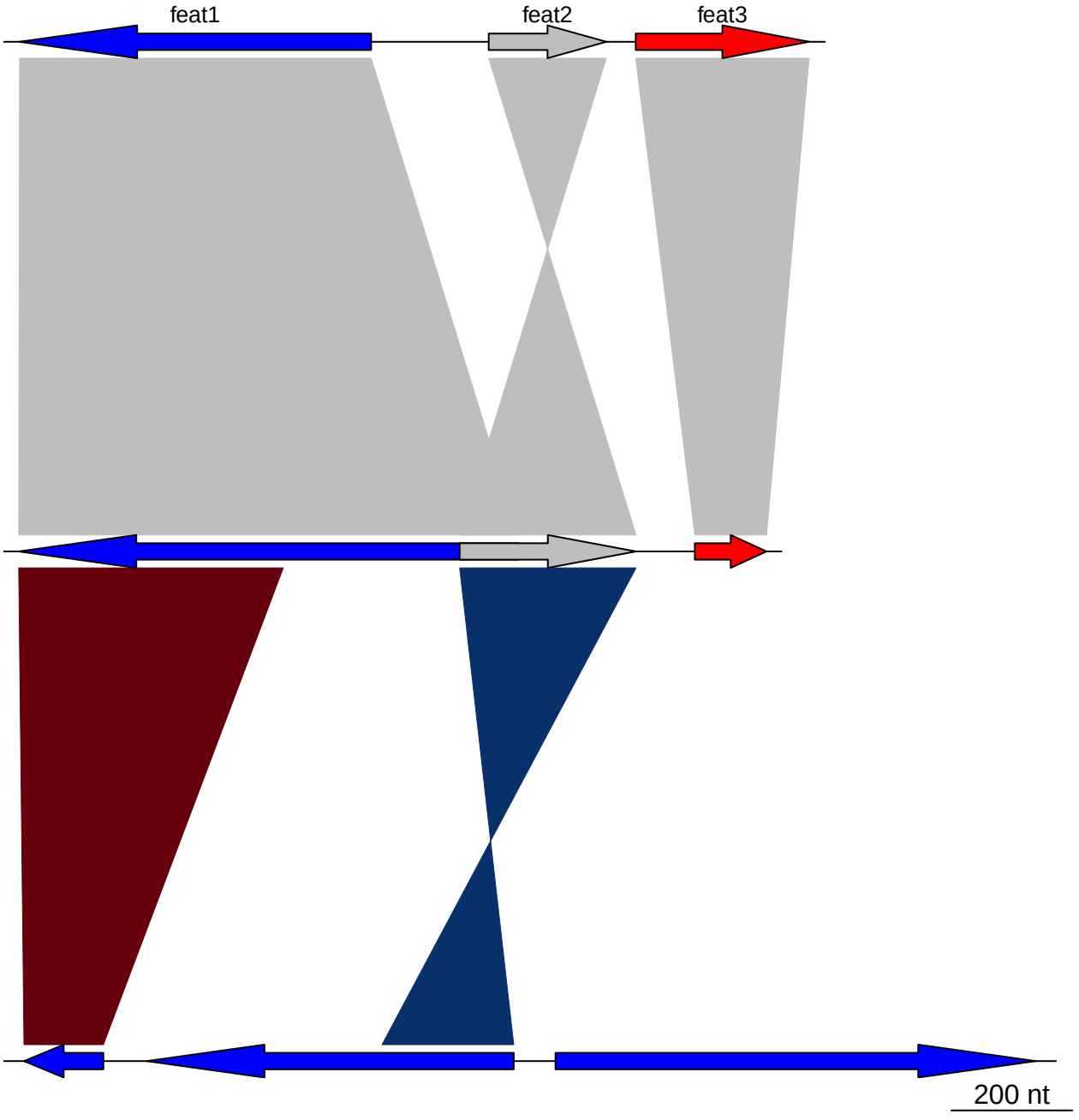
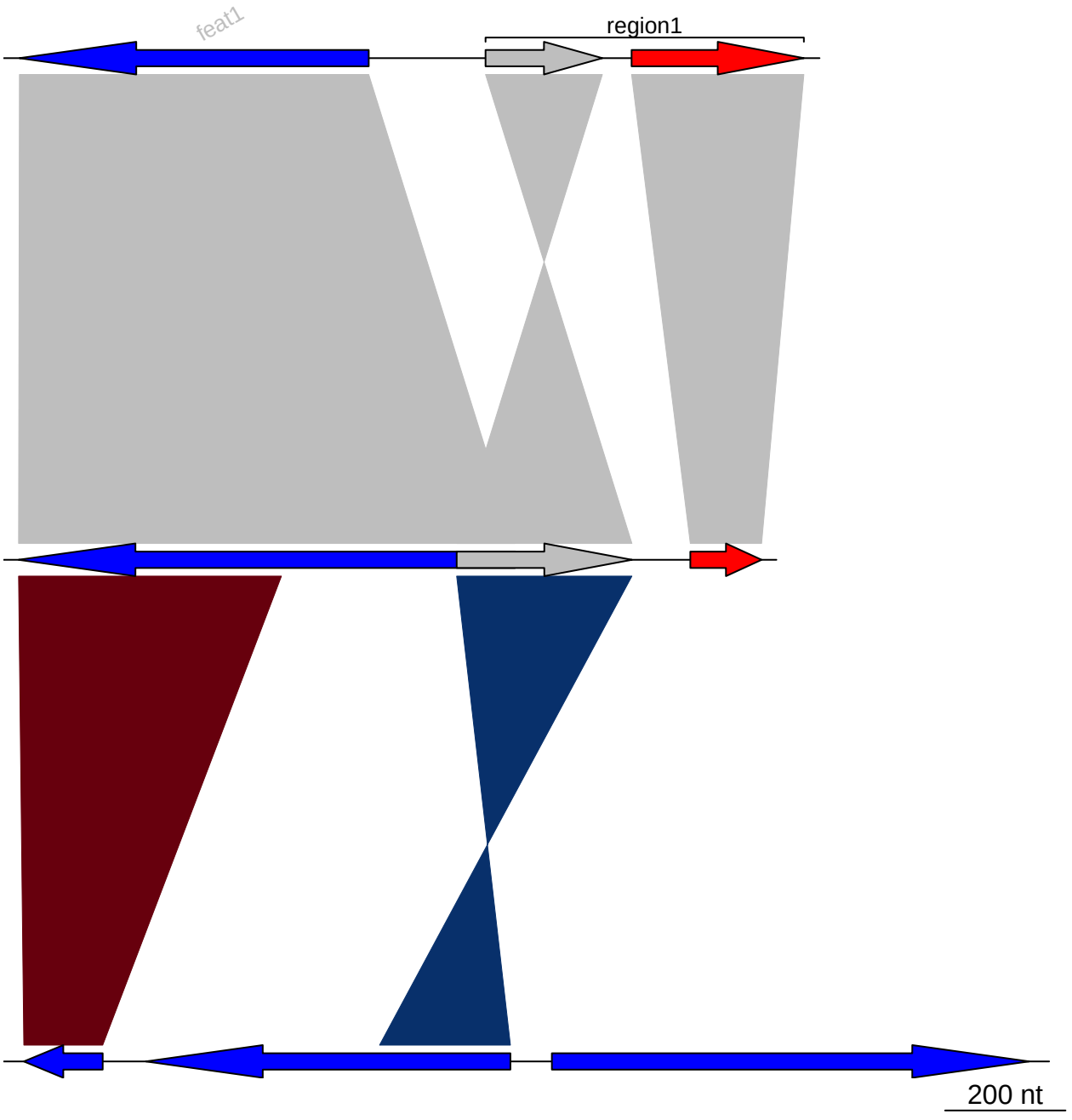
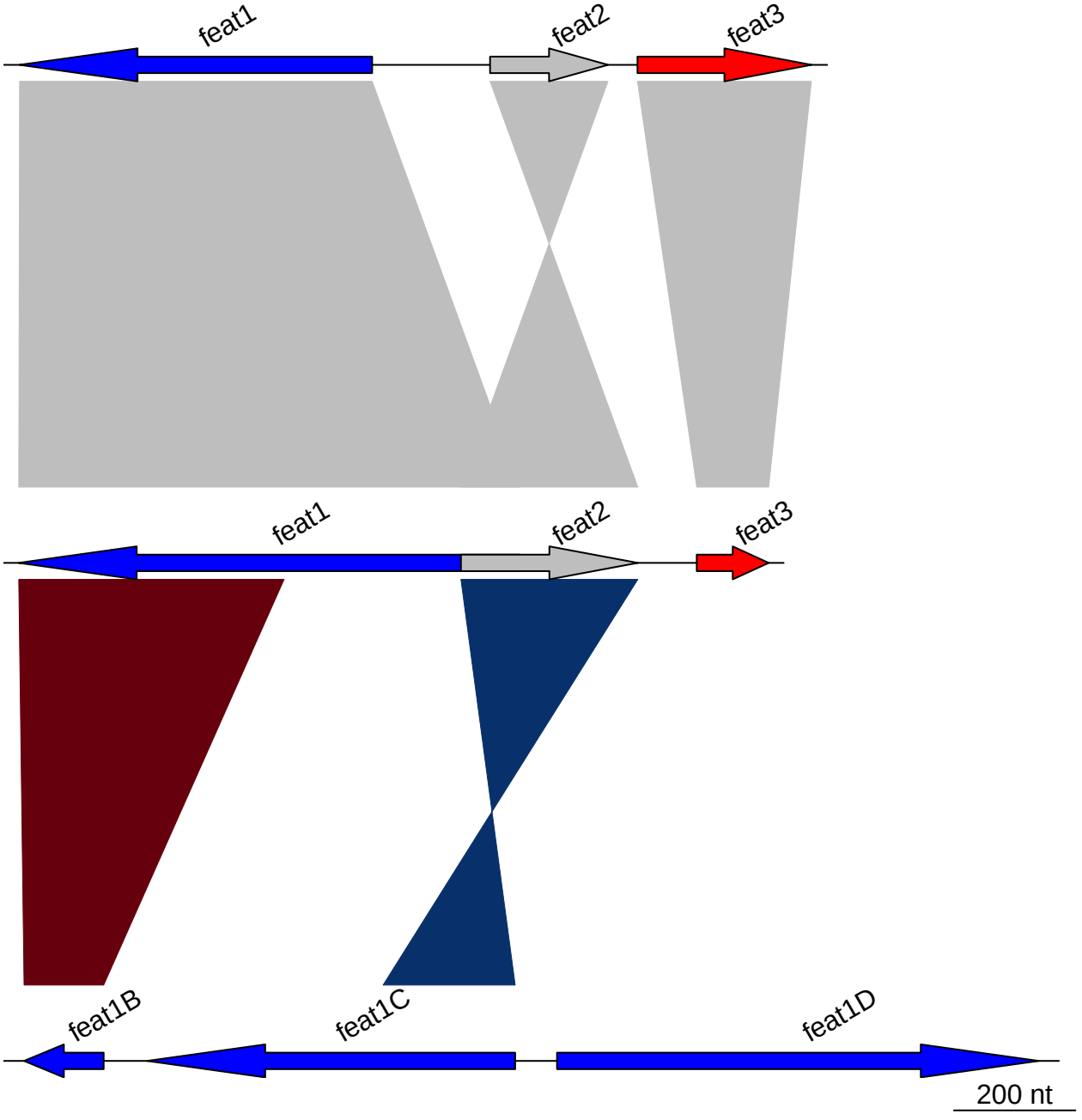


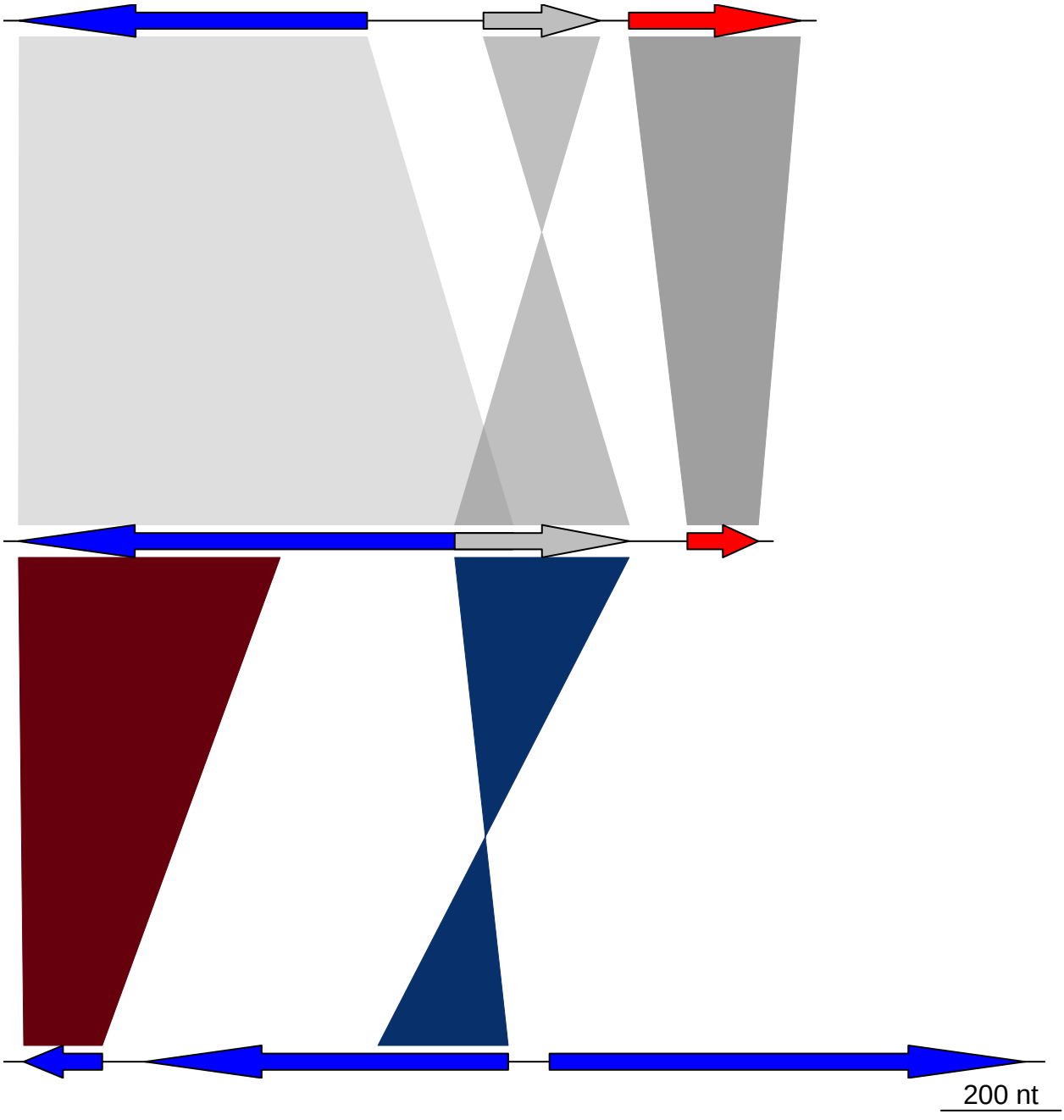
help("annotation")



help("annotation")



help("annotation")


