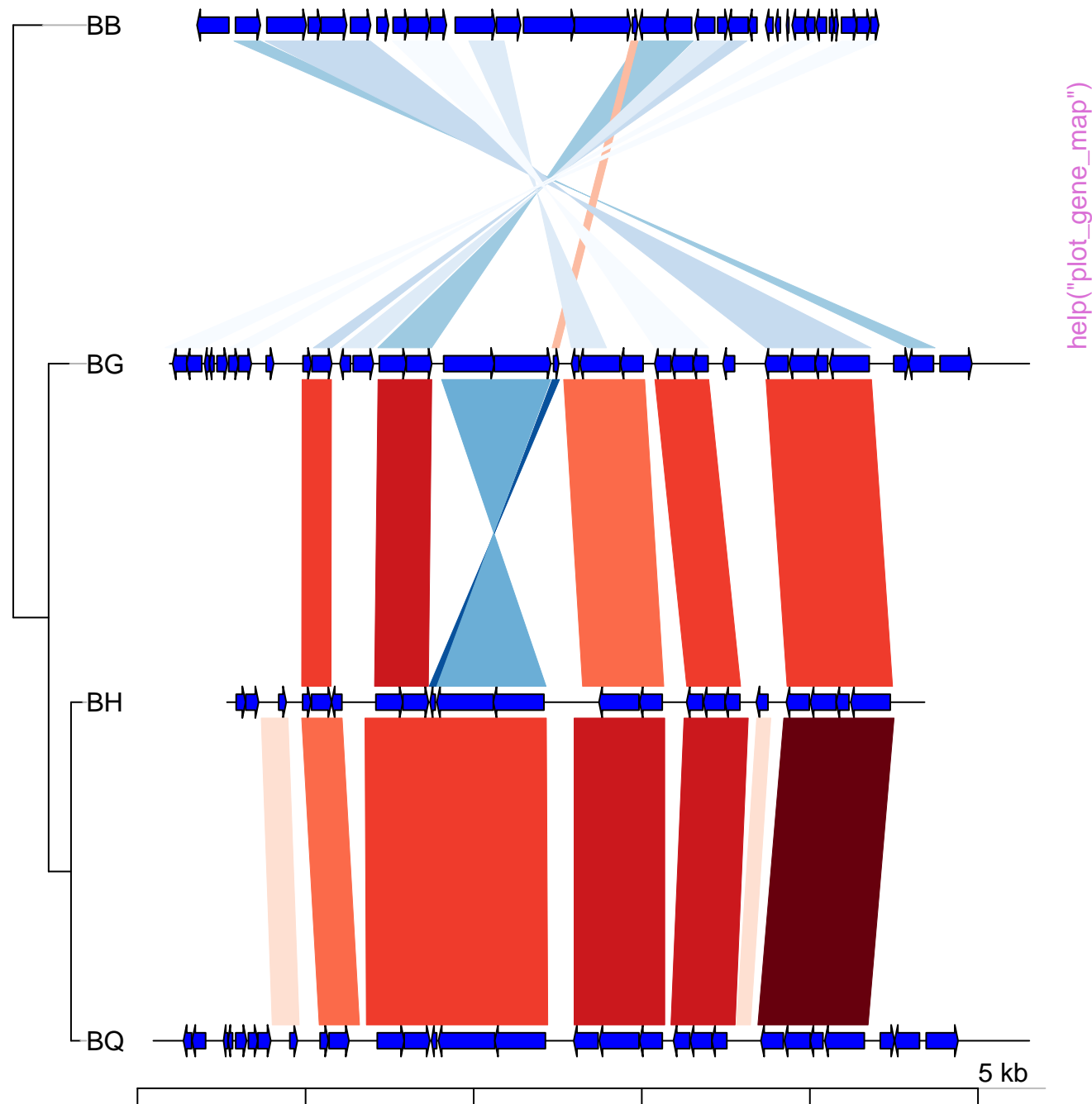
help("plot_gene_map")



