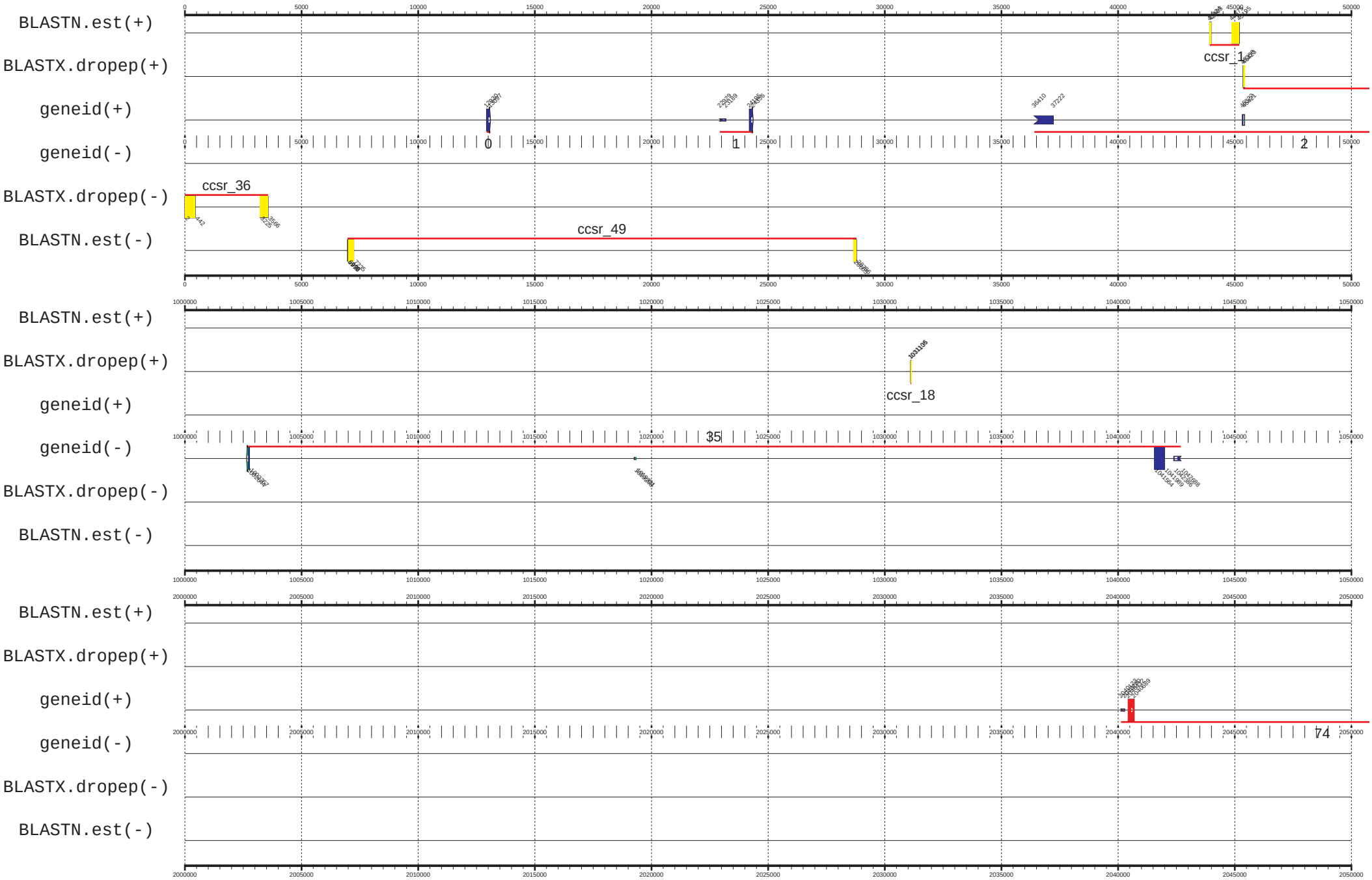
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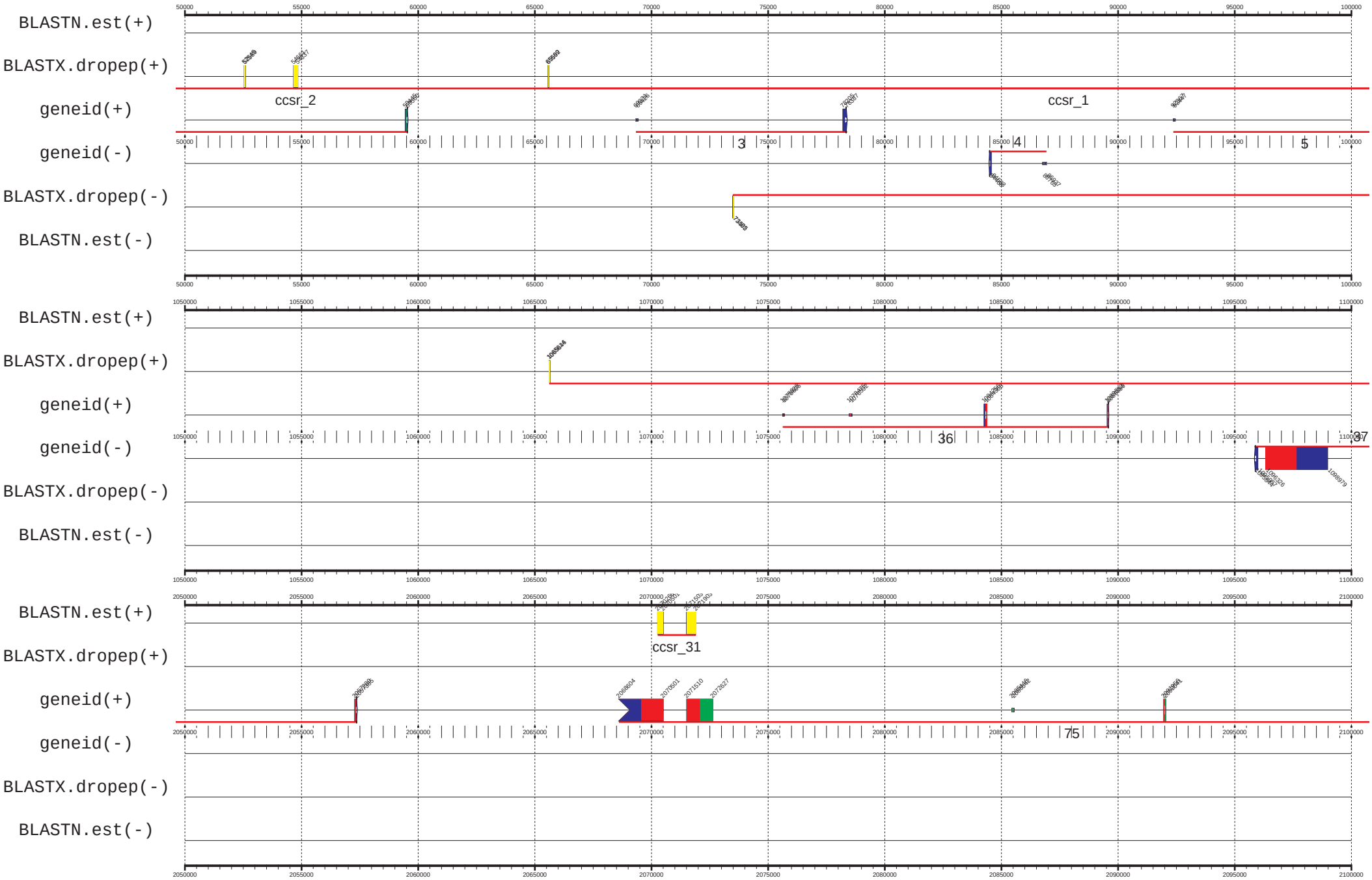
BLASTX.dropep(+)
geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
ccsr_1
ccsr_36
ccsr_49
ccsr_18
0
1
2
35
74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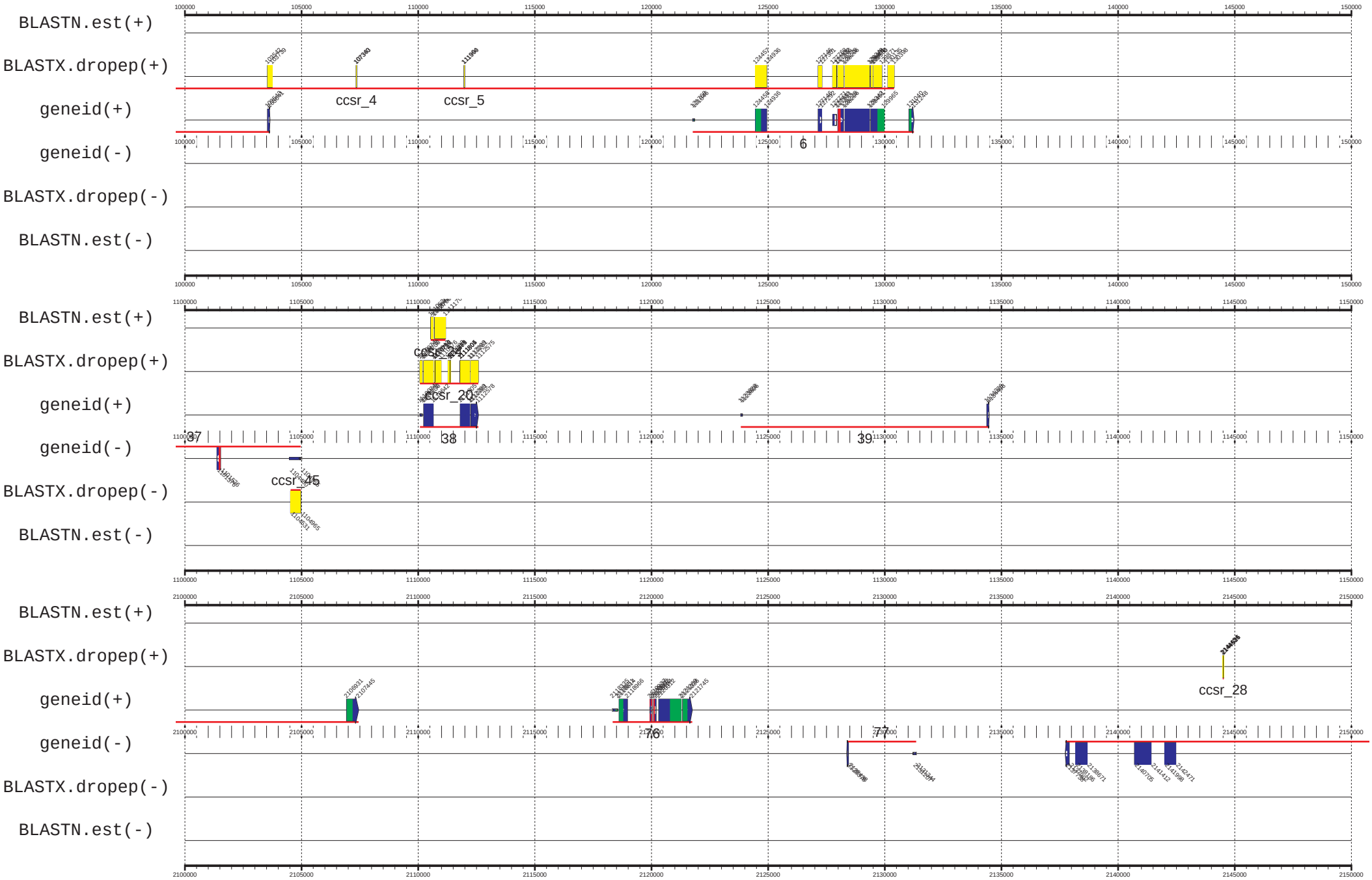
geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
ccsr_2
ccsr_1
3
4
5
36
37
ccsr_31
75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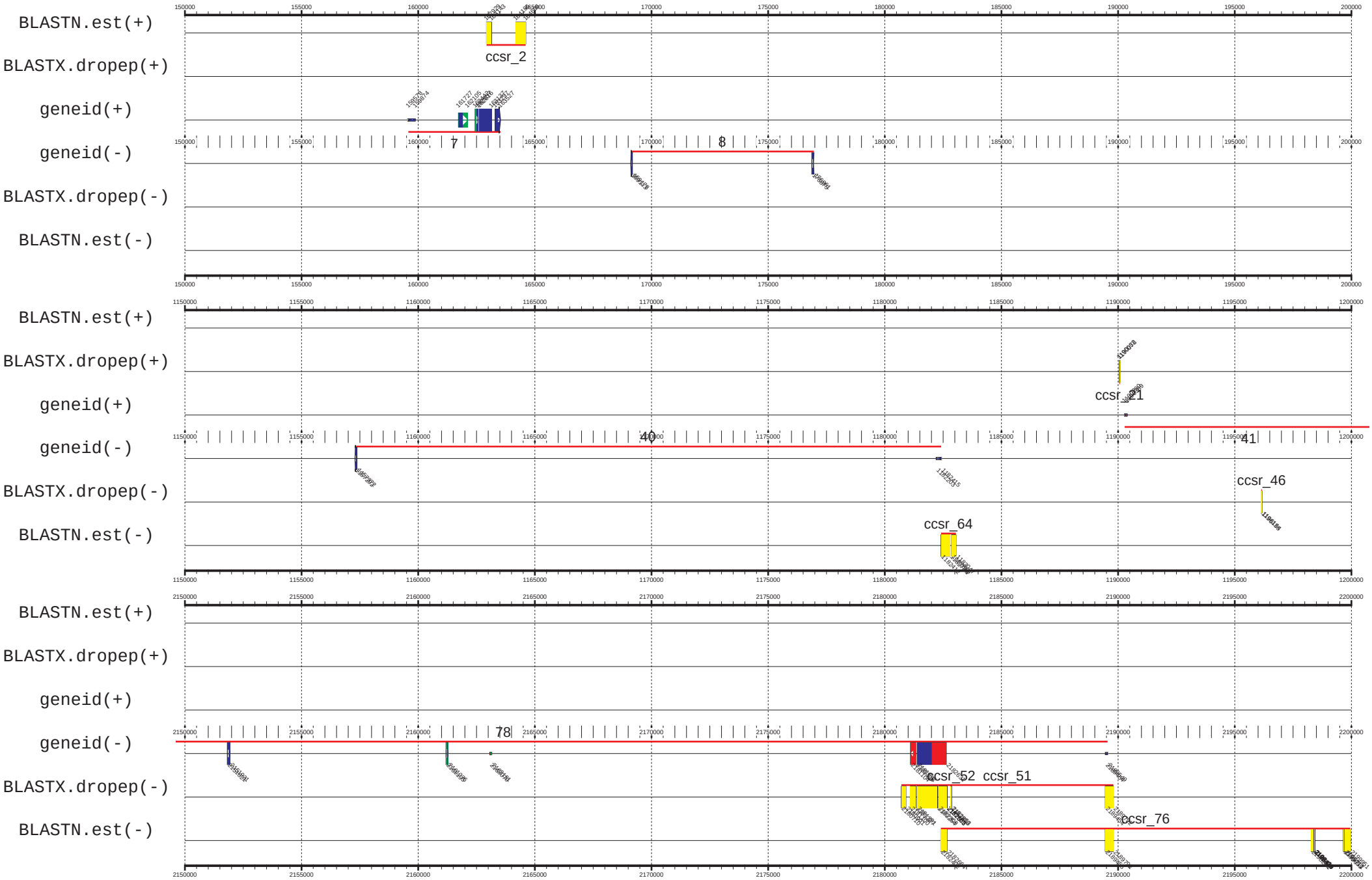
geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
ccsr_4
ccsr_5
6
ccsr_20
37
38
39
ccsr_45
76
77
ccsr_28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geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
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8
ccsr_21
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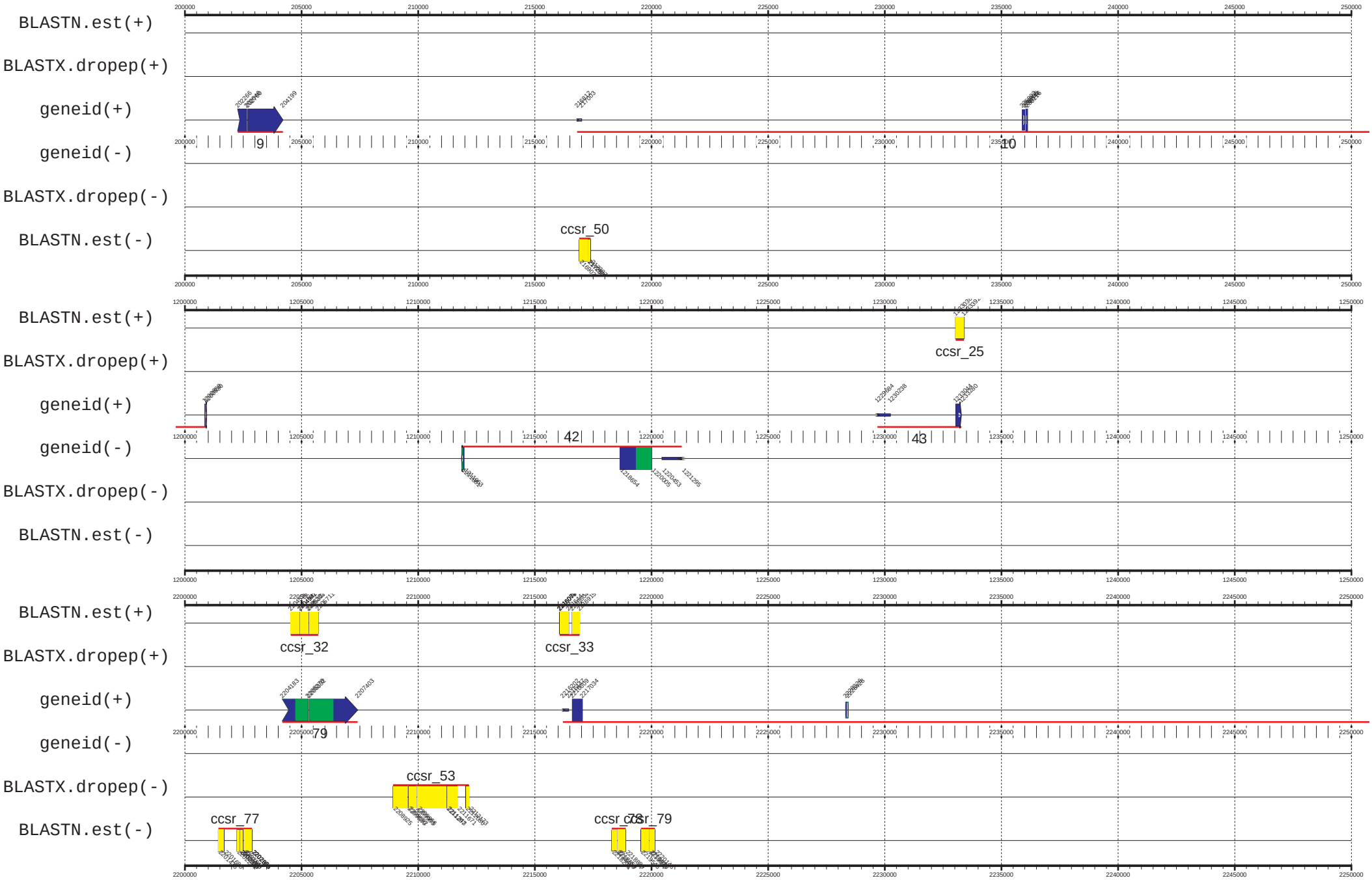
41
ccsr_46
ccsr_64
78
ccsr_52
ccsr_51
ccsr_76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geneid(+)
geneid(-)