help("apply_color_scheme")
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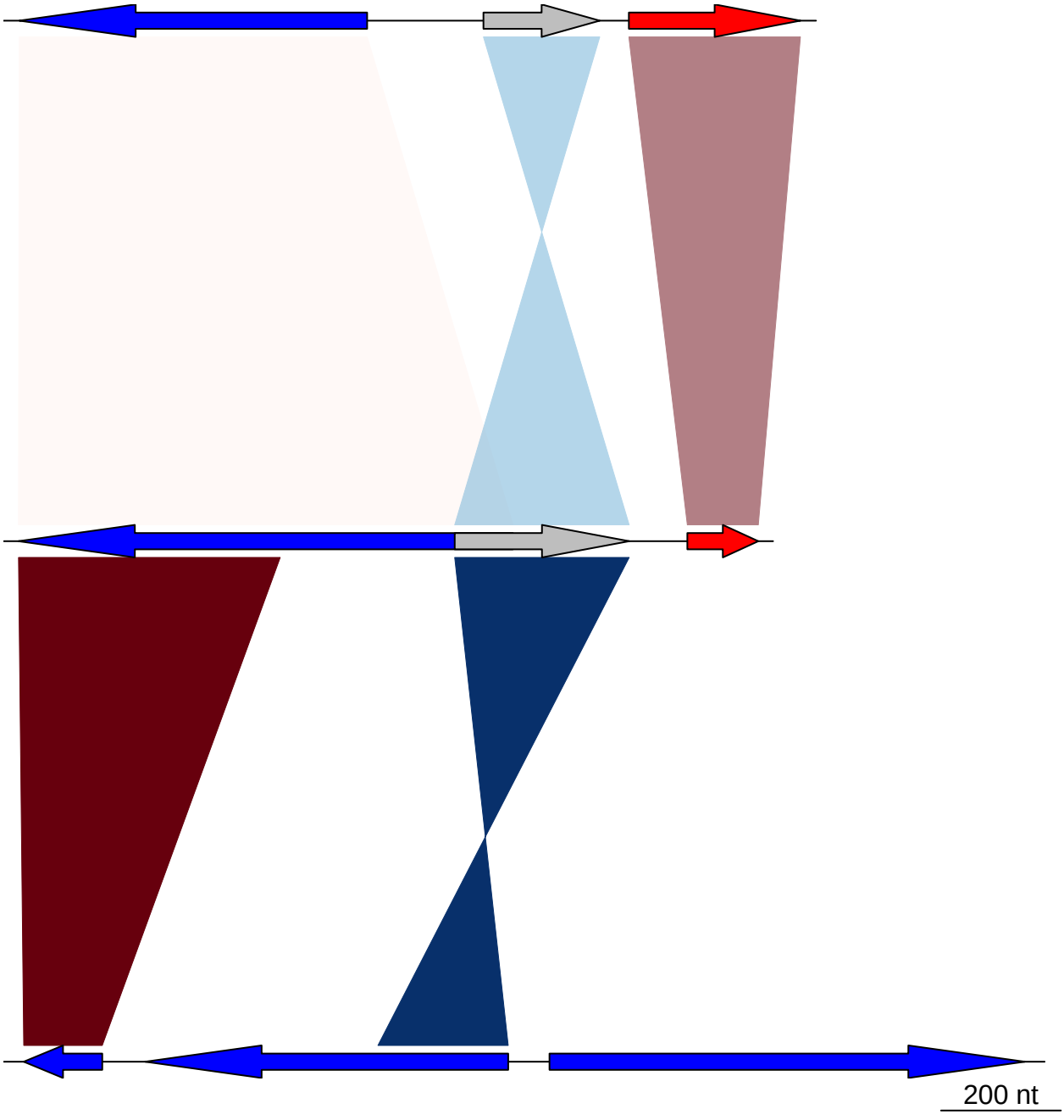
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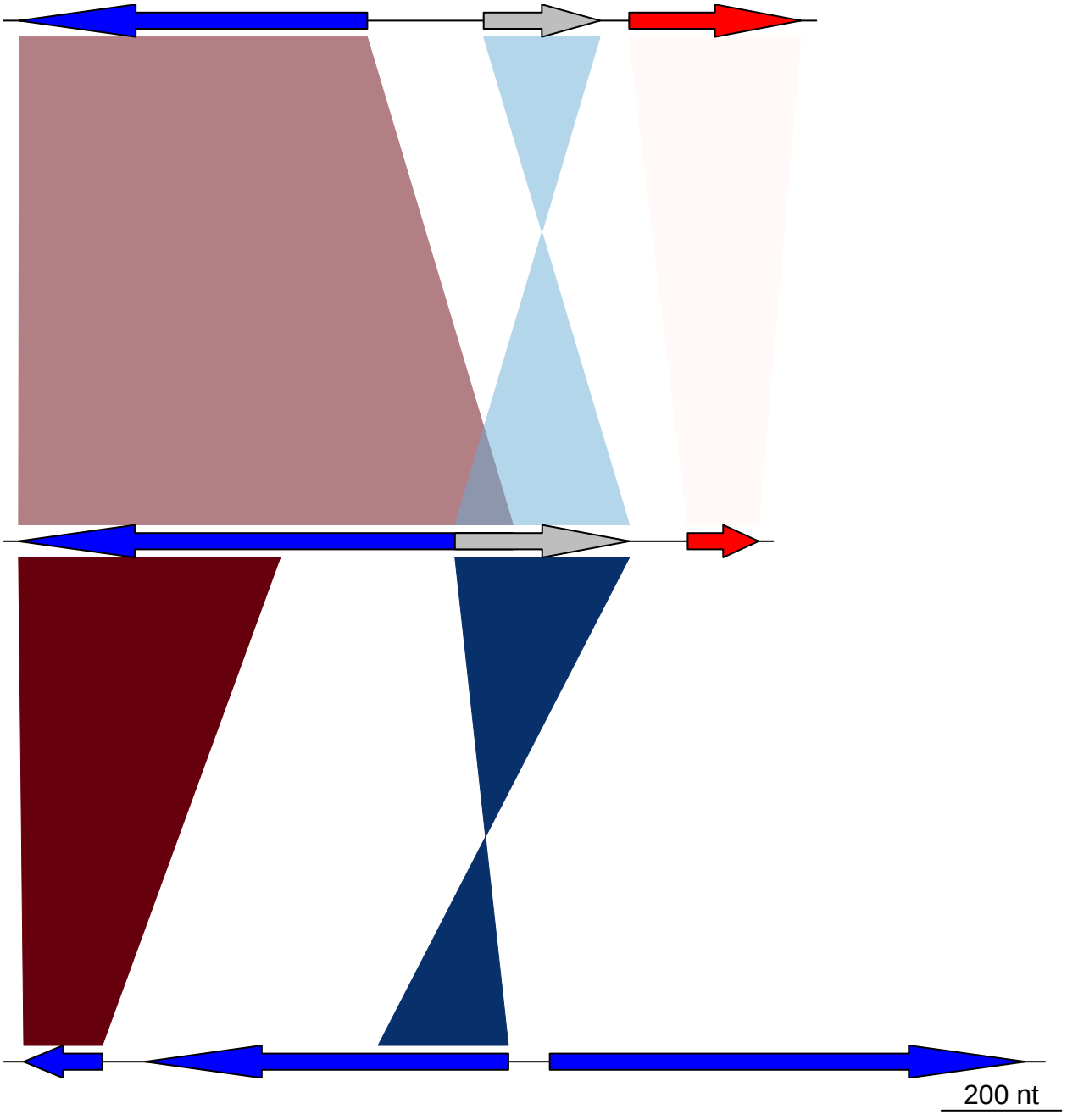
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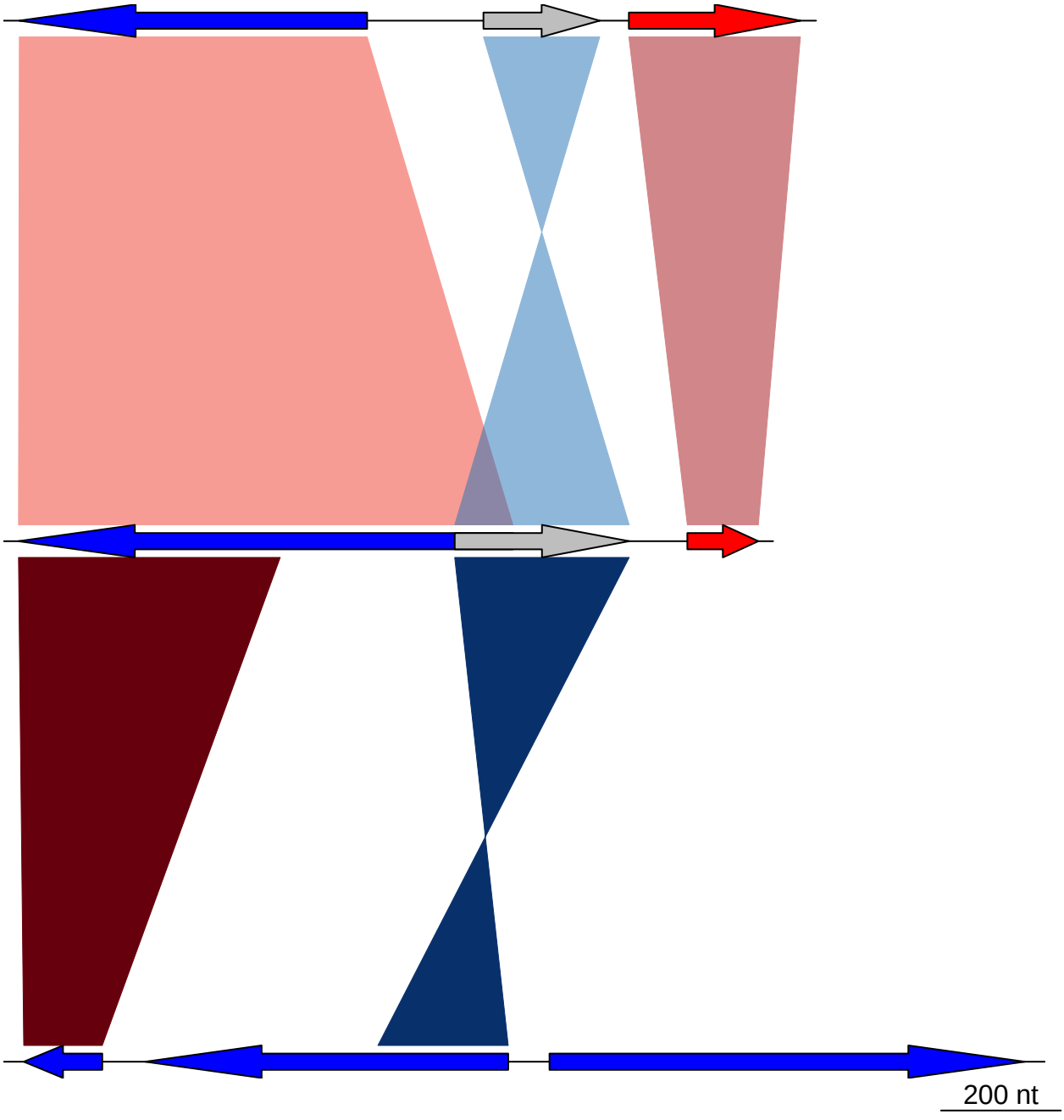
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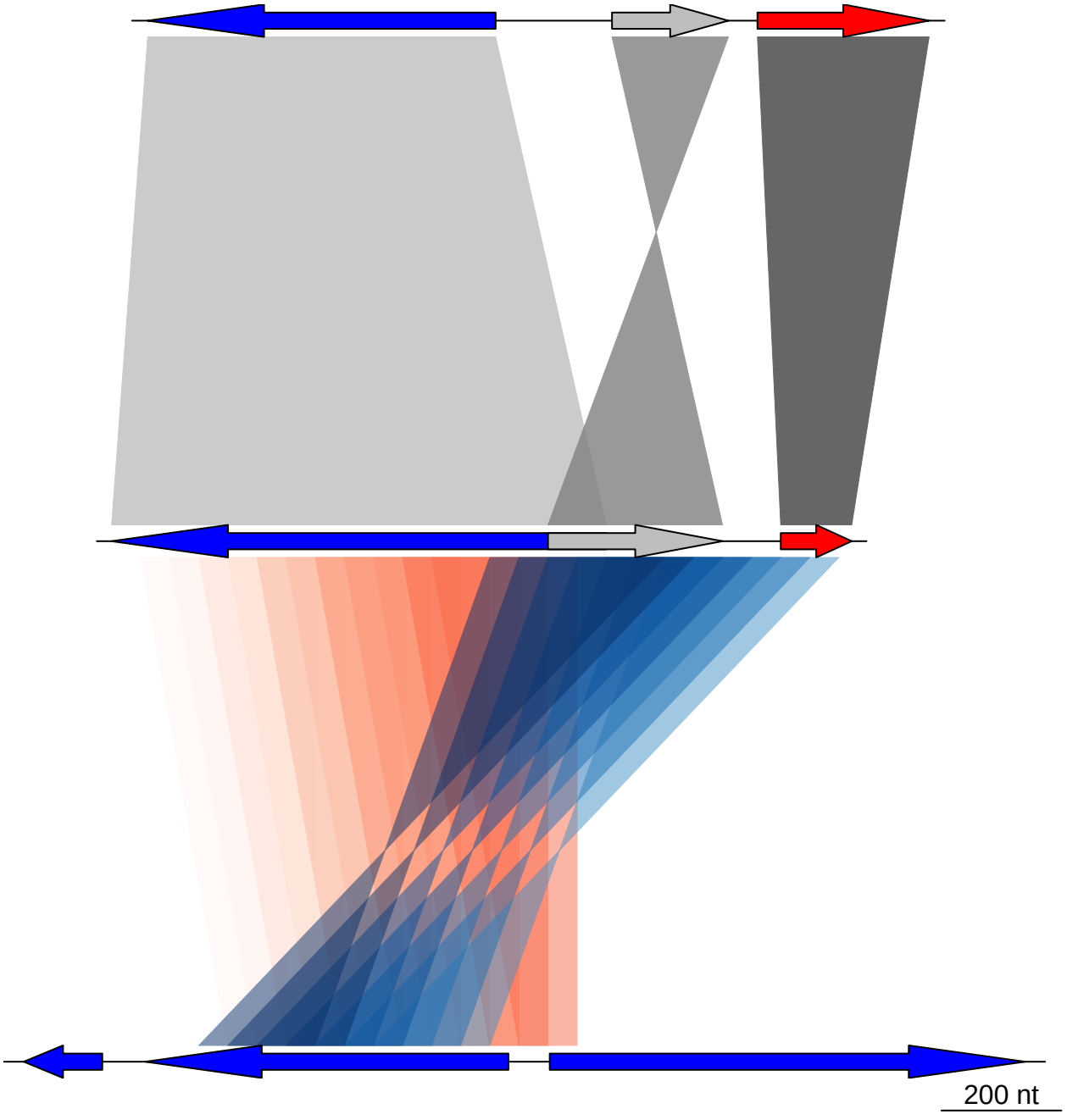
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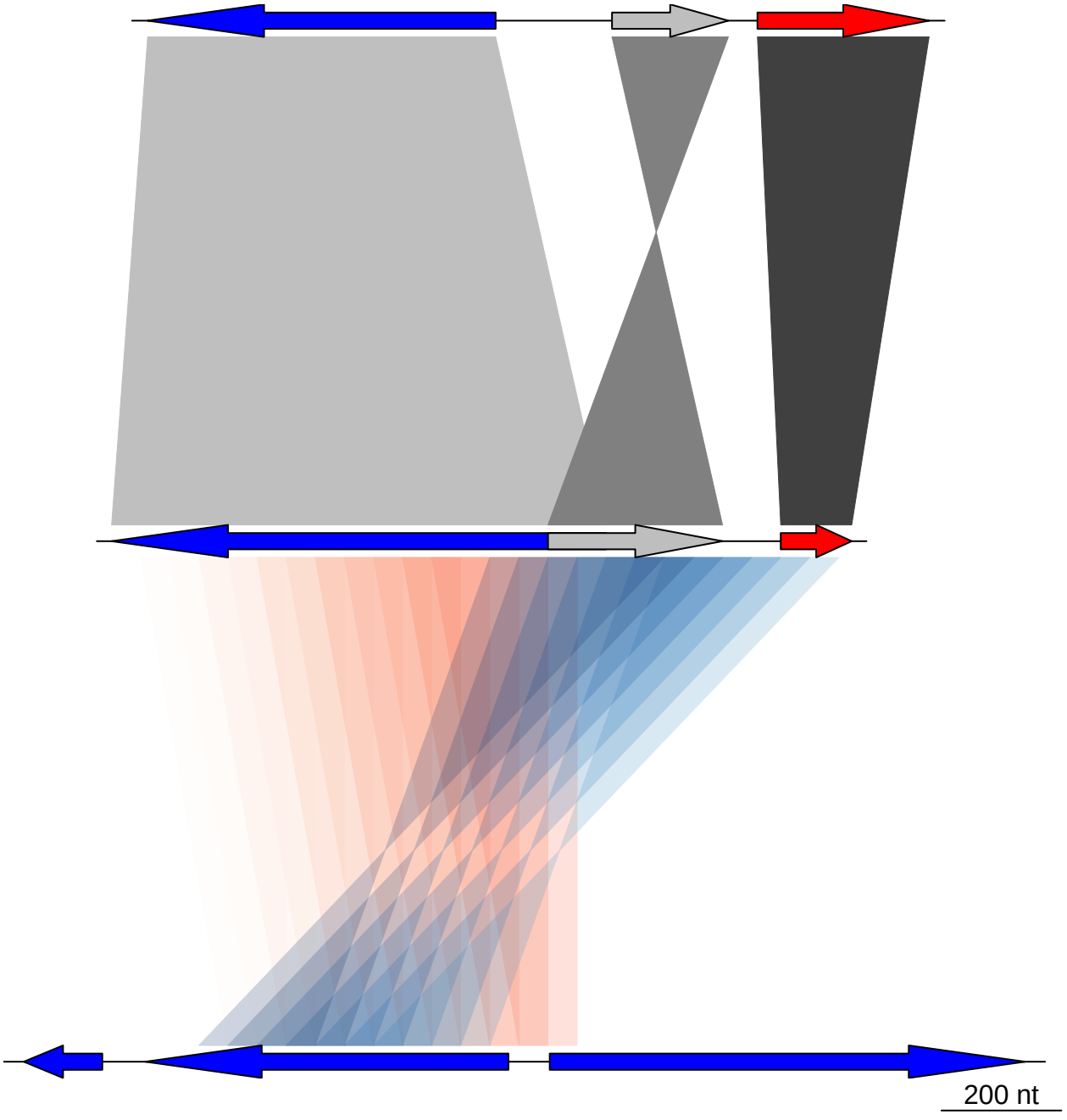
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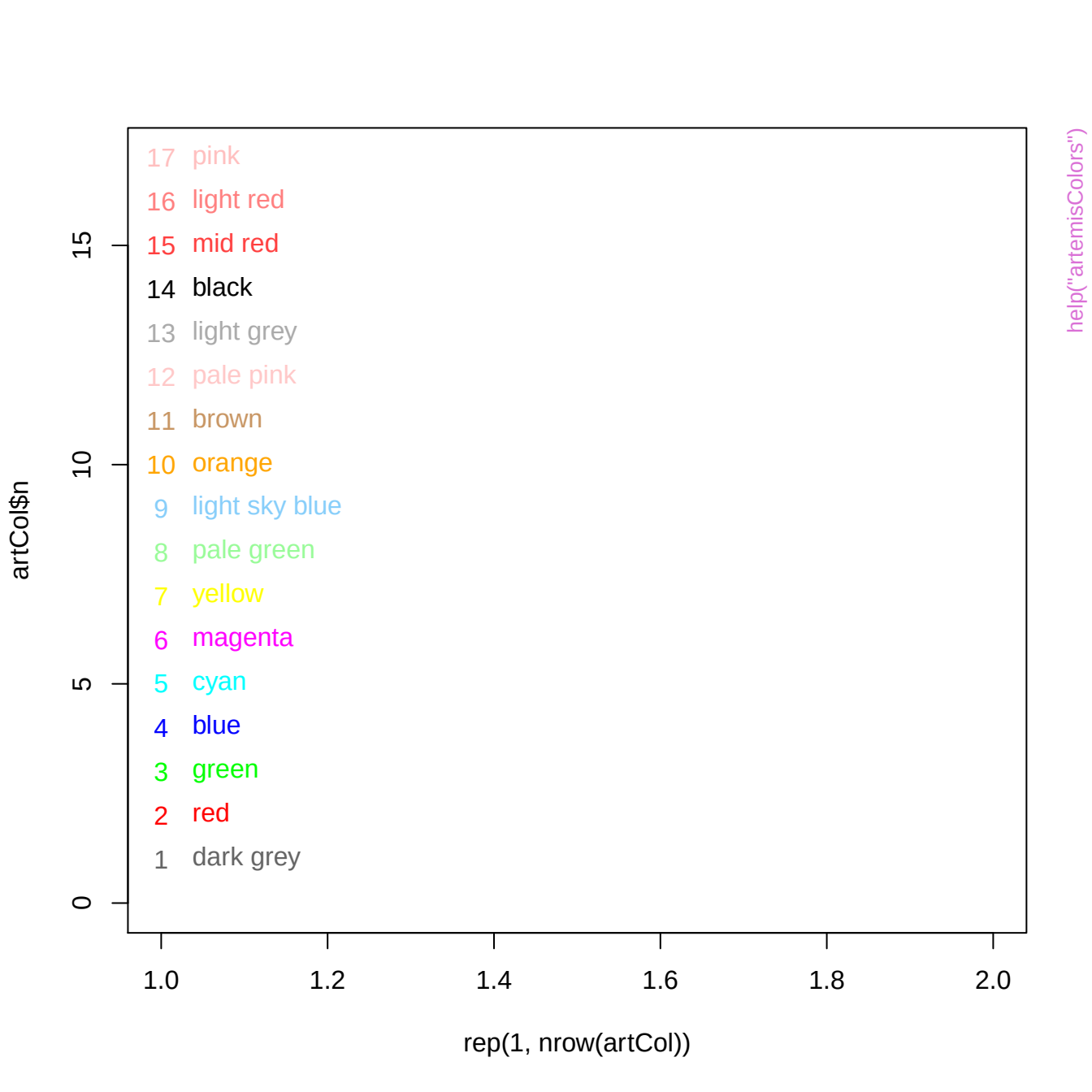
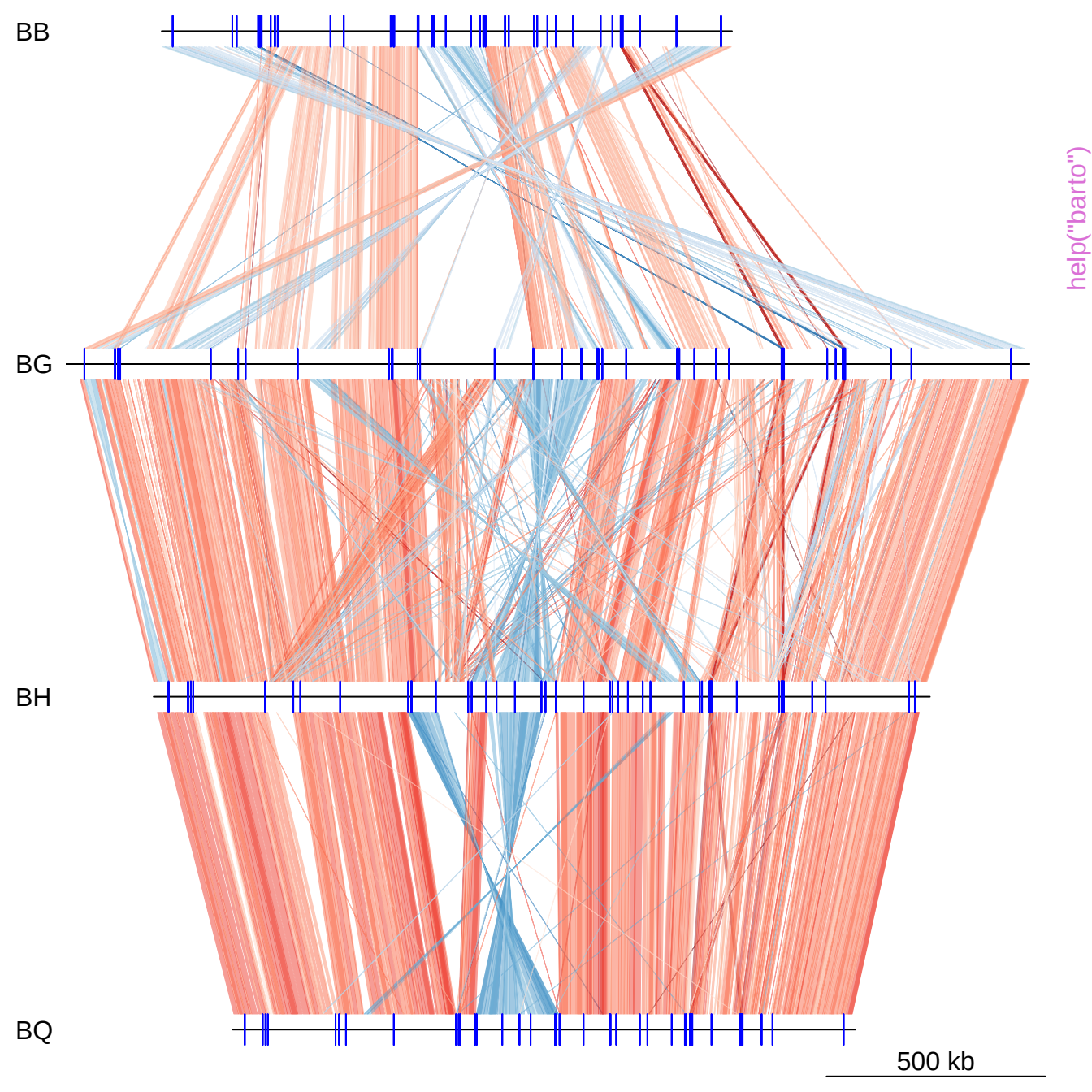

17 pink
16 light red
15 mid red
14 black
13 light grey
12 pale pink
11 brown
10 orange
9 light sky blue
8 pale green
7 yellow
6 magenta
5 cyan
4 blue
3 green
2 red
1 dark grey
0
5
10
15
artCol$n
1.0
1.2
1.4
1.6
1.8
2.0
rep(1, nrow(artCol))
help("artemisColors")


BB
BG
BH
BQ
help("barto")
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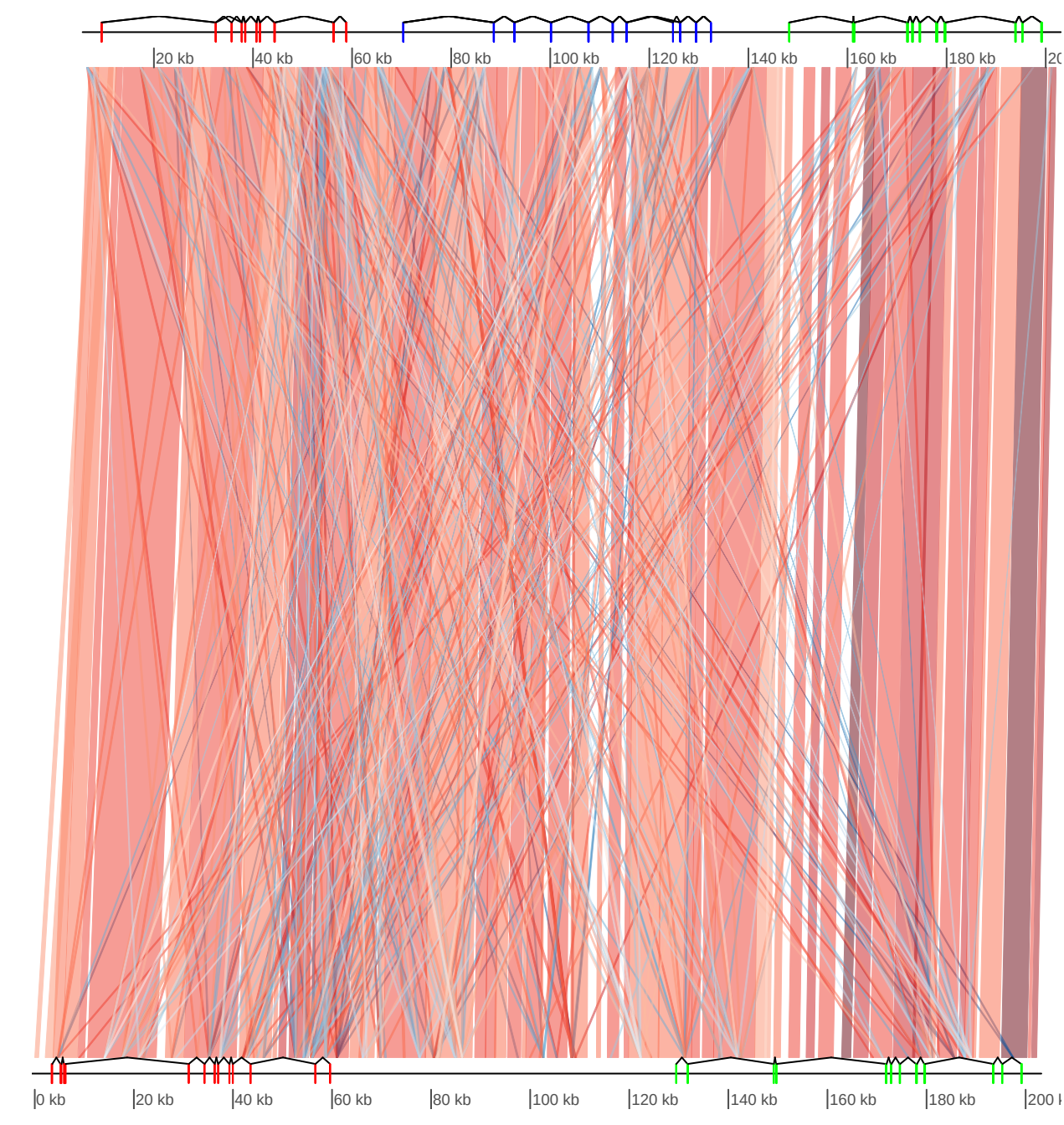
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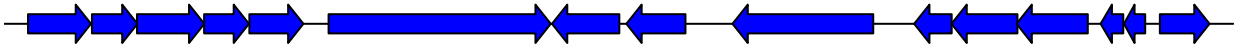
180 kb
help("chrY_subseg")

help("gene_types")