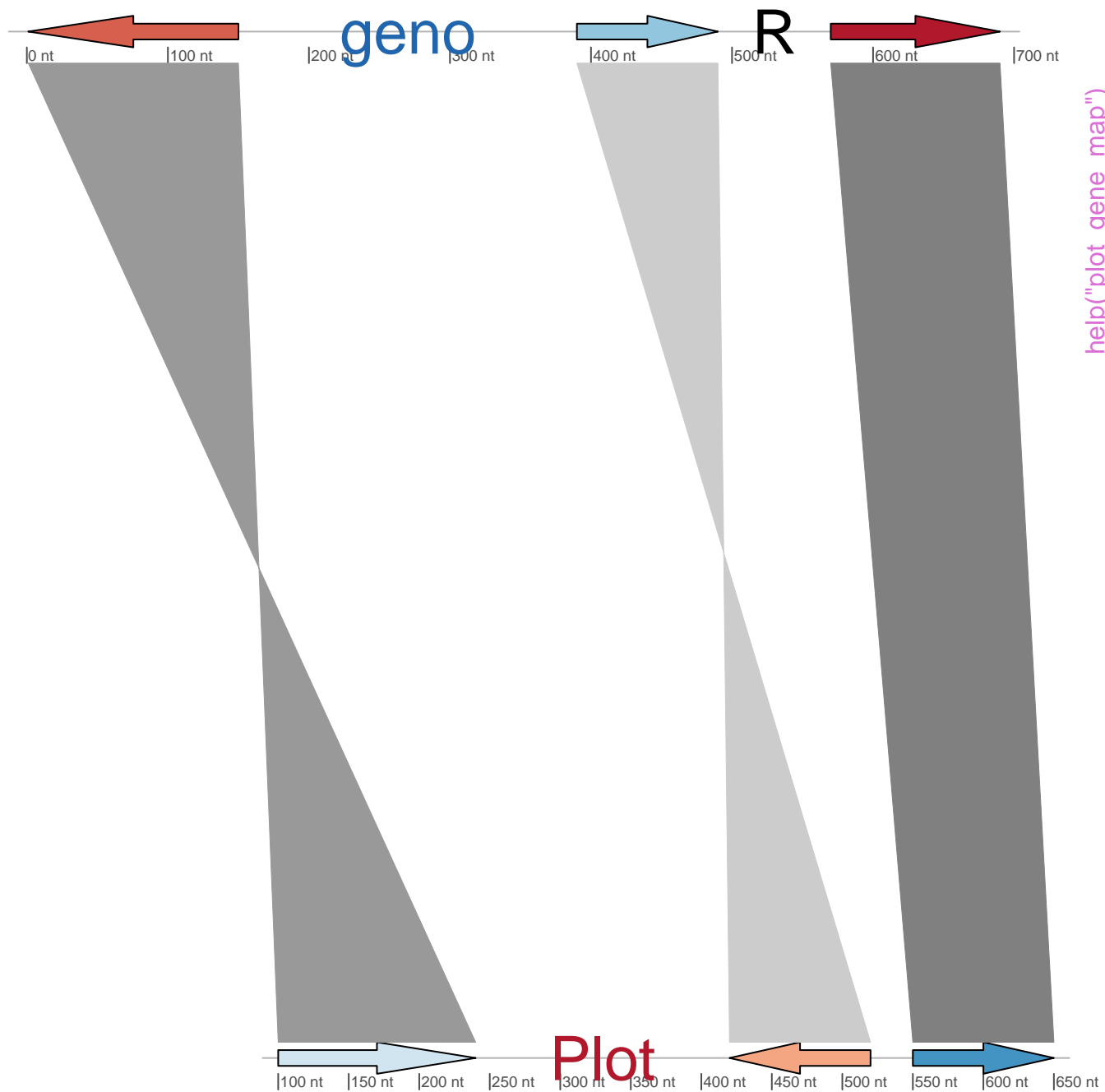








help("plot_gene_map")

