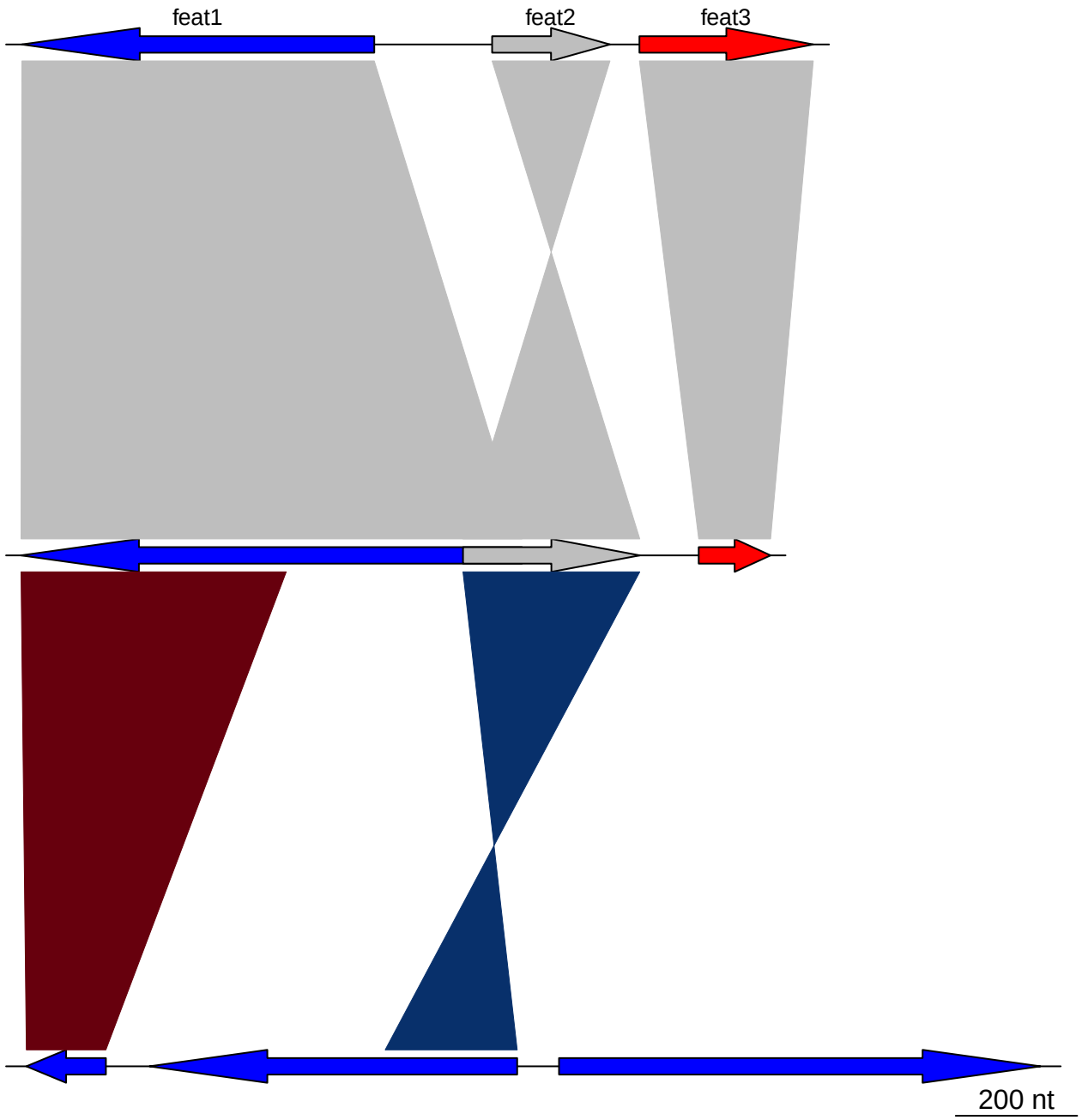

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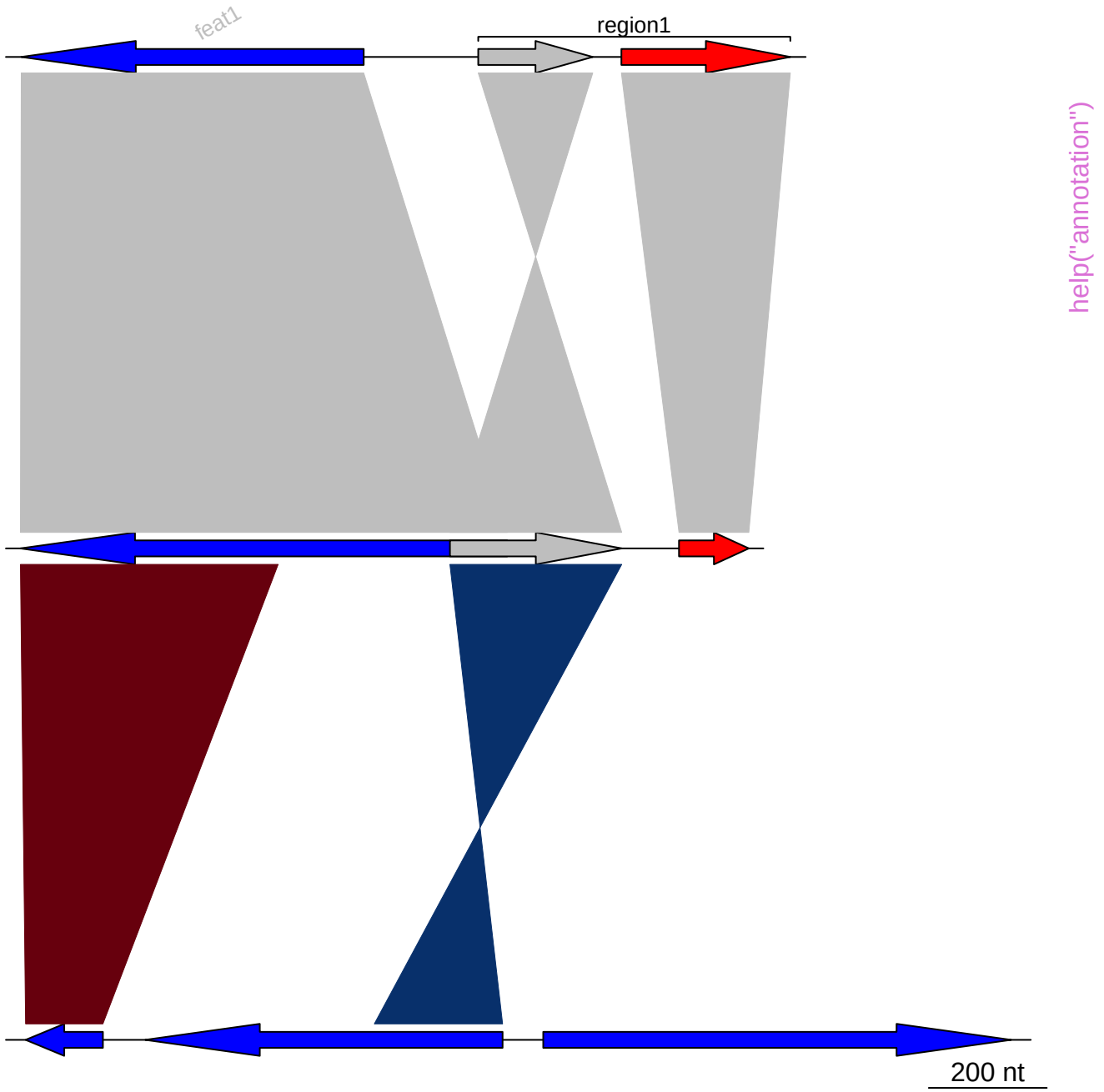
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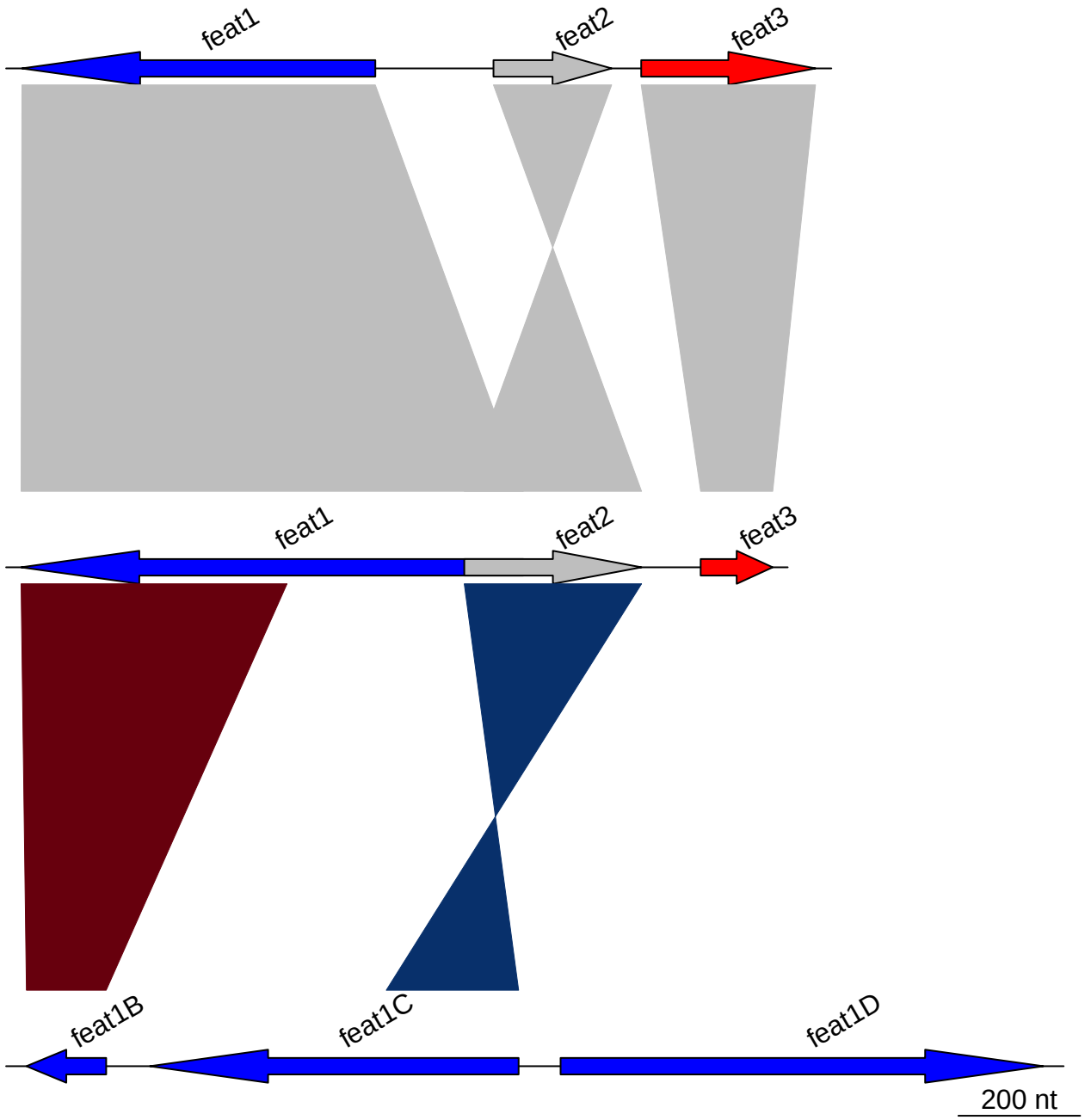
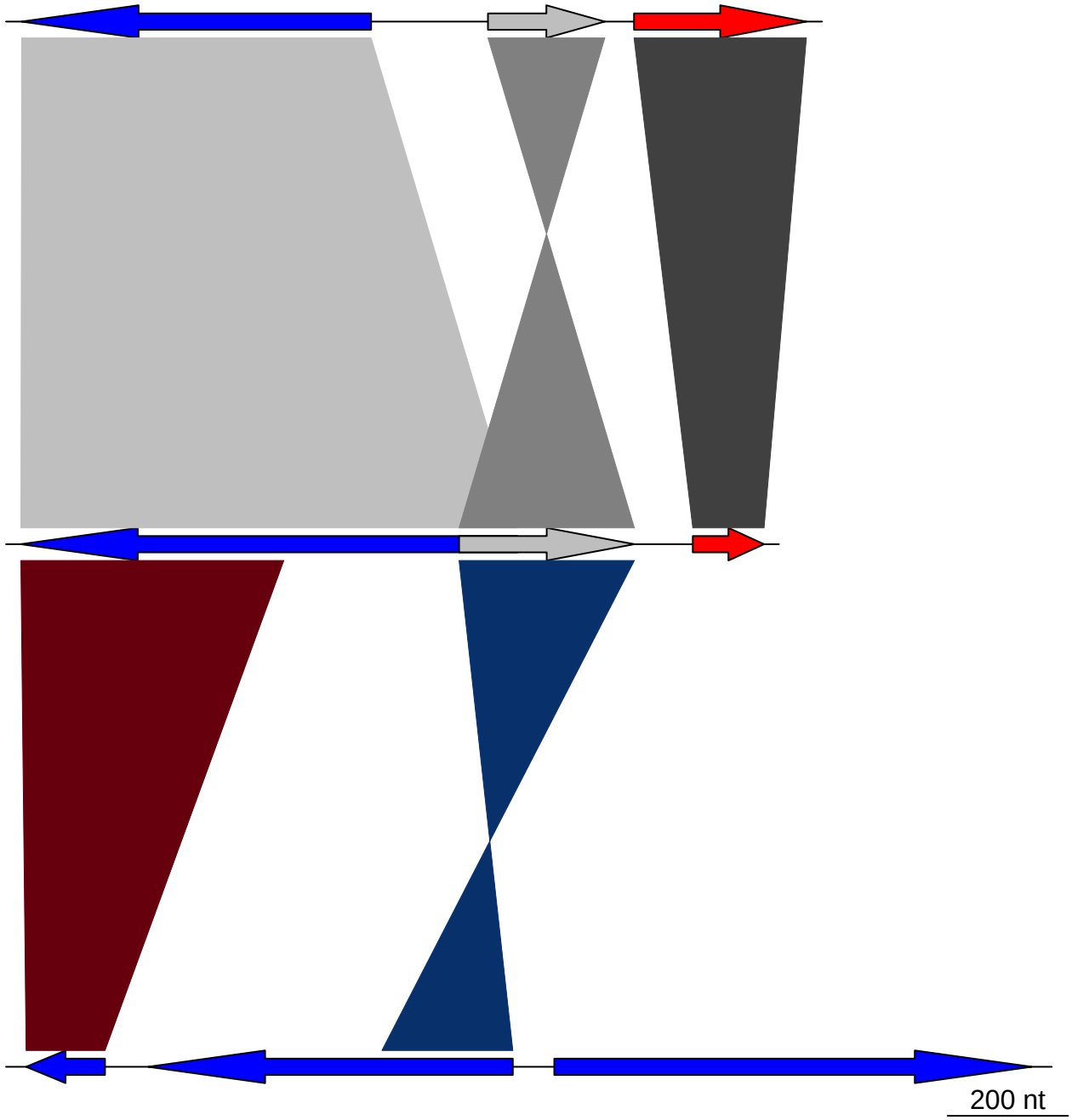