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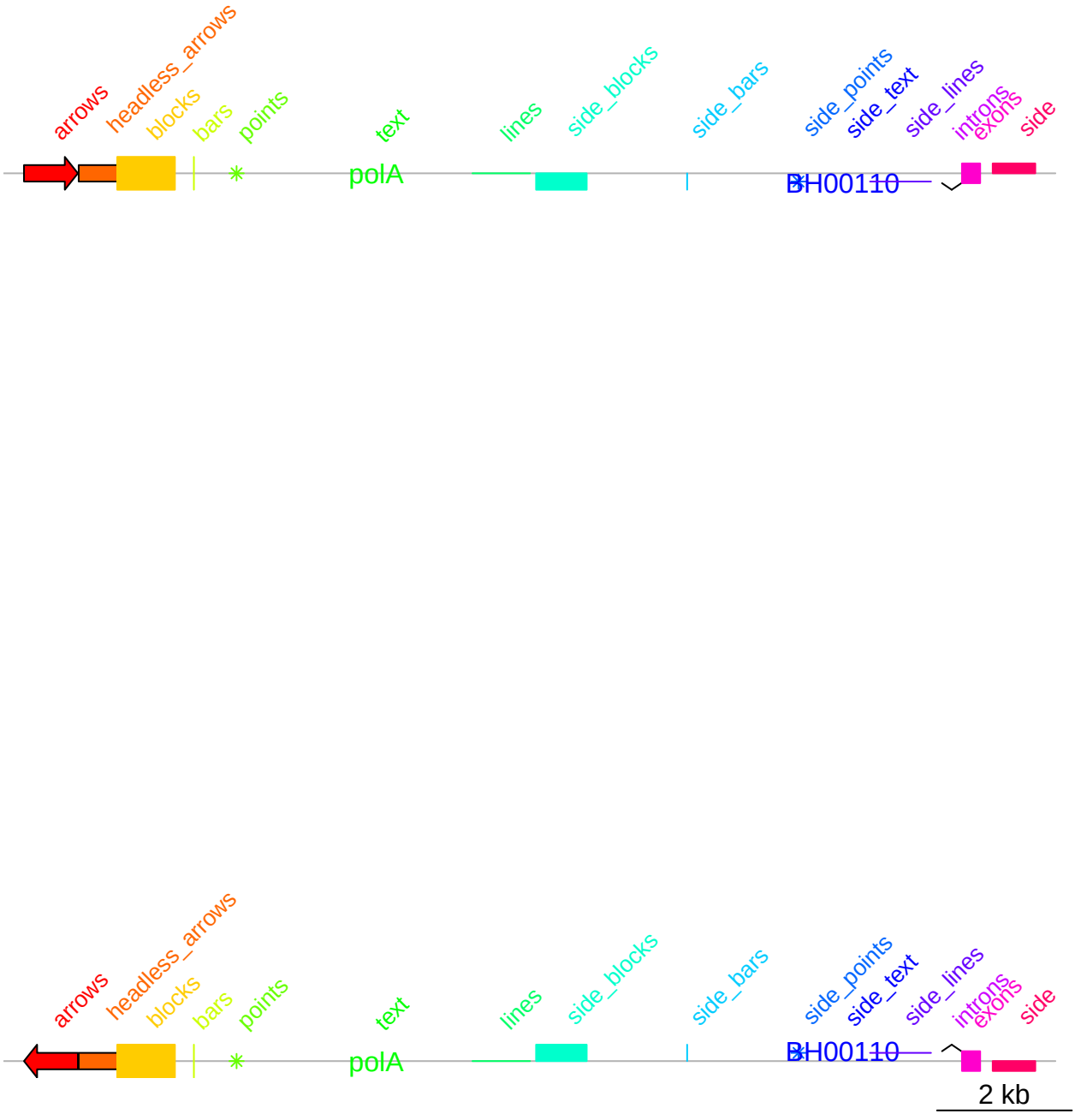
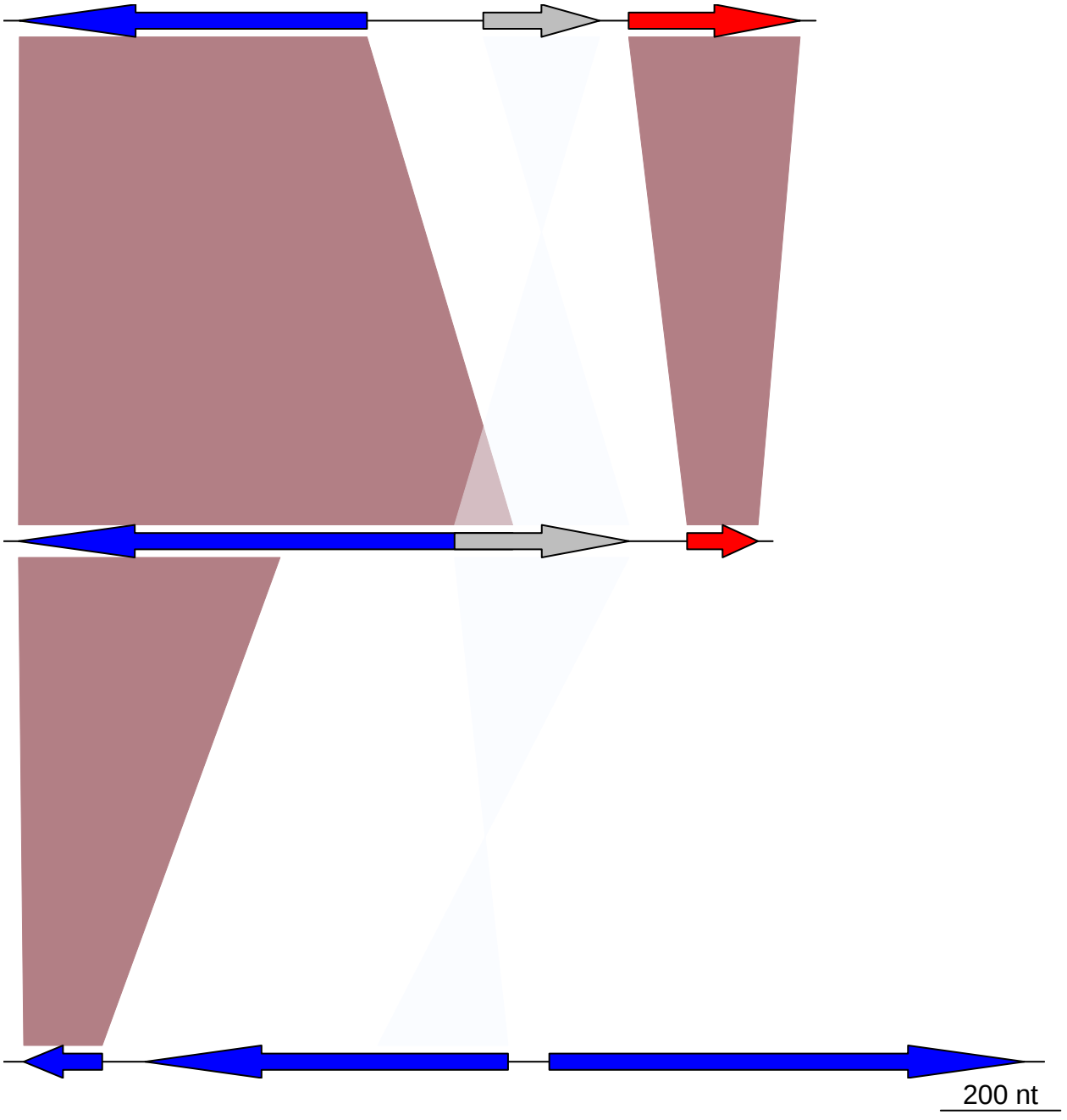
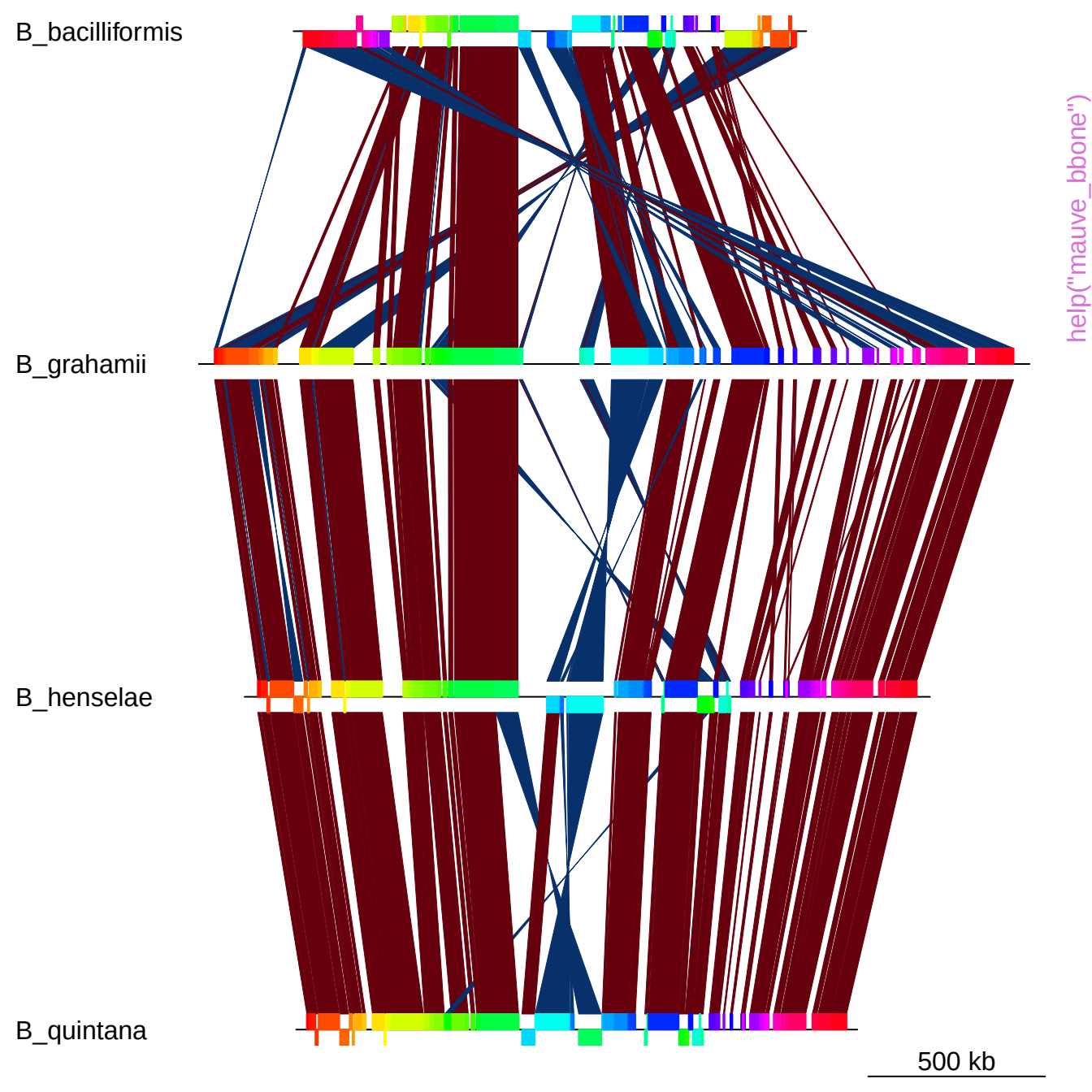
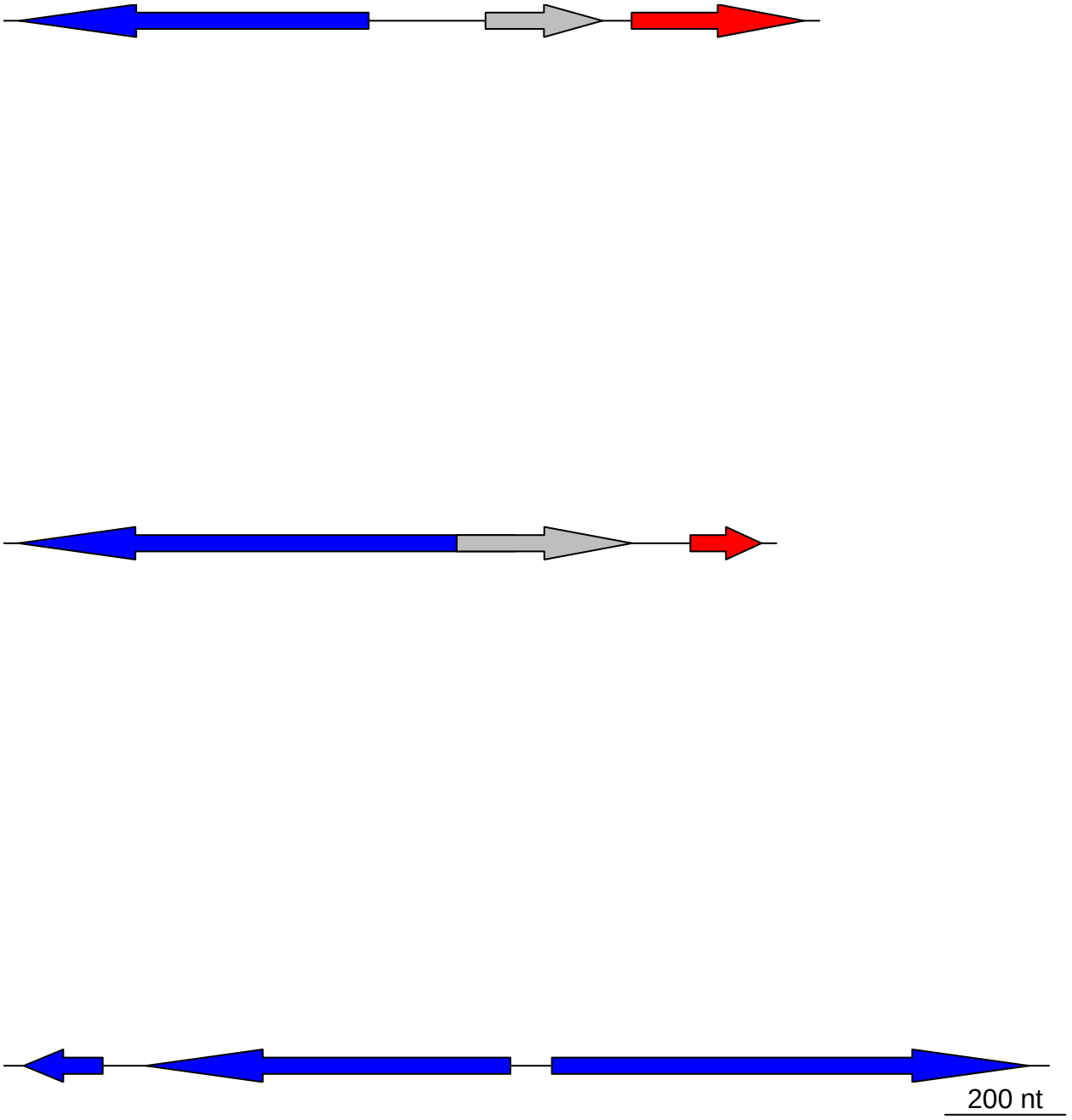

arrows
headless_arrows
blocks
bars
points
text
lines
side_blocks
side_bars
side_points
side_text
side_lines
introns
exons
side
polA
BH00110
help("gene_types")
2 kb



help("genoPlotR-package")


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


help("plot_gene_map")
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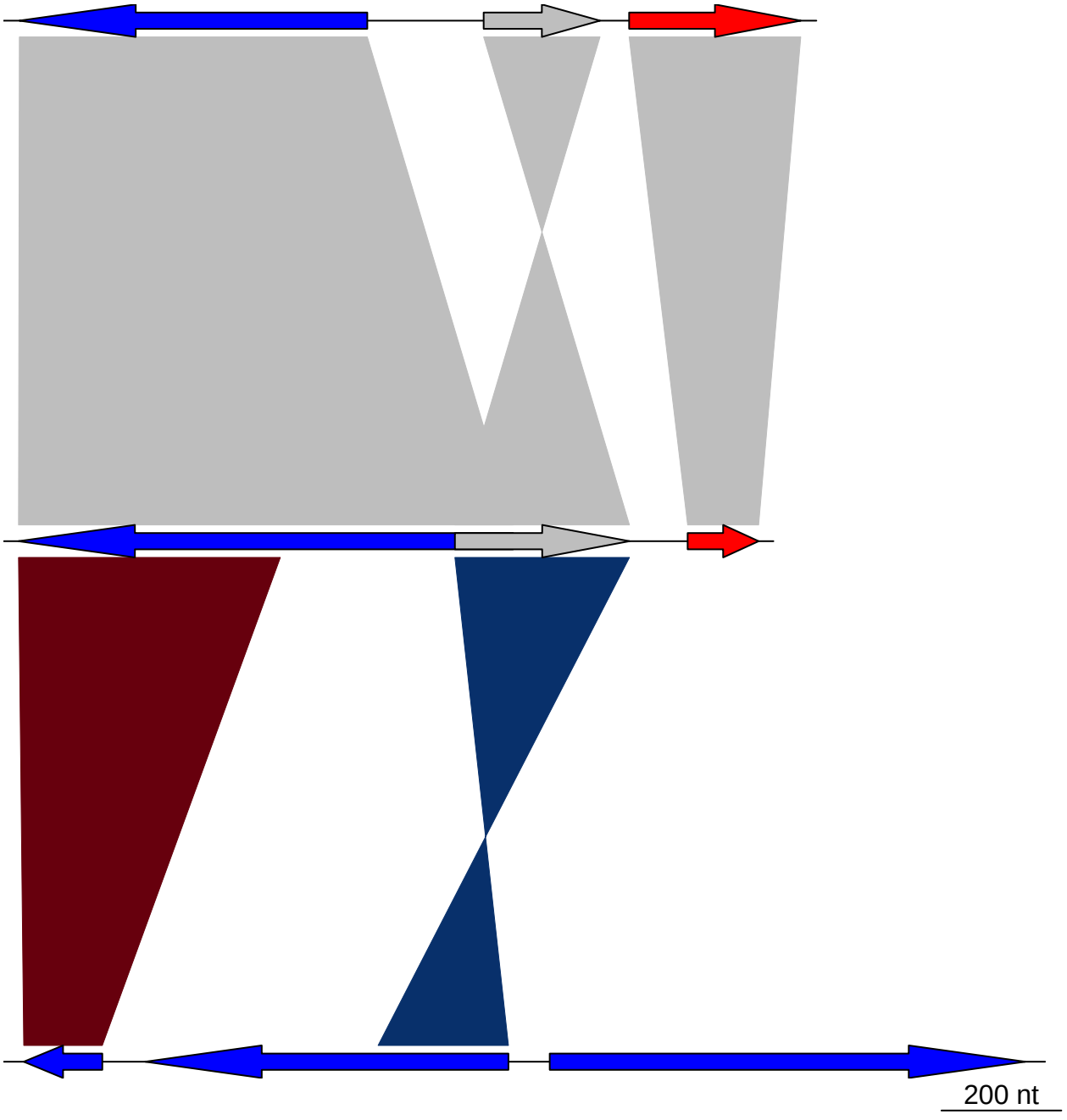
help("plot_gene_map")
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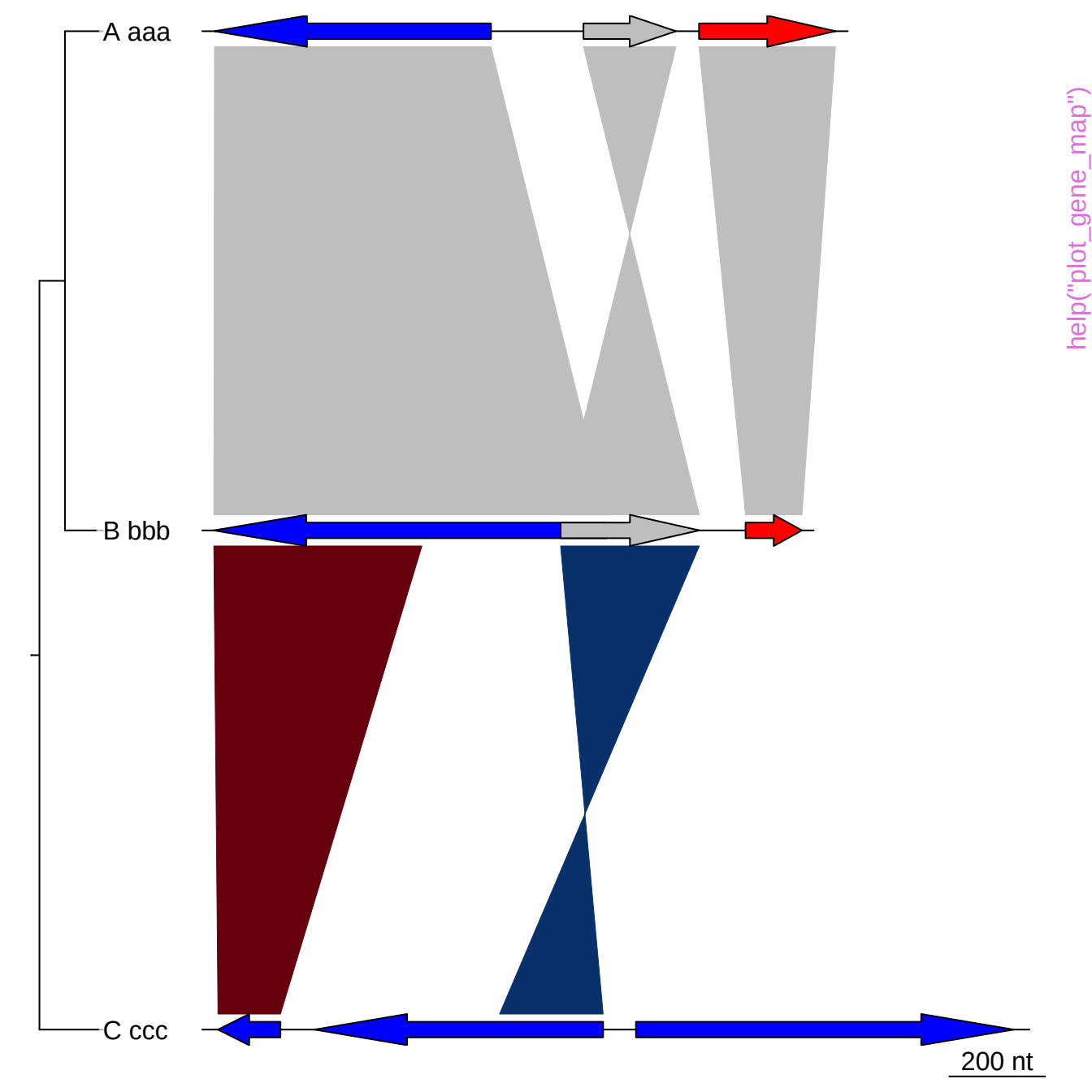
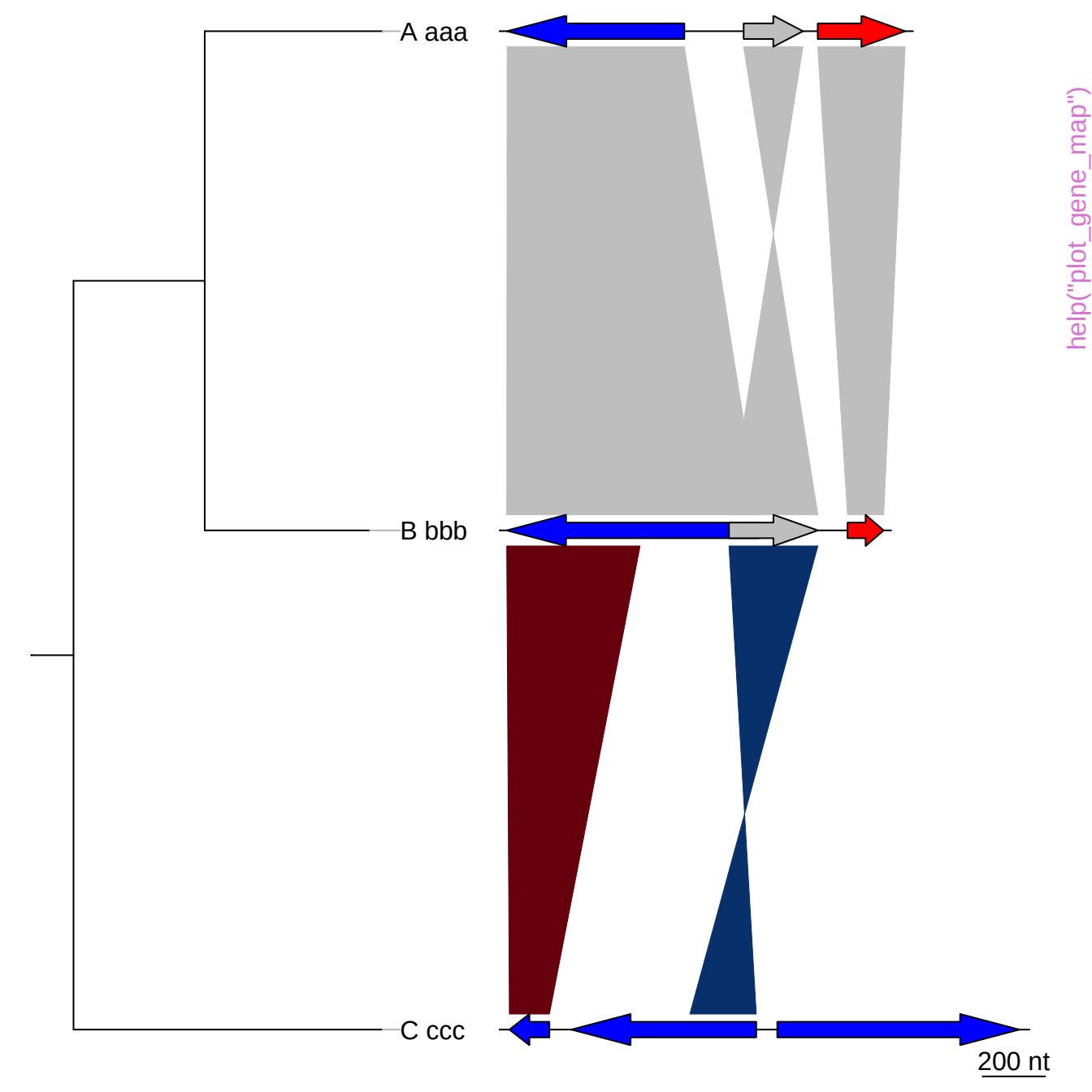
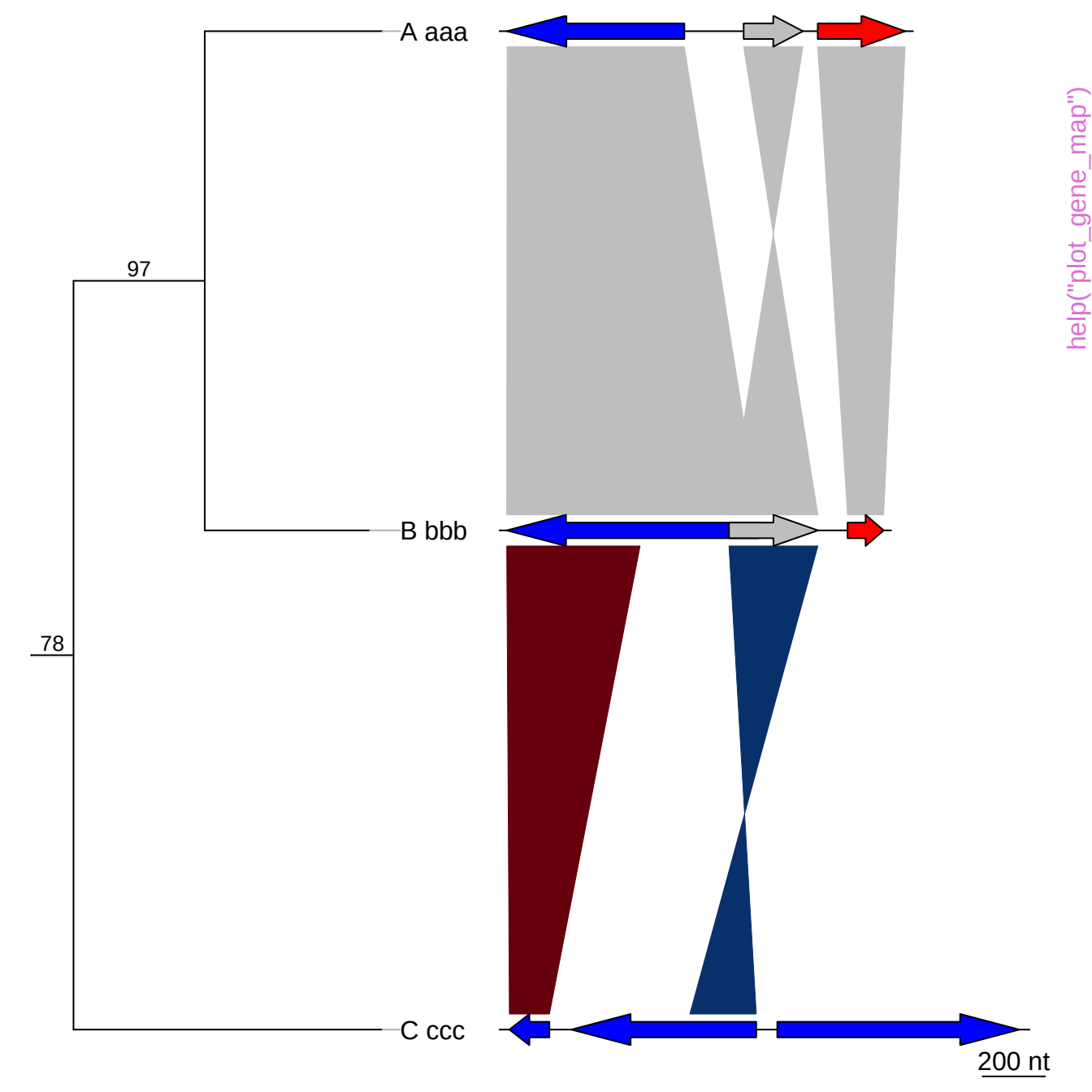
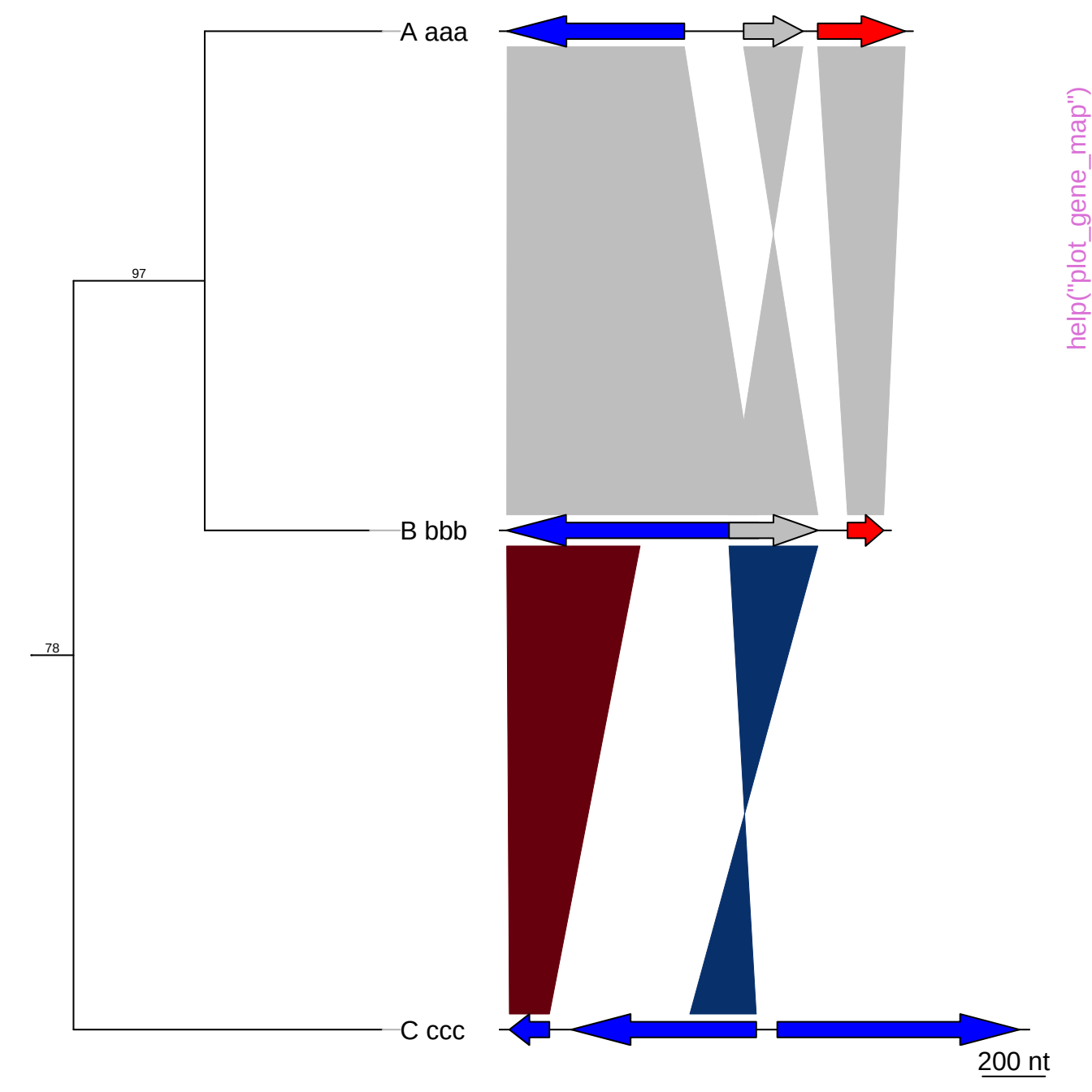
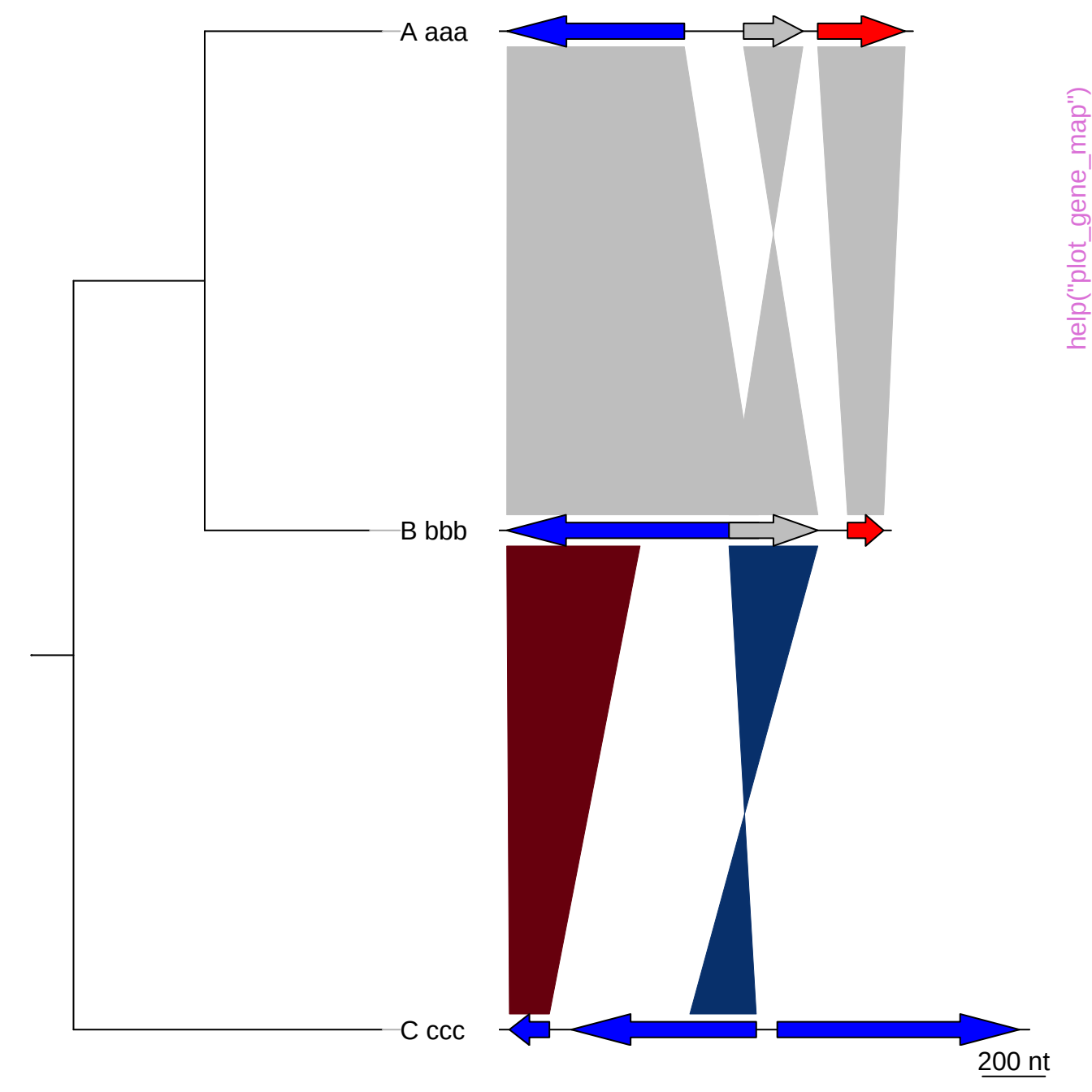
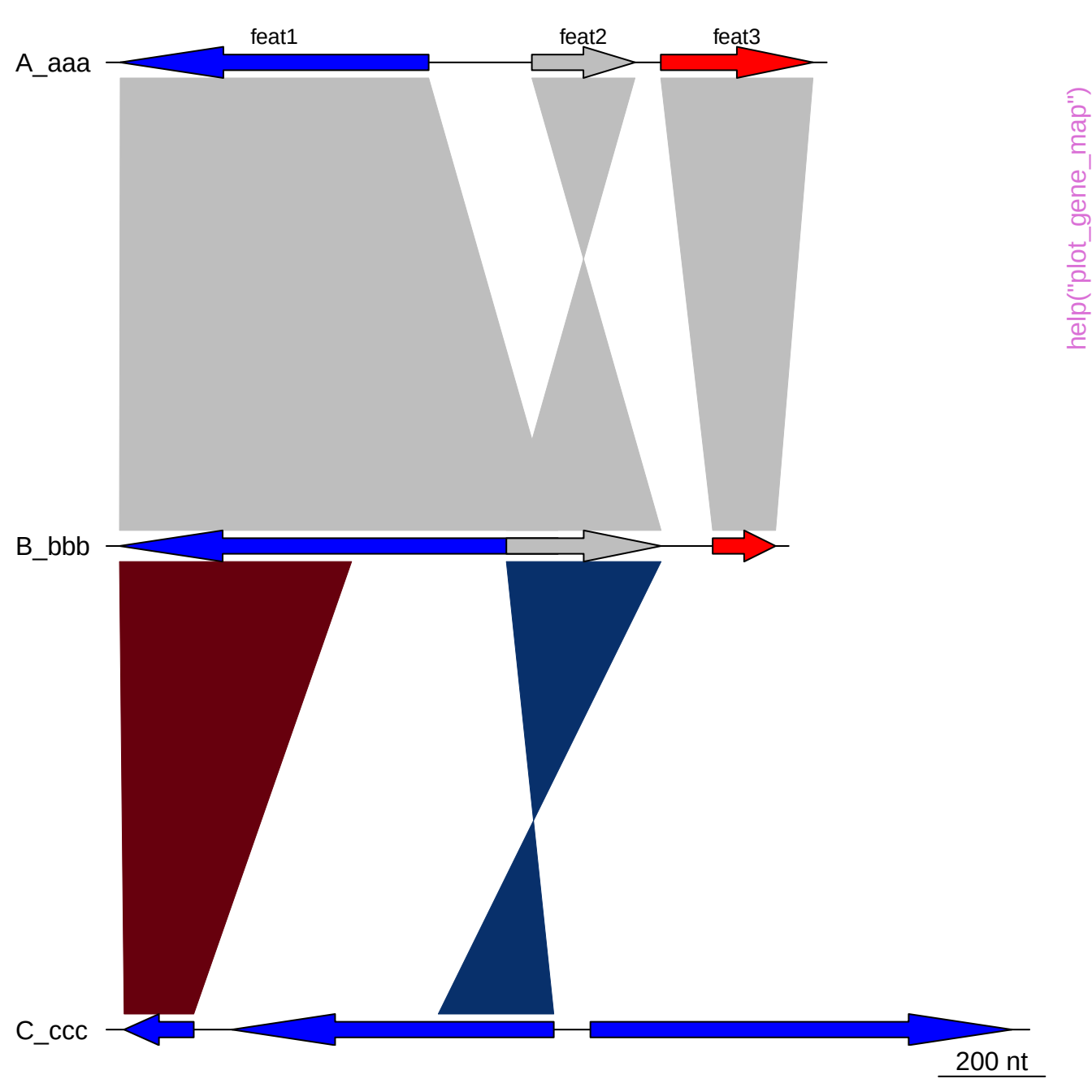