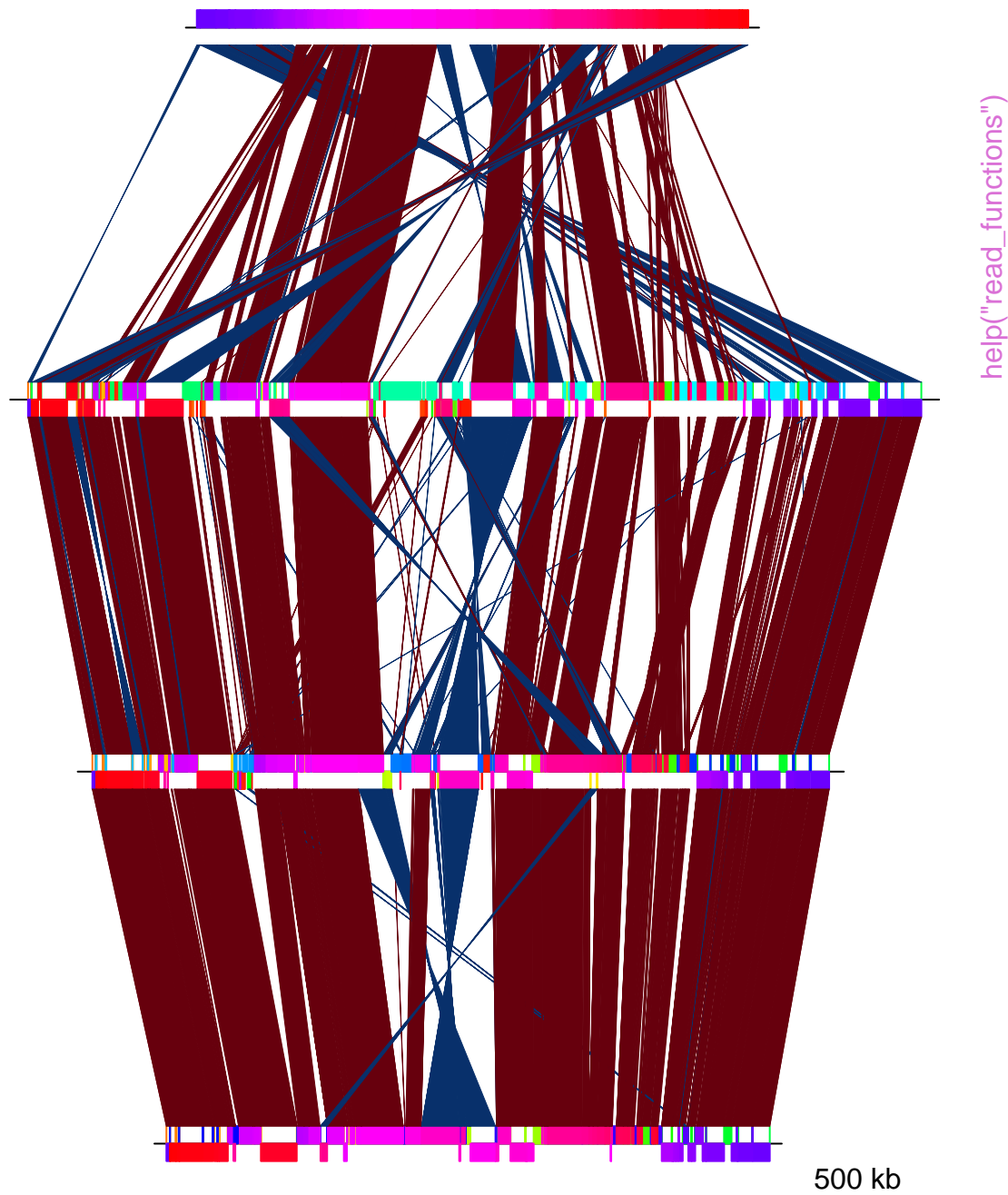


B_bacilliformis

B_grahamii

B_henselae

B_quintana



B_bacilliformis