help("annotation")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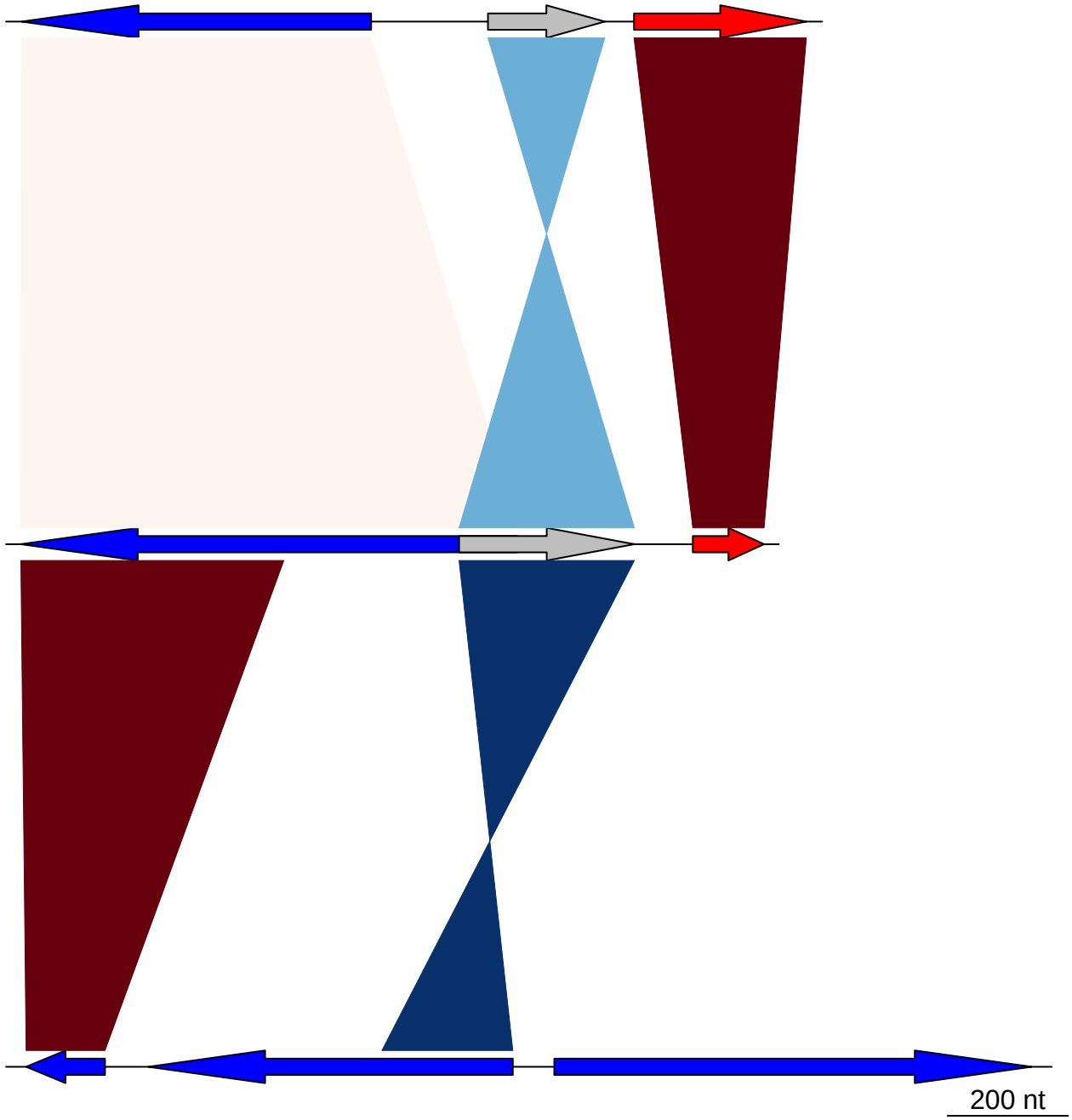
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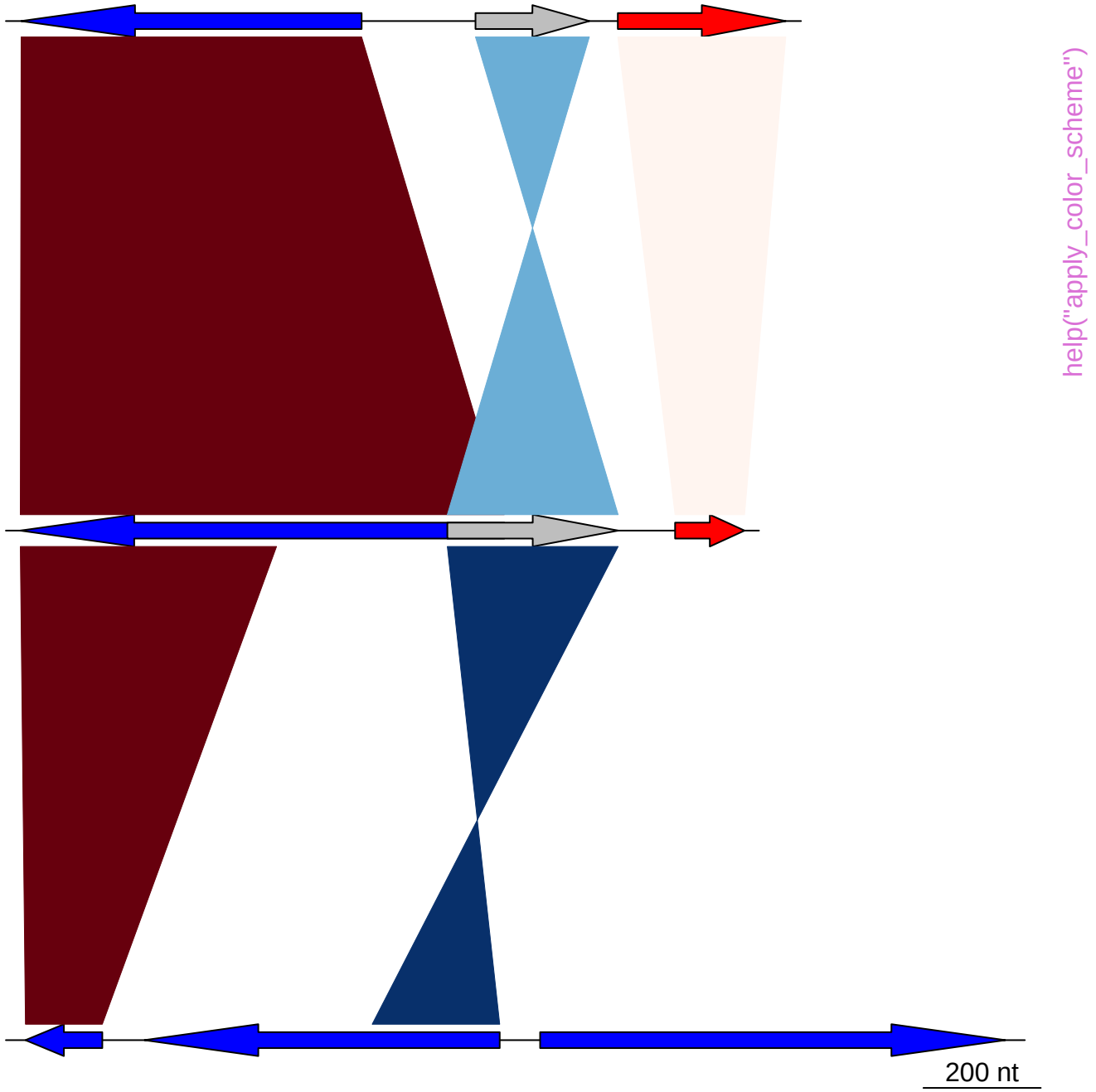
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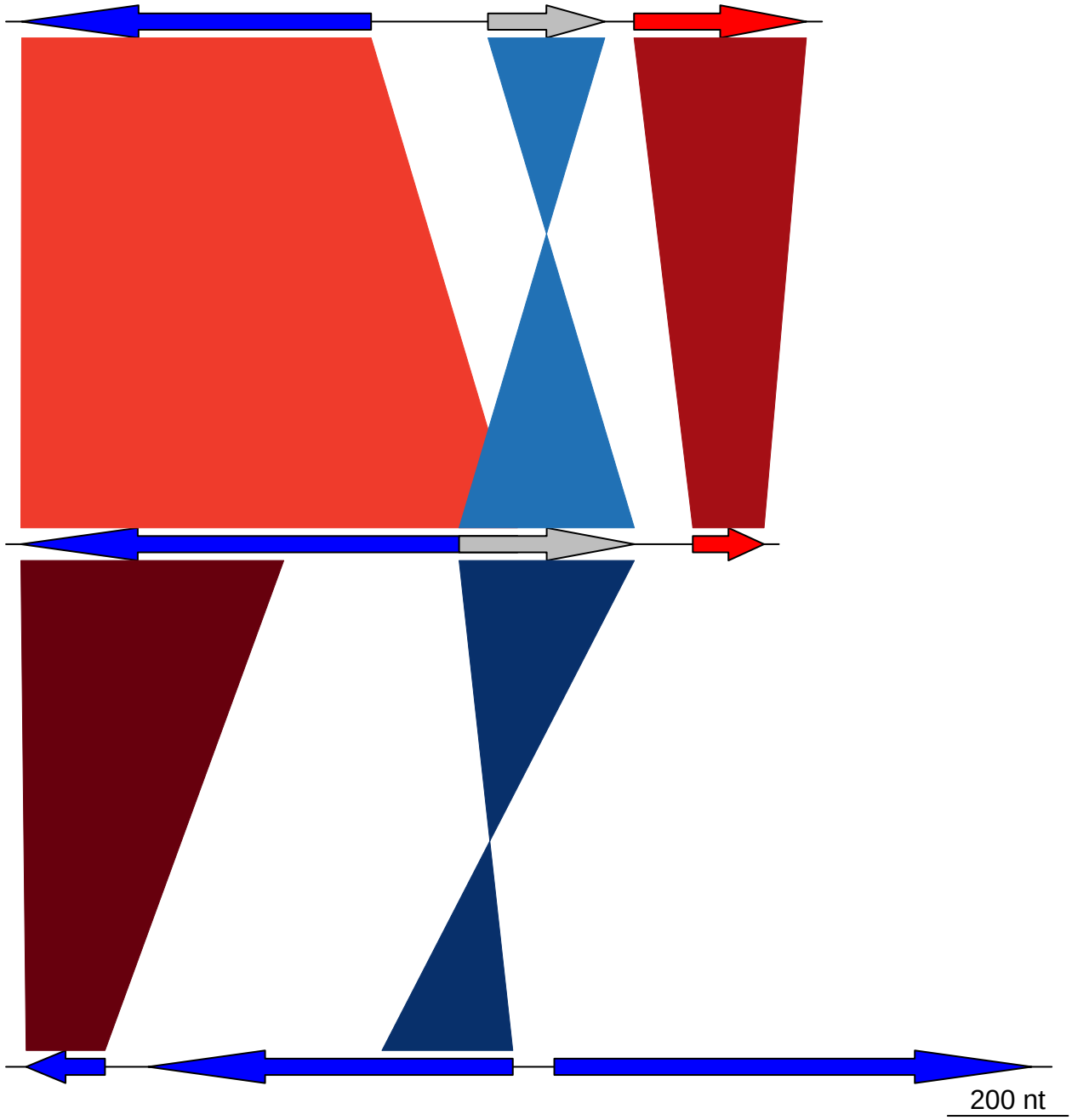
help("apply_color_scheme")
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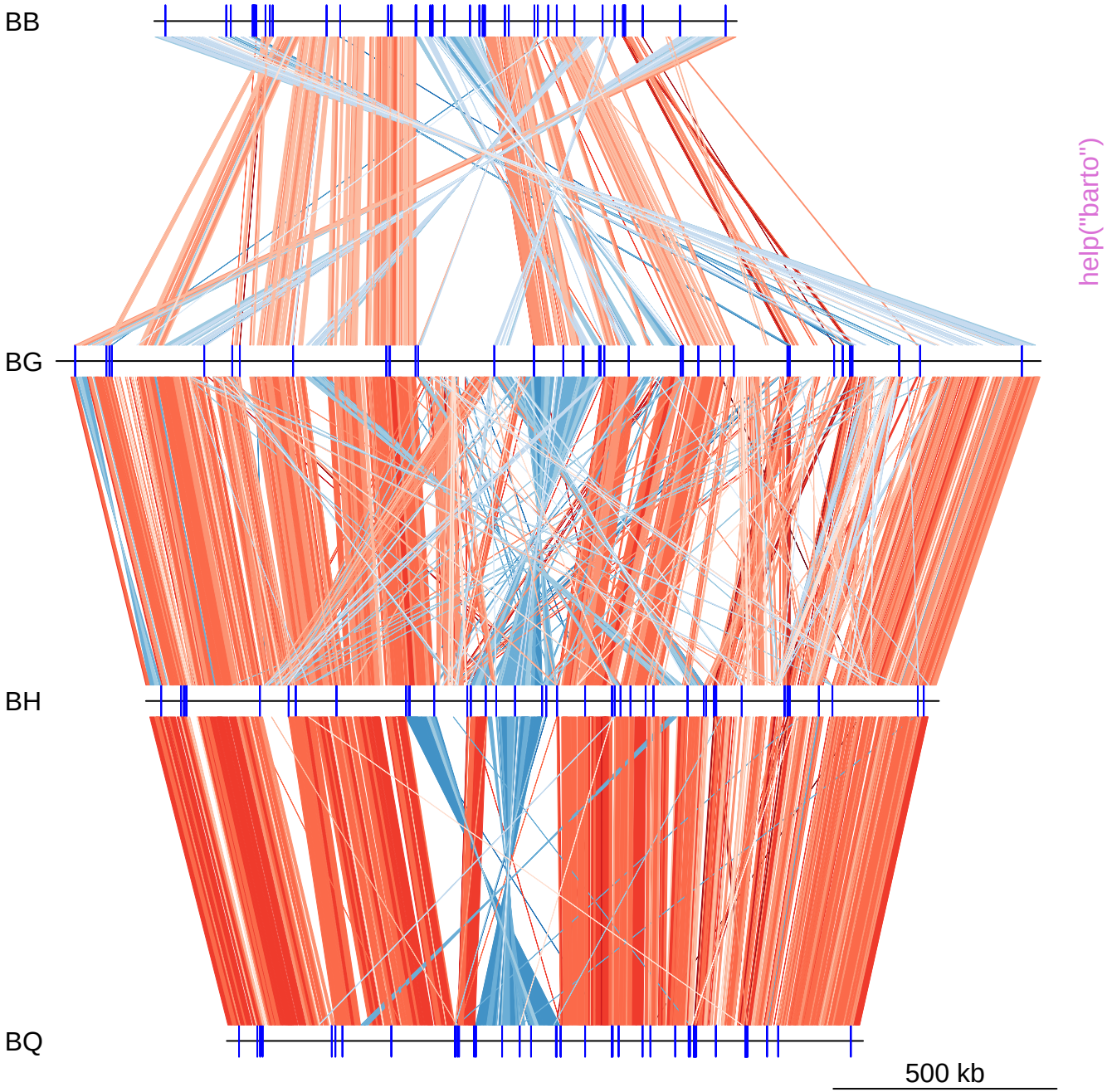
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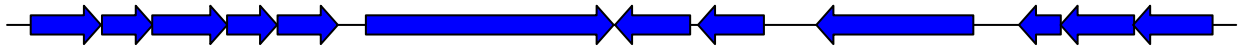
BG
BH
BQ
500 kb
help("barto")

help("gene_types")