A aaa
B bbb
C ccc
200 nt
help("plot_gene_map")


A aaa
B bbb
C ccc
200 nt
help("plot_gene_map")


A aaa
B bbb
C ccc
97
78
200 nt
help("plot_gene_map")


A aaa
B bbb
C ccc
97
78
200 nt
help("plot_gene_map")


A aaa
B bbb
C ccc
200 nt
help("plot_gene_map")


feat1
feat2
feat3
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B_bbb
C_ccc
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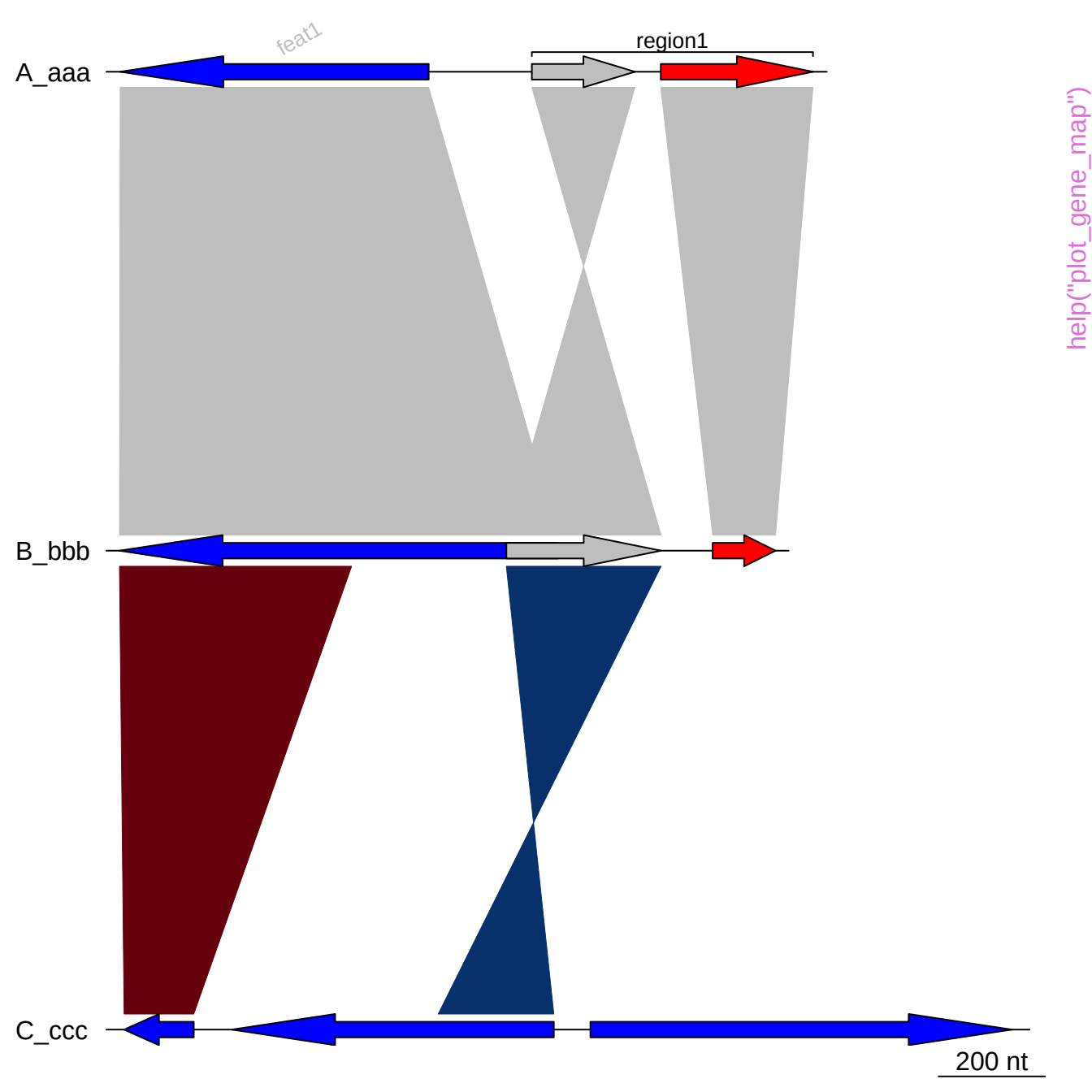
feat1
region1
A_aaa
B_bbb
C_ccc
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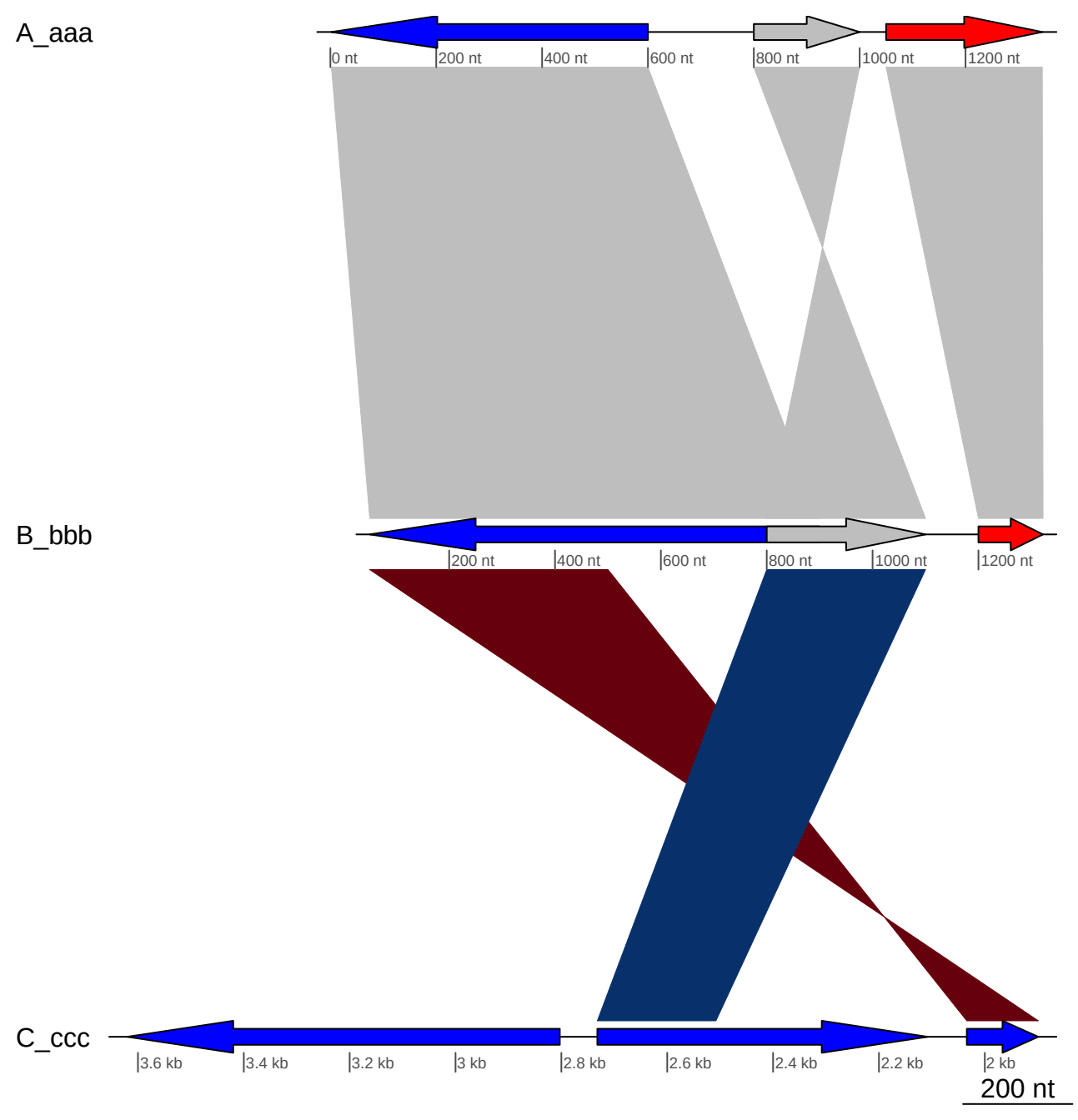
help("plot_gene_map")


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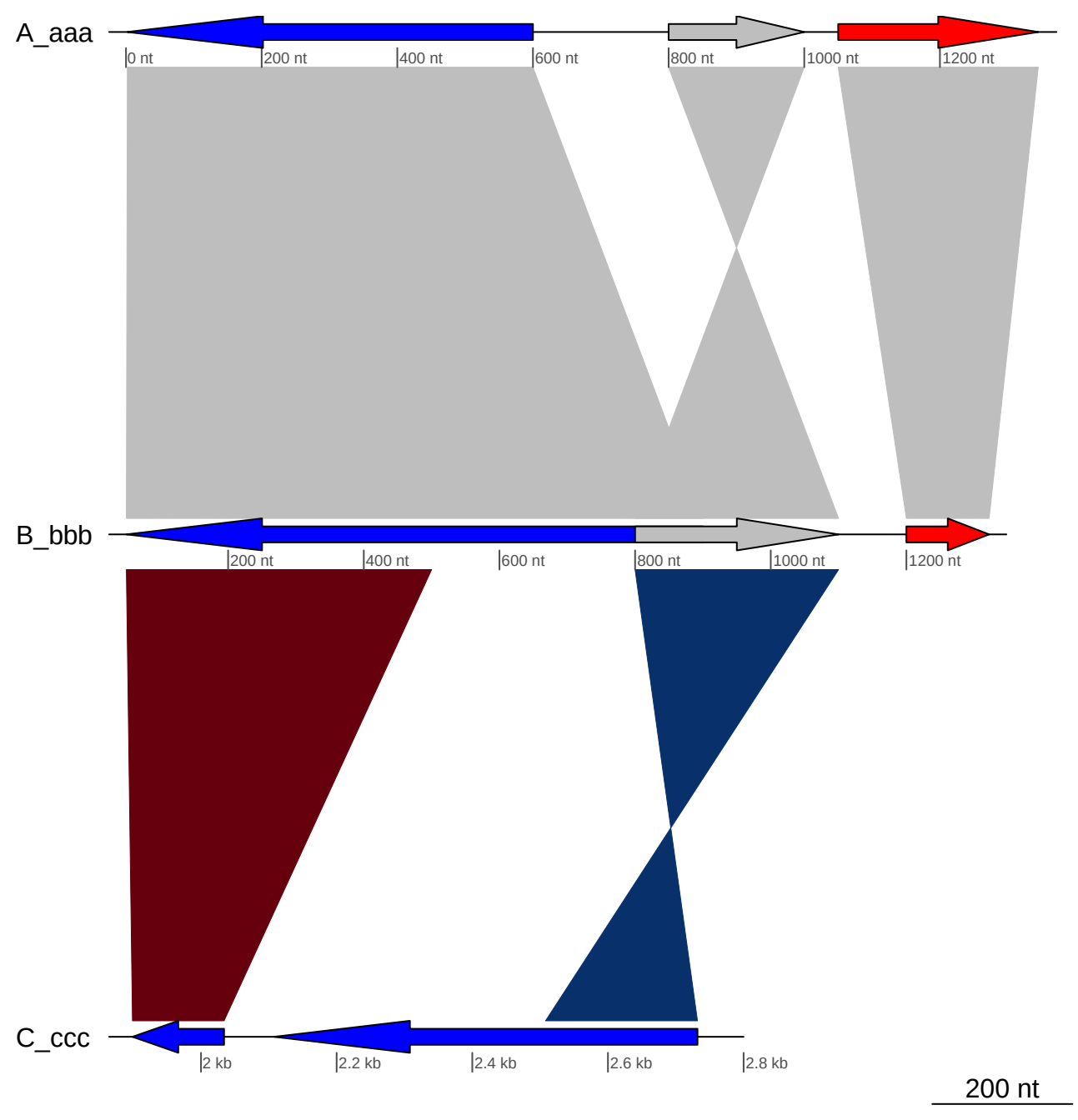
help("plot_gene_map")