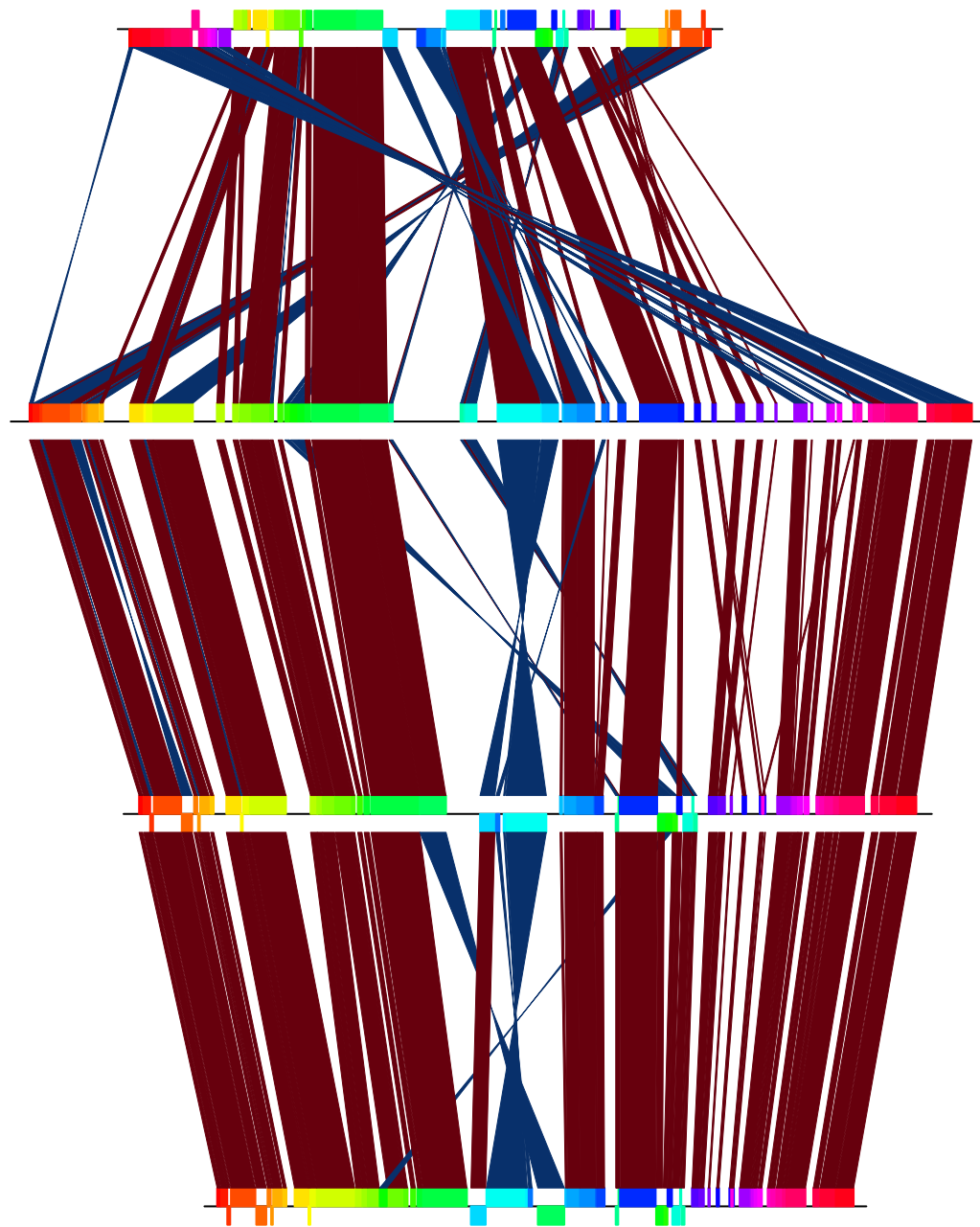
B_grahamii

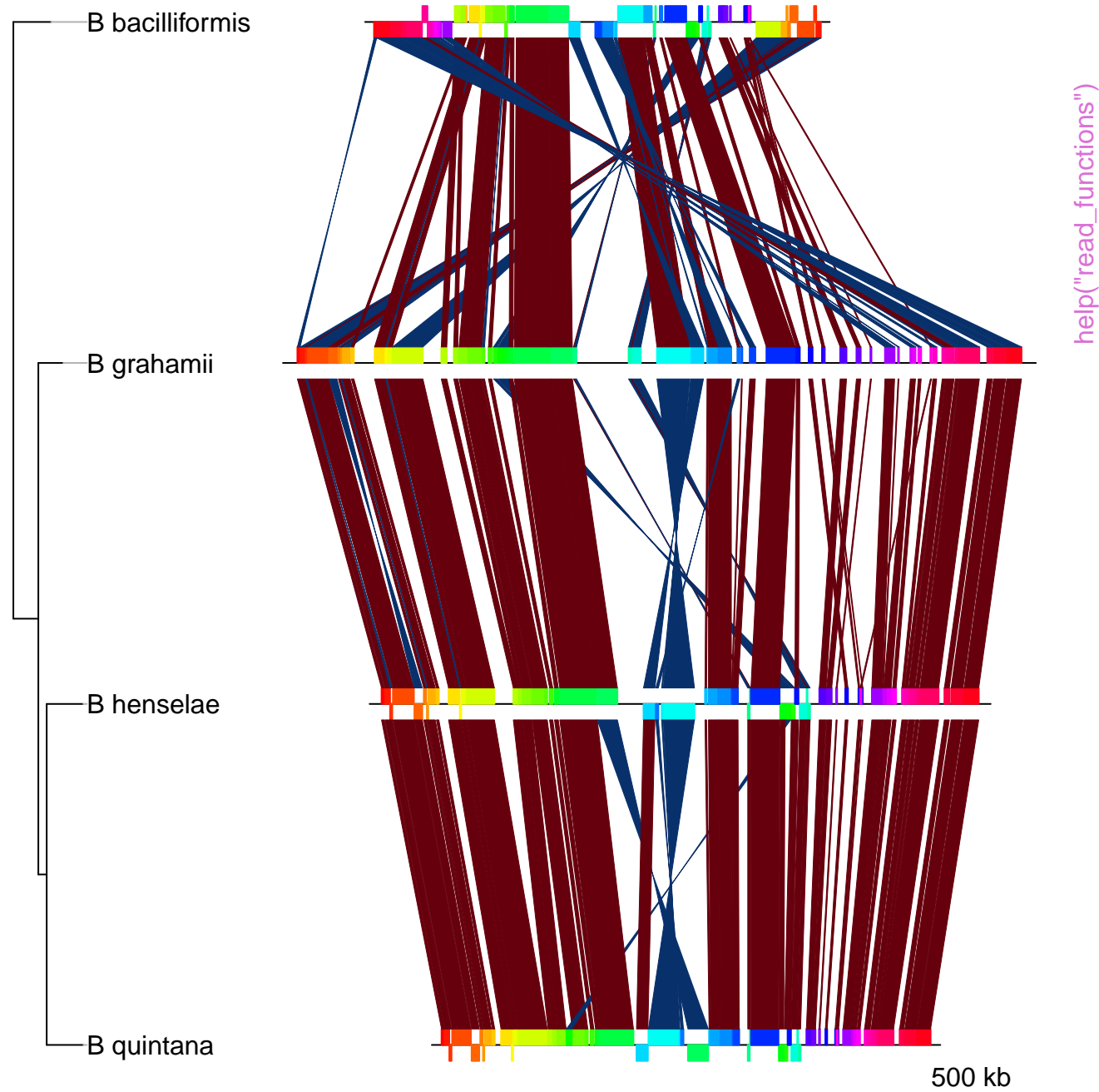
B_henselae