2 kb

BH00080
BH00090
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BH00080
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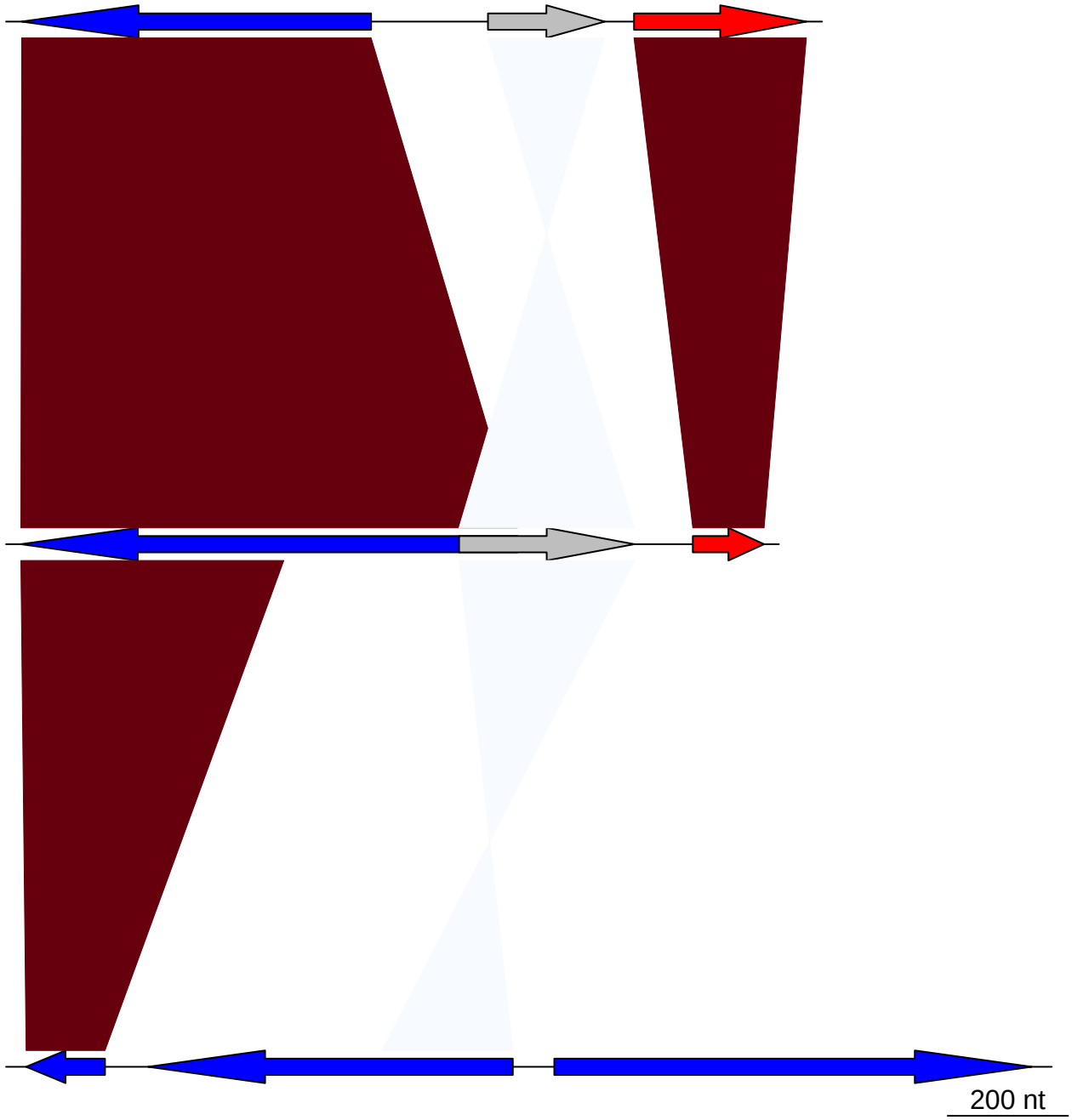
2 kb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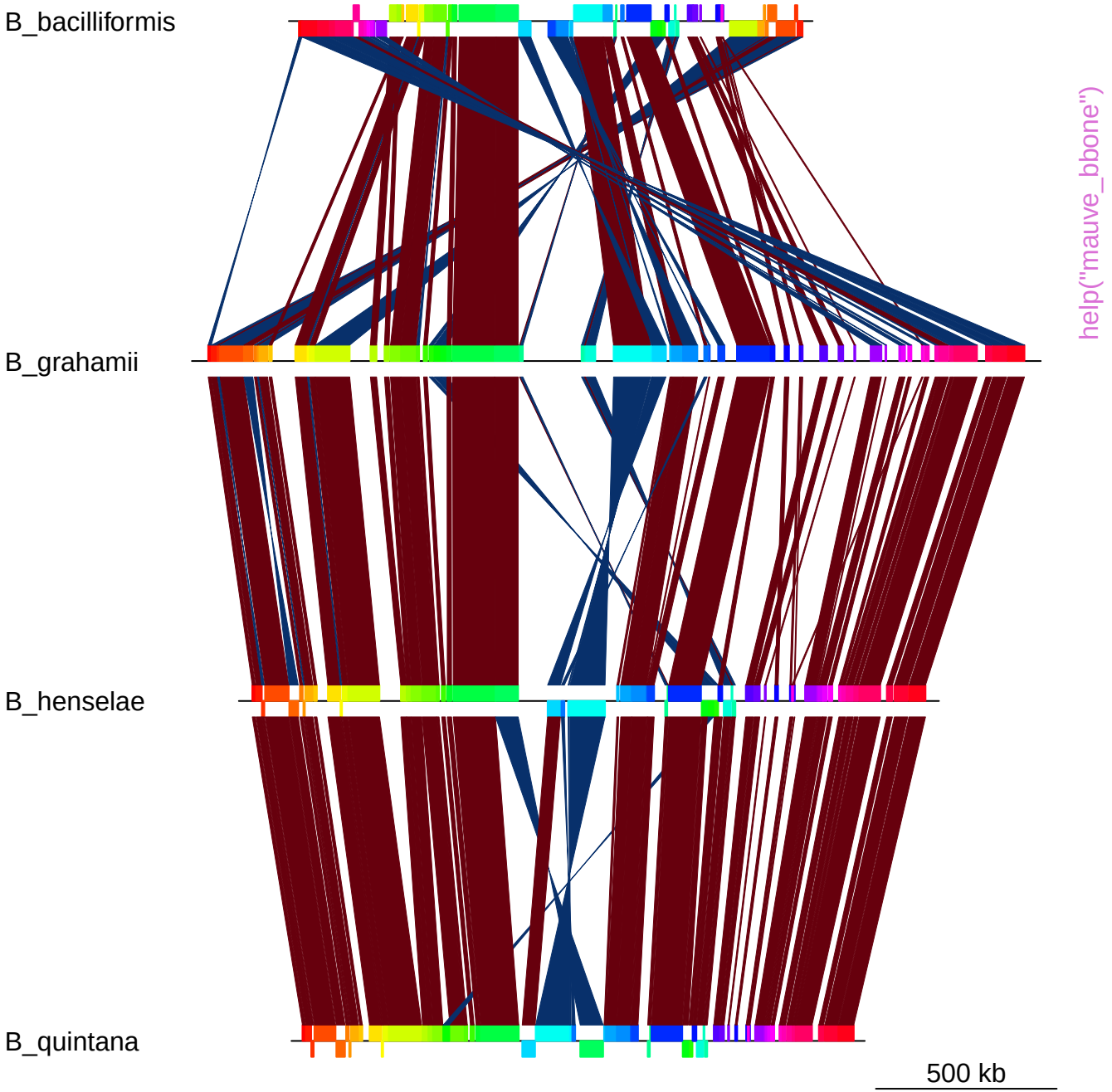
B_bacilliformis
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B_henselae
B_quintana
500 kb
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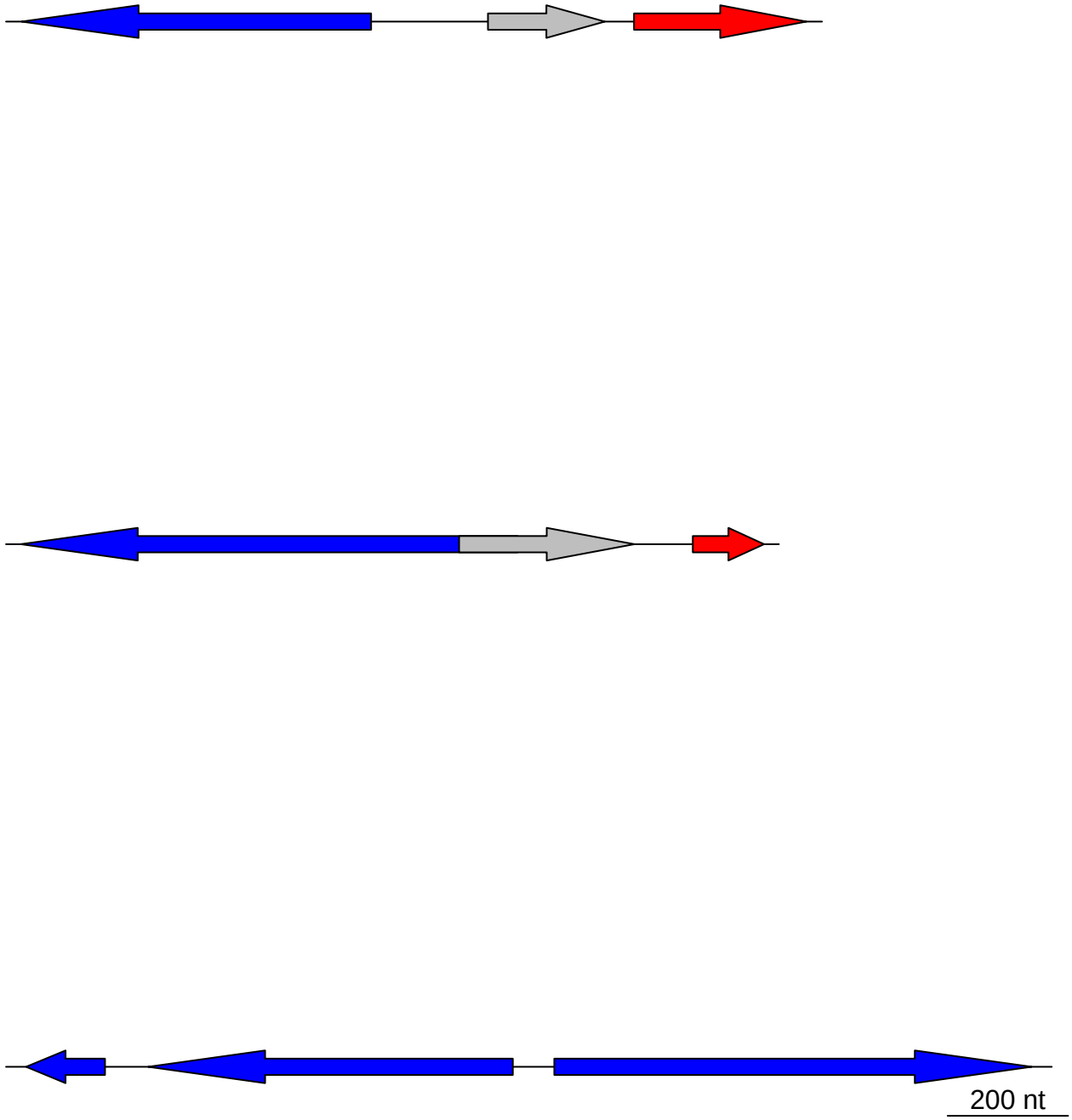
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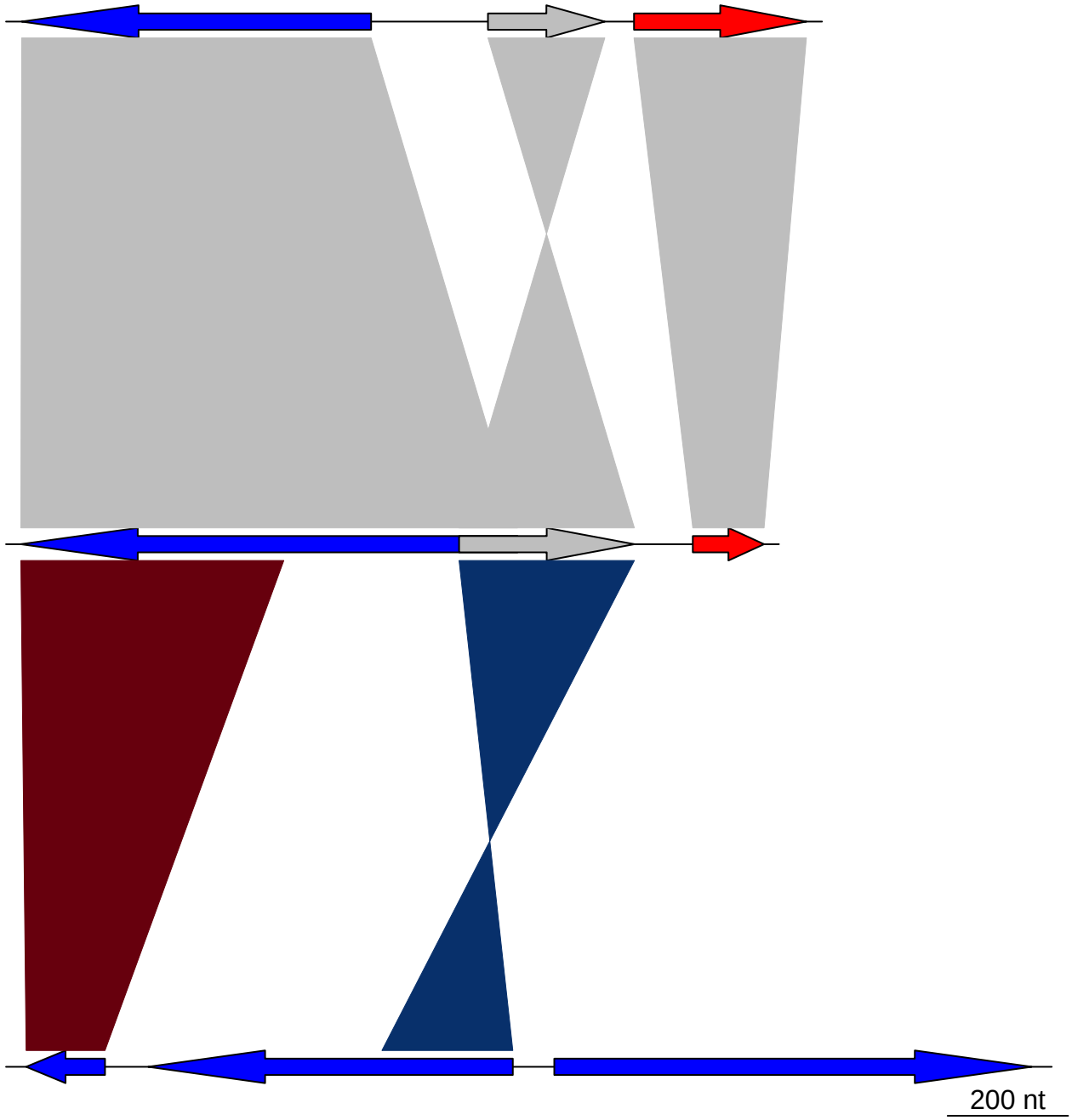
help("plot_gene_map")
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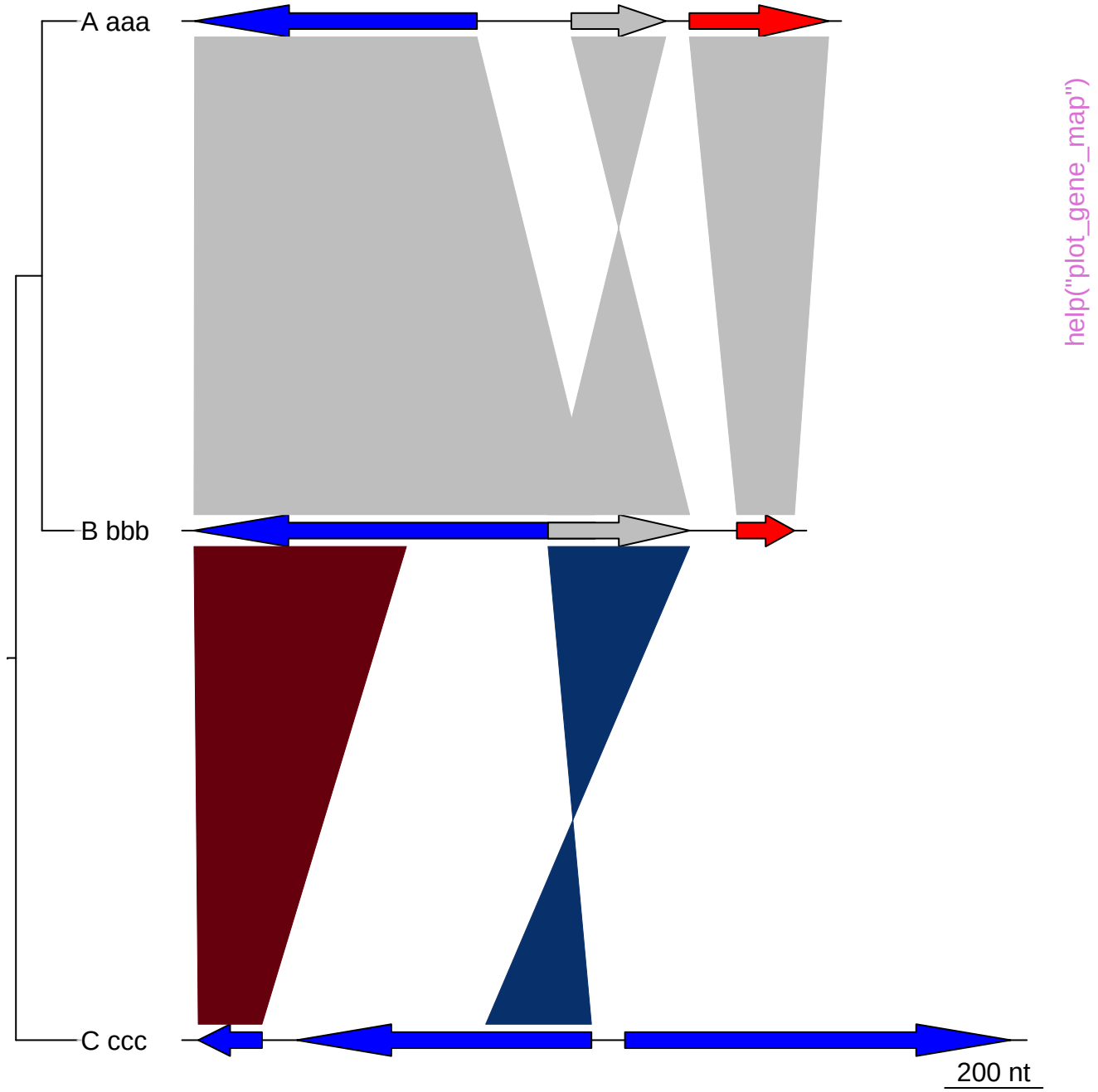
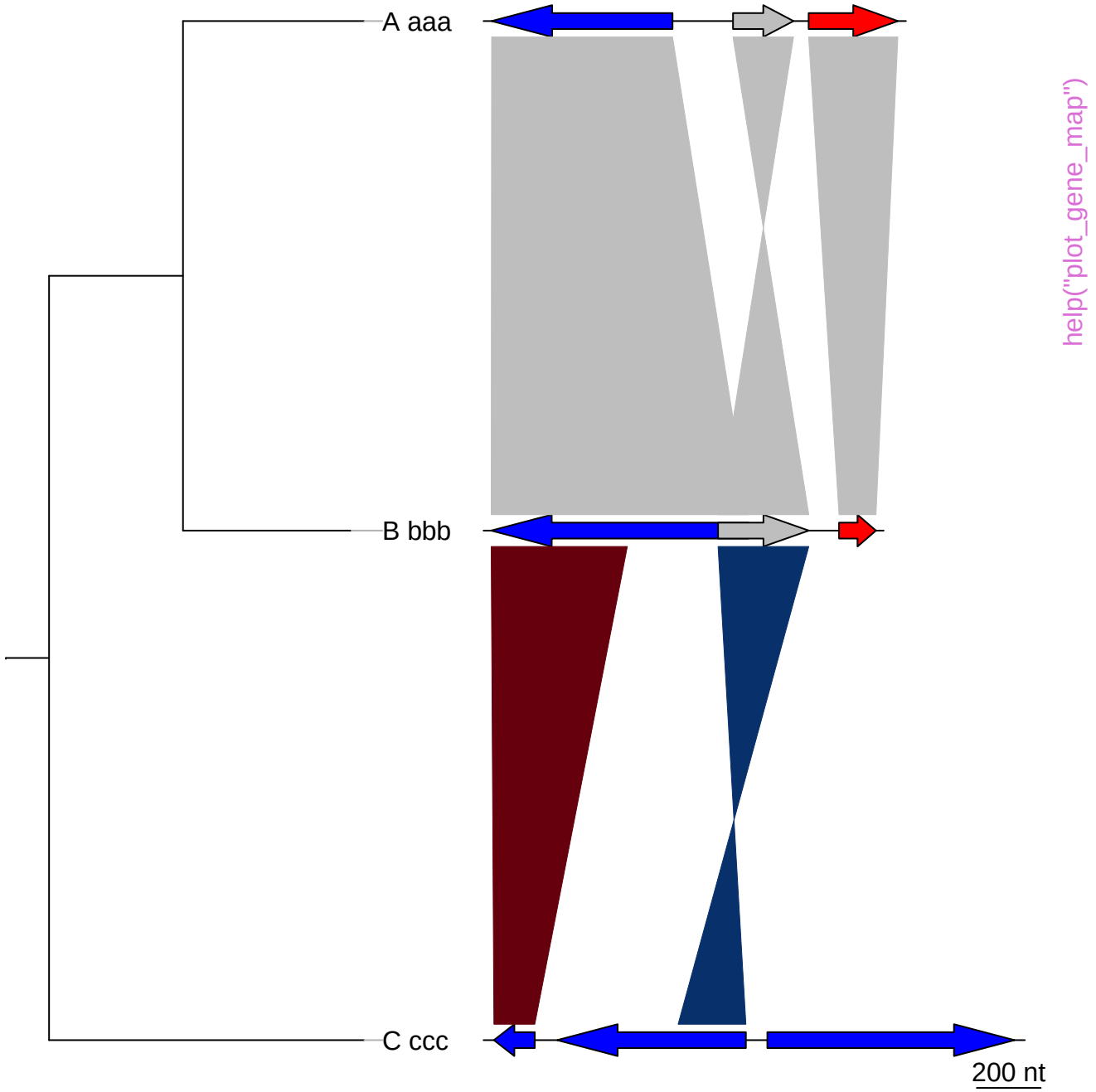

A aaa
B bbb
C ccc
200 nt
help("plot_gene_map")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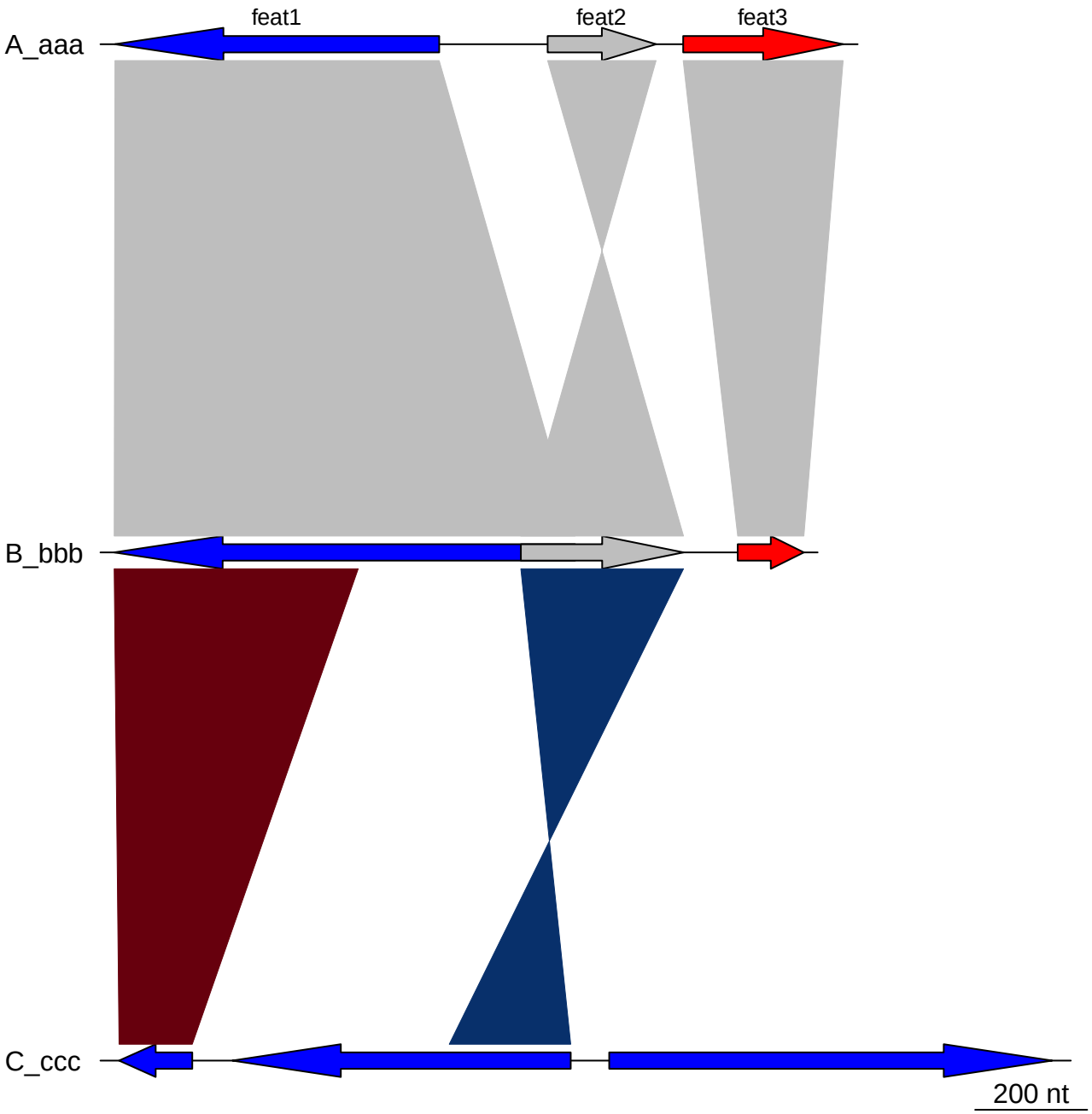
B bbb
C ccc
200 nt
help("plot_gene_map")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feat2
feat3
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B_bbb
C_ccc
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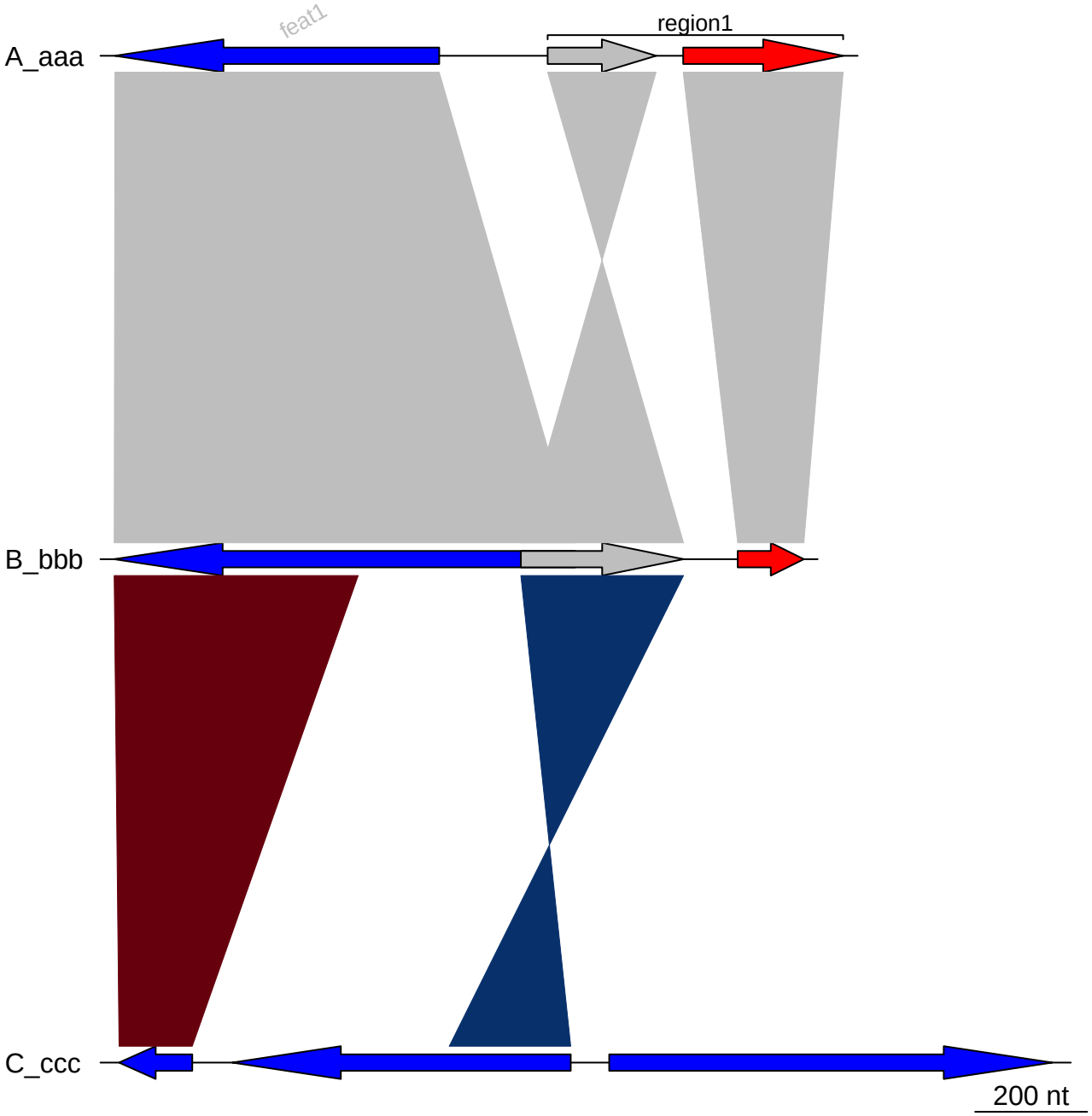
help("plot_gene_map")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region1
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B_bbb
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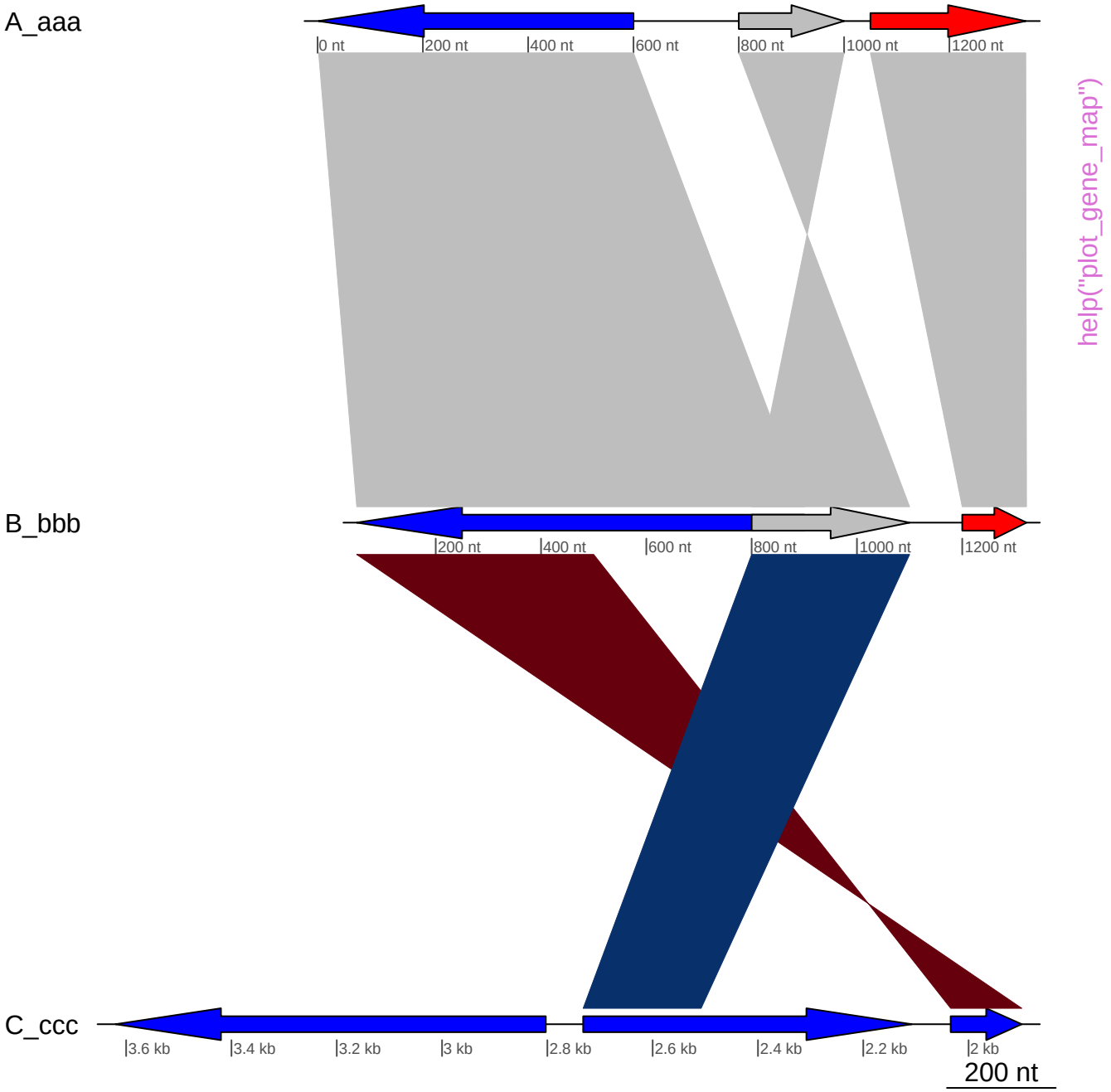
help("plot_gene_map")