BLASTX.dropep(-)
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9
10
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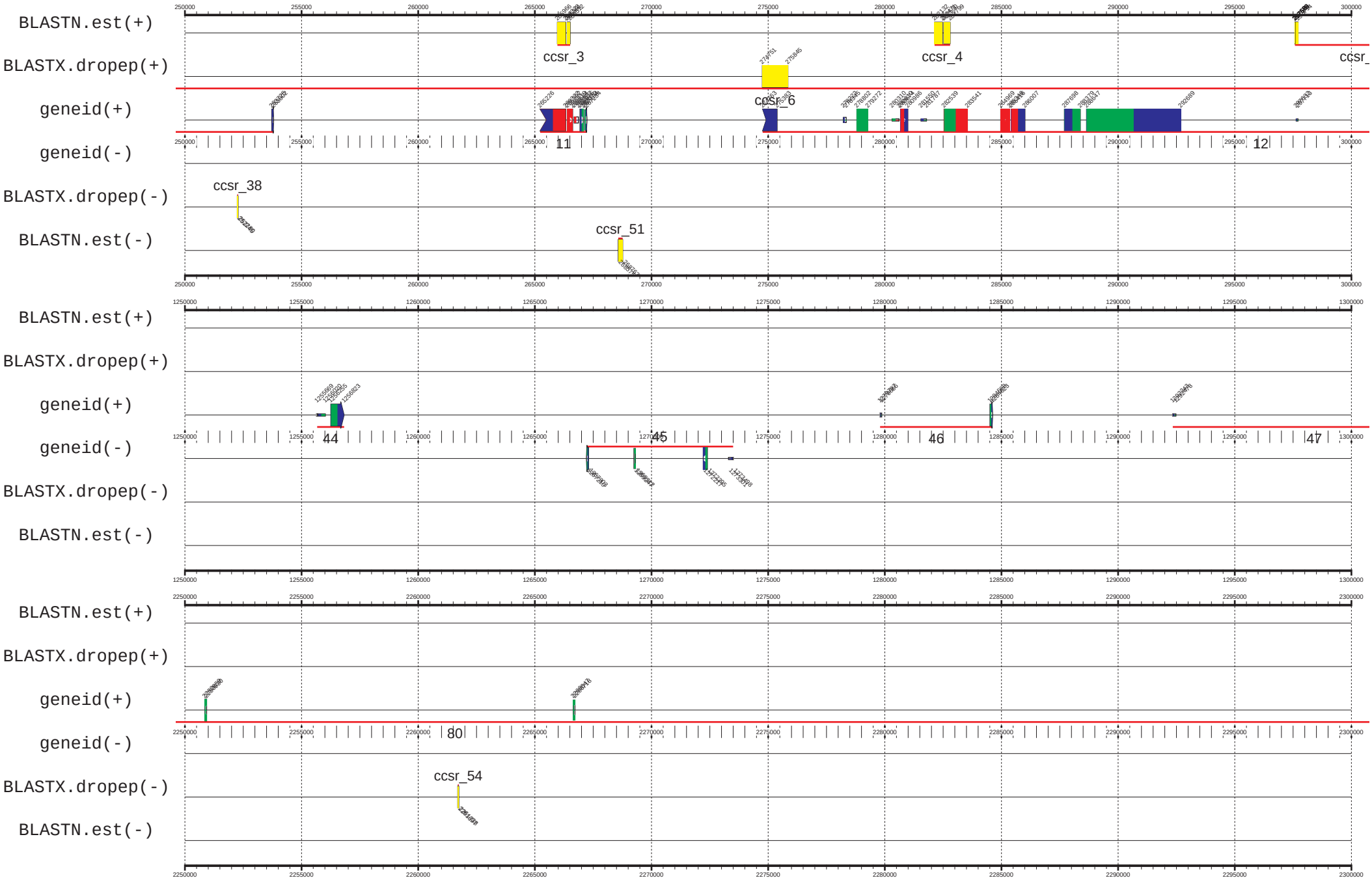
ccsr_25
42
43
ccsr_32
ccsr_33
79
ccsr_53
ccsr_77
ccsr_78
ccsr_79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geneid(+)
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ccsr_3
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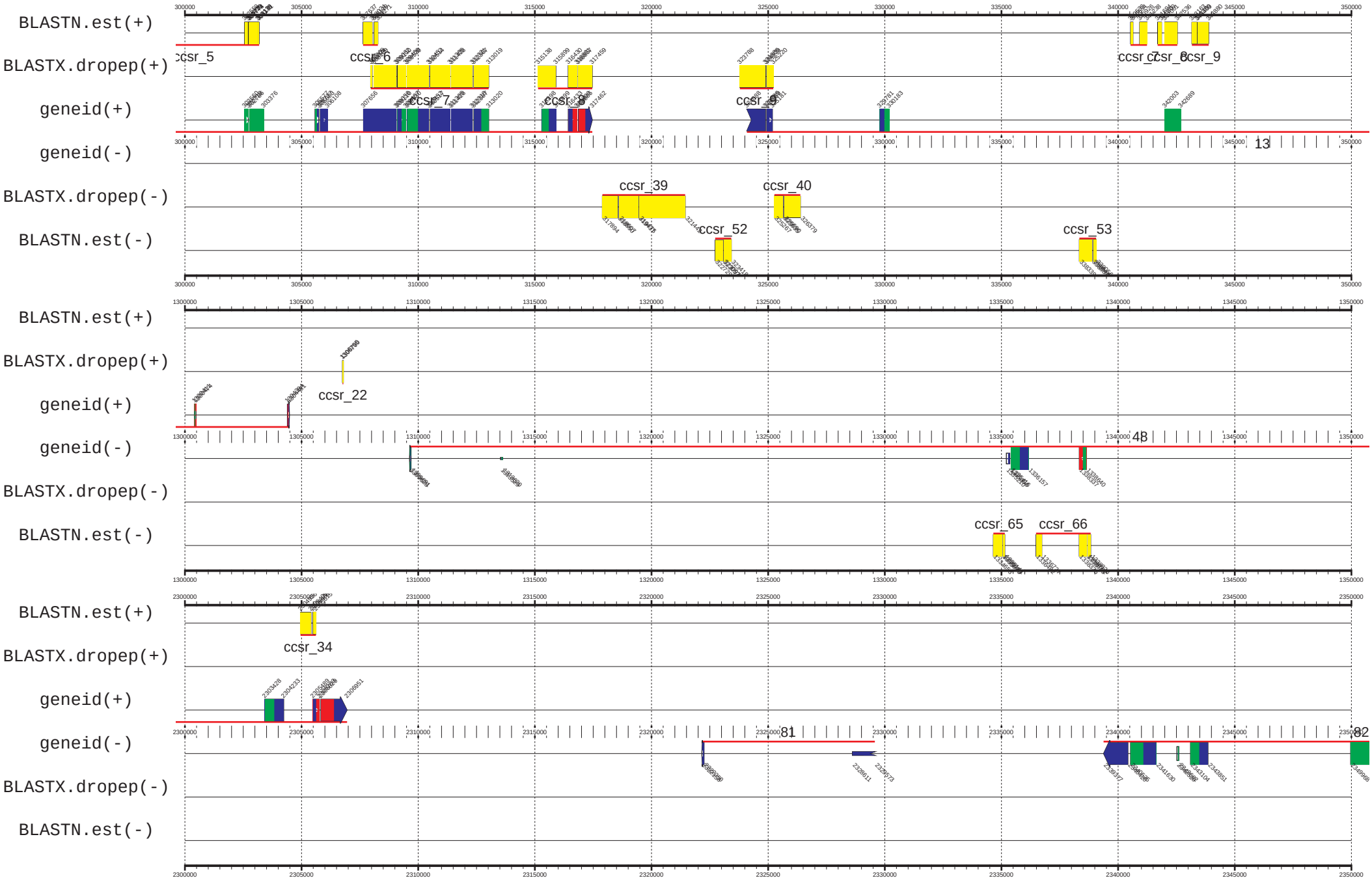
11
12
44
45
46
47
80
ccsr_54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geneid(+)