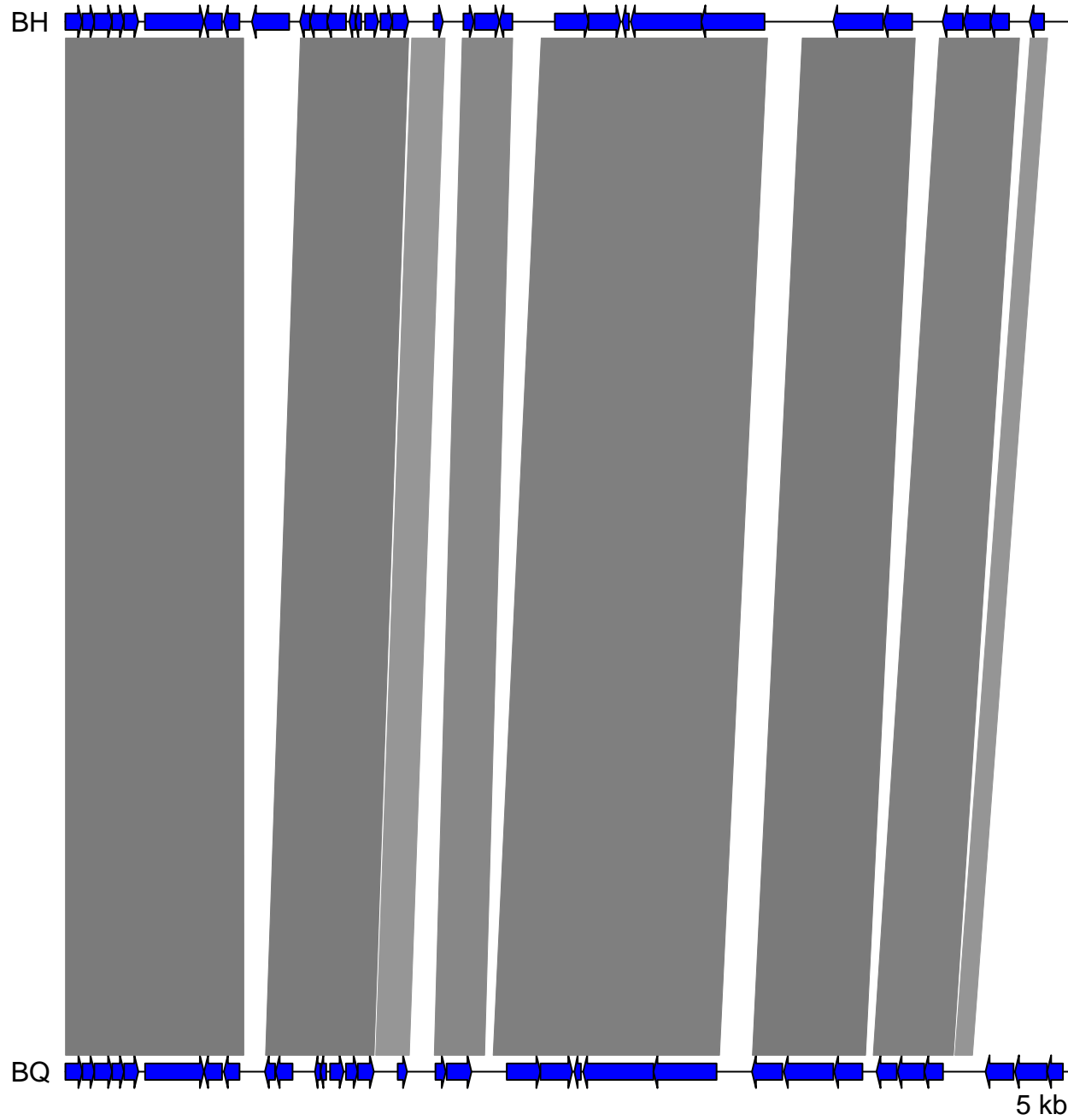
B_quintana

help("read_functions")

500 kb







help("read_functions")