A_aaa
0 nt
200 nt
400 nt
600 nt
800 nt
1000 nt
1200 nt
help("plot_gene_map")
B_bbb
200 nt
400 nt
600 nt
800 nt
1000 nt
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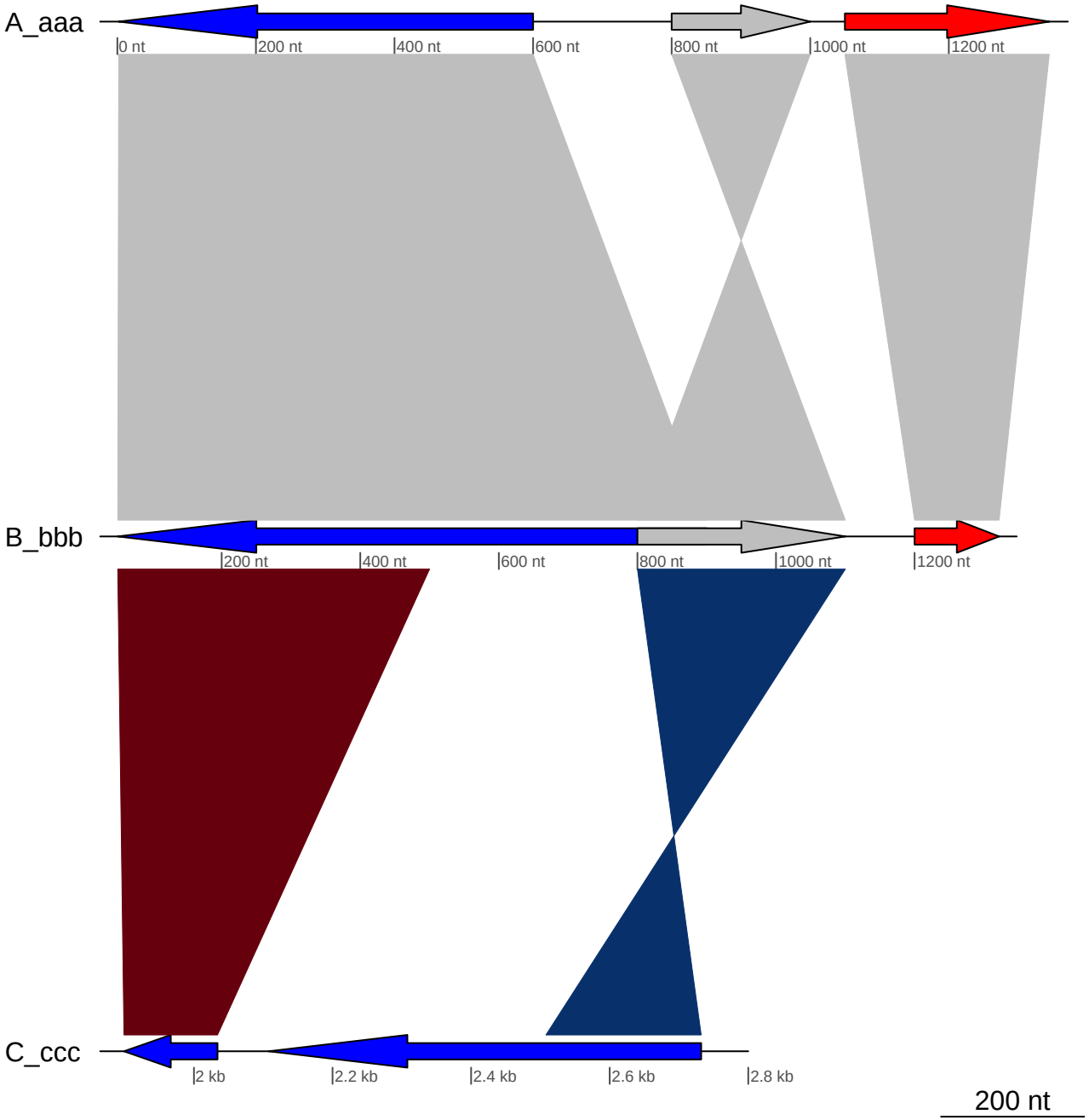
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1200 nt
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200 nt
400 nt
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C_ccc
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2.2 kb
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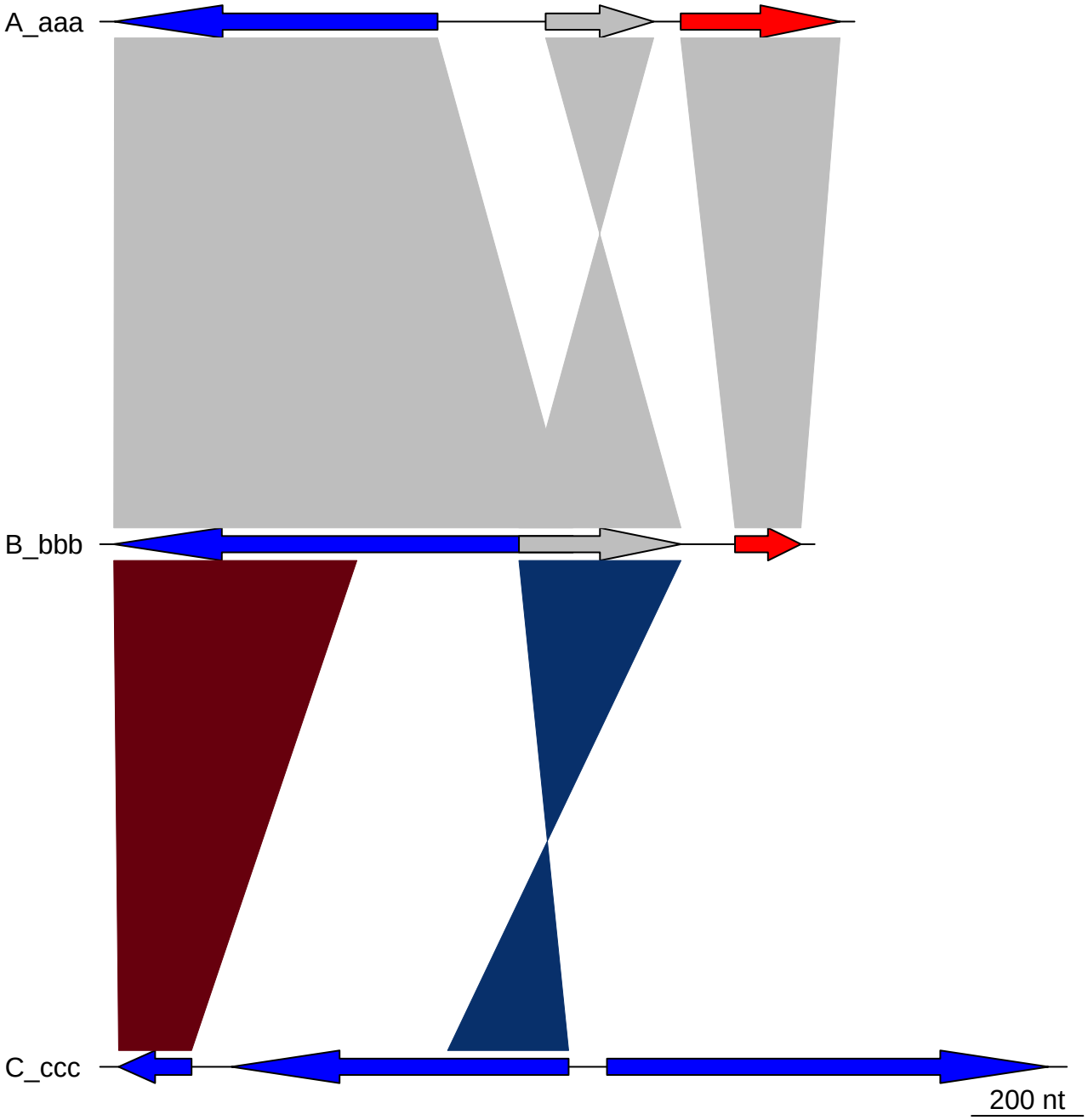
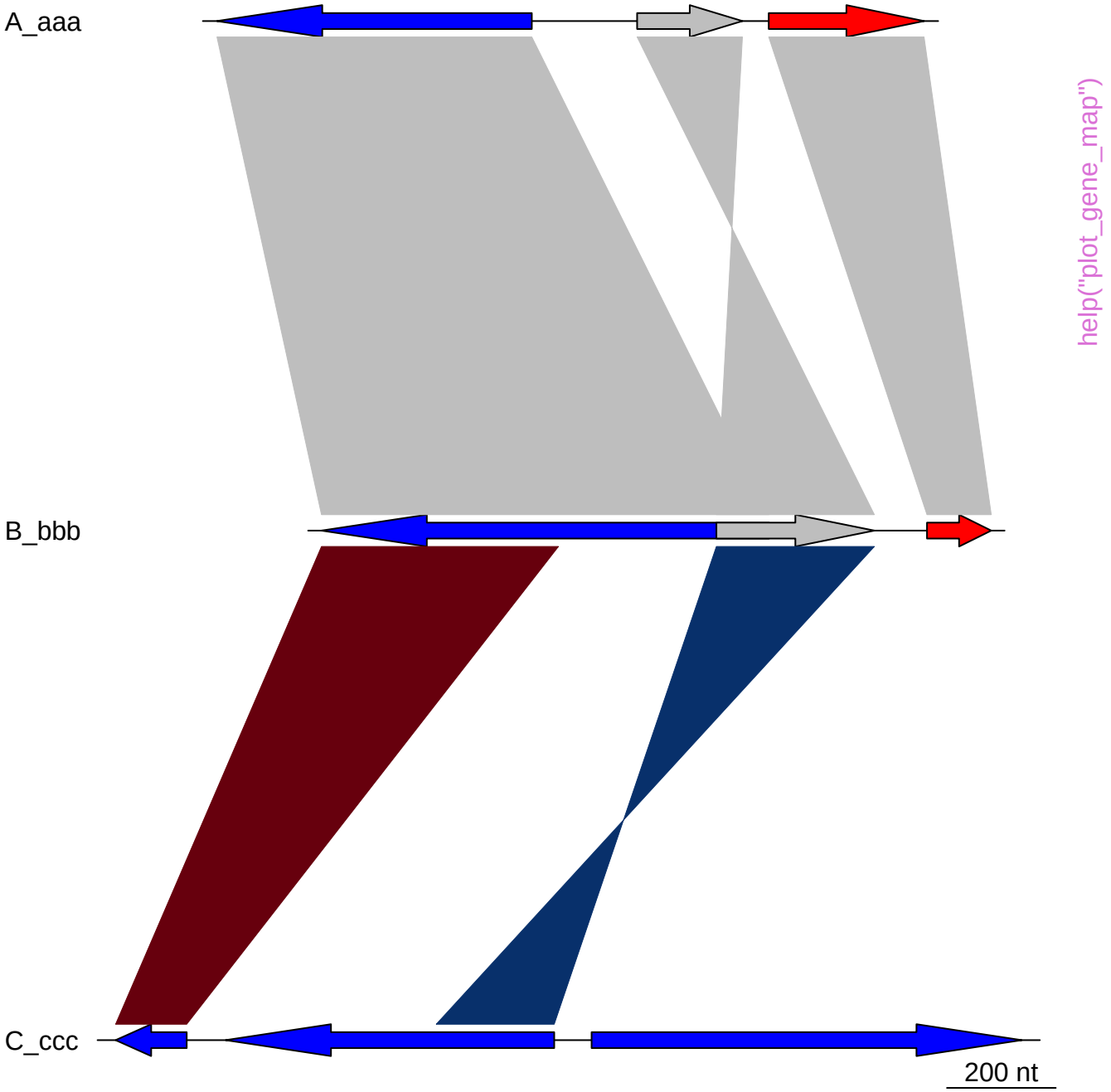
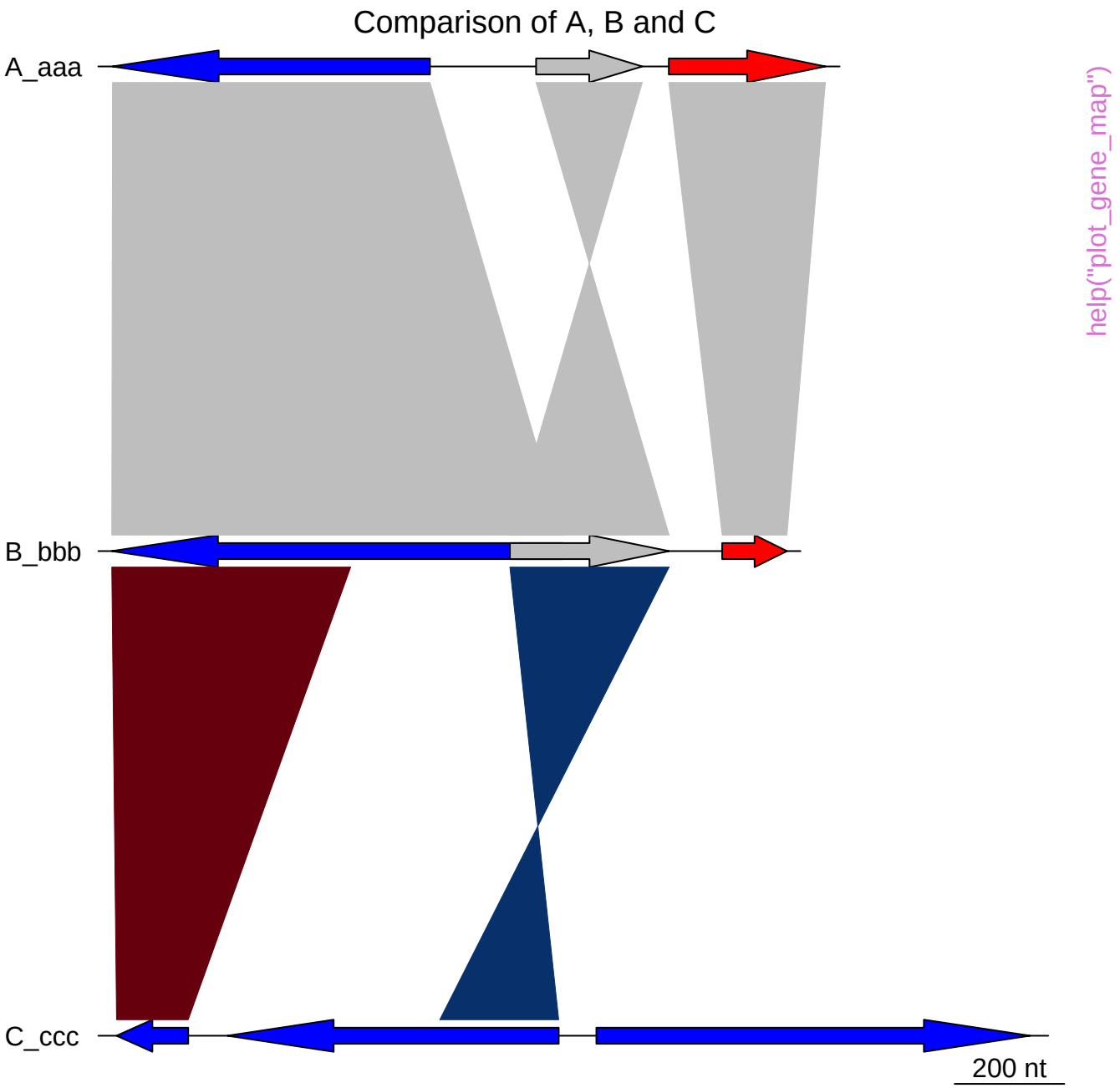
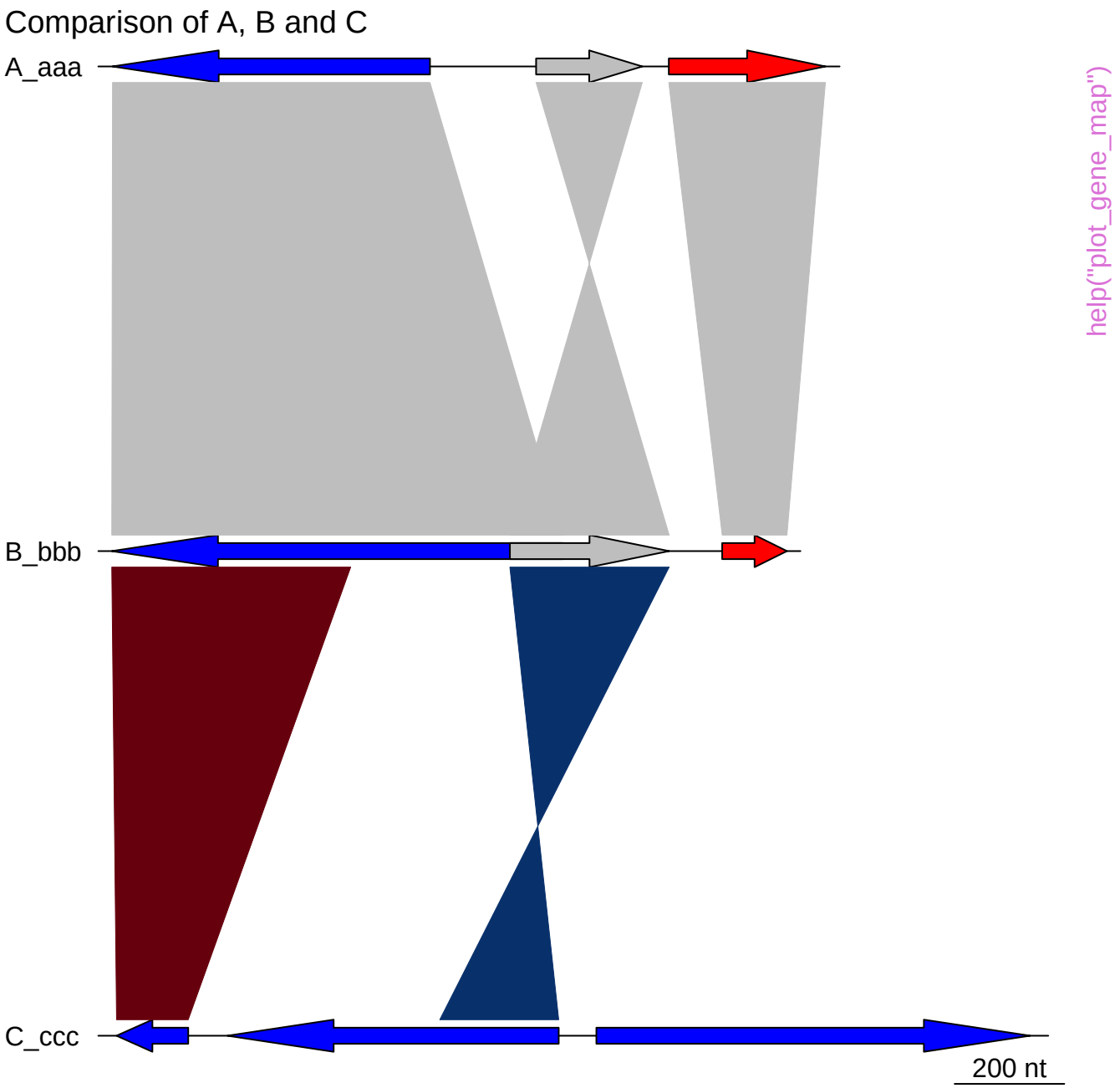
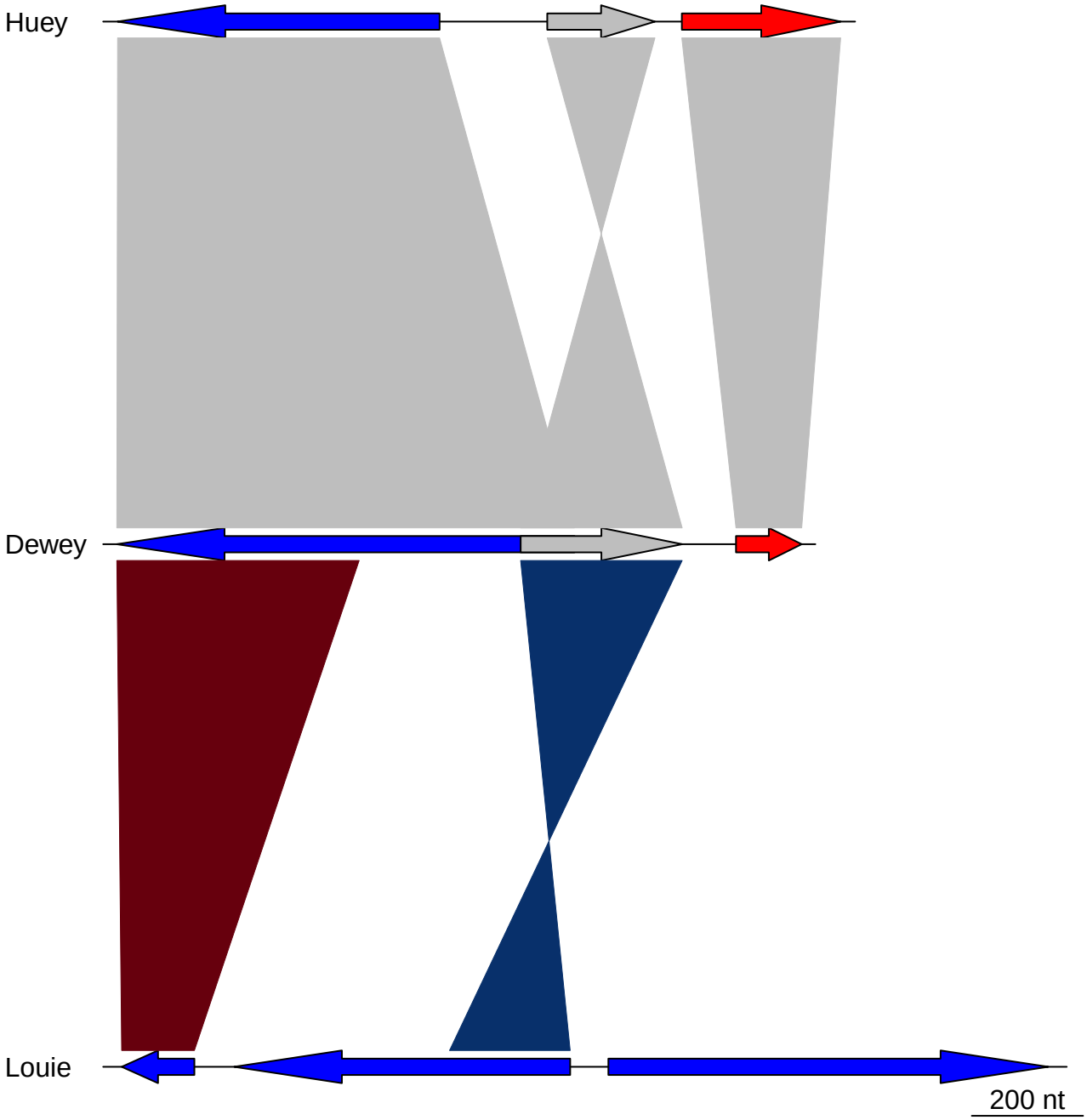
help("plot_gene_map")