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BLASTX.dropep(-)
BLASTN.est(-)
ccsr_5
ccsr_6
ccsr_7
ccsr_8
ccsr_9
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ccsr_52
ccsr_53
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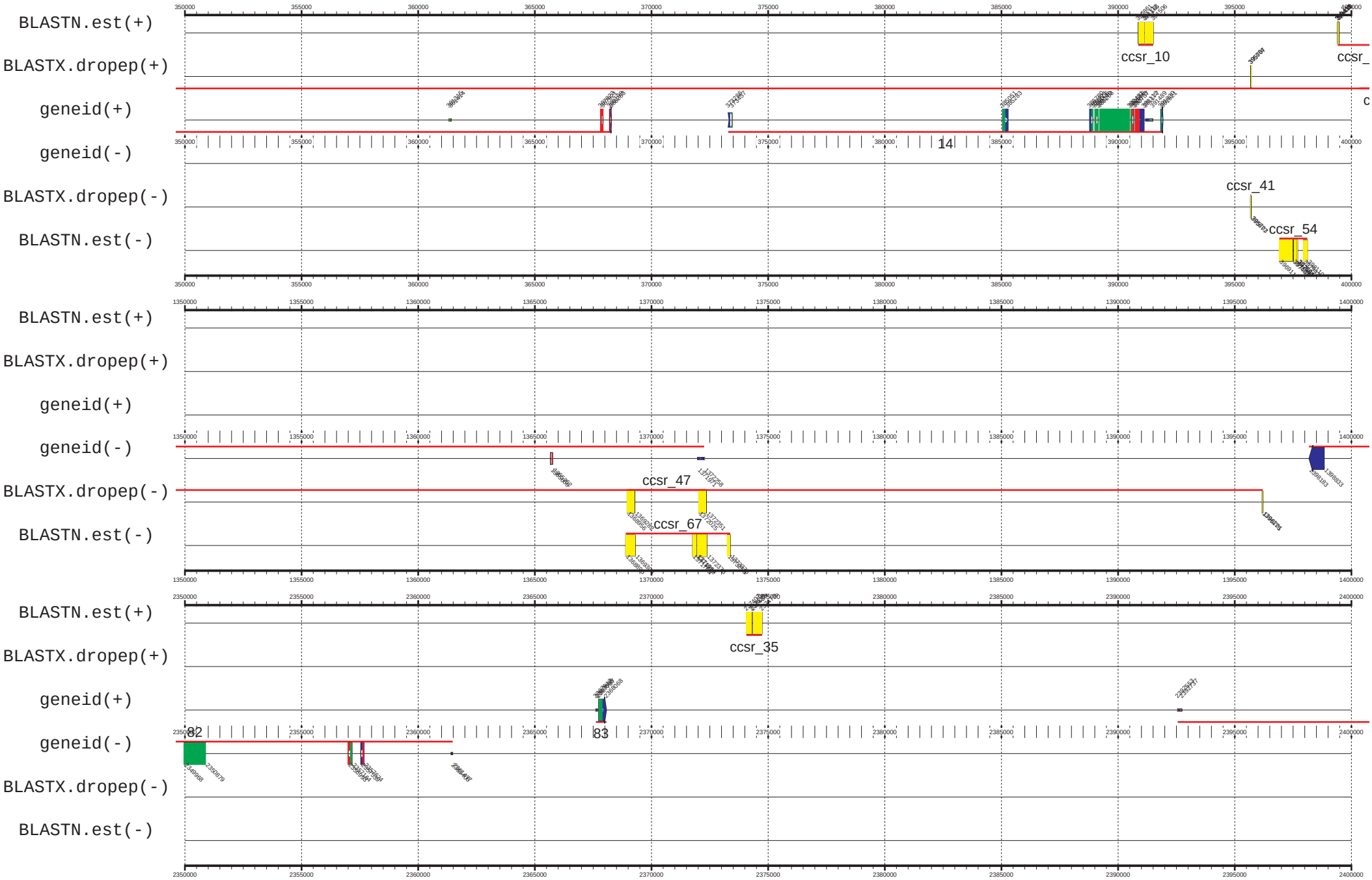
ccsr_22
48
ccsr_65
ccsr_66
ccsr_34
81
82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geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
ccsr_10