A_aaa
0 nt
200 nt
400 nt
600 nt
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B_bbb
200 nt
400 nt
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C_ccc
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2.2 kb
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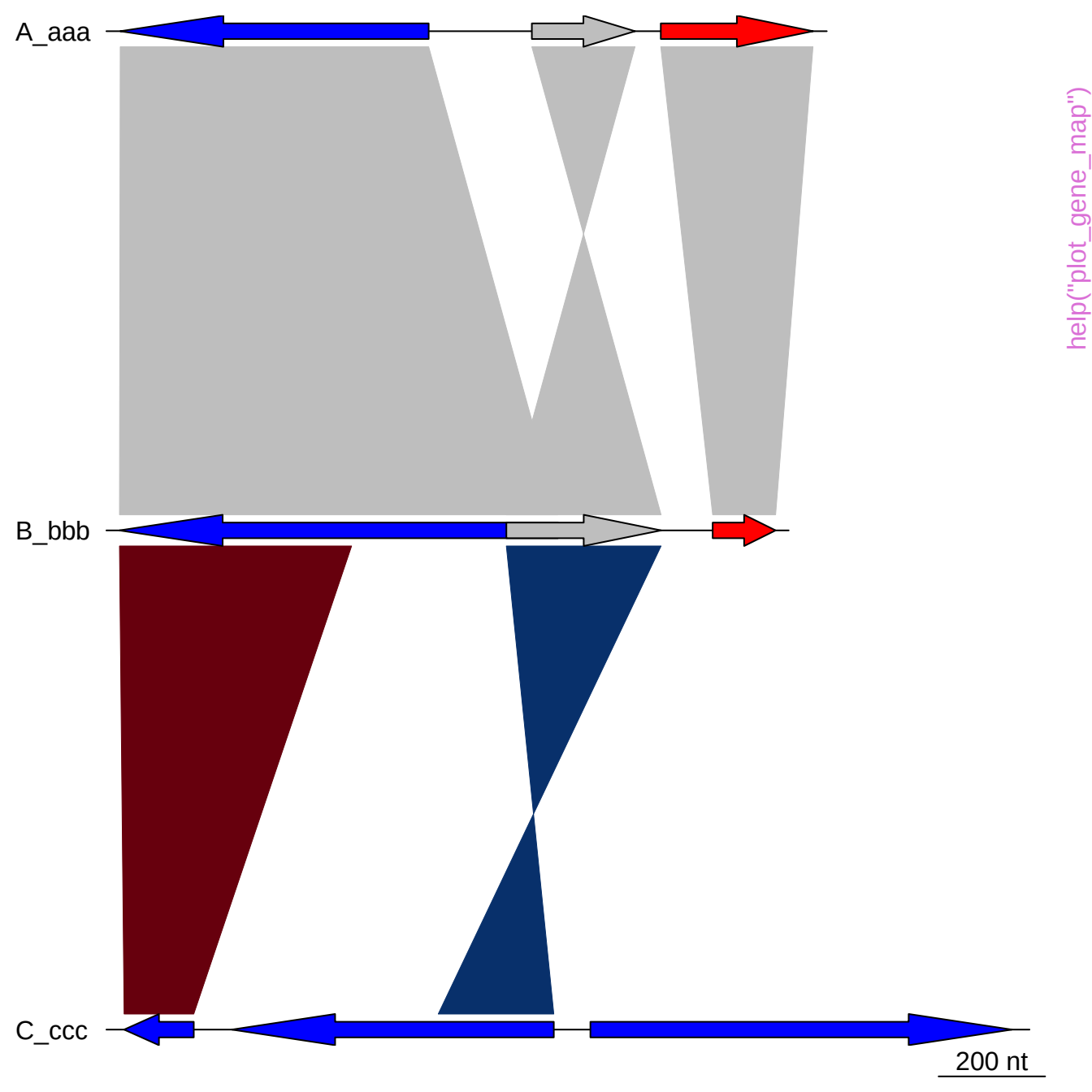
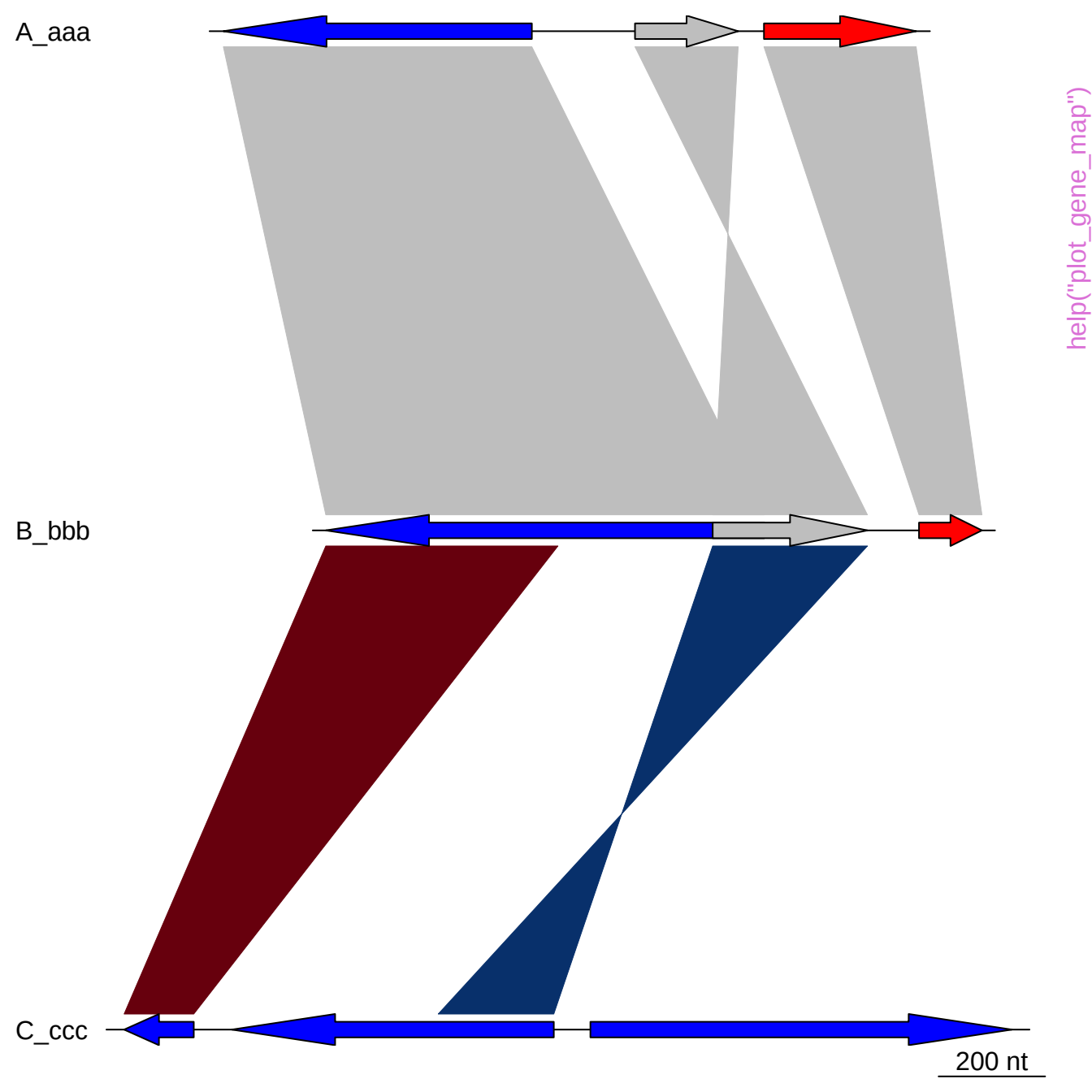
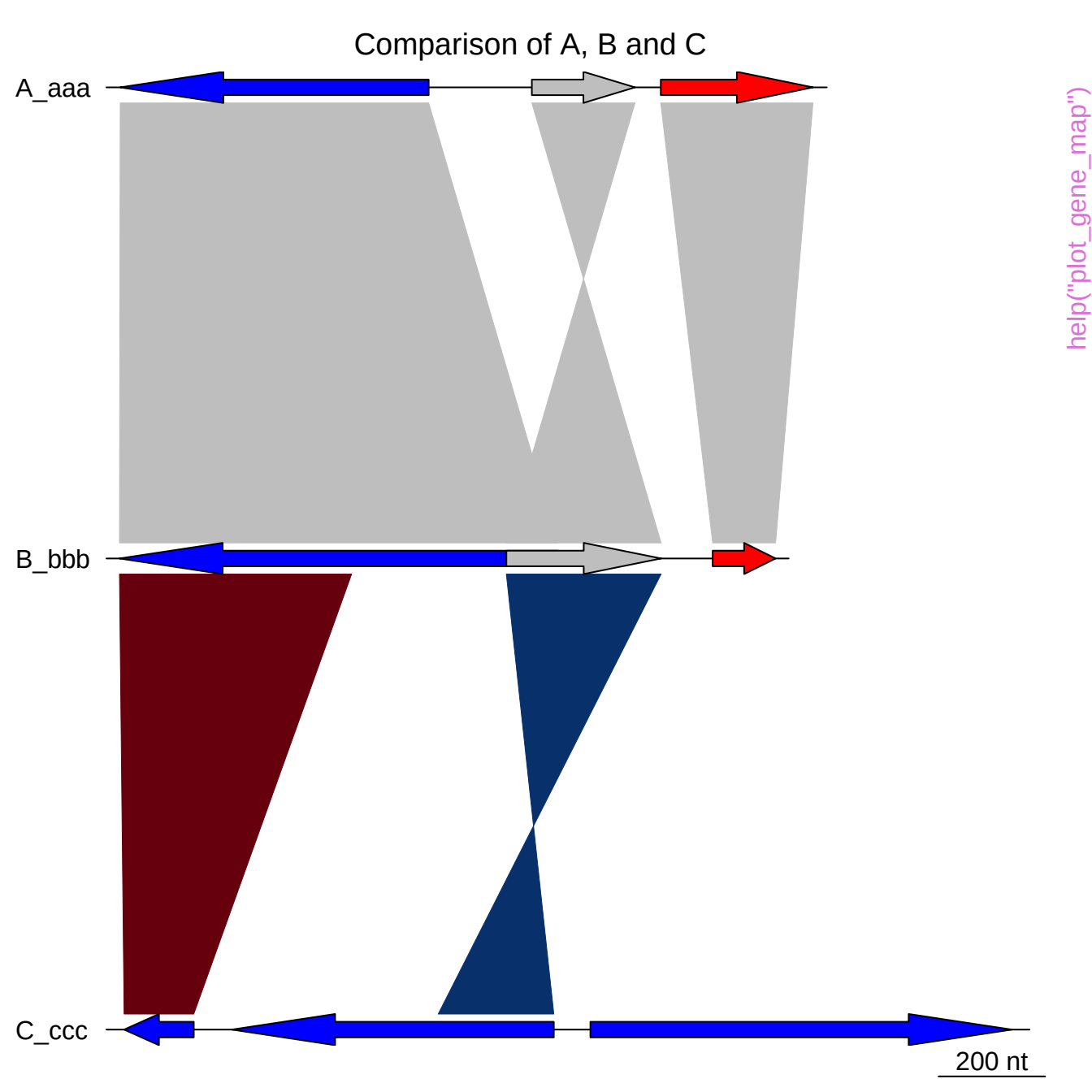
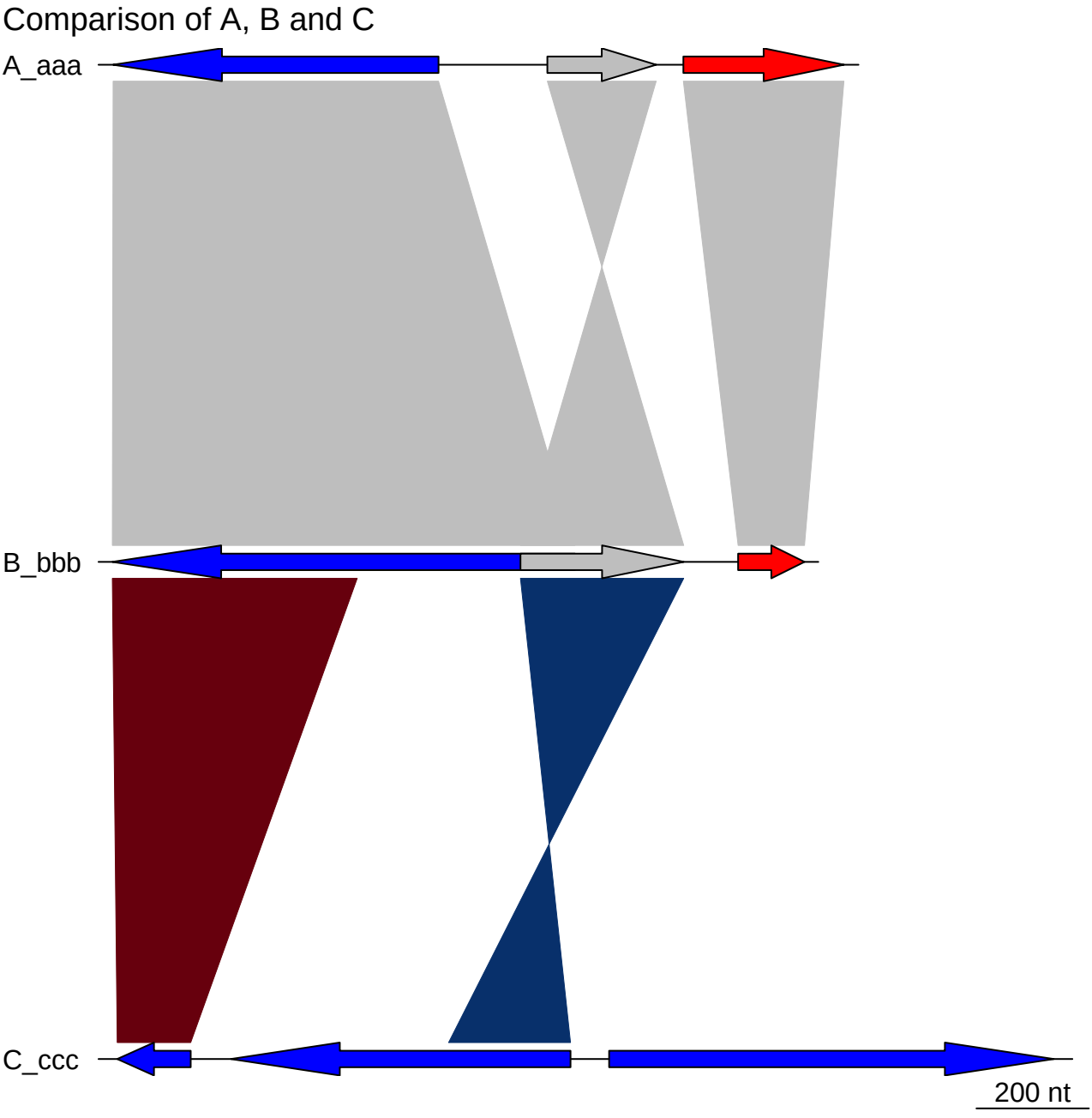
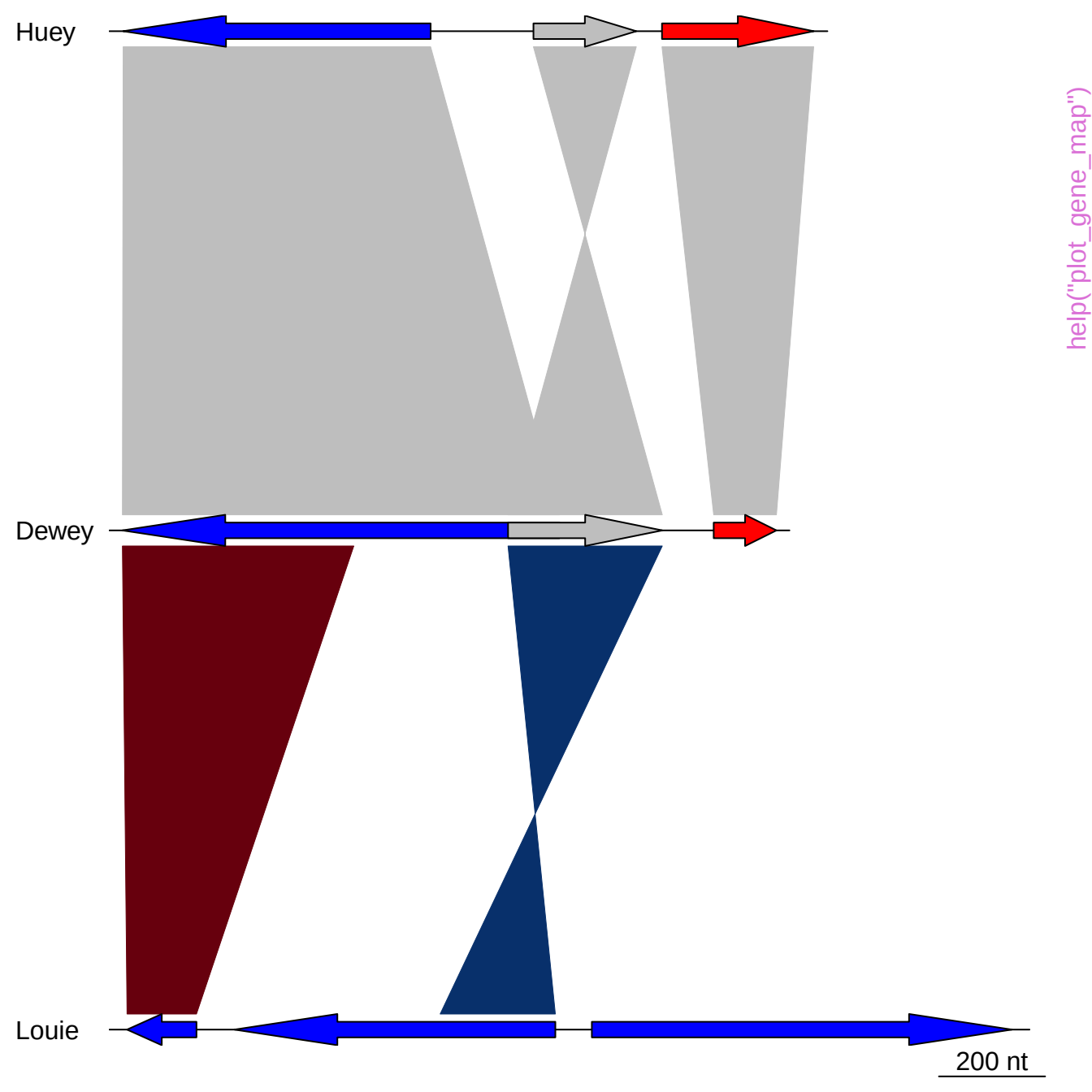

A_aaa
B_bbb
C_ccc
200 nt
help("plot_gene_map")


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
200 nt
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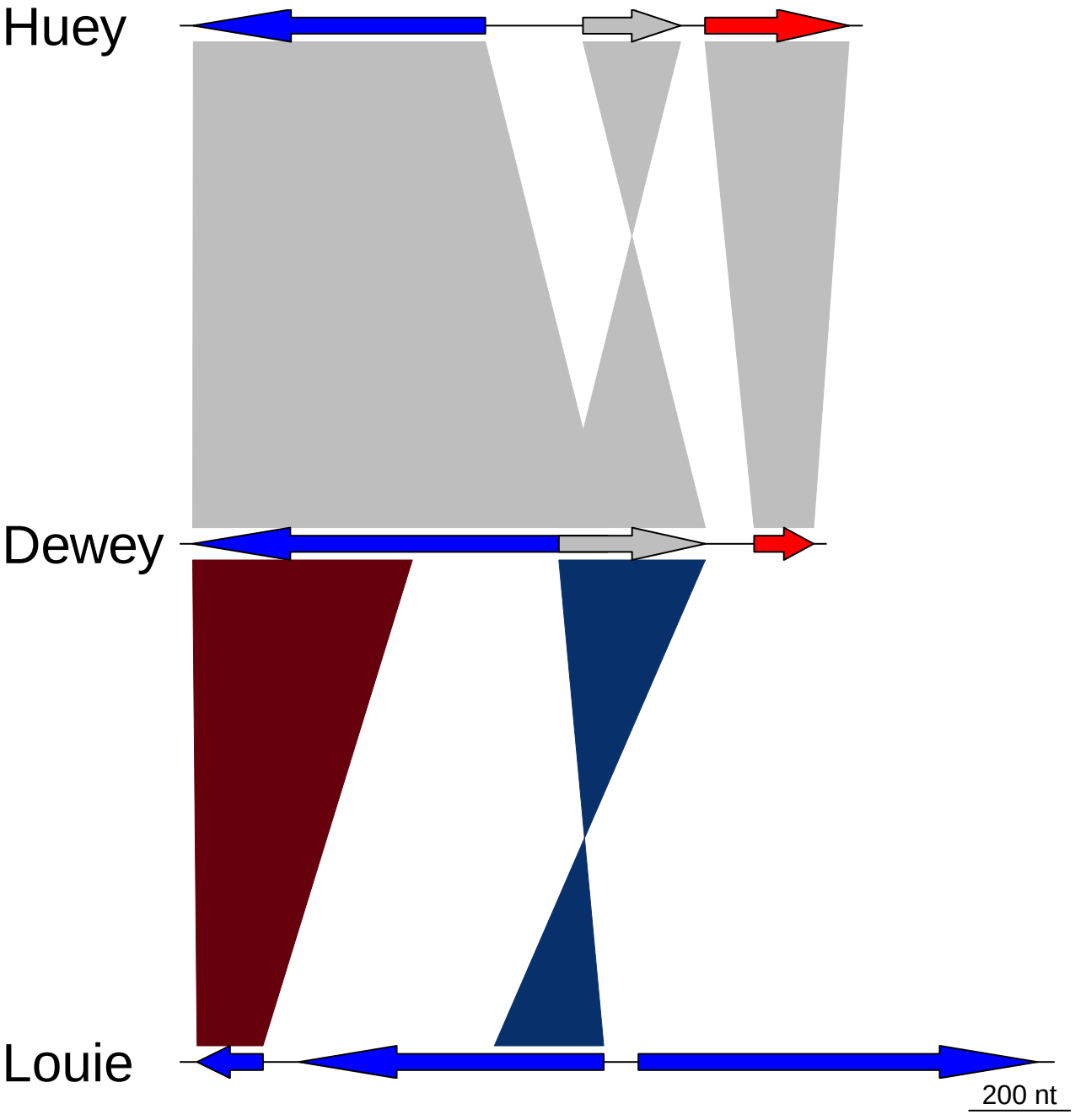
Huey
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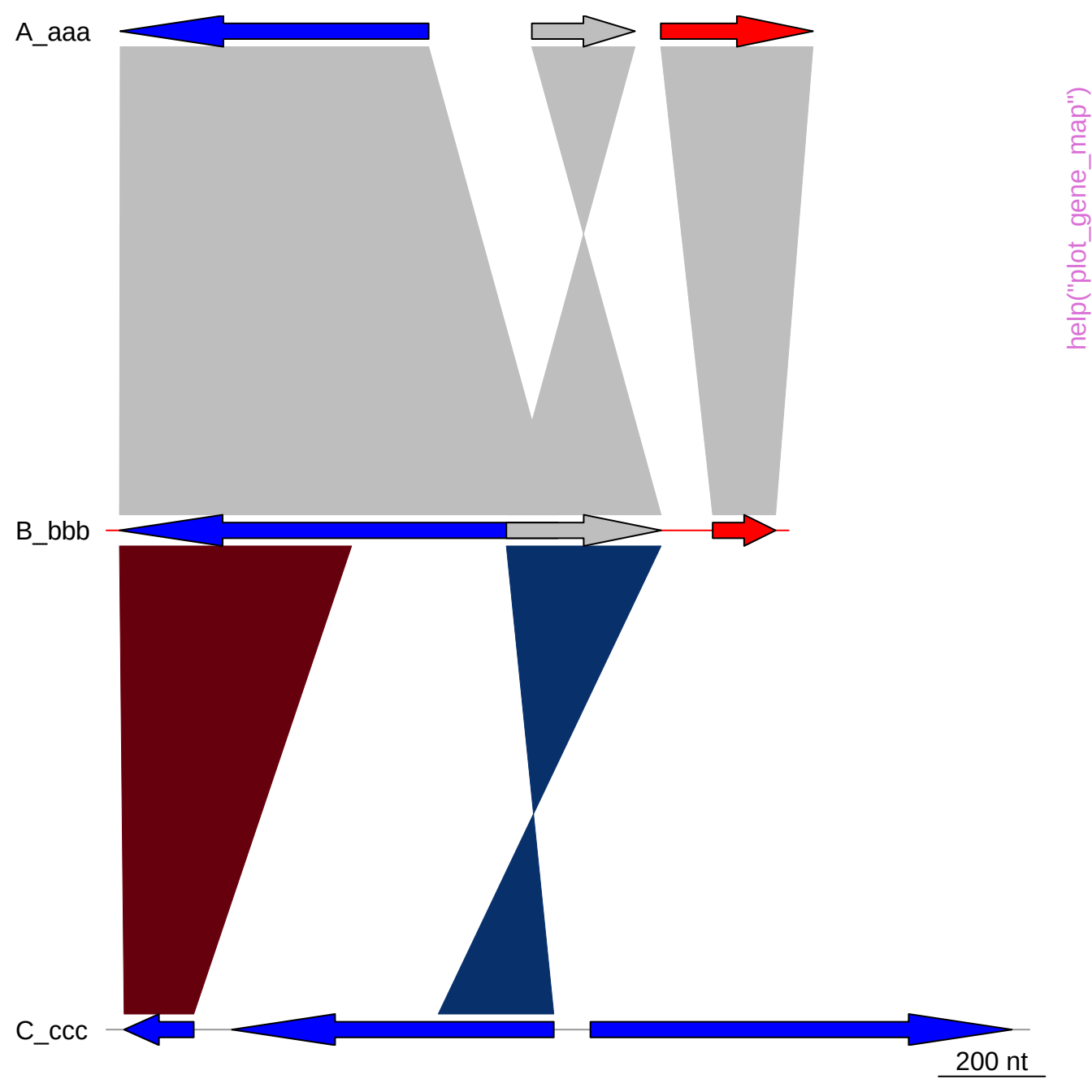
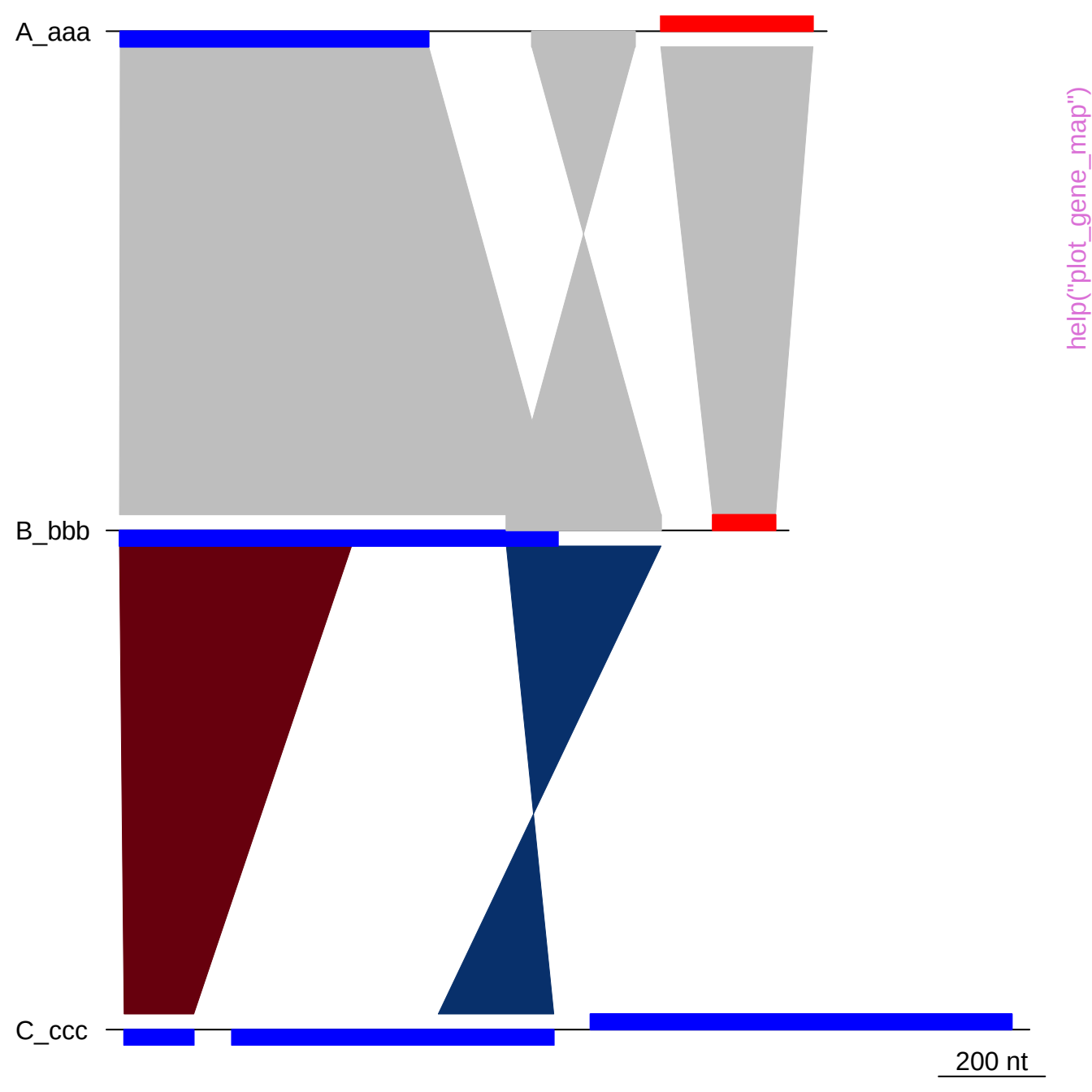

A_aaa
B_bbb
C_ccc
200 nt
help("plot_gene_map")