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
A_aaa
B_bbb
C_ccc
200 nt
help("plot_gene_map")


Comparison of A, B and C
A_aaa
B_bbb
C_ccc
200 nt
help("plot_gene_map")


Huey
Dewey
Louie
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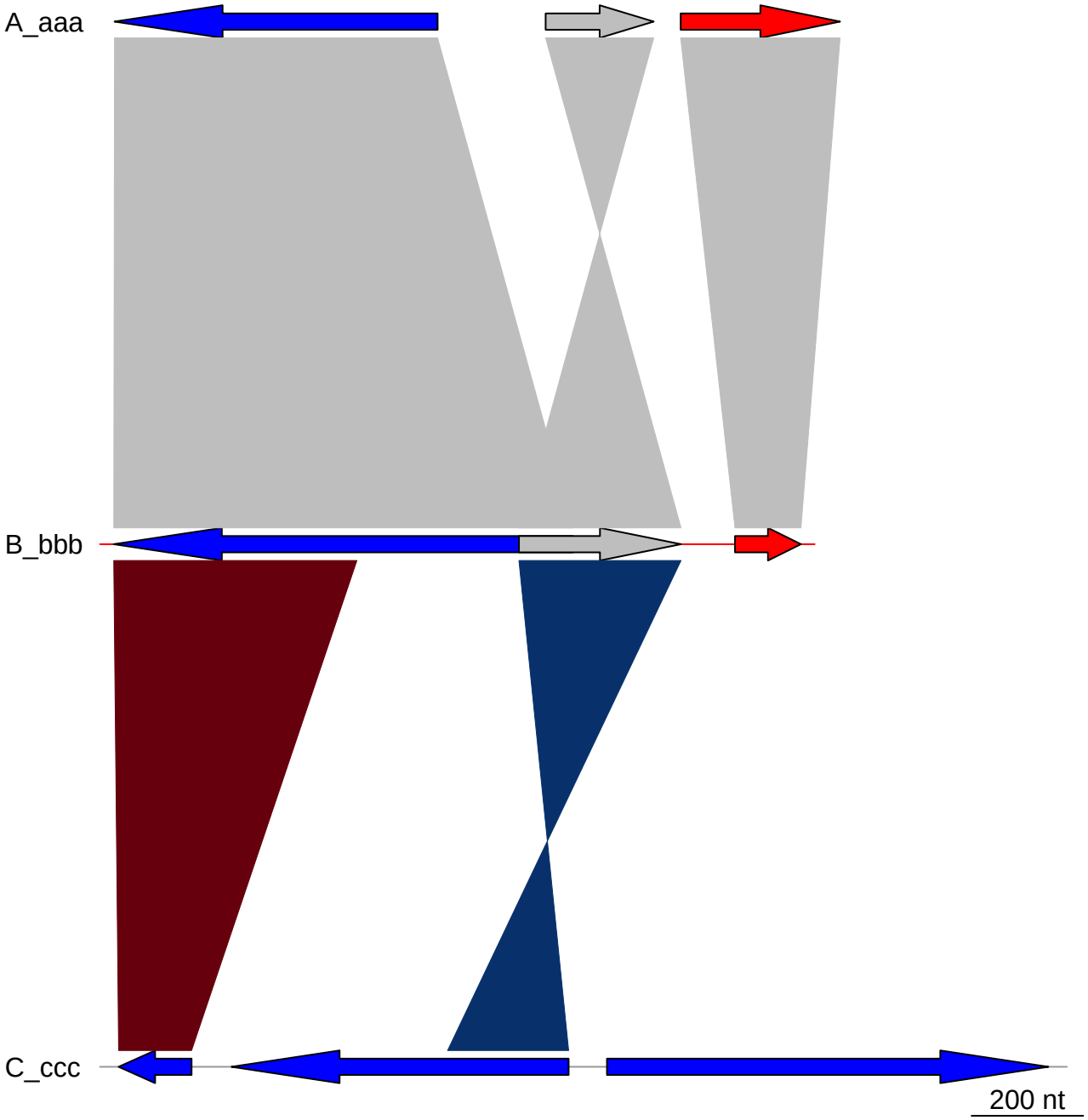
help("plot_gene_map")