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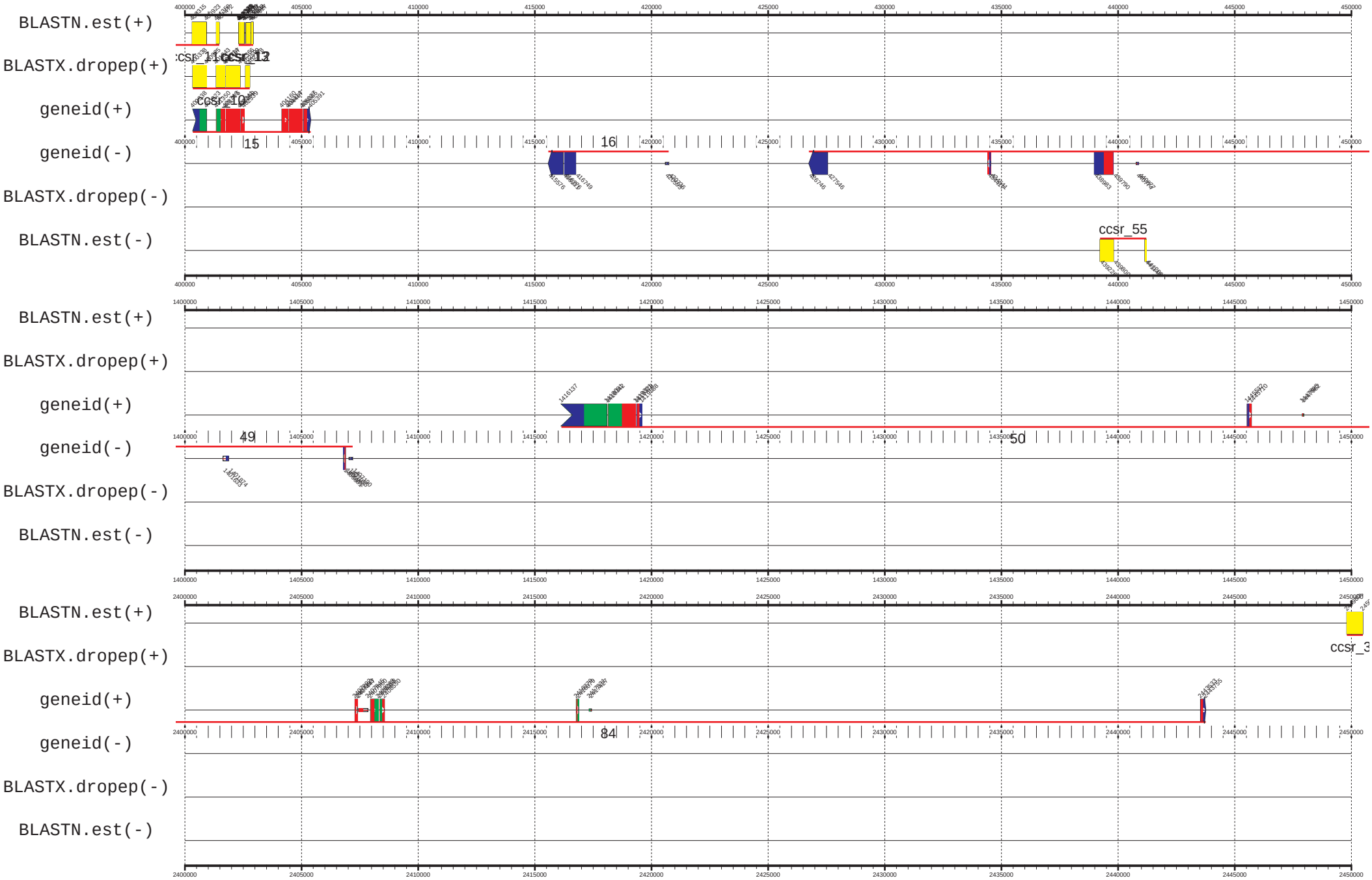
ccsr_35
14
82
83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geneid(+)
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BLASTX.dropep(-)
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ccsr_11
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ccsr_10
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16
ccsr_55
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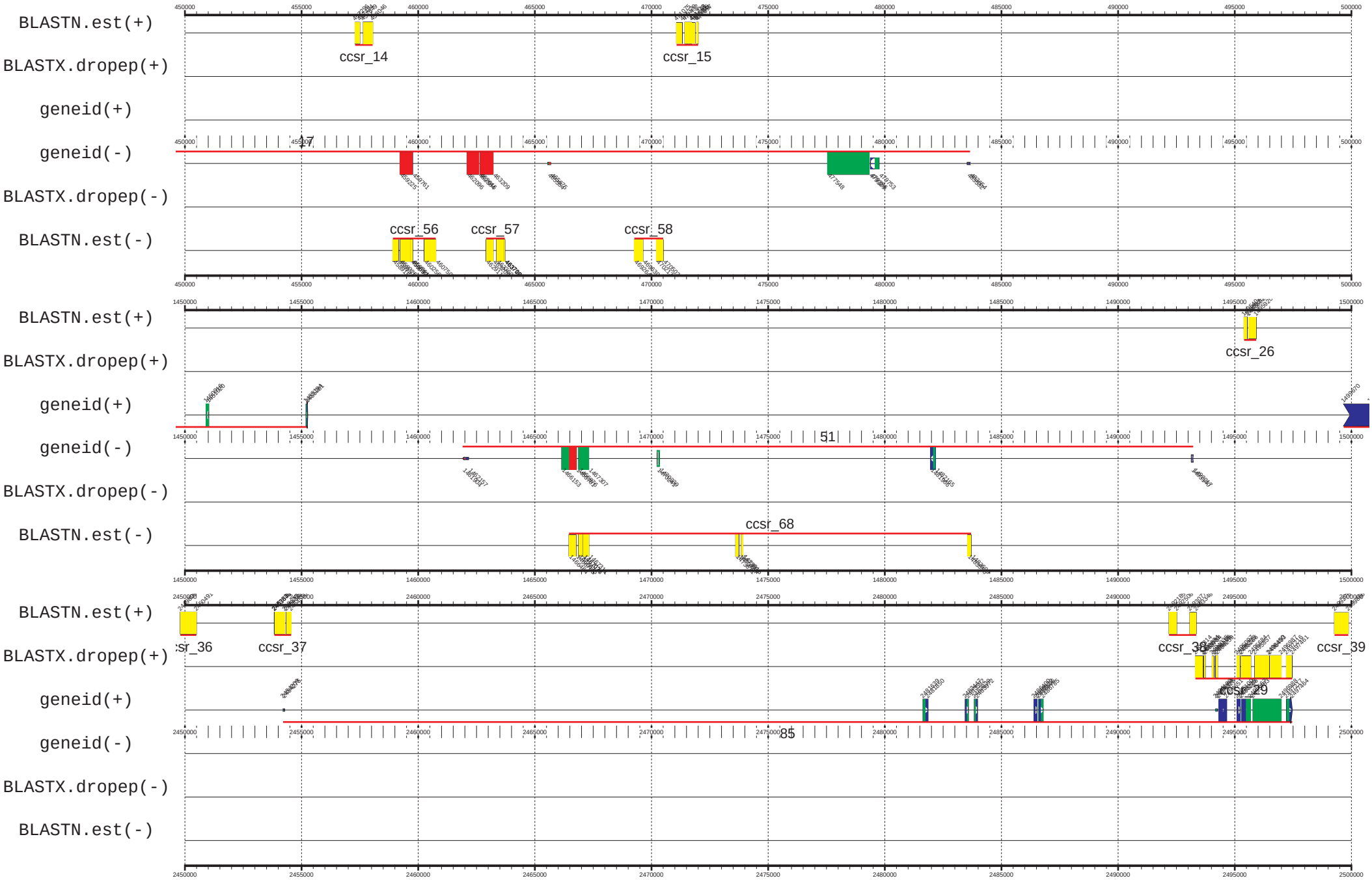
50
84