B_bacilliformis

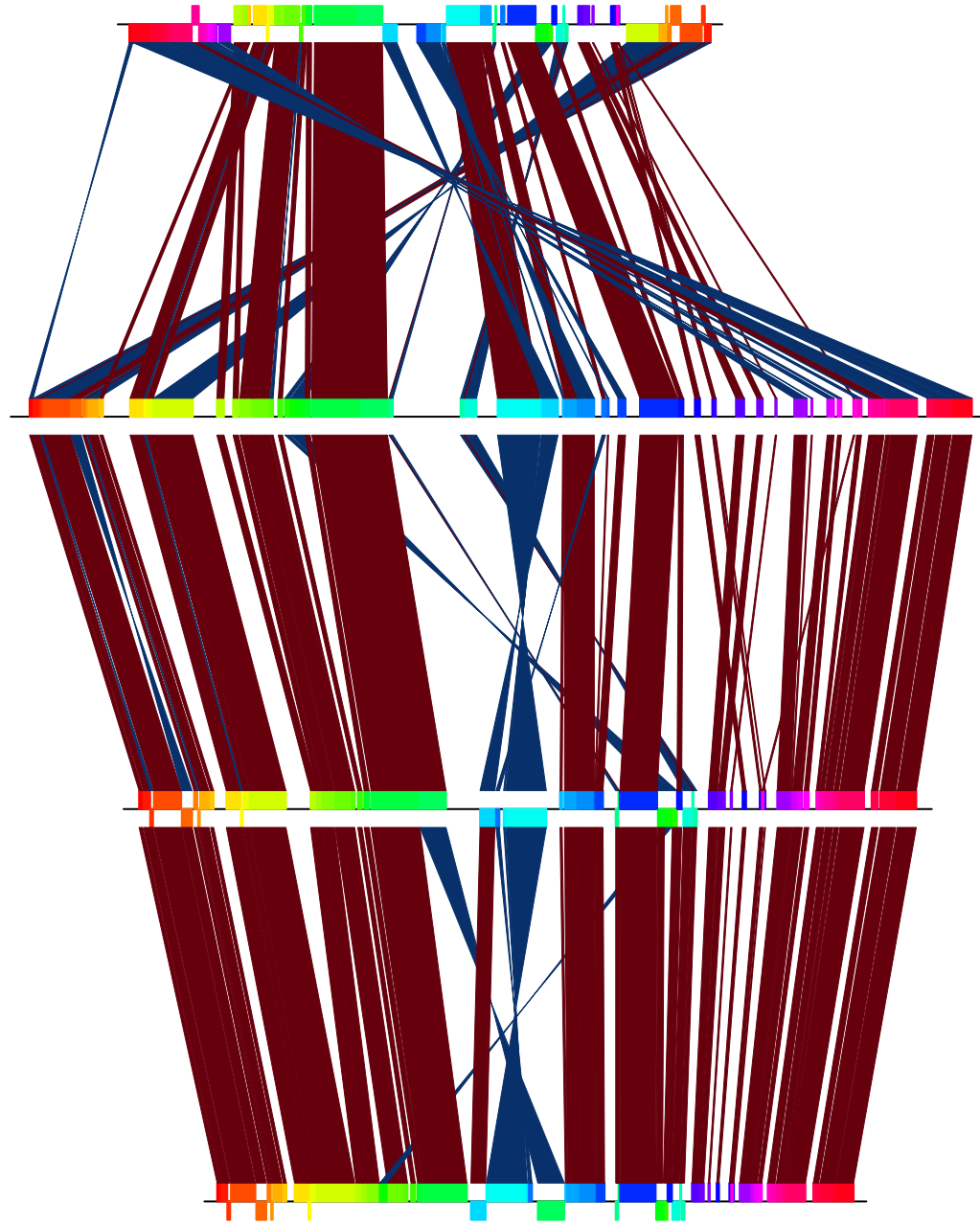
B_grahamii

B_henselae

B_quintana

help("reverse")