A_aaa
B_bbb
C_ccc
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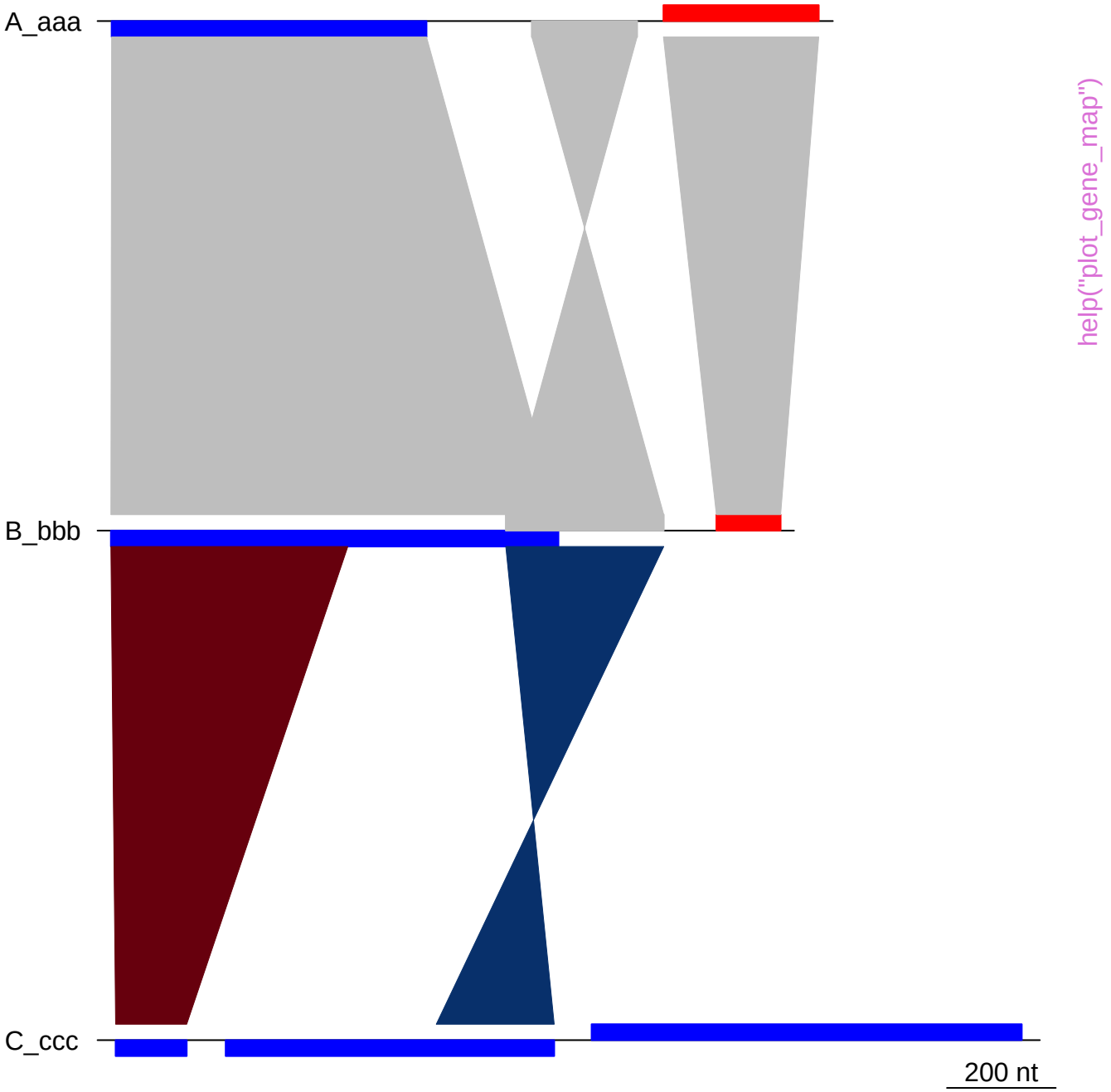
help("plot_gene_map")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B_bbb
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200 nt
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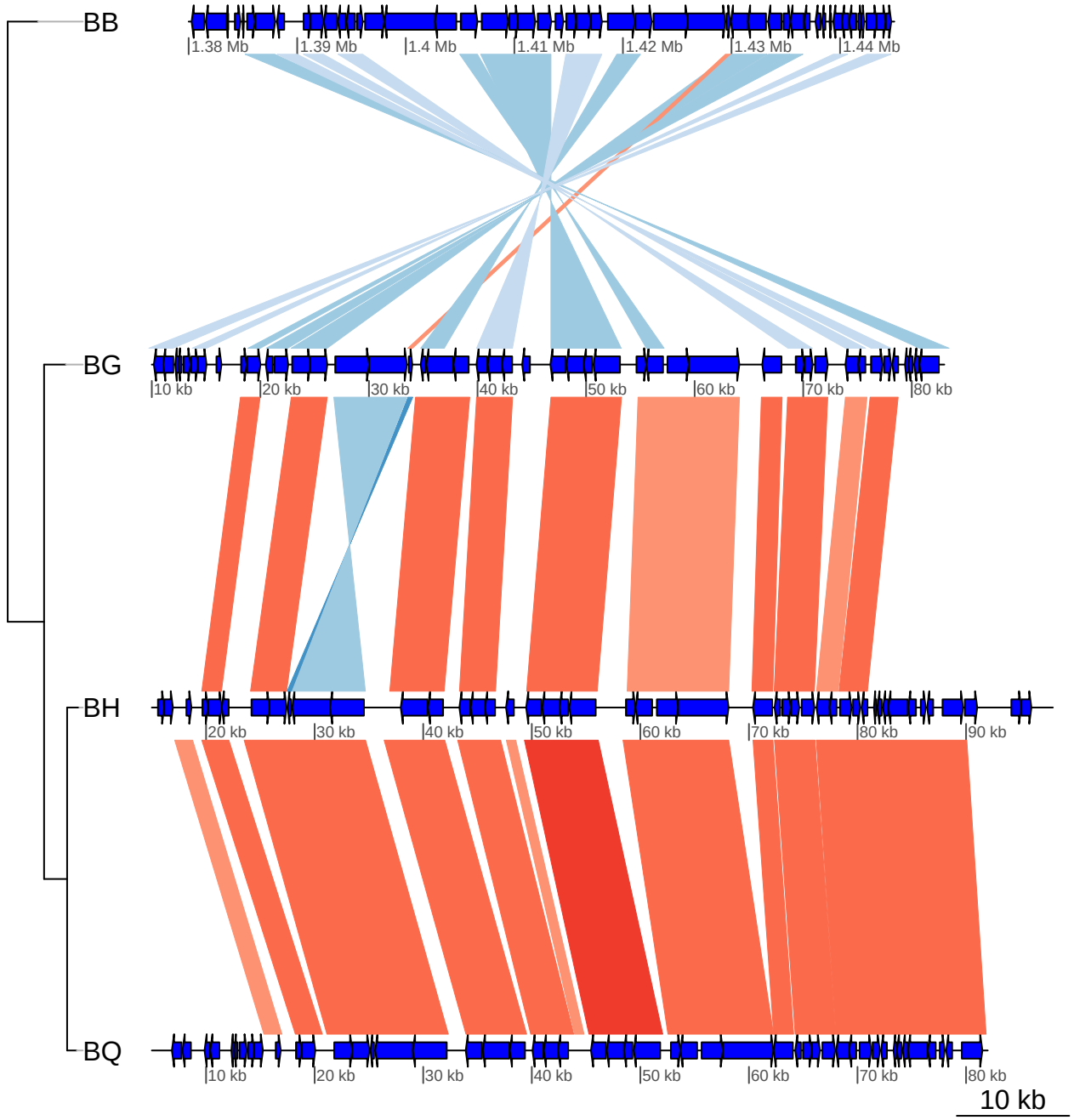
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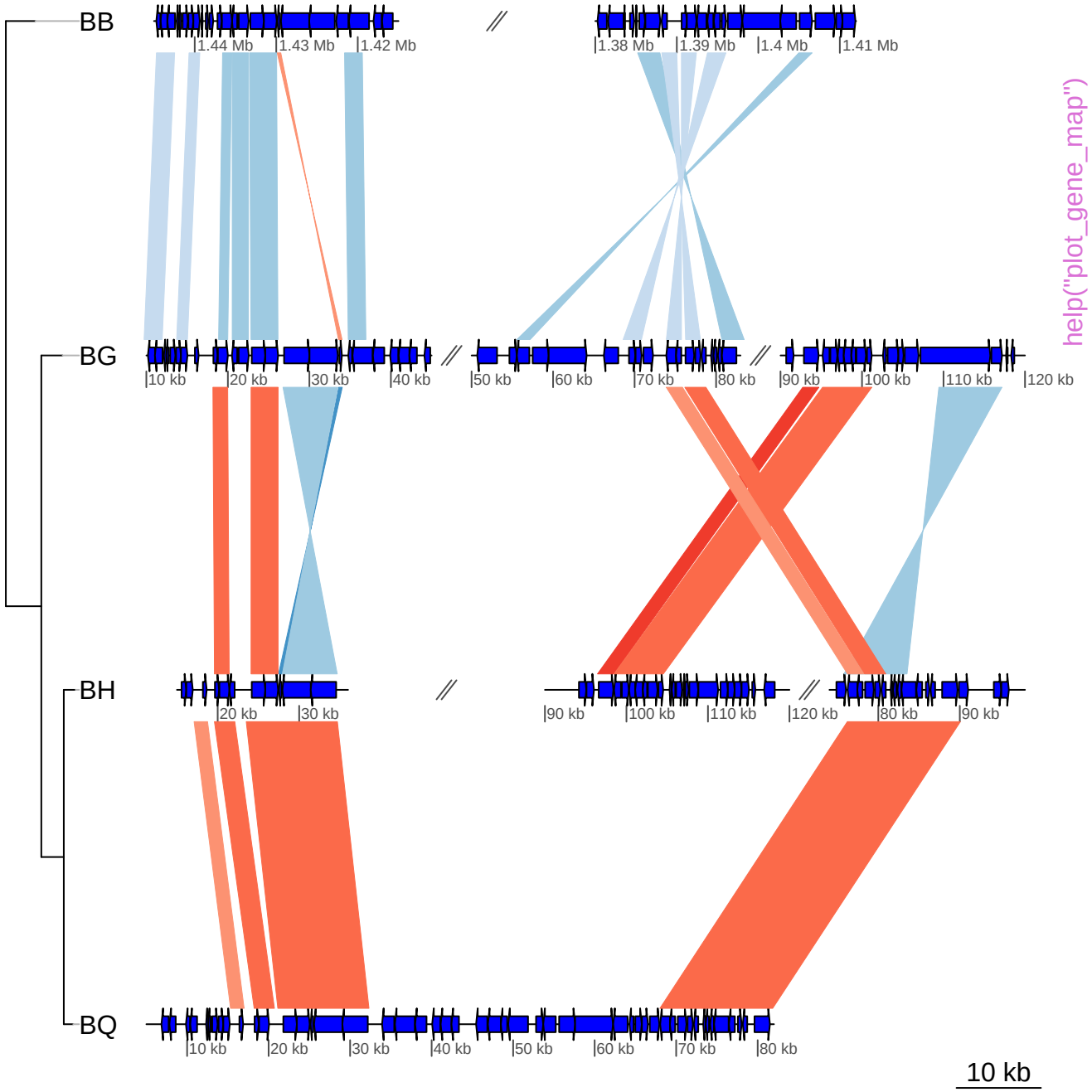
help("plot_gene_map")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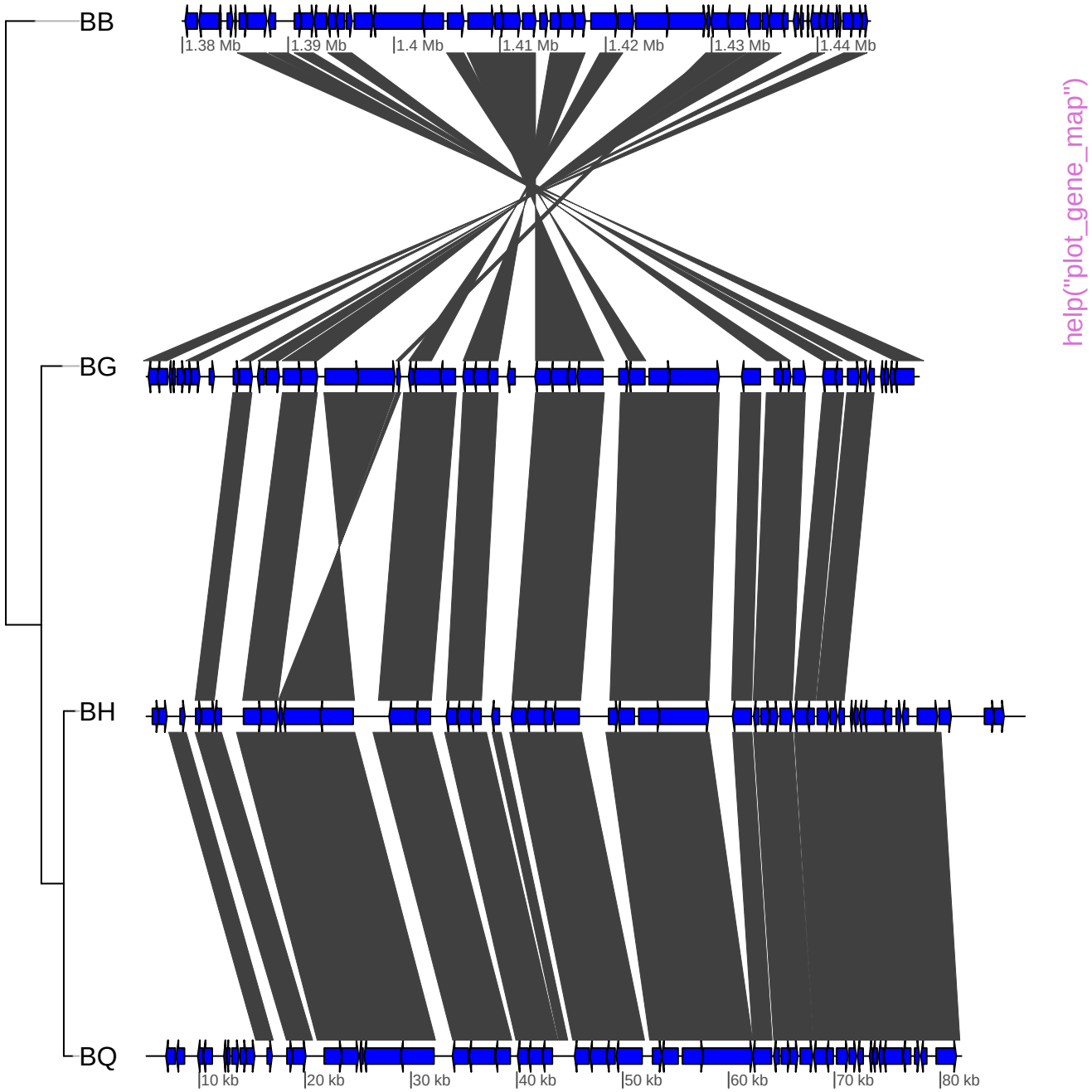
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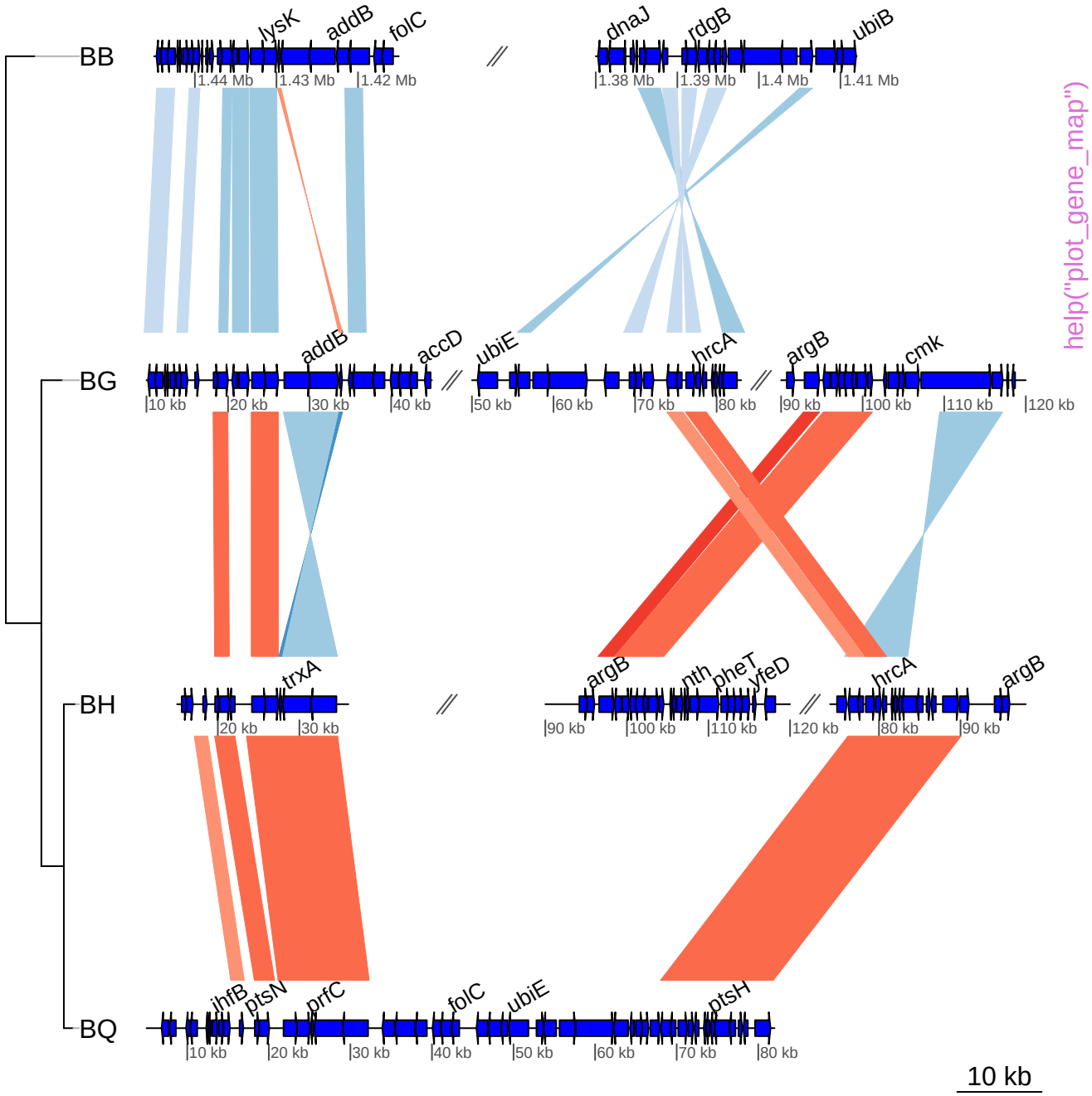
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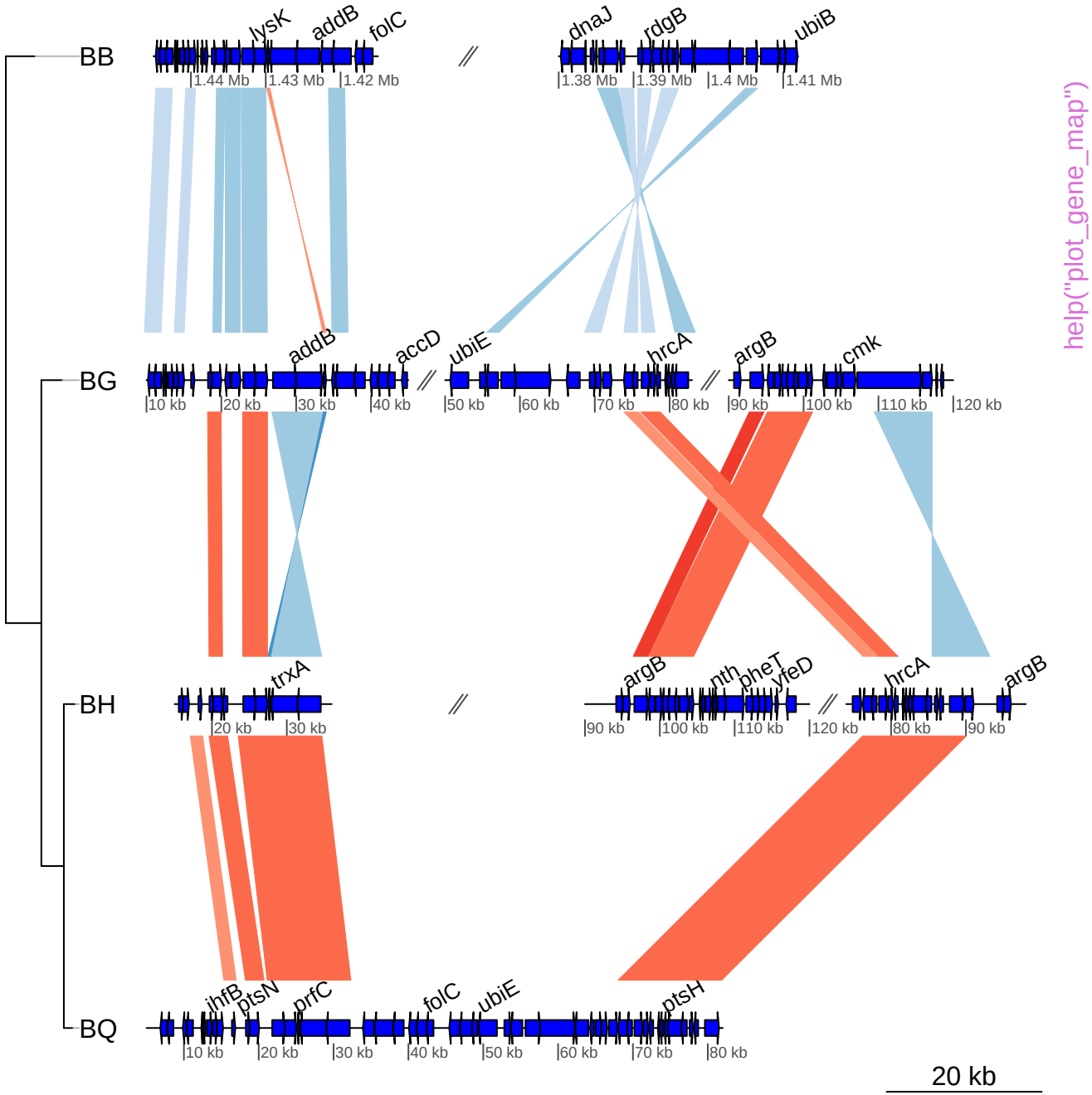
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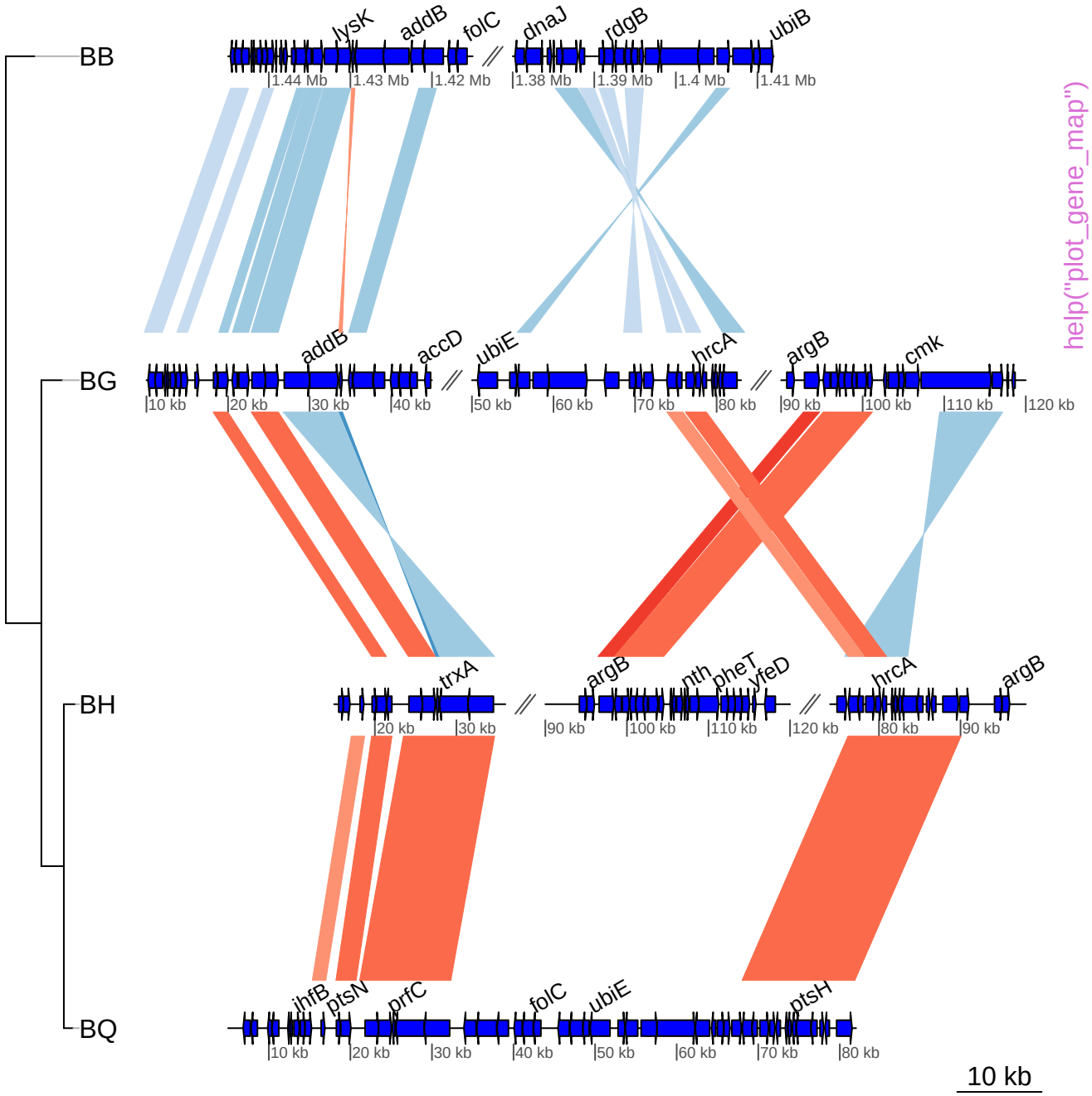
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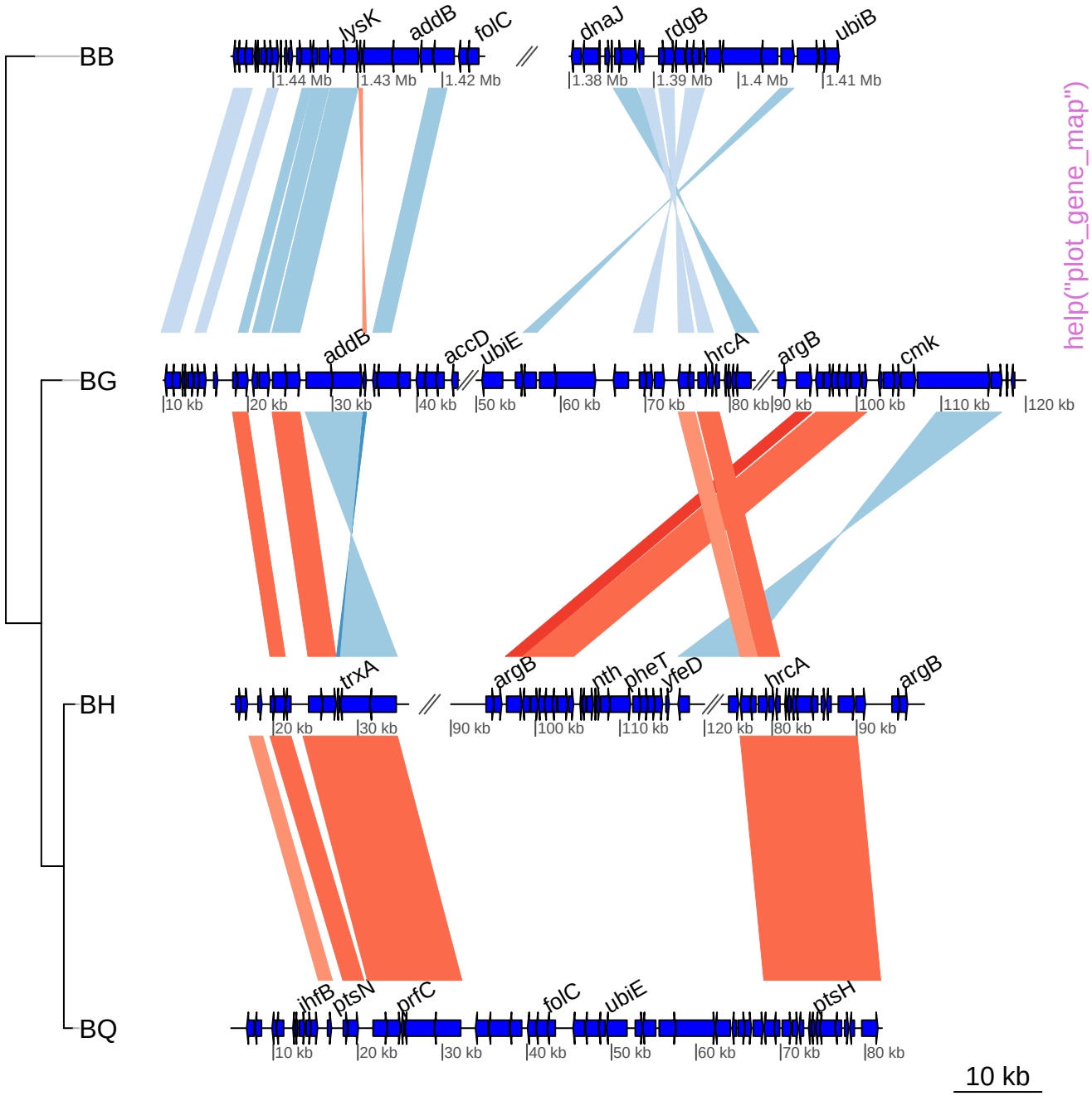
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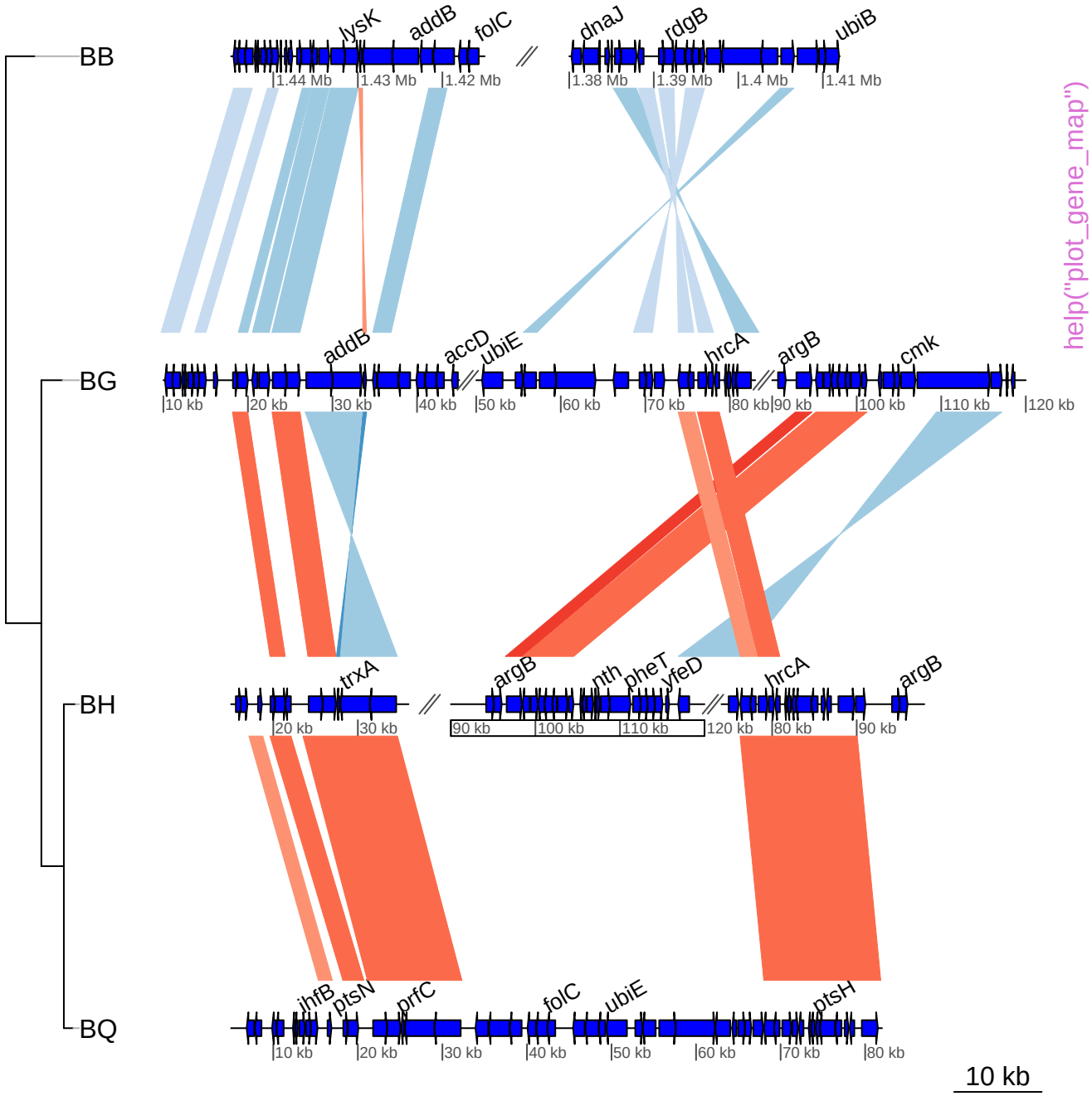
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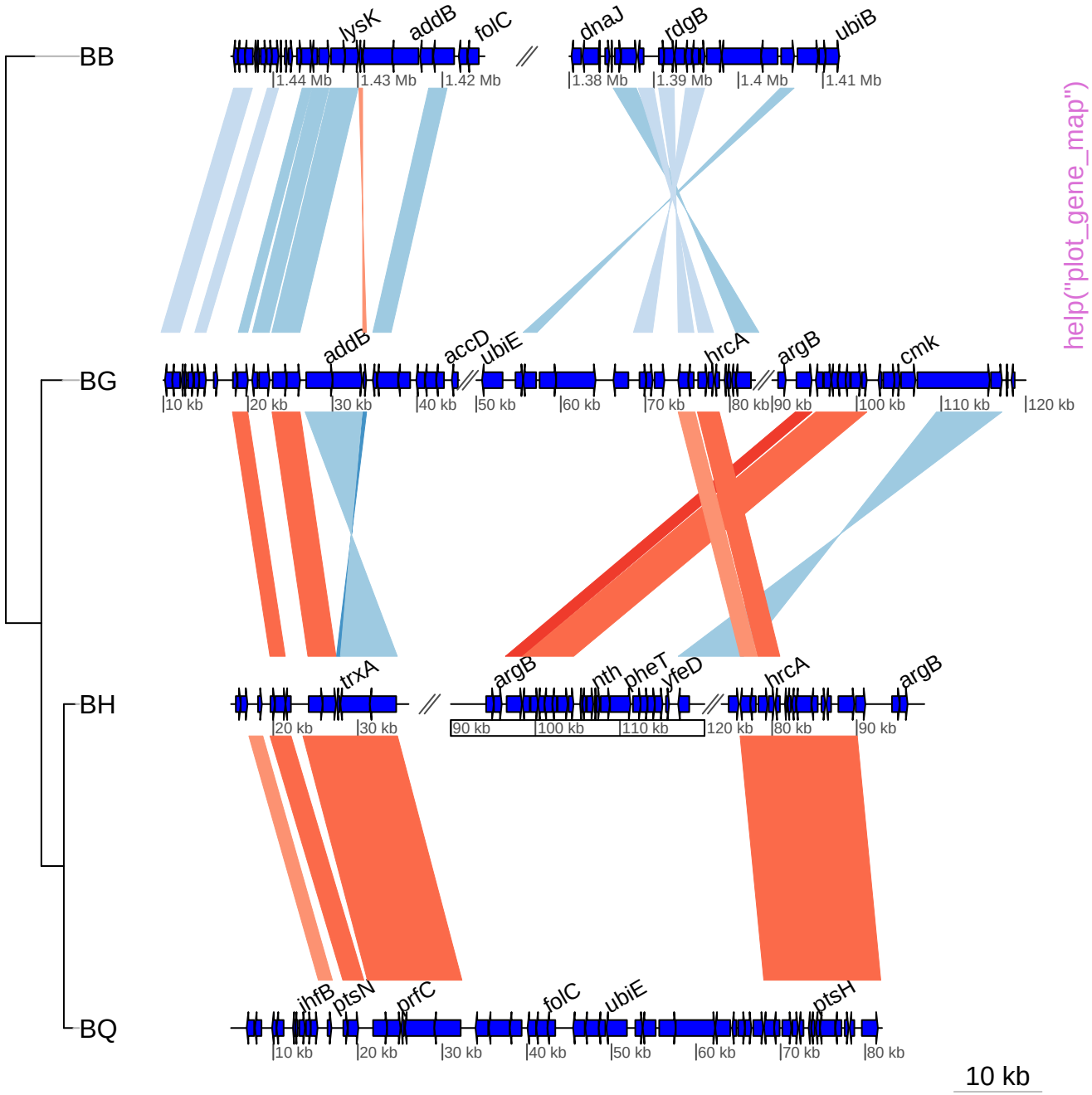
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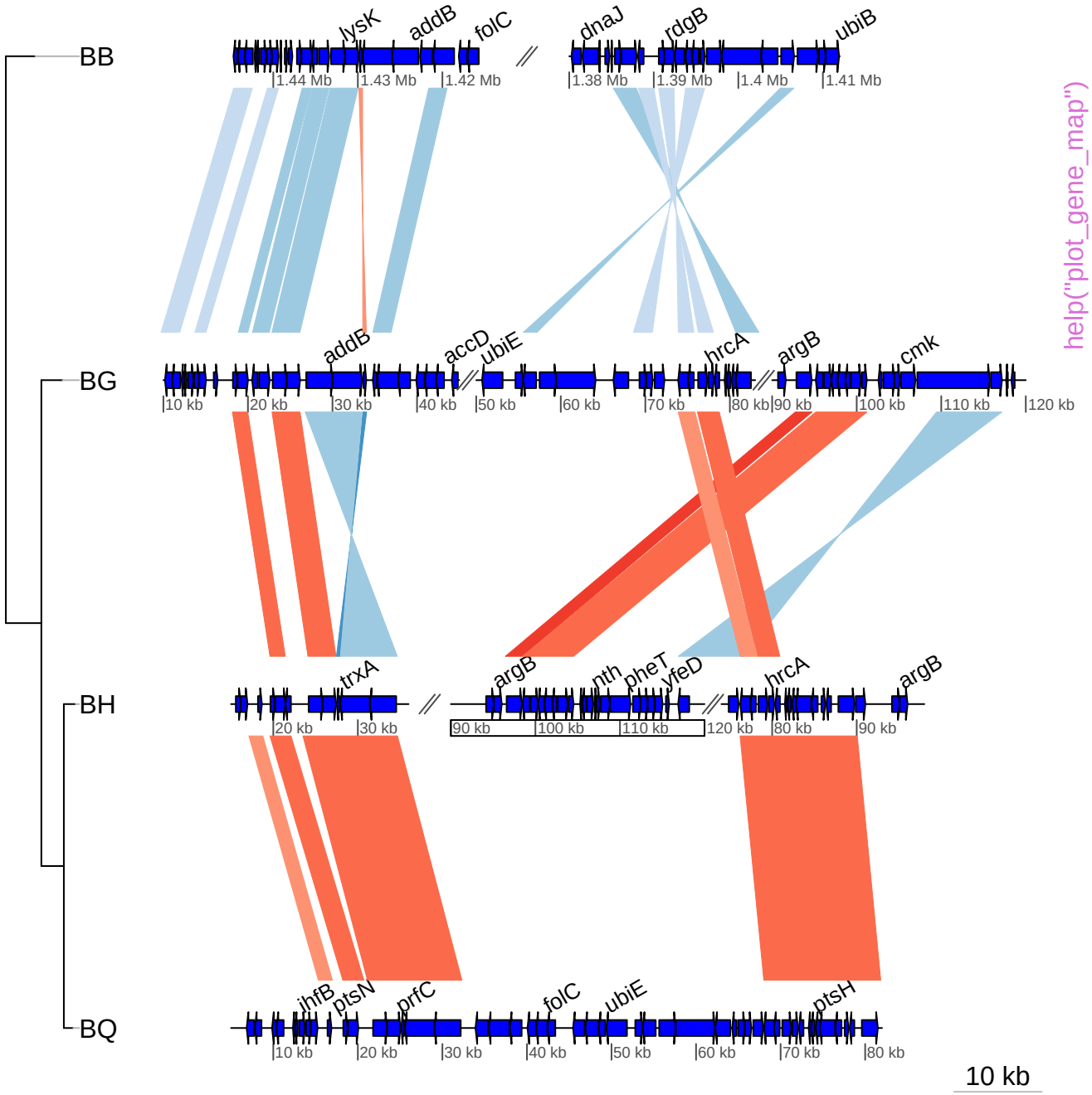
help("plot_gene_map")