ccsr_3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BLASTX.dropep(+)
geneid(+)
geneid(-)
BLASTX.dropep(-)
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ccsr_14
ccsr_15
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ccsr_58
ccsr_26
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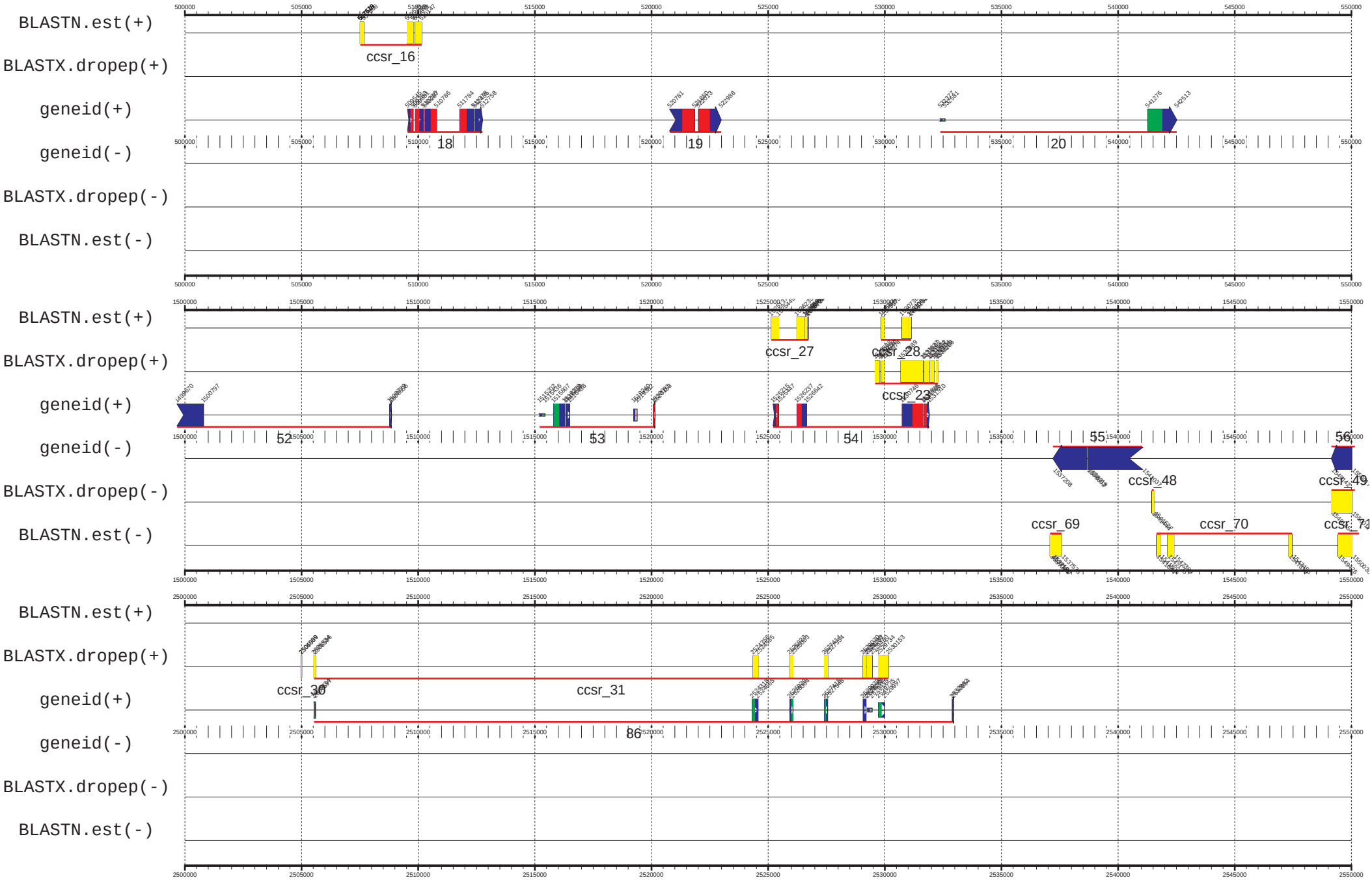
ccsr_36
ccsr_37
ccsr_38
ccsr_39
ccsr_29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BLASTX.dropep(+)
geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
ccsr_16
18
19
20
ccsr_27