A_aaa
B_bbb
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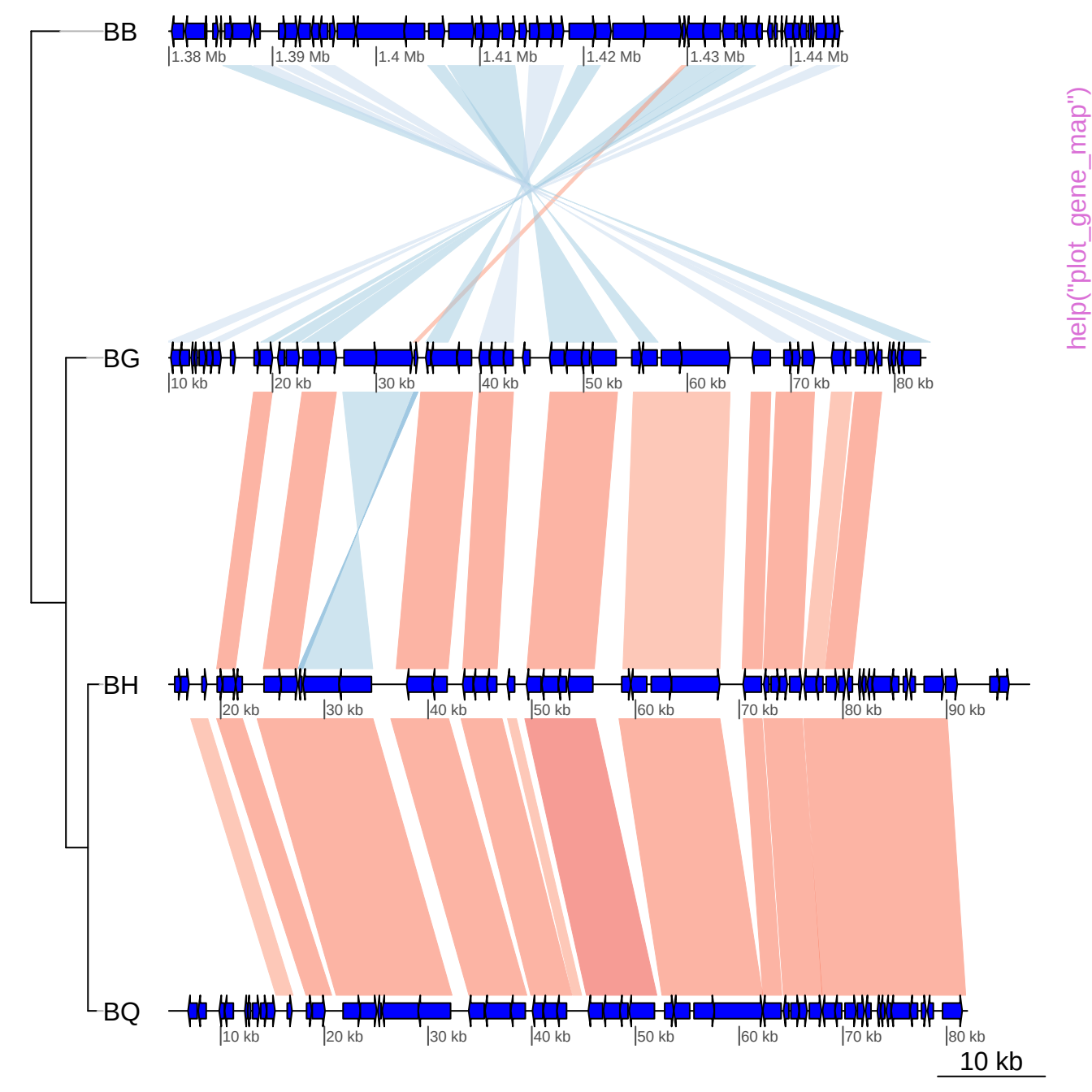
help("plot_gene_map")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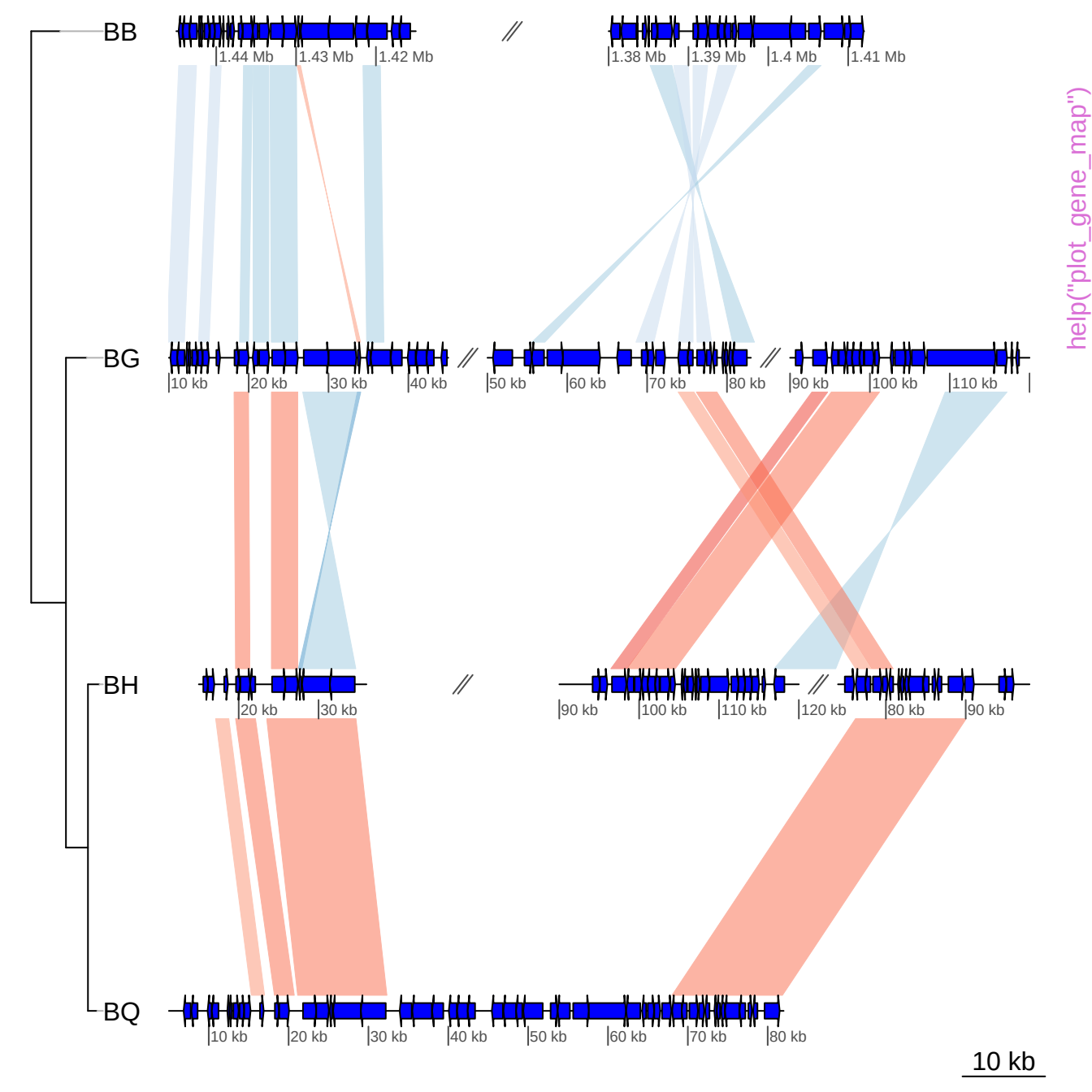
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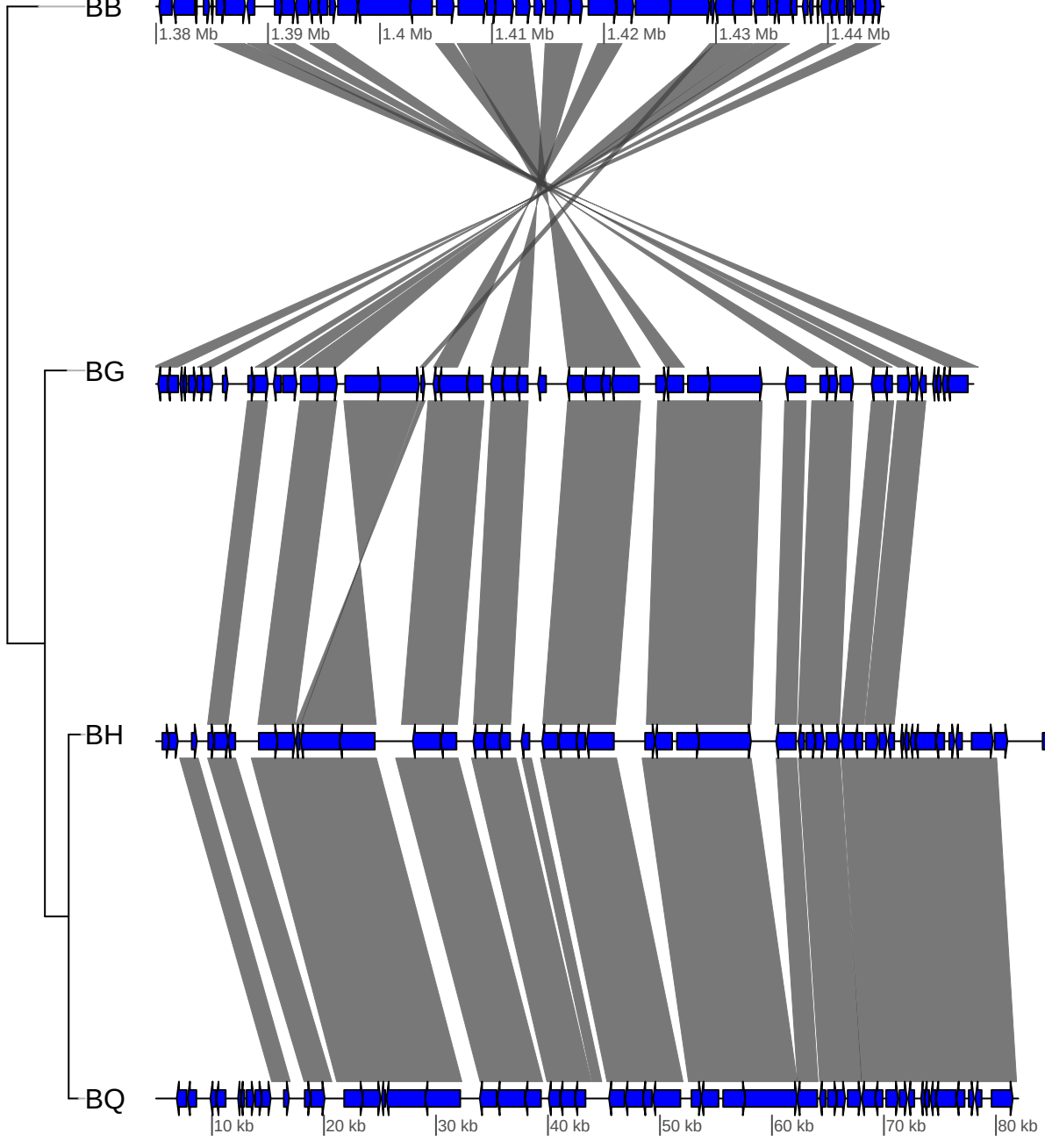
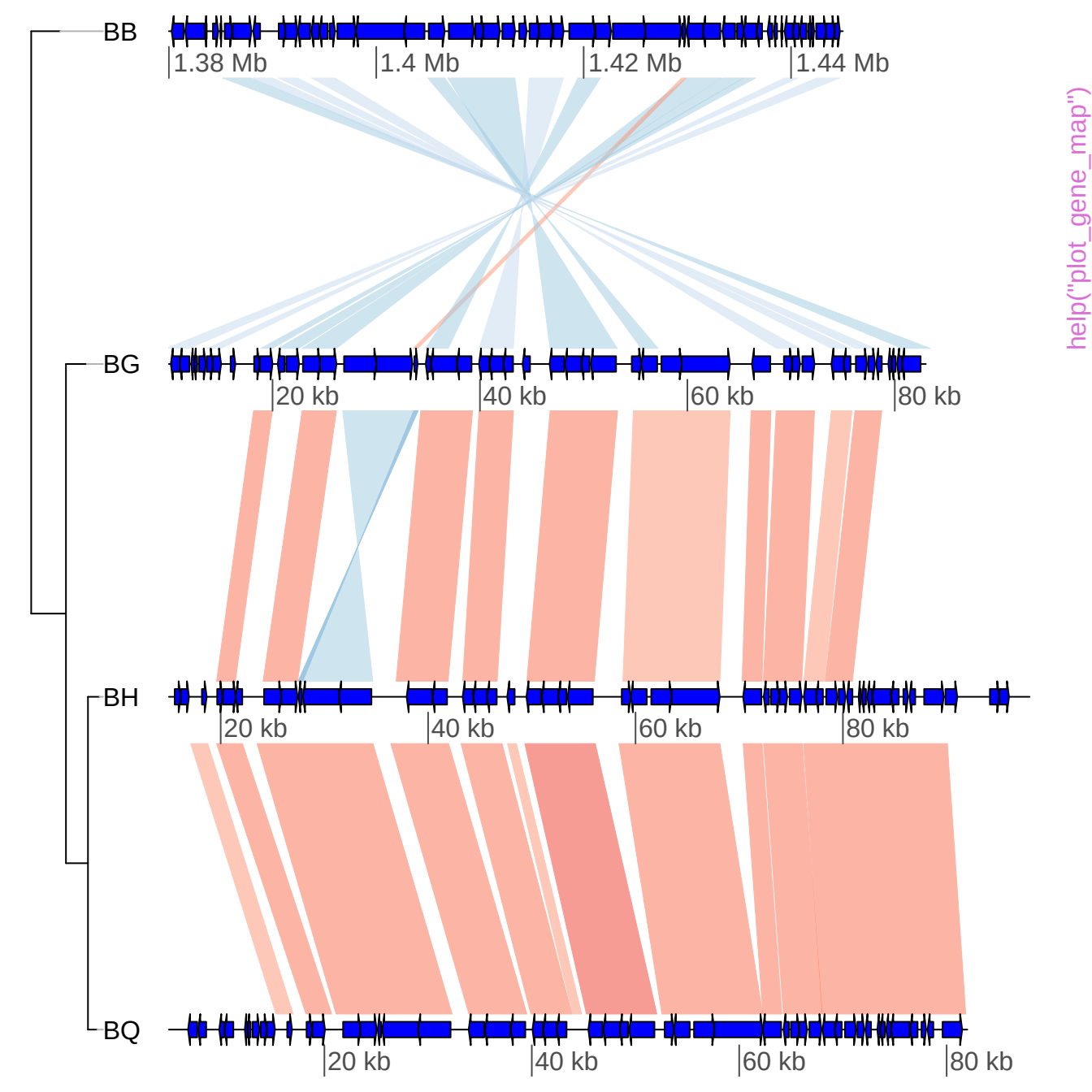


help("plot_gene_map")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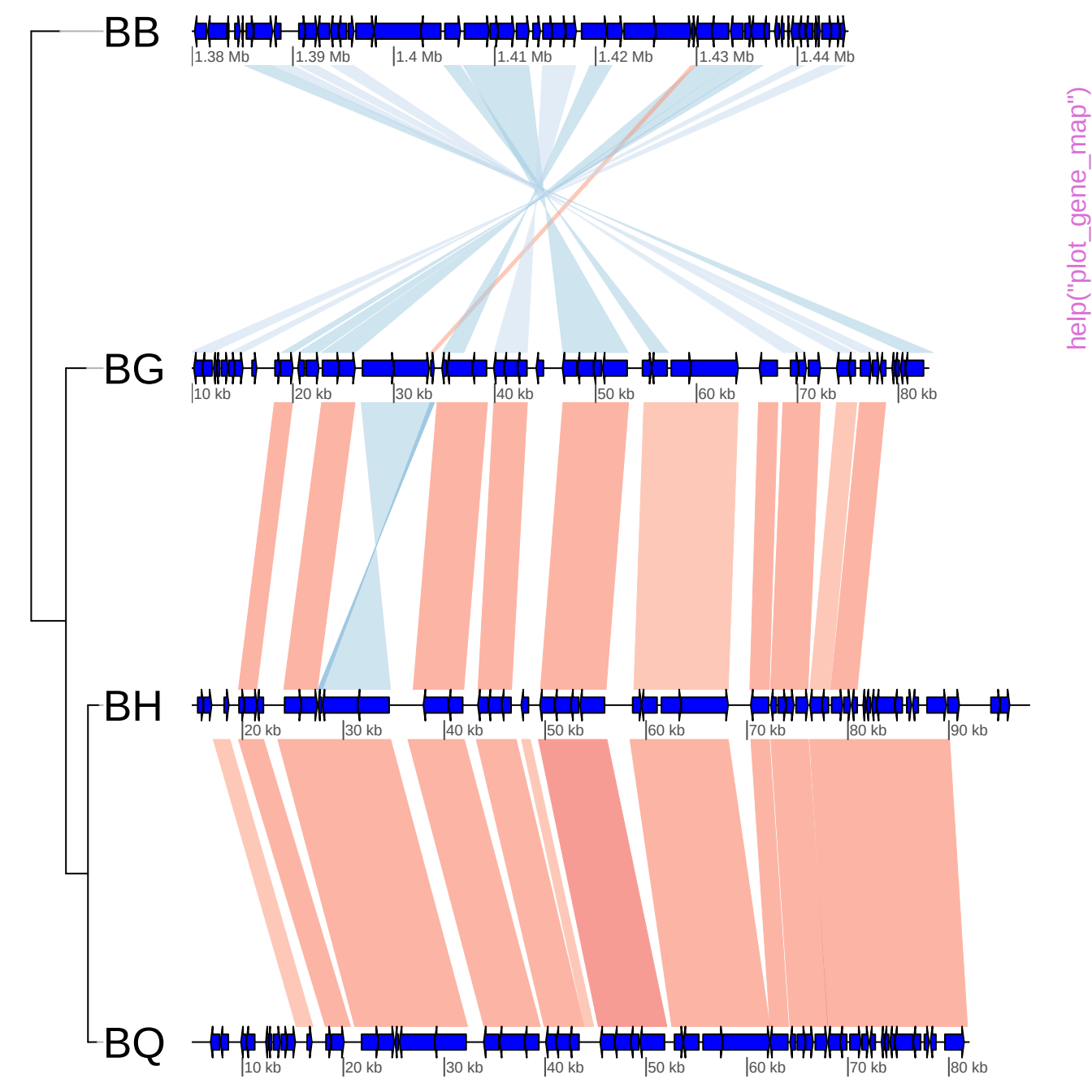
help("plot_gene_map")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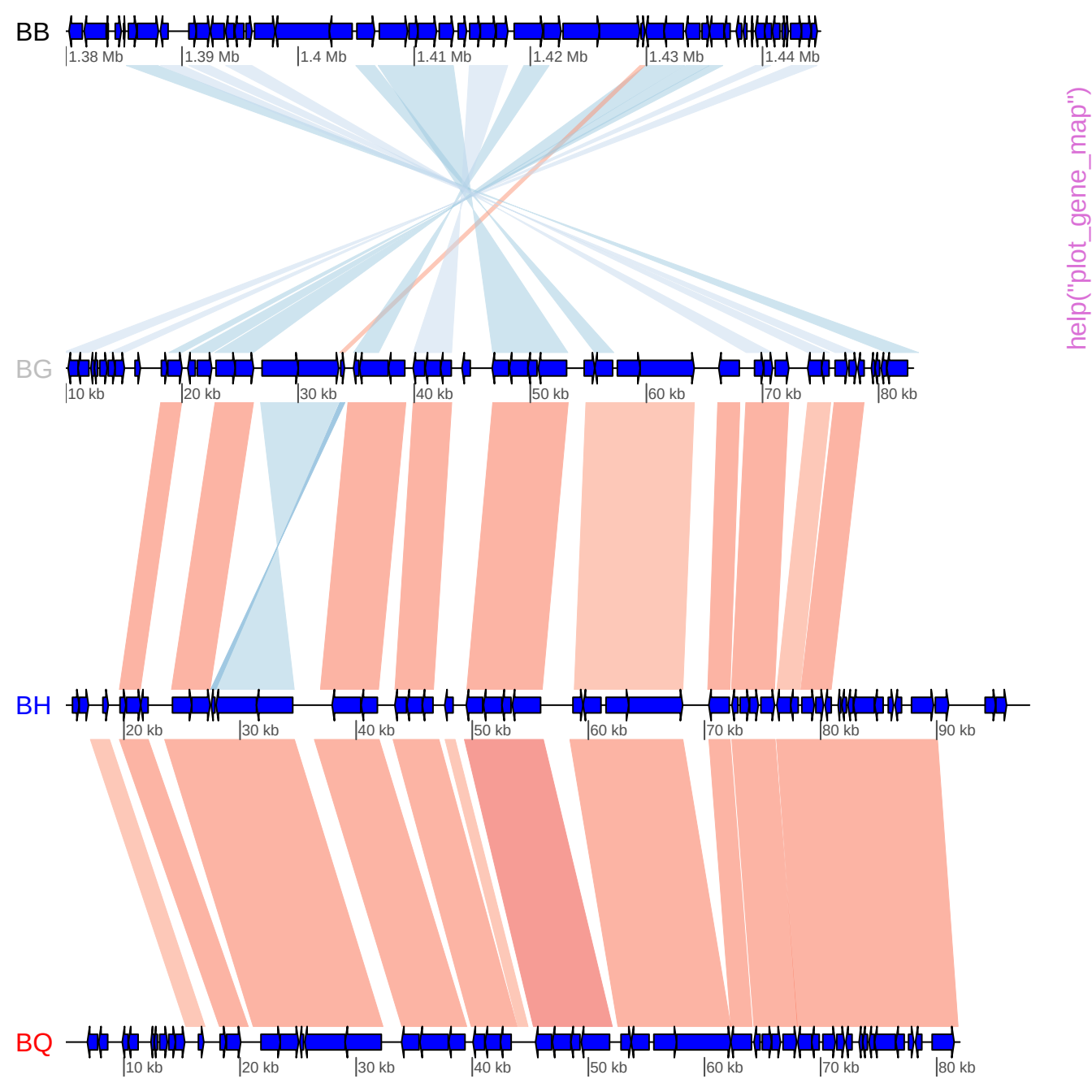
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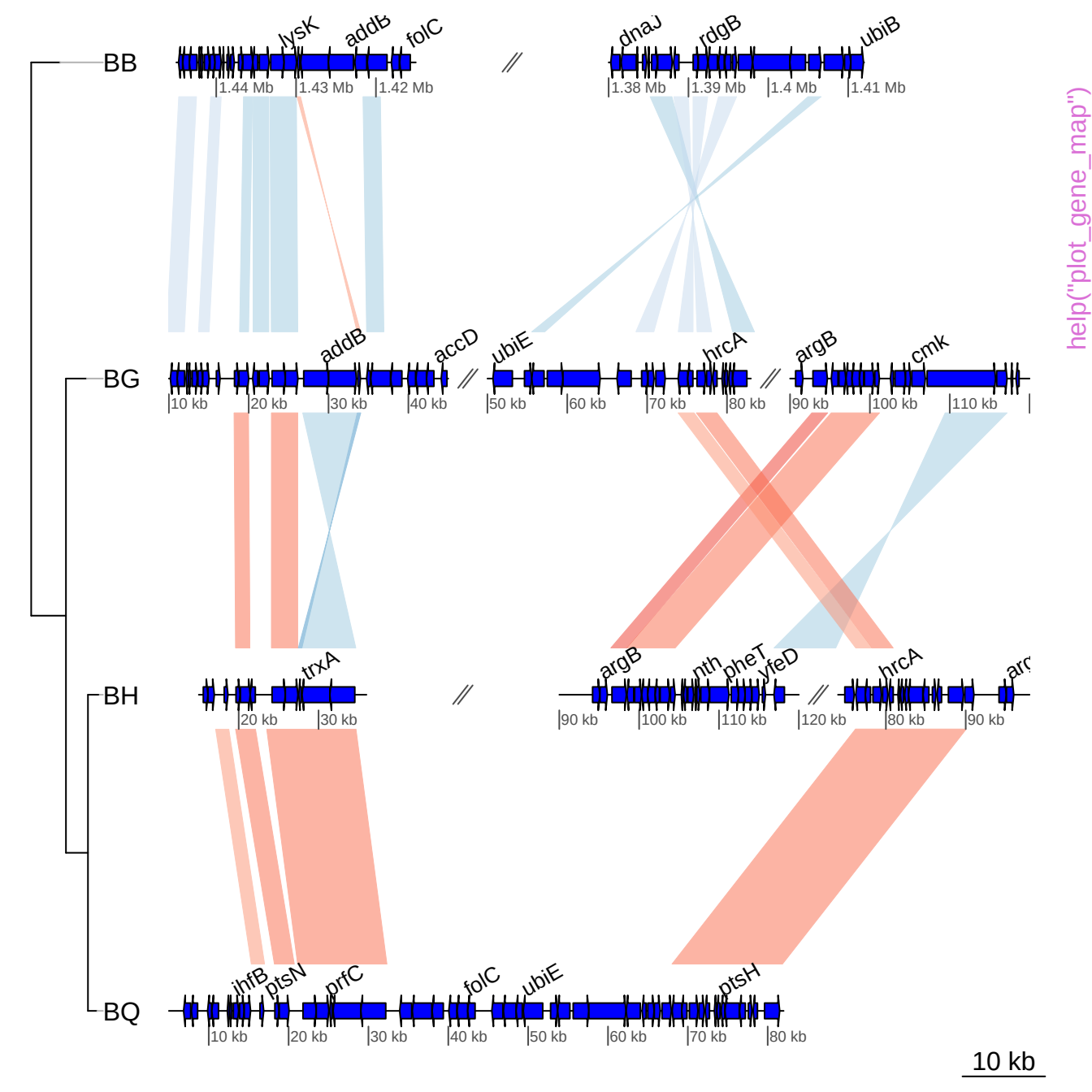
help("plot_gene_map")


BB
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rdgB
ubiB
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accD
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hrcA
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BQ
ihfB
ptsN
prfC
folC
ubiE
ptsH
10 kb
20 kb
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70 kb
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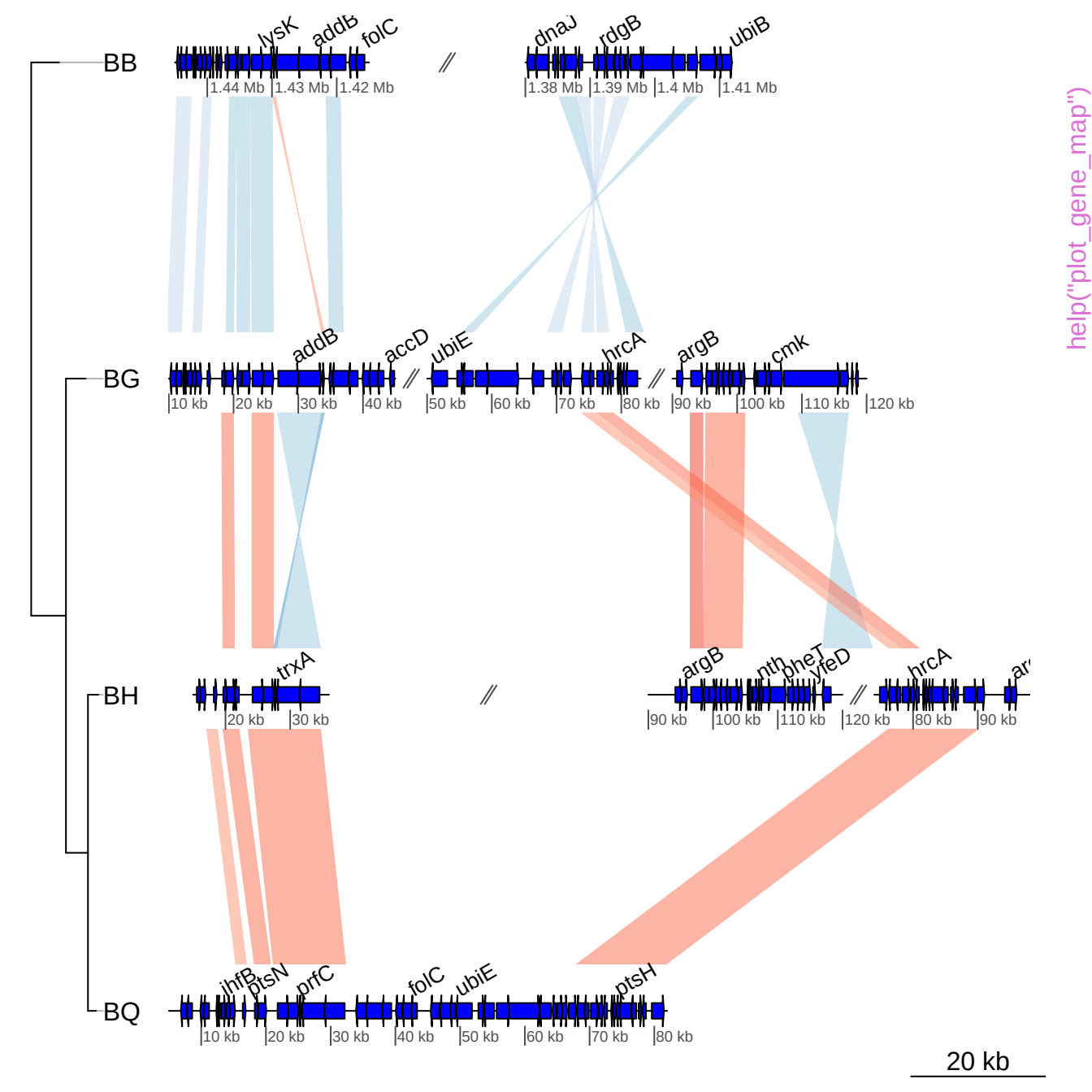
10 kb
help("plot_gene_map")