BB
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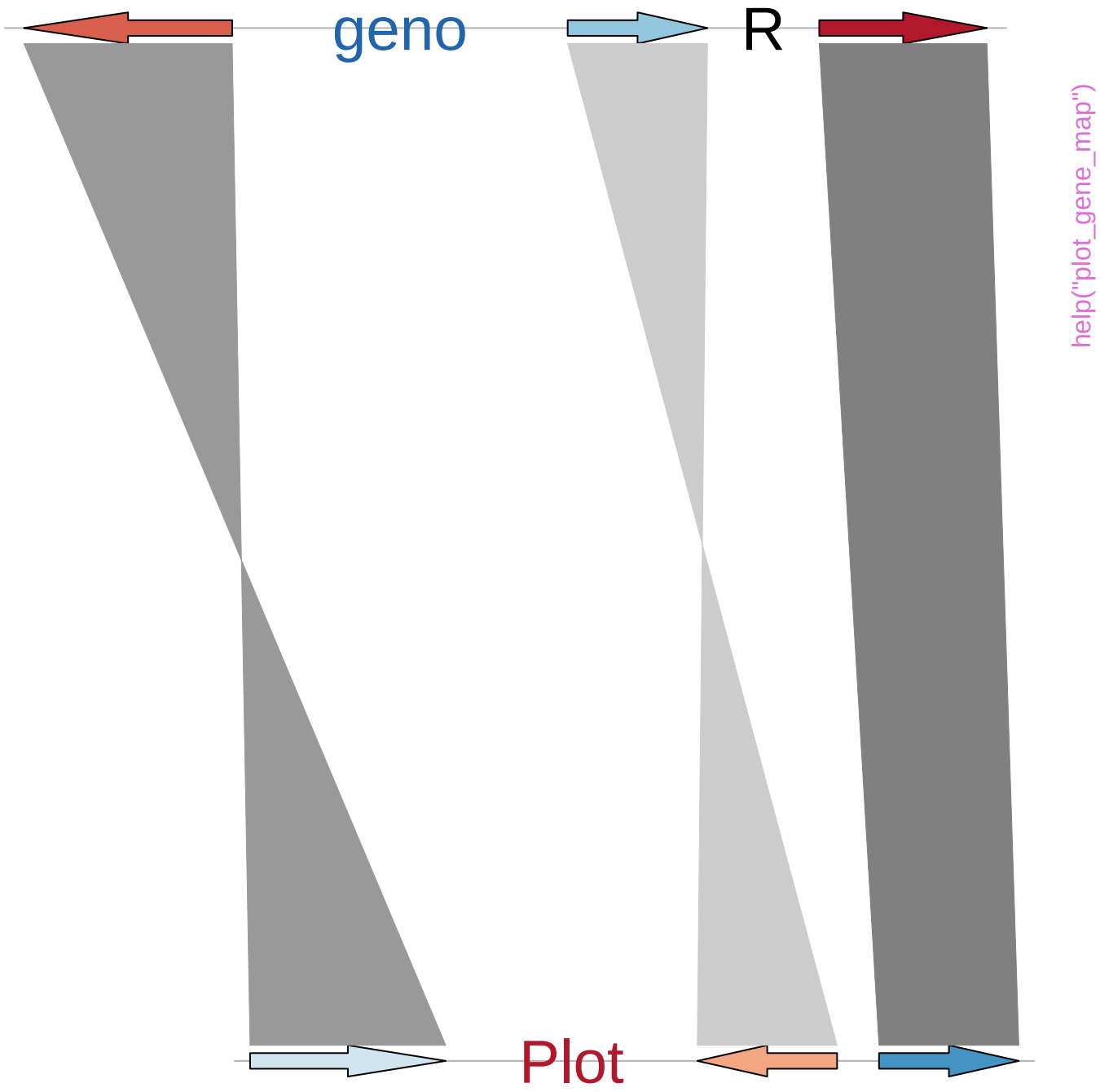
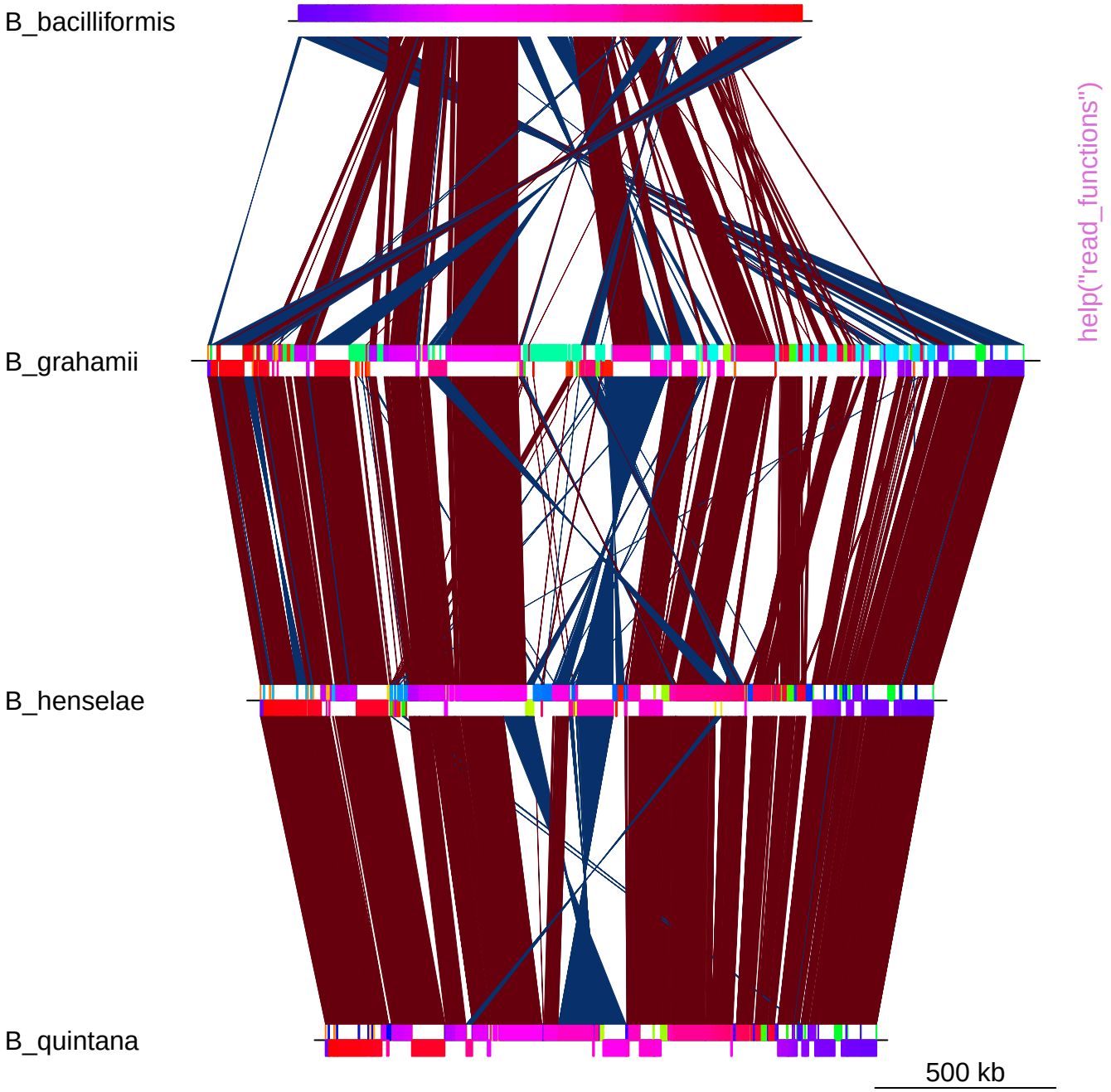
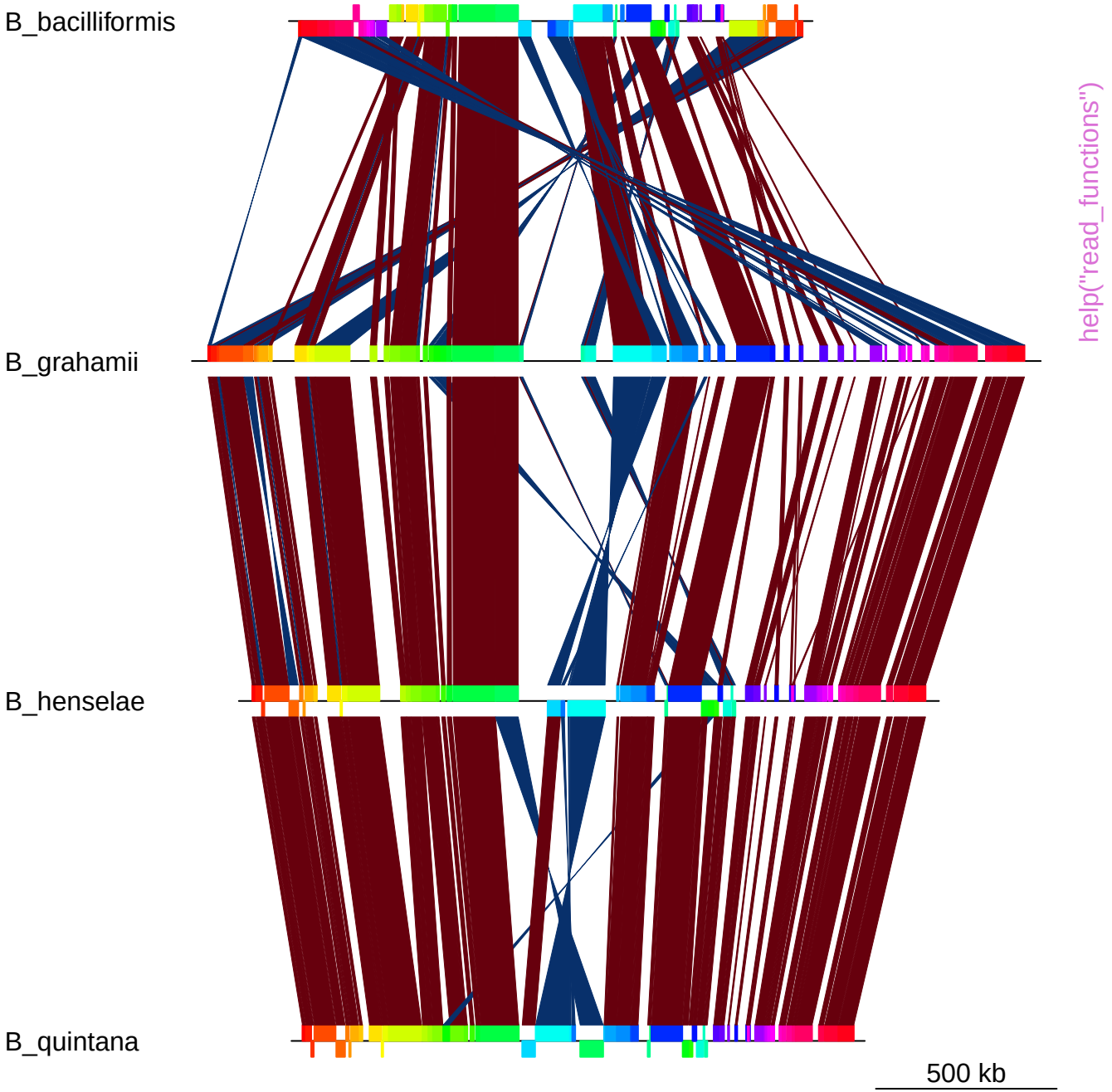
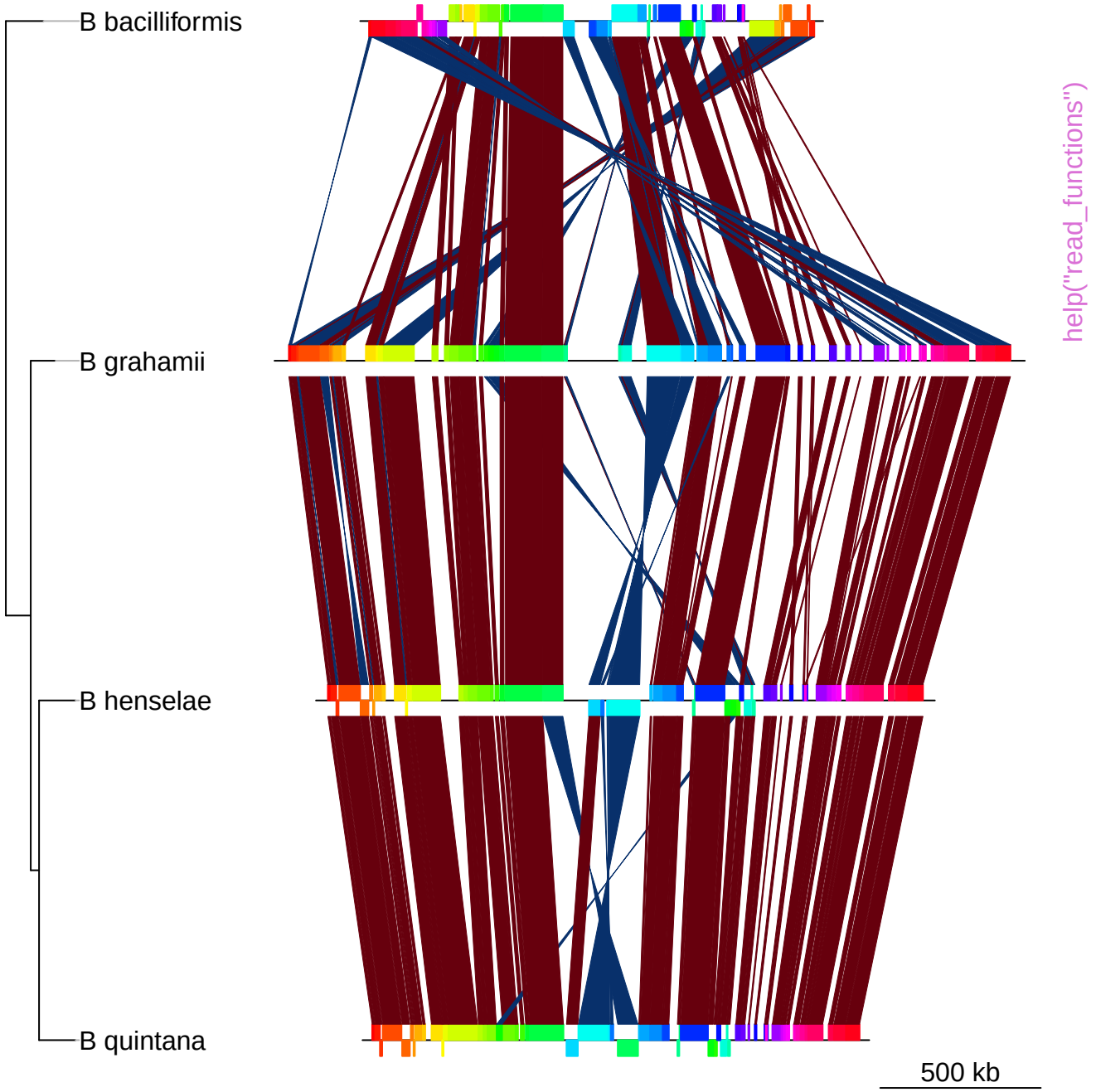
120 kb
BH
trxA
argB
nth
pheT
yfeD
hrcA
argB
20 kb
30 kb
90 kb
100 kb
110 kb
120 kb
80 kb
90 kb
BQ
ihfB
ptsN
prfC
folC
ubiE
ptsH
10 kb
20 kb
30 kb
40 kb
50 kb
60 kb
70 kb
80 kb
10 kb
help("plot_gene_map")