500 kb



B_bacilliformis

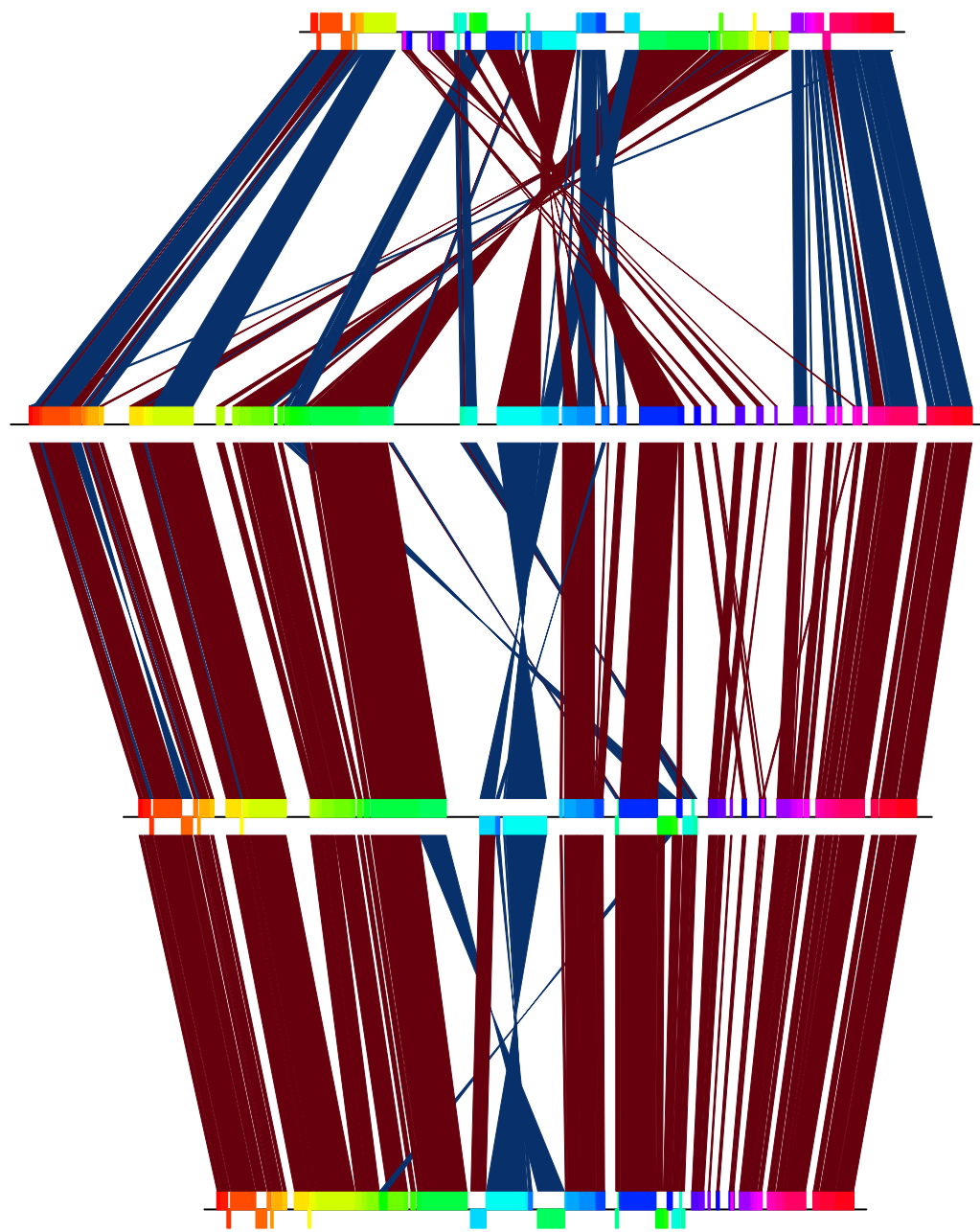
B_grahamii