BB
lysK
addB
folC
1.44 Mb
1.43 Mb
1.42 Mb
dnaJ
rdgB
ubiB
1.38 Mb
1.39 Mb
1.4 Mb
1.41 Mb
BG
addB
accD
ubiE
hrcA
argB
cmk
10 kb
20 kb
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90 kb
100 kb
110 kb
120 kb
BH
trxA
20 kb
30 kb
argB
nth
pheT
ylfeD
hrcA
90 kb
100 kb
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120 kb
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BQ
ihfB
ptsN
prfC
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ptsH
10 kb
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40 kb
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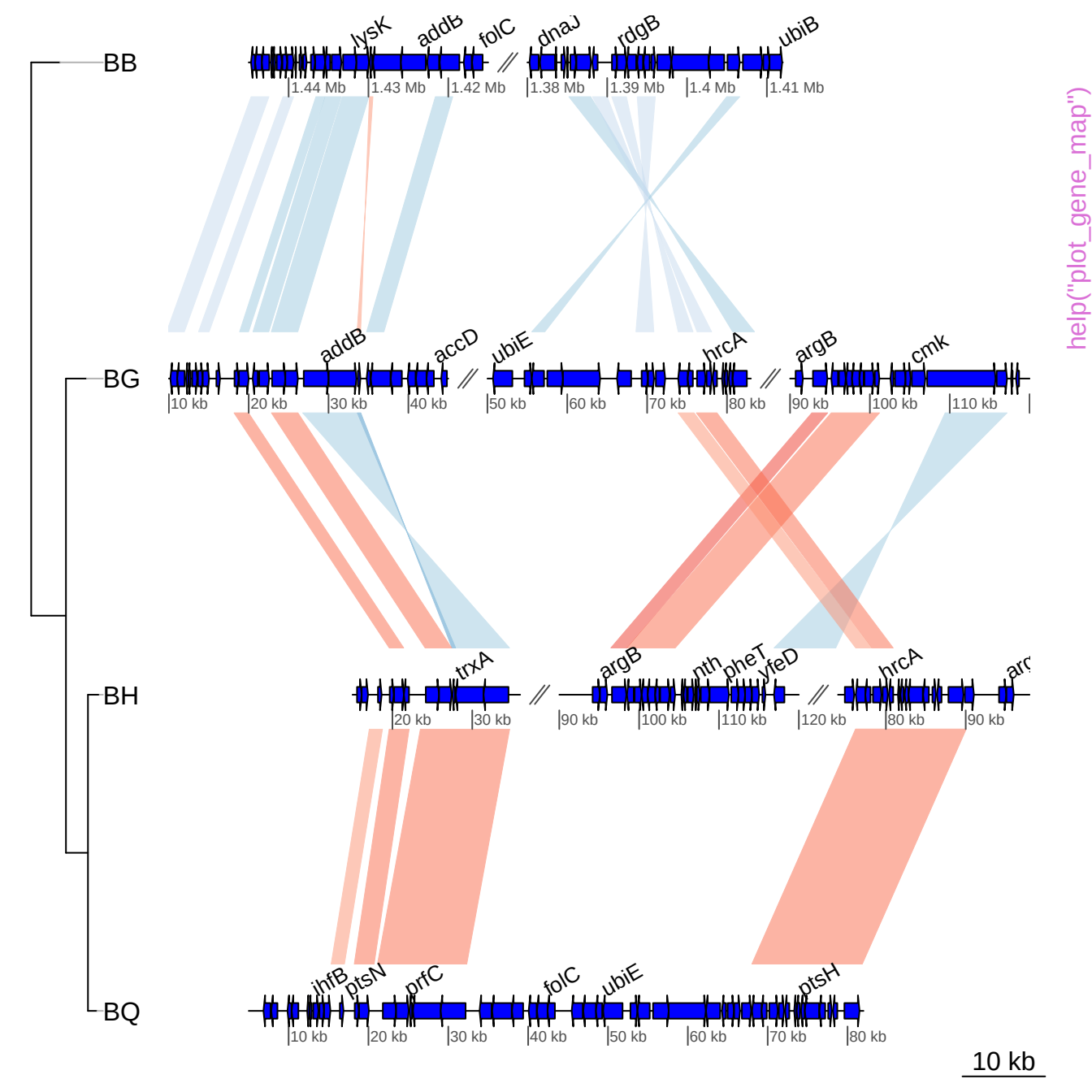
60 kb
70 kb
80 kb
20 kb
help("plot_gene_map")


BB
lysK
addB
folC
dnaJ
rdgB
ubiB
1.44 Mb
1.43 Mb
1.42 Mb
1.38 Mb
1.39 Mb
1.4 Mb
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BG
addB
accD
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hrcA
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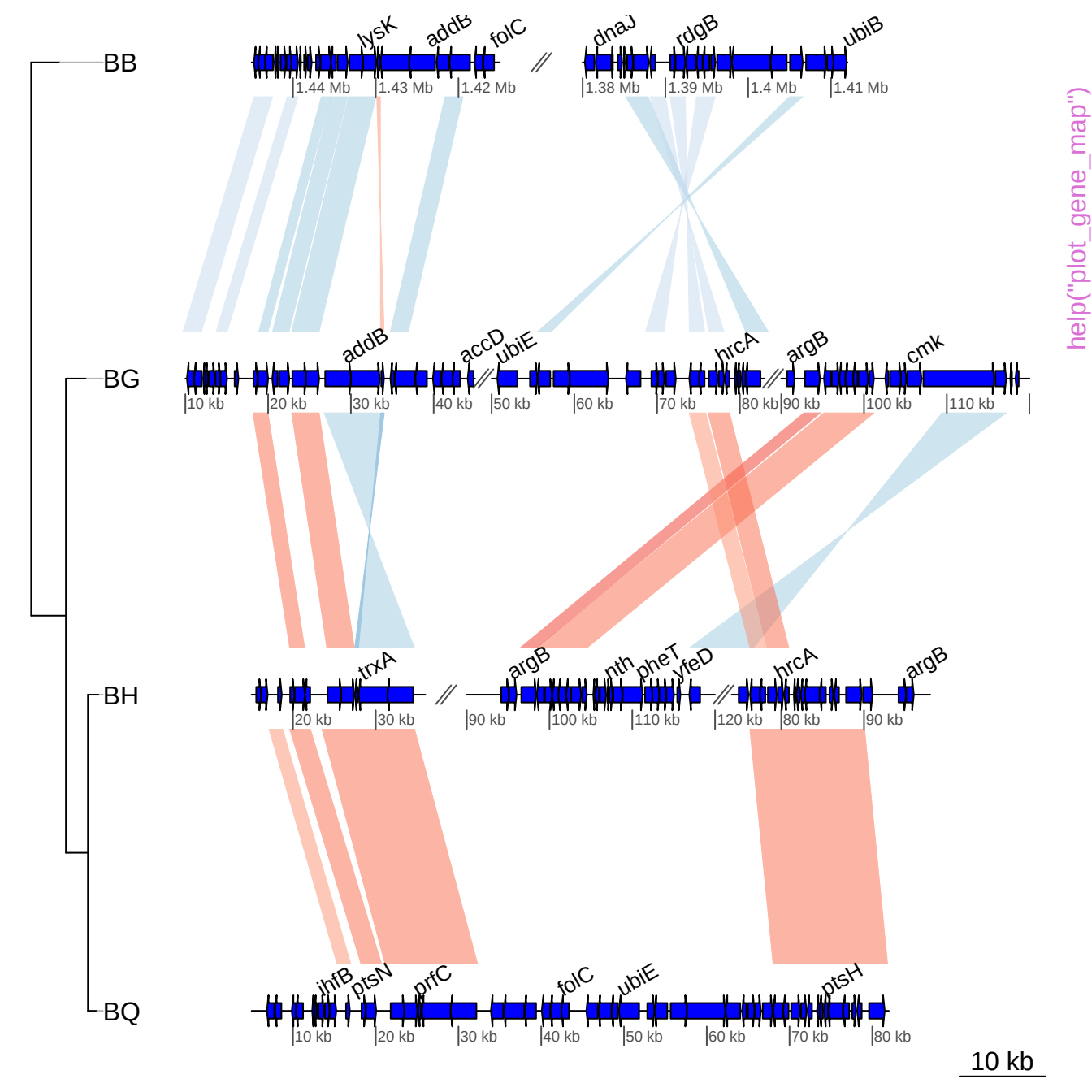
help("plot_gene_map")


BB
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ubiB
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BG
addB
accD
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argB
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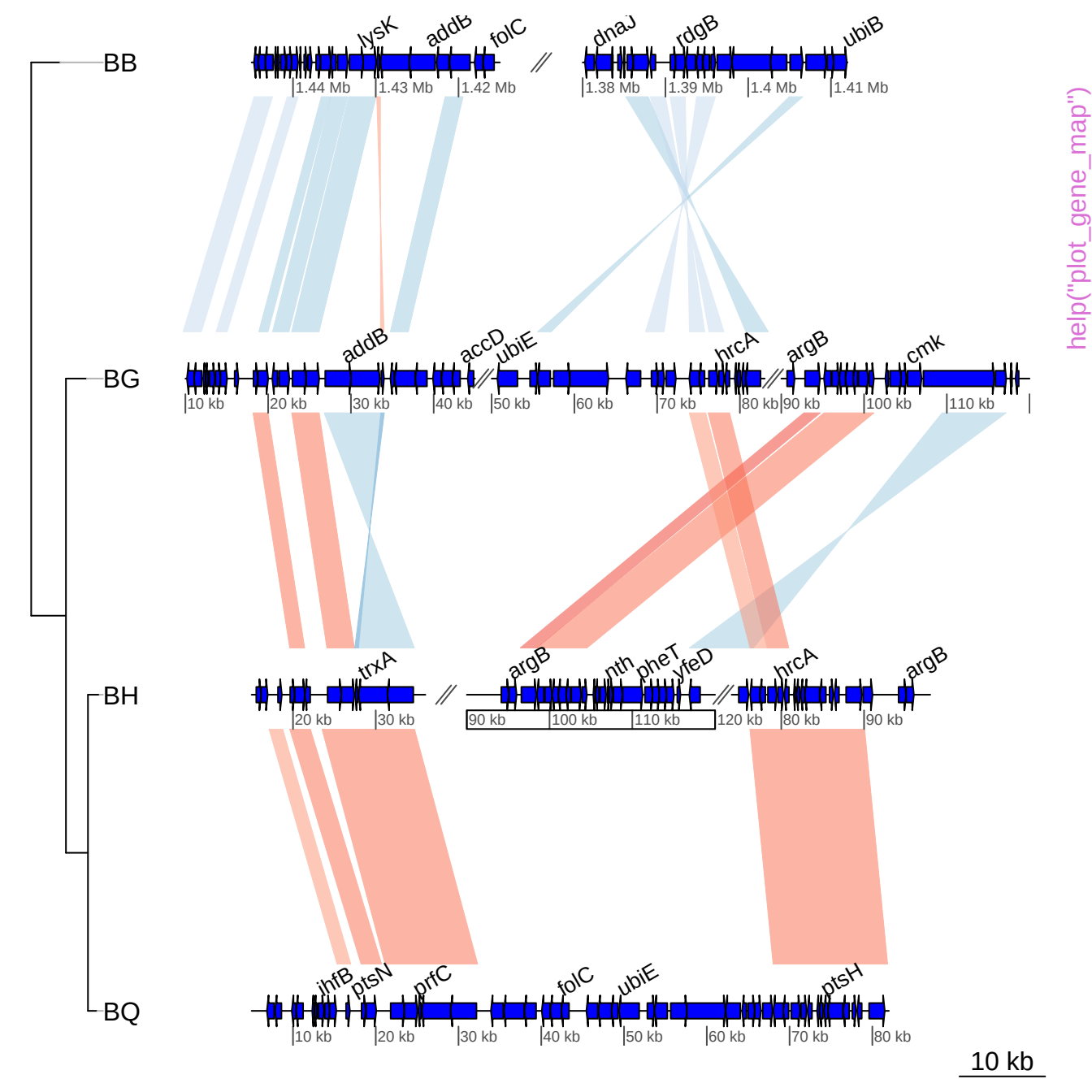
help("plot_gene_map")


BB
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dnaJ
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ubiB
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1.43 Mb
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ihfB
ptsN
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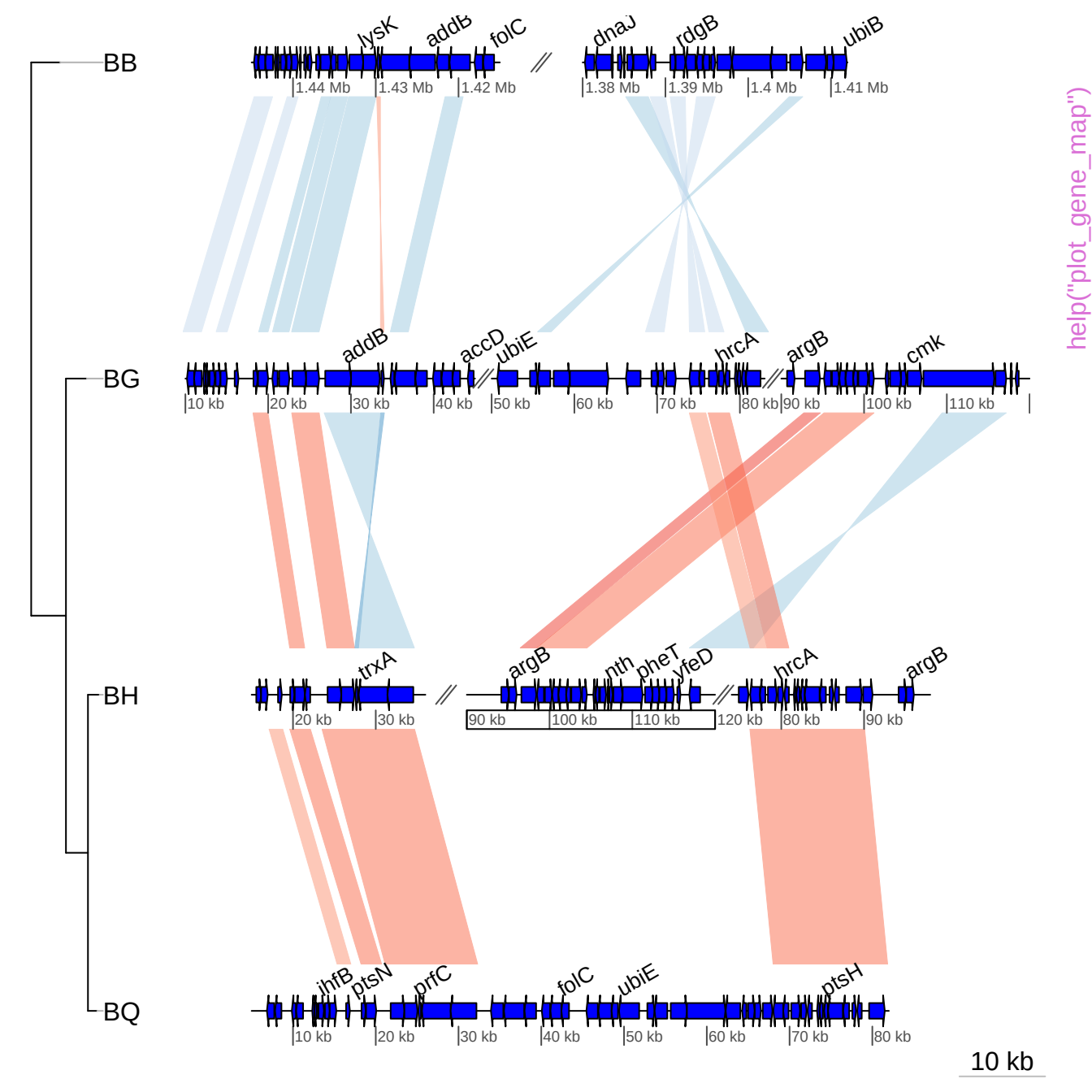
help("plot_gene_map")


BB
lysK
addB
folC
dnaJ
rdgB
ubiB
1.44 Mb
1.43 Mb
1.42 Mb
1.38 Mb
1.39 Mb
1.4 Mb
1.41 Mb
BG
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accD
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argB
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ptsN
prfC
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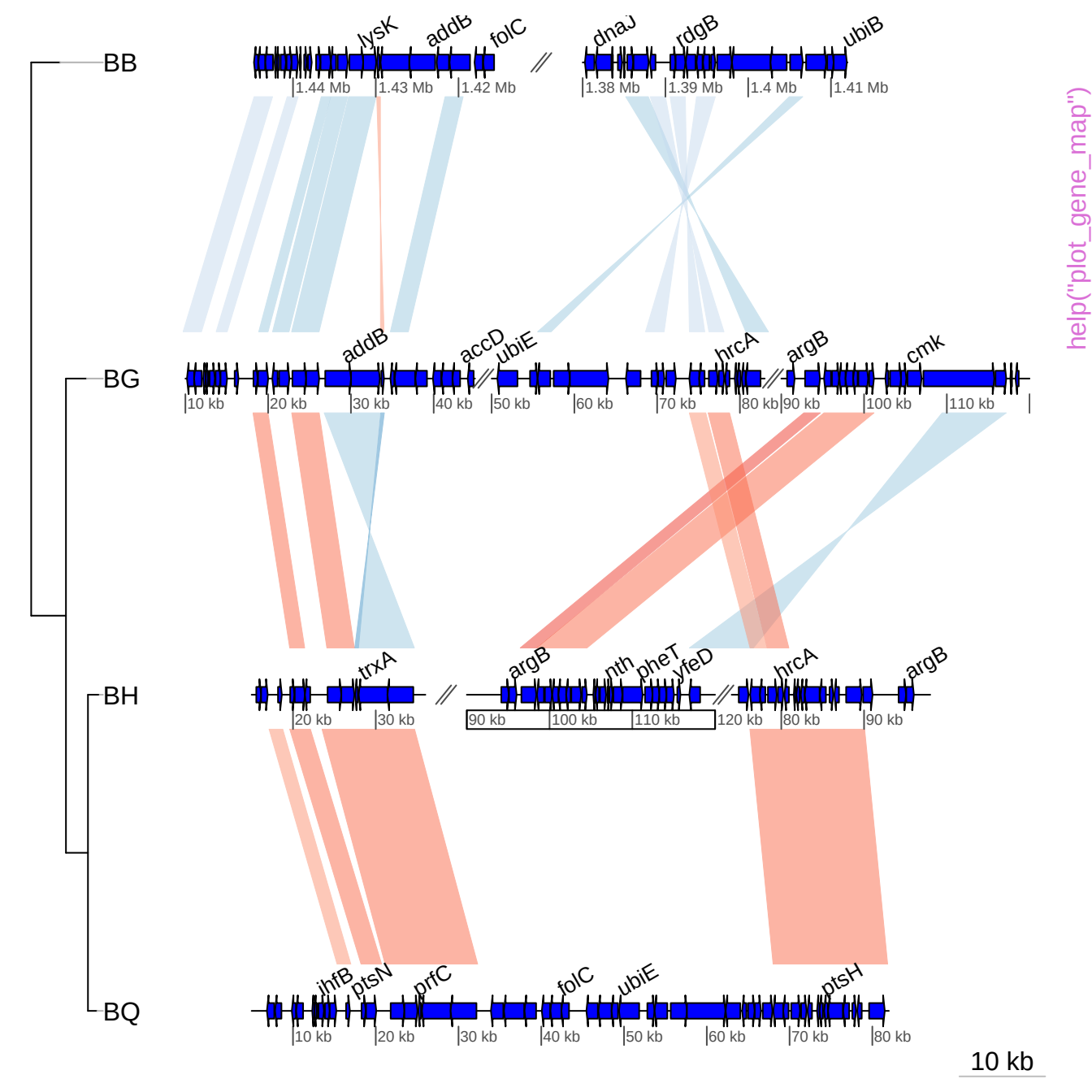
help("plot_gene_map")