geno
R
Plot
help("plot_gene_map")


B_bacilliformis
B_grahamii
B_henselae
B_quintana
500 kb
help("read_functions")

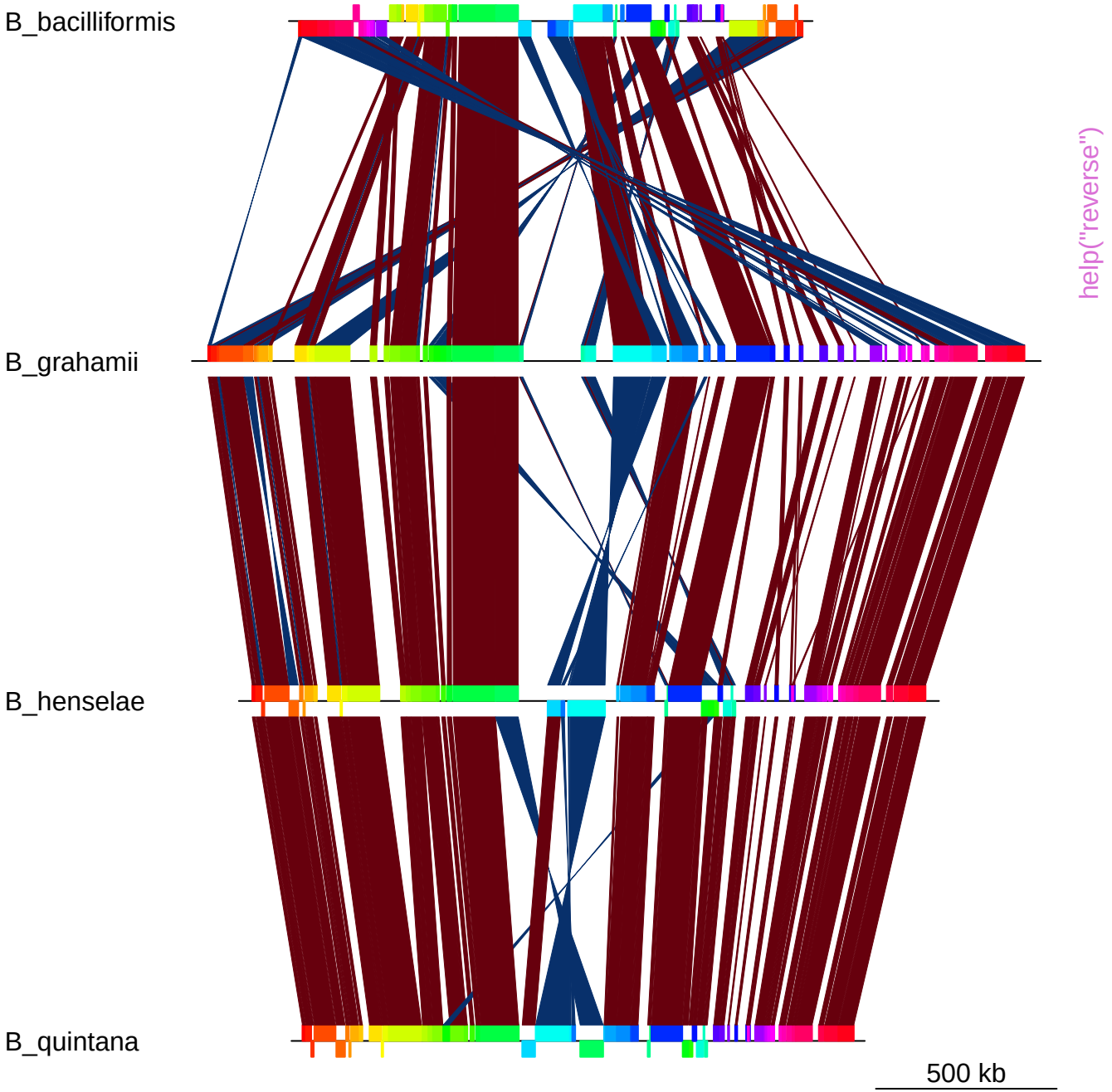

B_bacilliformis
B_grahamii
B_henselae
B_quintana
500 kb
help("read_functions")

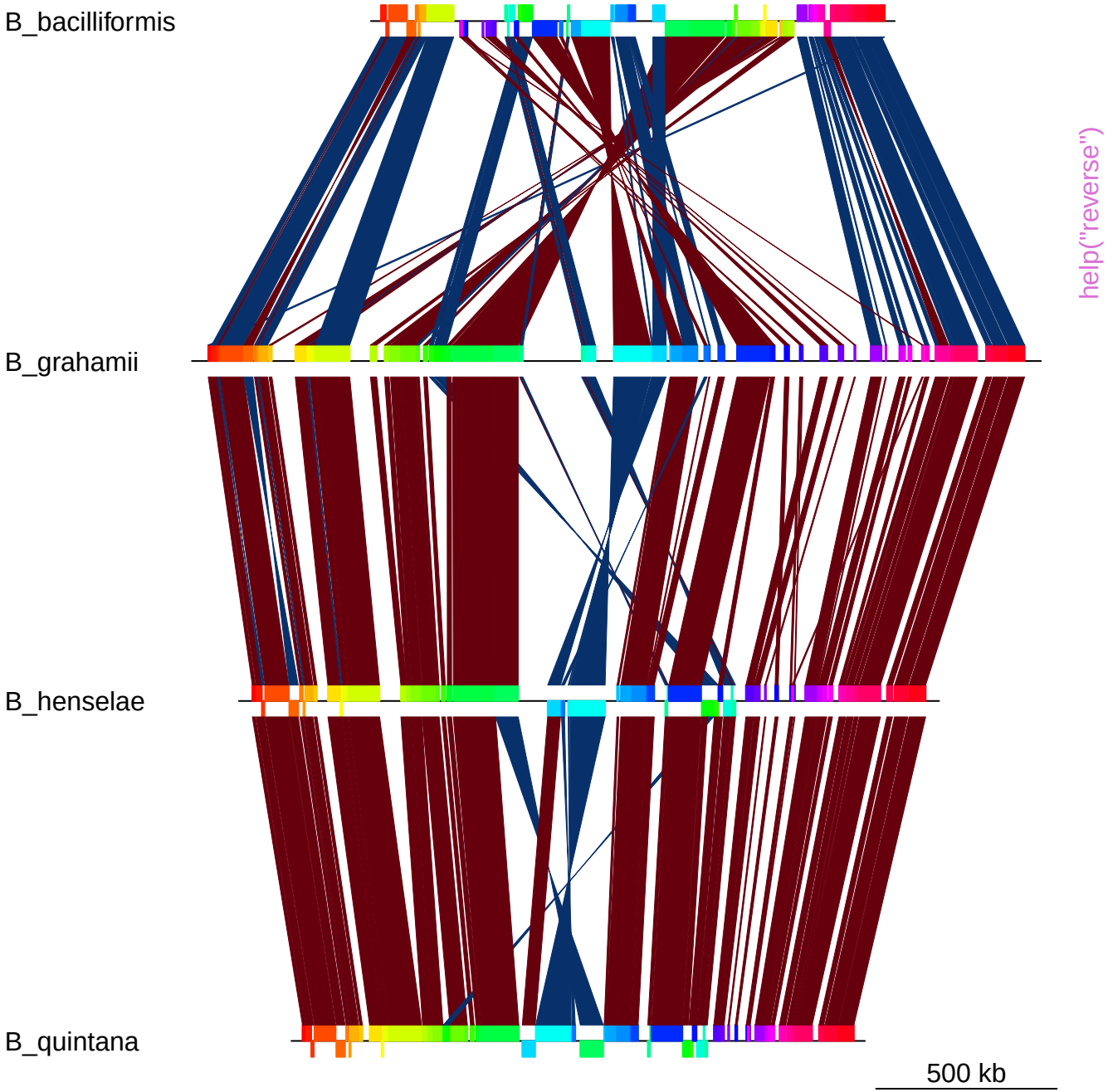

B bacilliformis
B grahamii
B henselae
B quintana
500 kb
help("read_functions")

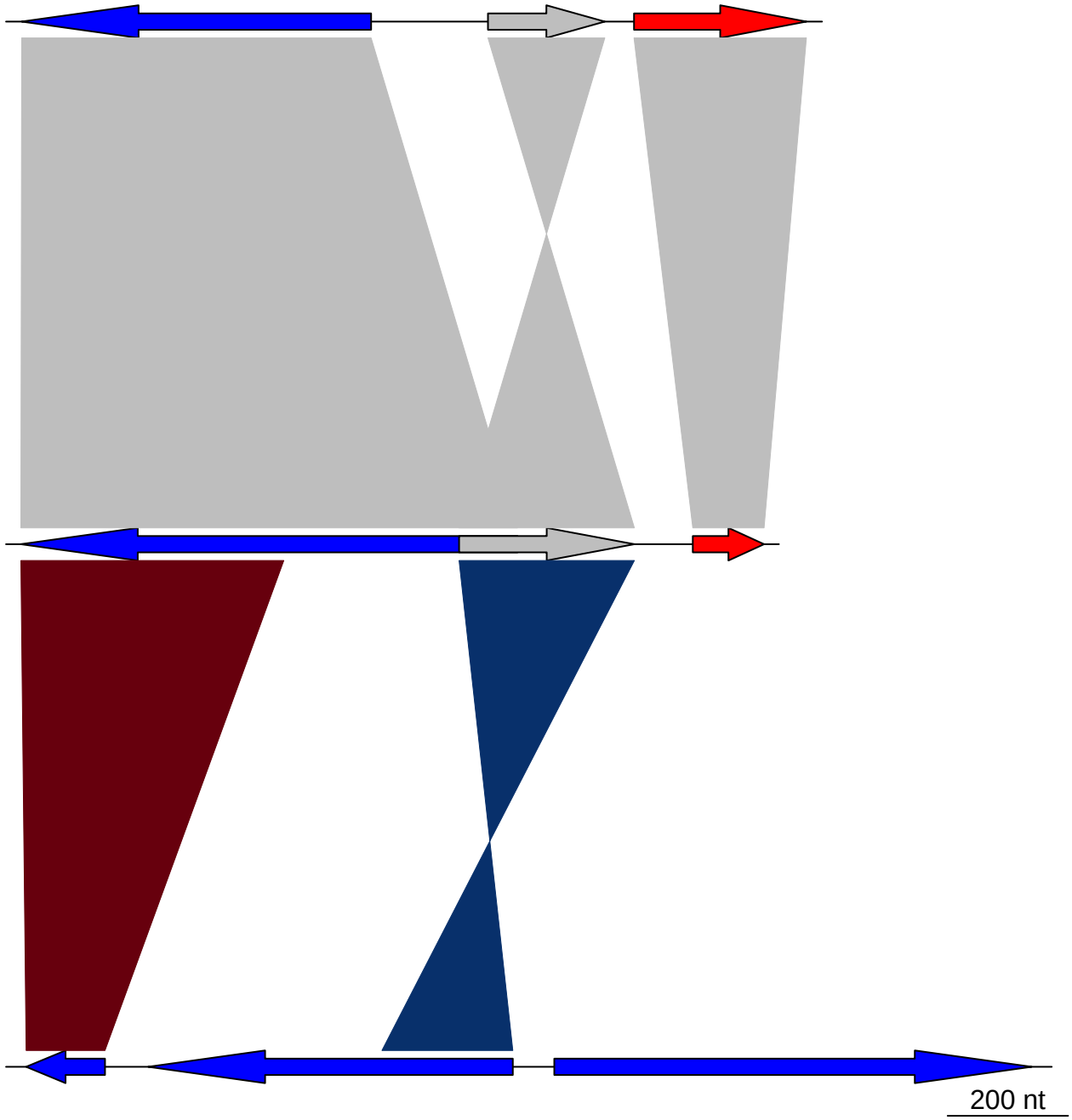
help("read_functions")

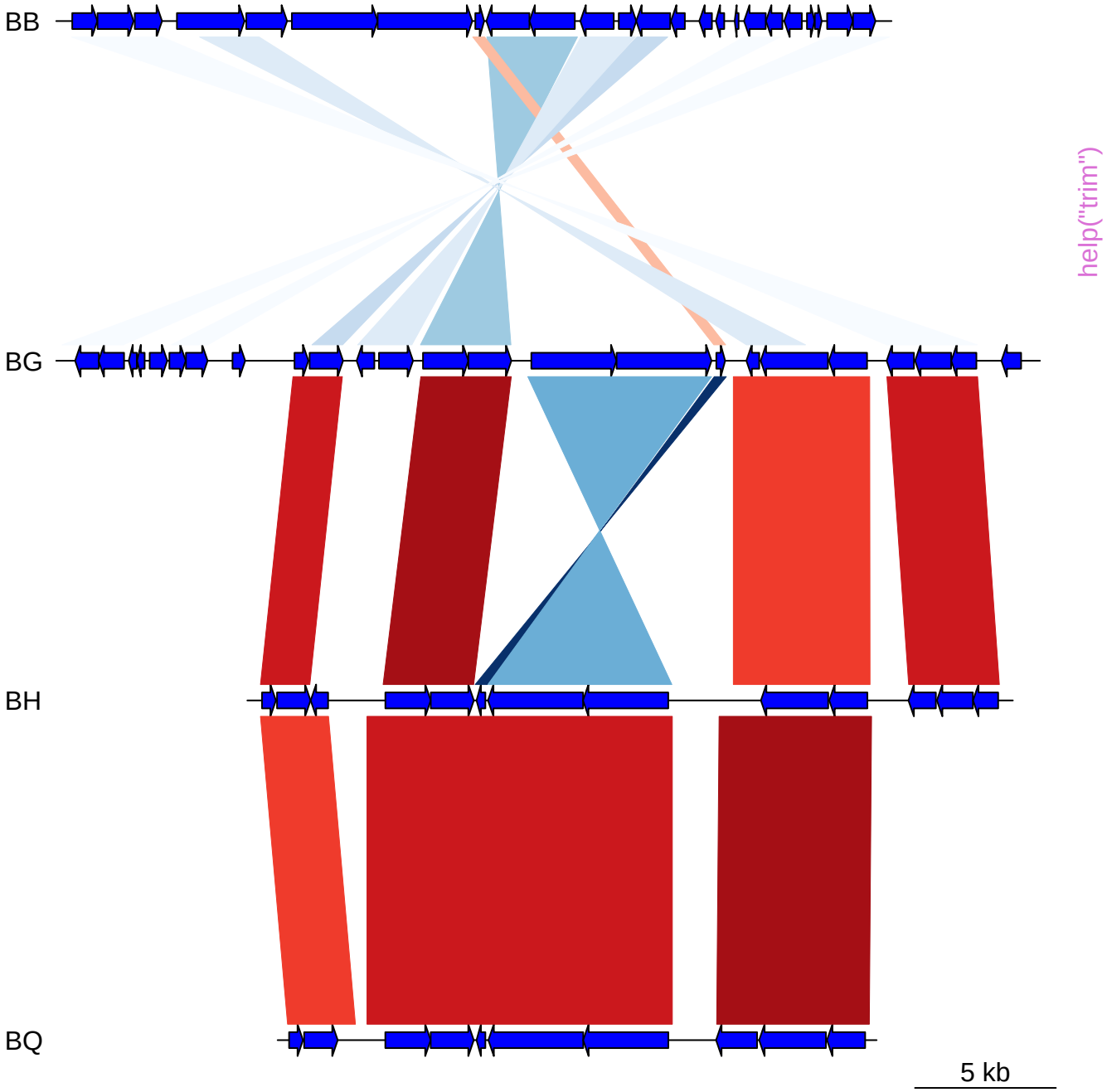
BQ

5 kb


B_bacilliformis
B_grahamii
B_henselae
B_quintana
500 kb
help("reverse")


B_bacilliformis
B_grahamii
B_henselae
B_quintana
help("reverse")
500 kb


help("three_genes")
200 nt


BB
BG
BH
BQ
help("trim")
5 kb