ccsr_28
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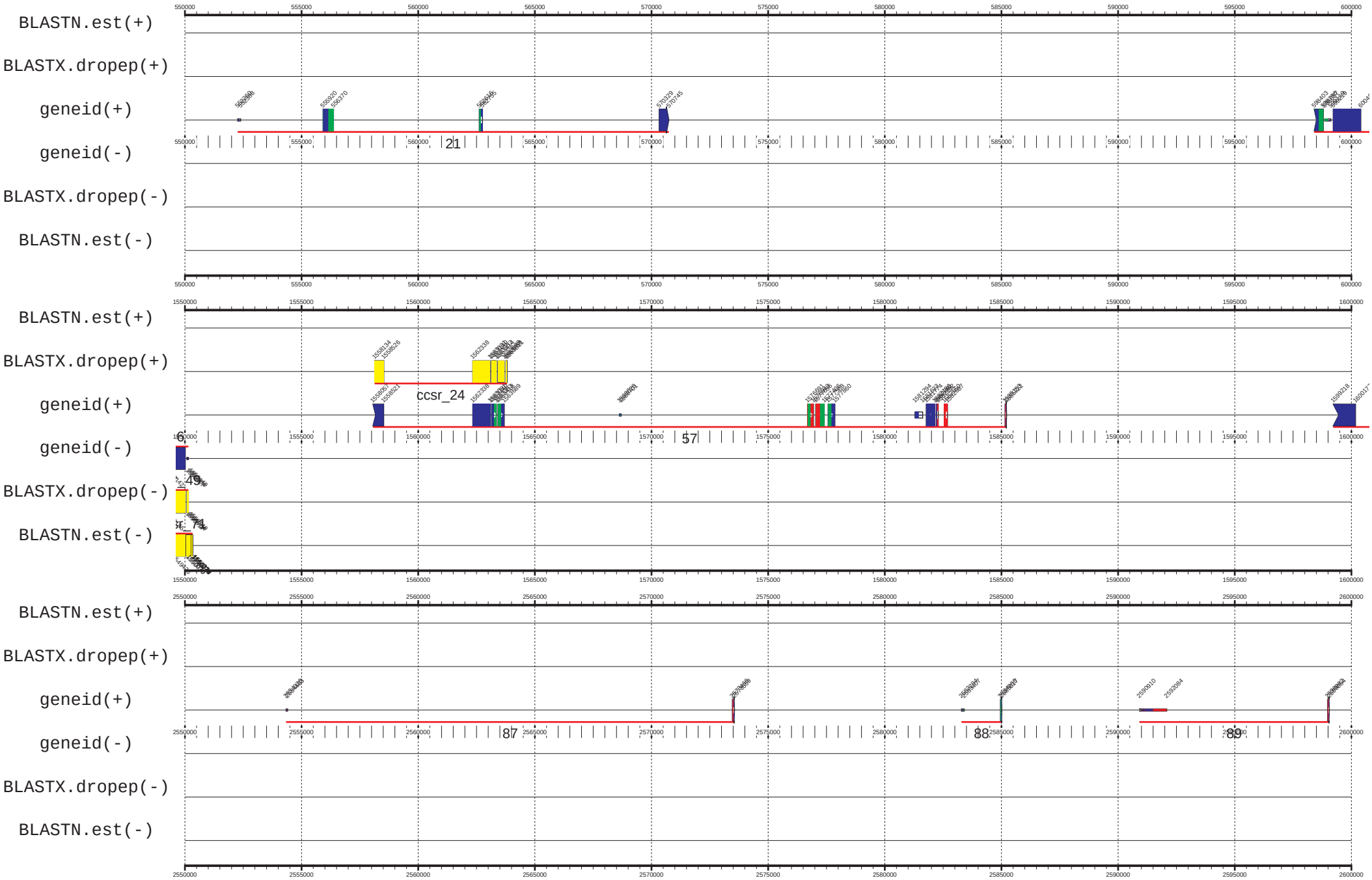
54
55
56
ccsr_48
ccsr_49
ccsr_69
ccsr_70
ccsr_30
ccsr_31
86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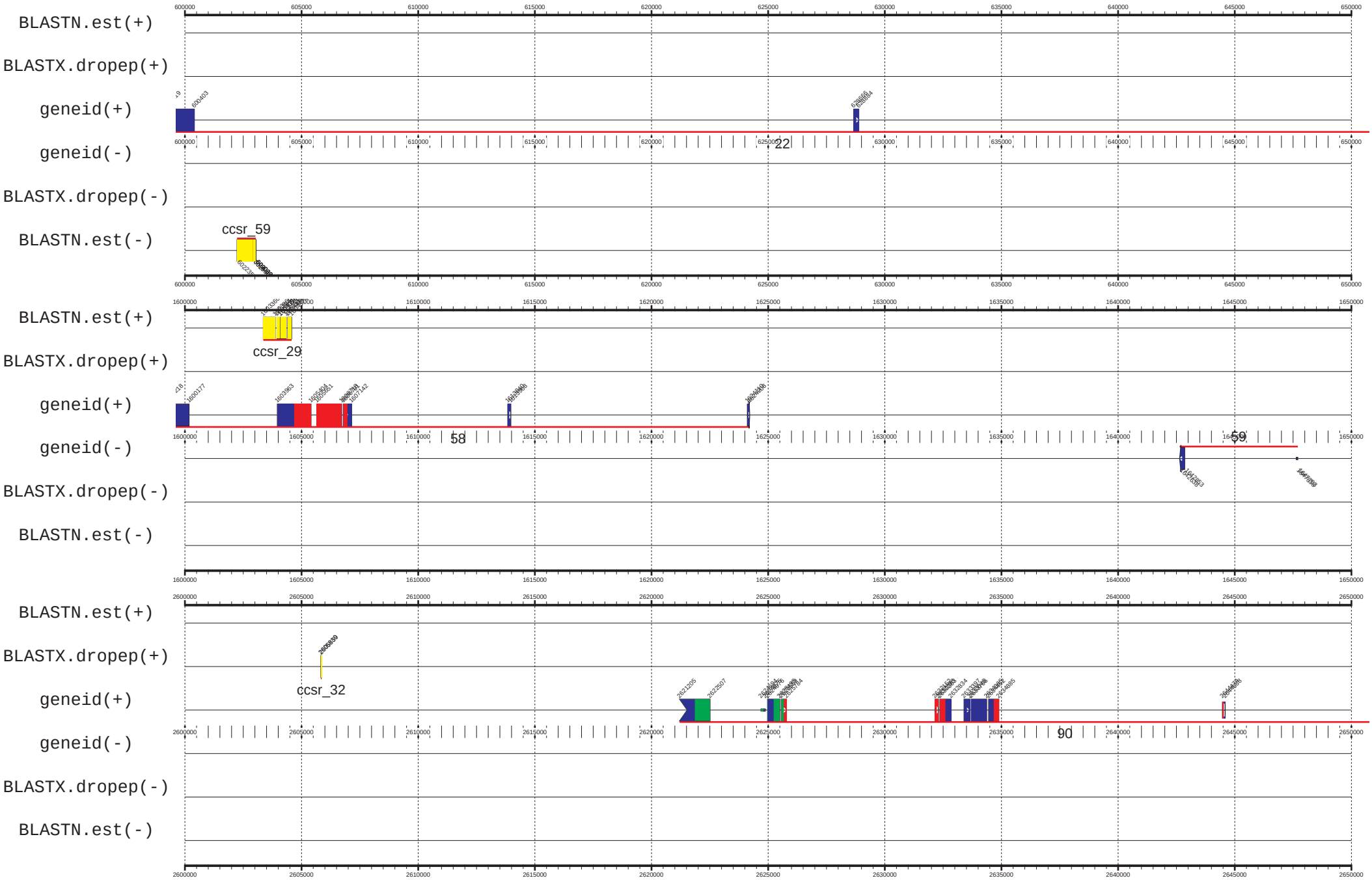
BLASTX.dropep(+)
geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
21
ccsr_24
57
49
87
88
89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BLASTX.dropep(+)
geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