BB
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addB
folC
dnaJ
rdgB
ubiB
1.44 Mb
1.43 Mb
1.42 Mb
1.38 Mb
1.39 Mb
1.4 Mb
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BG
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accD
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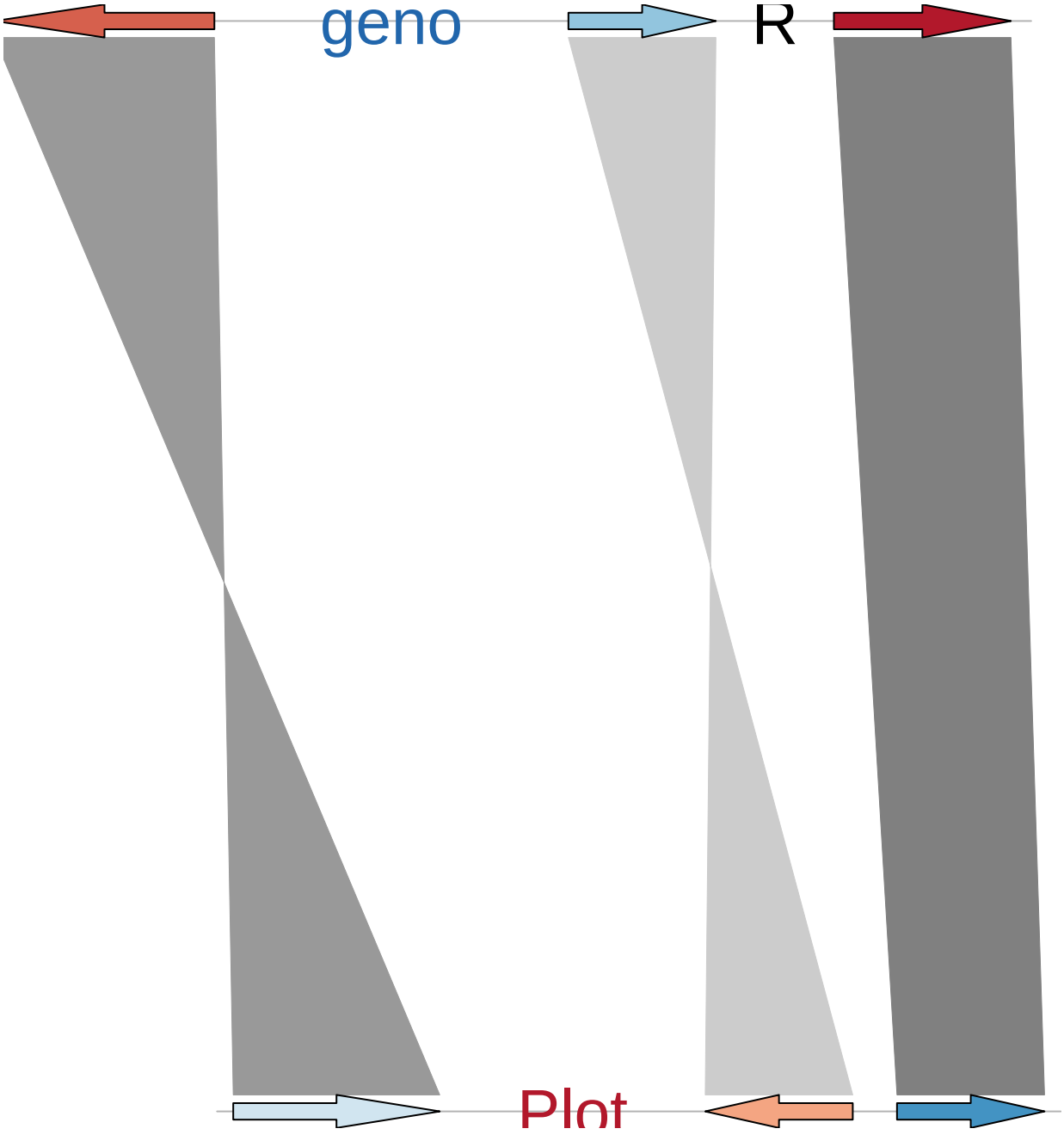
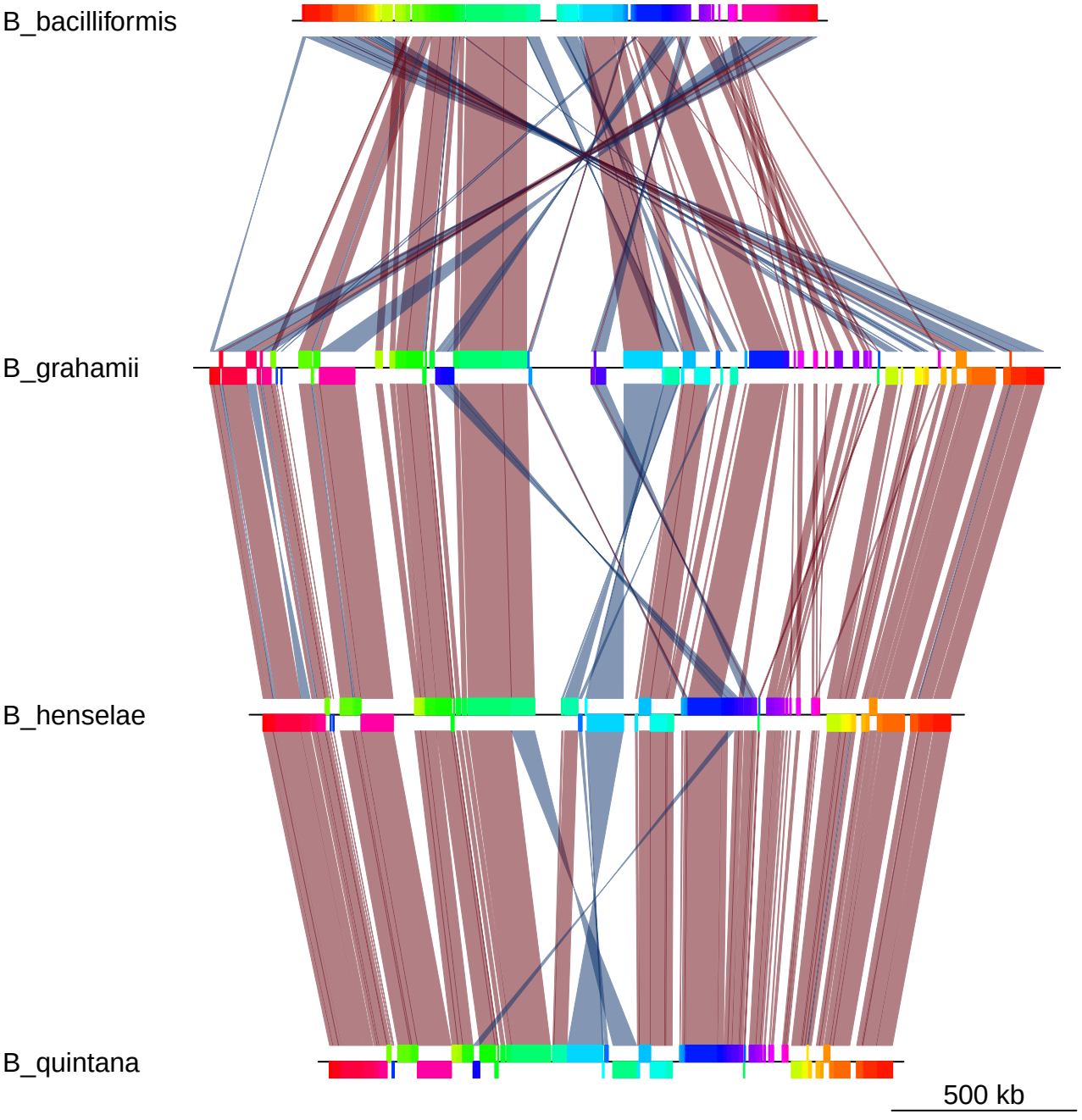
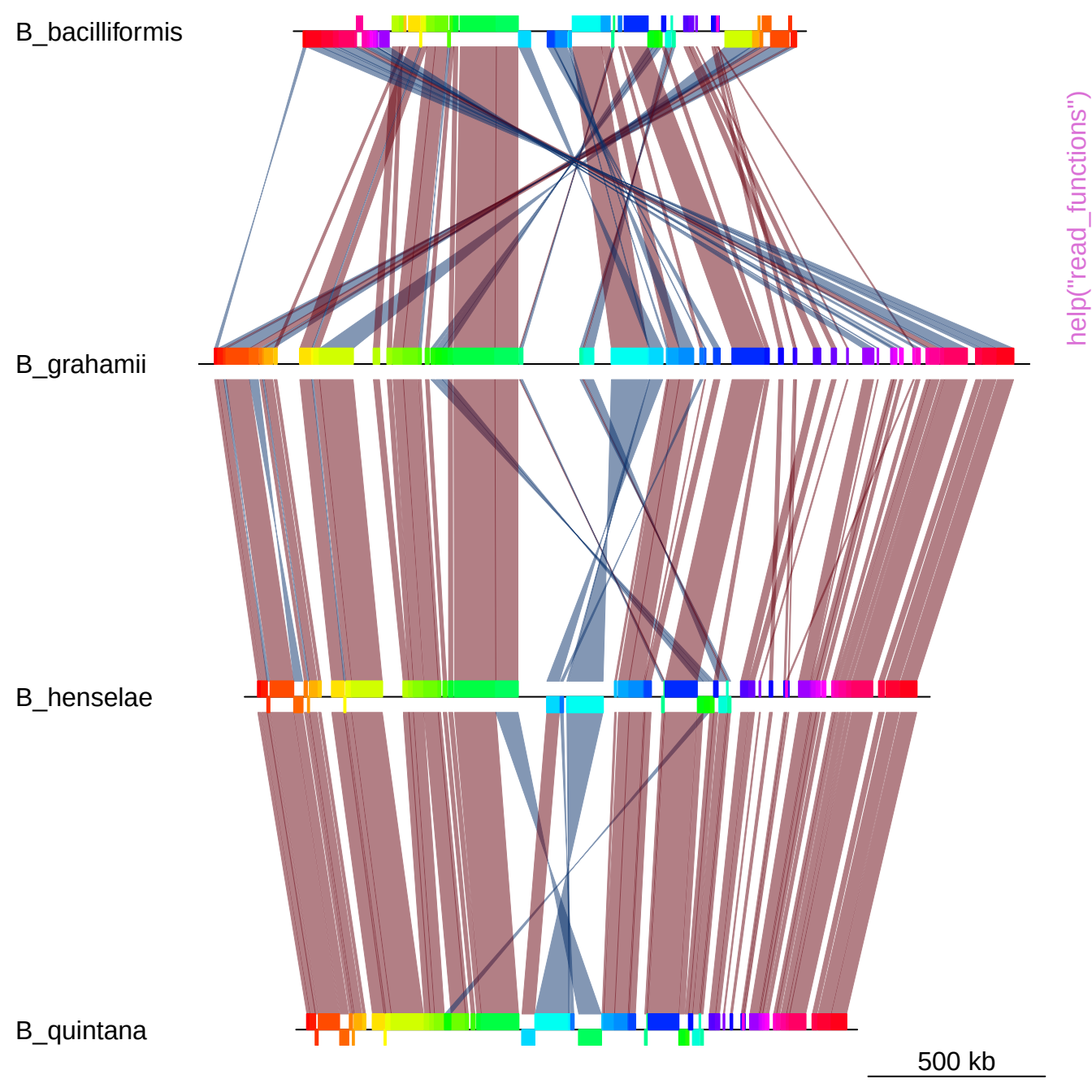
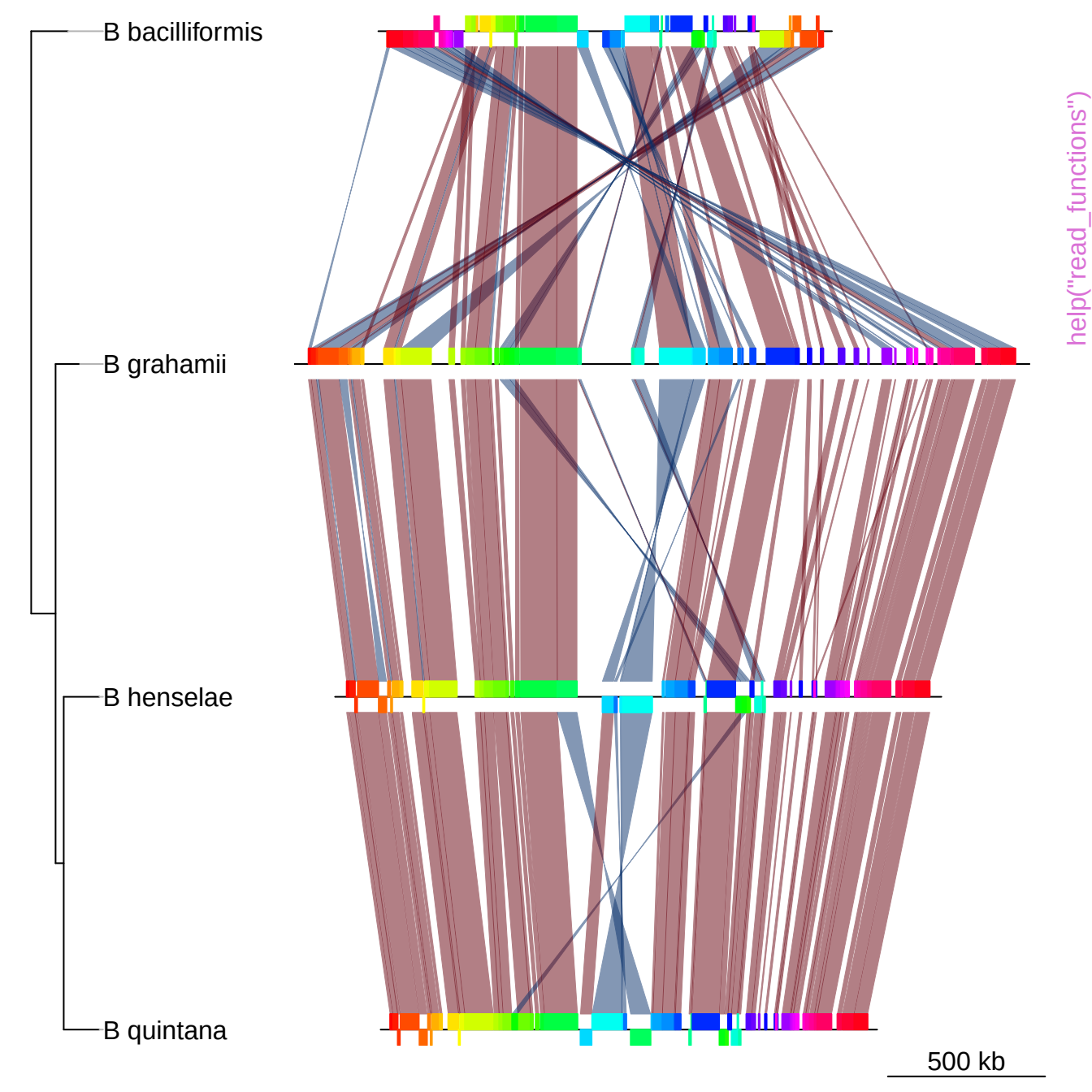
help("plot_gene_map")

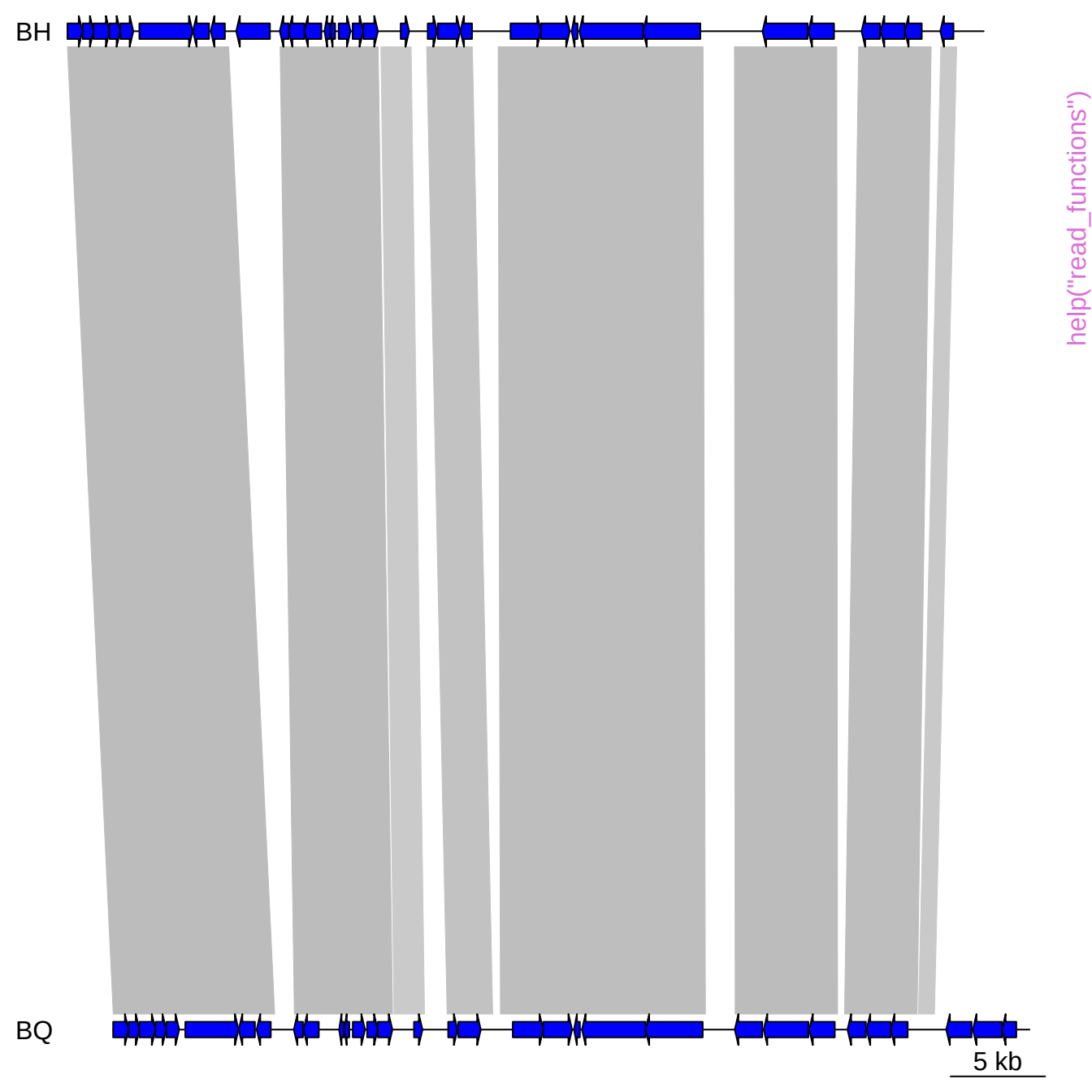


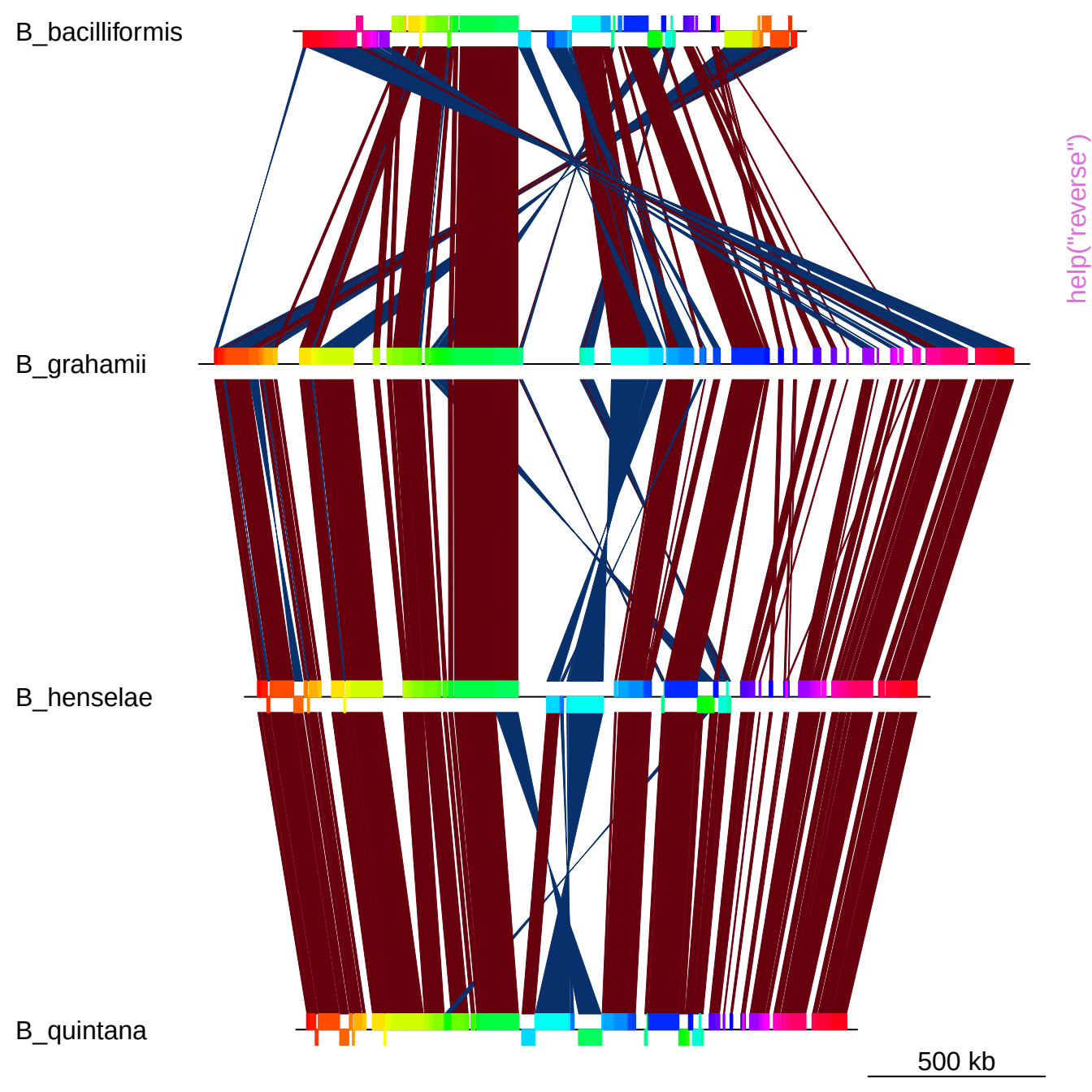
help("plot_gene_map")

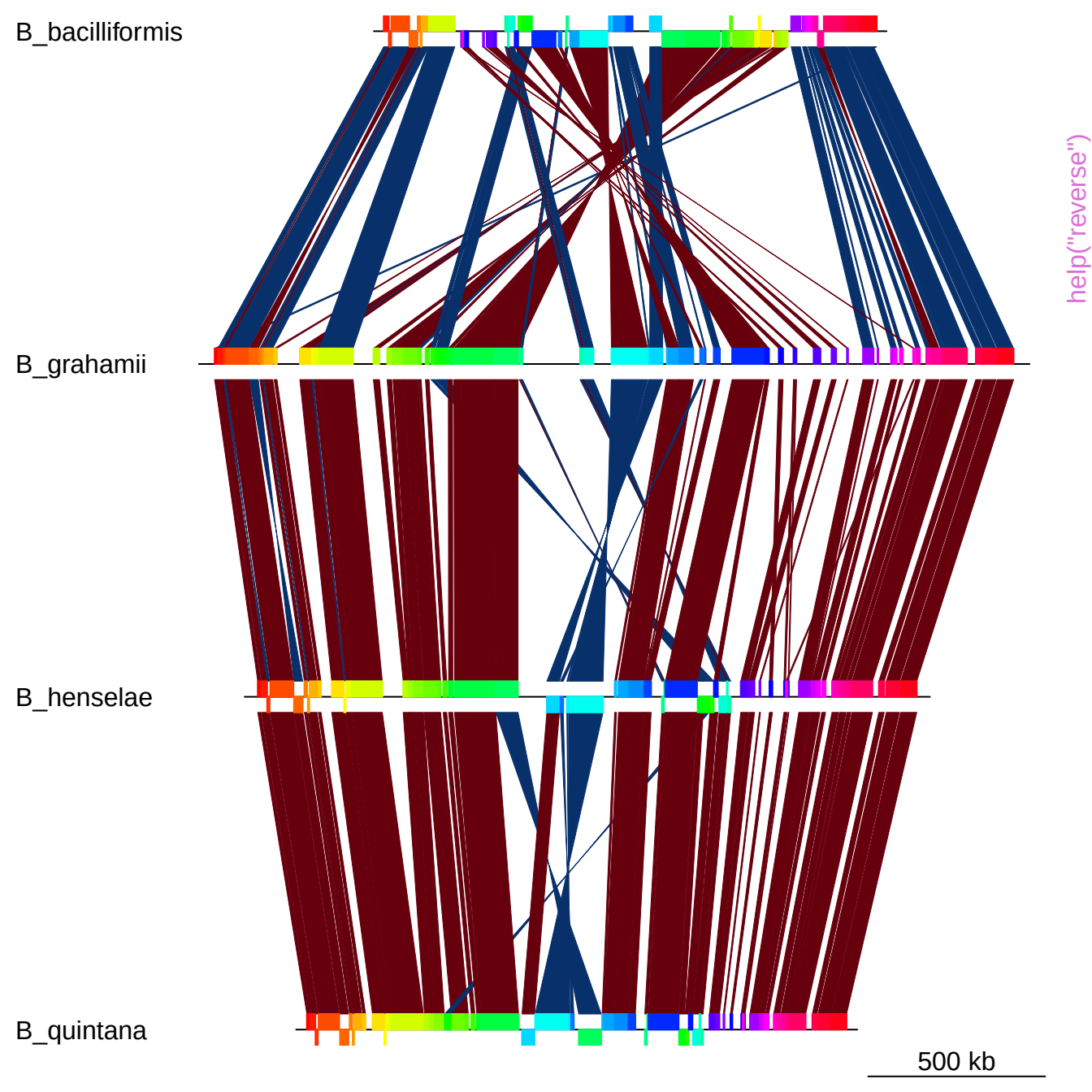

B_bacilliformis
B_grahamii
B_henselae
B_quintana
500 kb
help("read_functions")

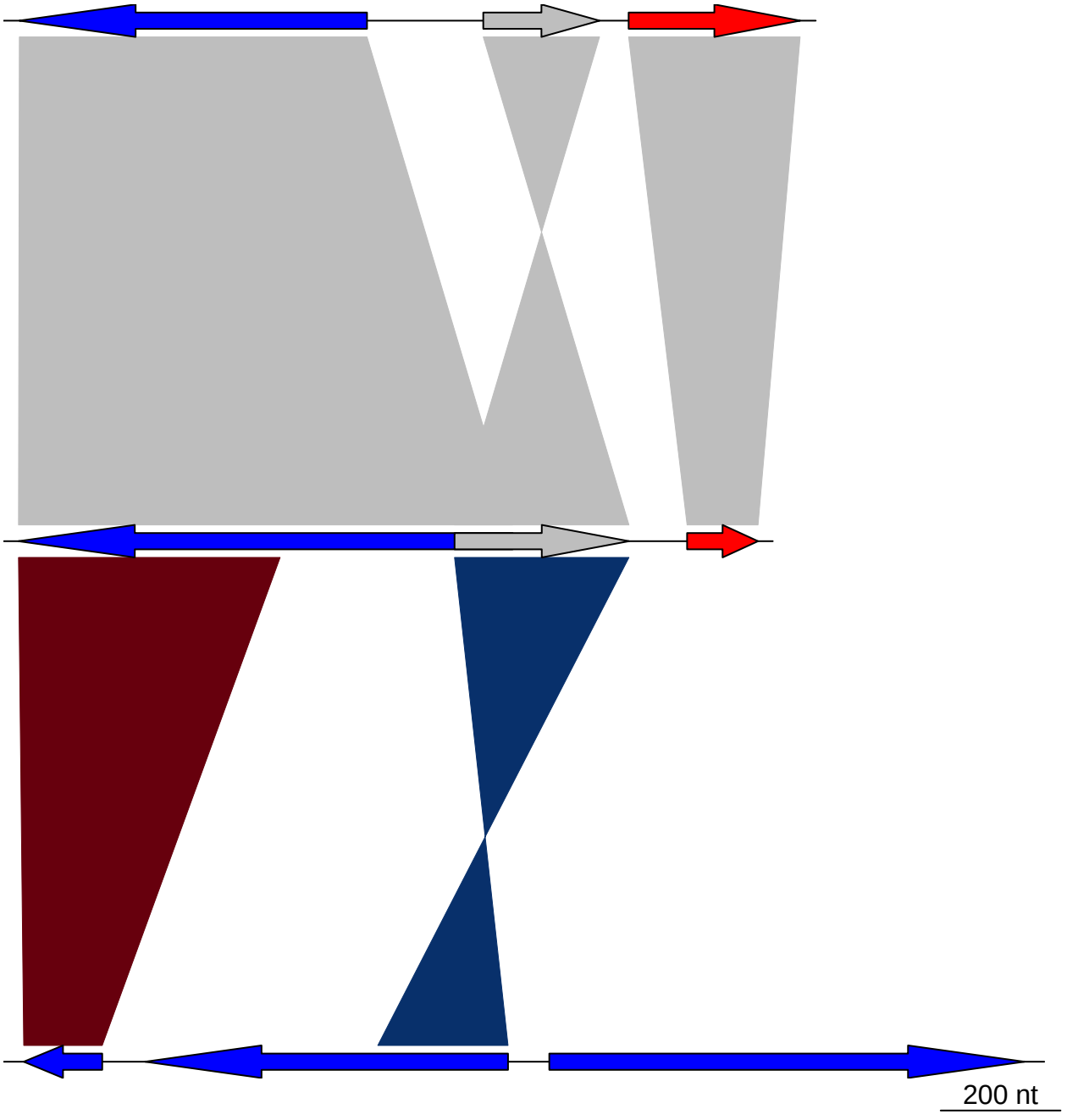

B_bacilliformis
B_grahamii
B_henselae
B_quintana
500 kb
help("read_functions")

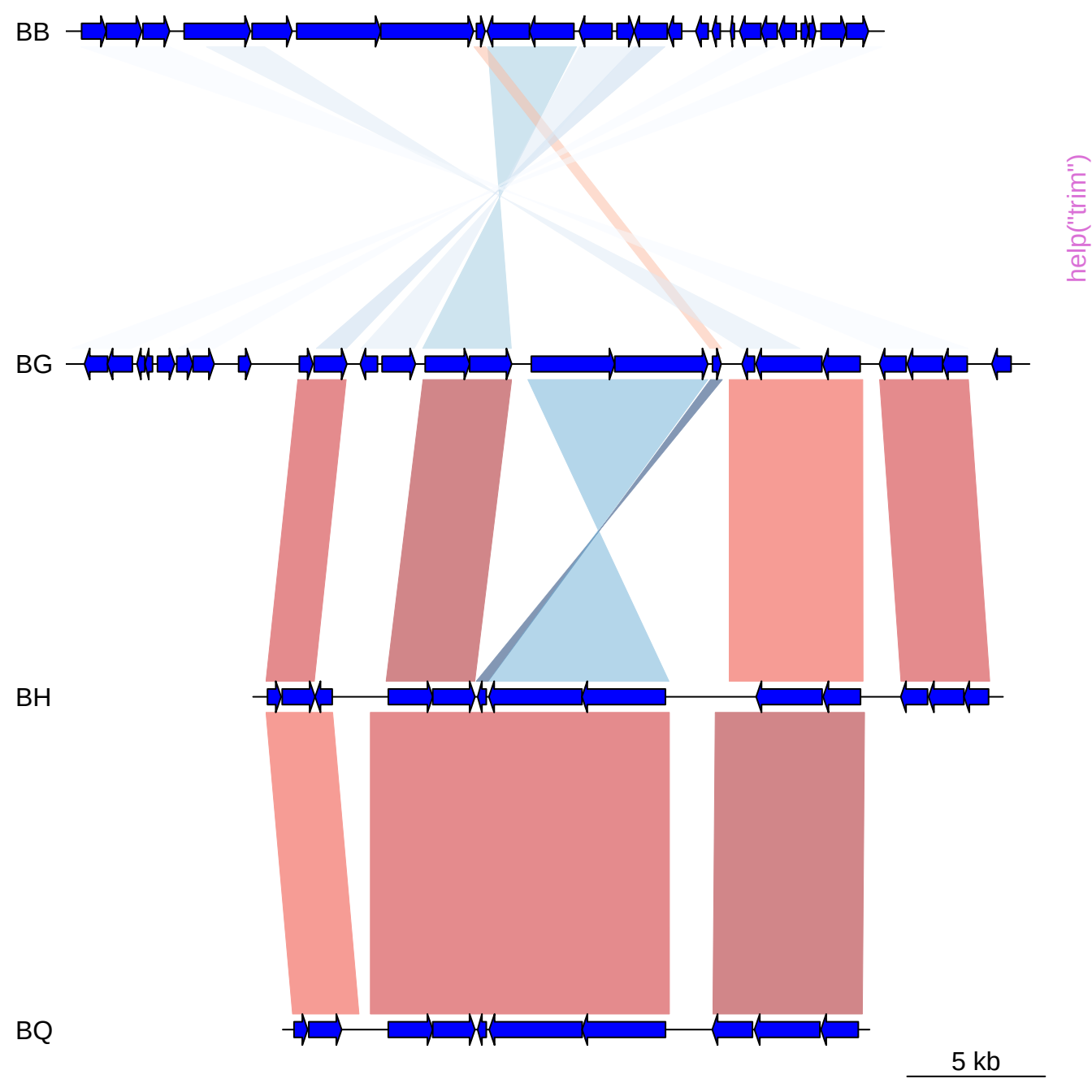

B bacilliformis
B grahamii
B henselae
B quintana
500 kb
help("read_functions")


BH
BQ
5 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")
200 nt


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