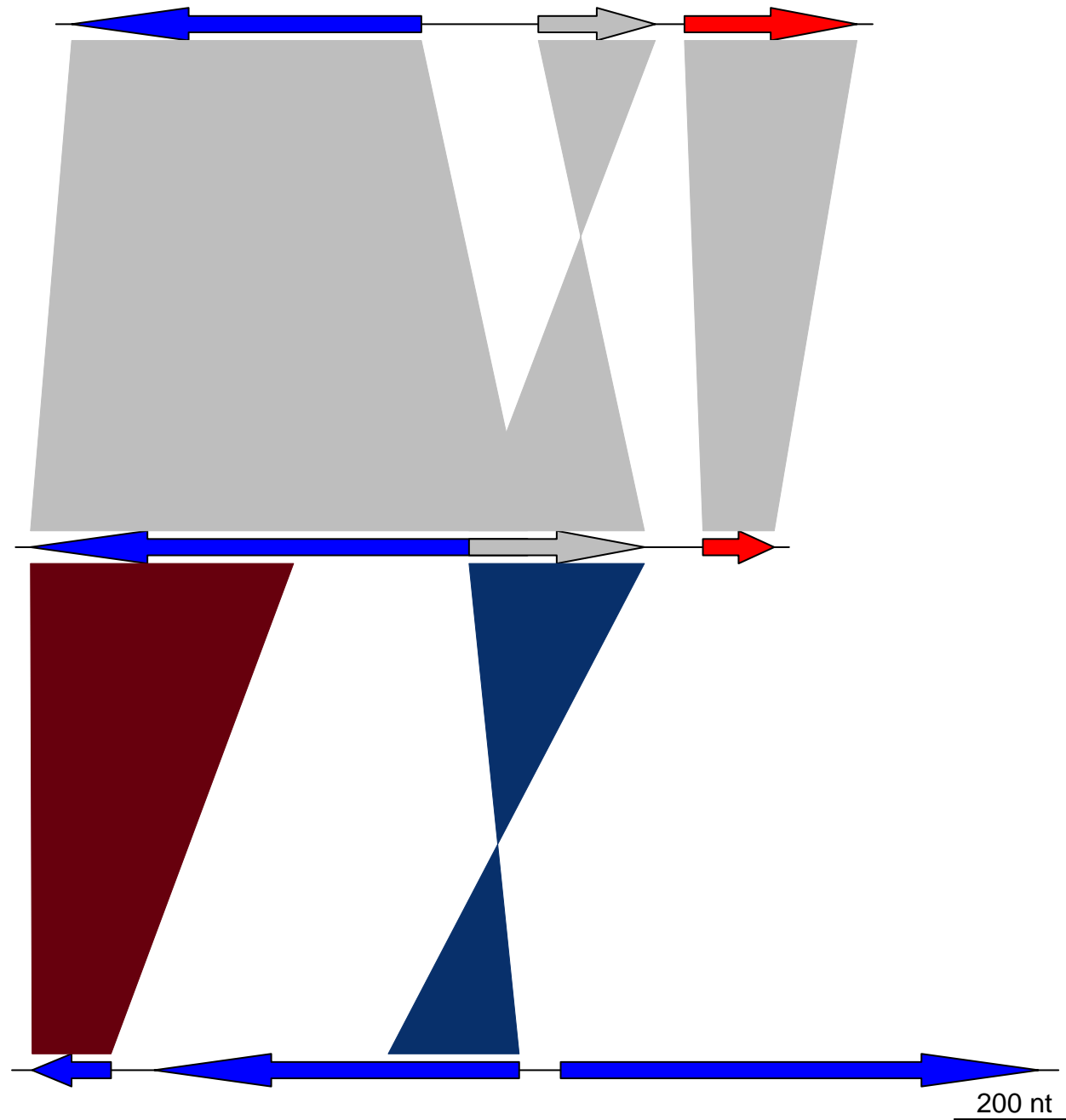
B_henselae

B_quintana

help("reverse")

500 kb





help("three_genes")