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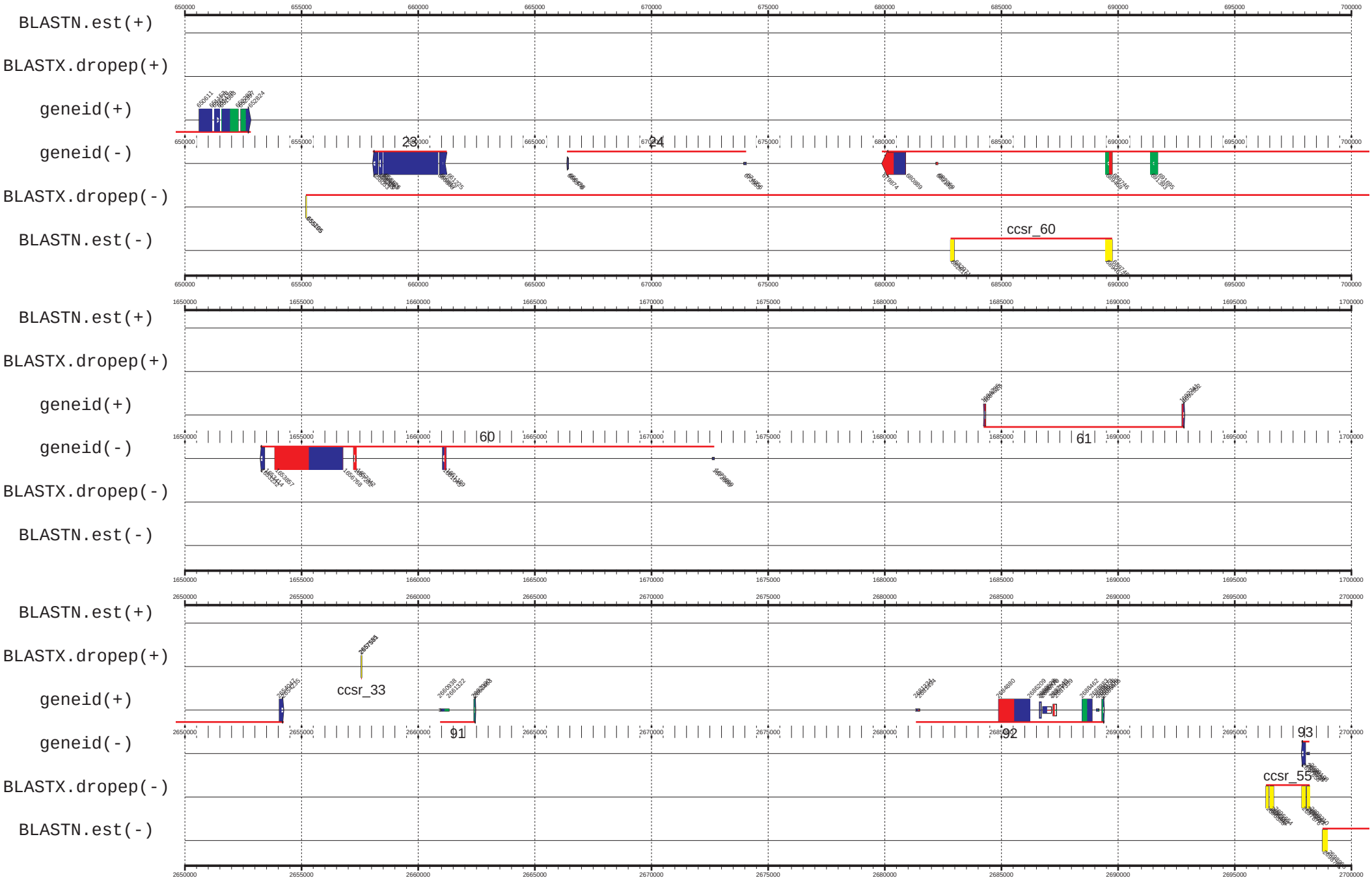
ccsr_29
ccsr_32
22
58
59
90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geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
650000
655000
660000
665000
670000
675000
680000
685000
690000
695000
700000
23
24
ccsr_60
1650000
1655000
1660000
1665000
1670000
1675000
1680000
1685000
1690000
1695000
1700000
60
61
2650000
2655000
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2700000
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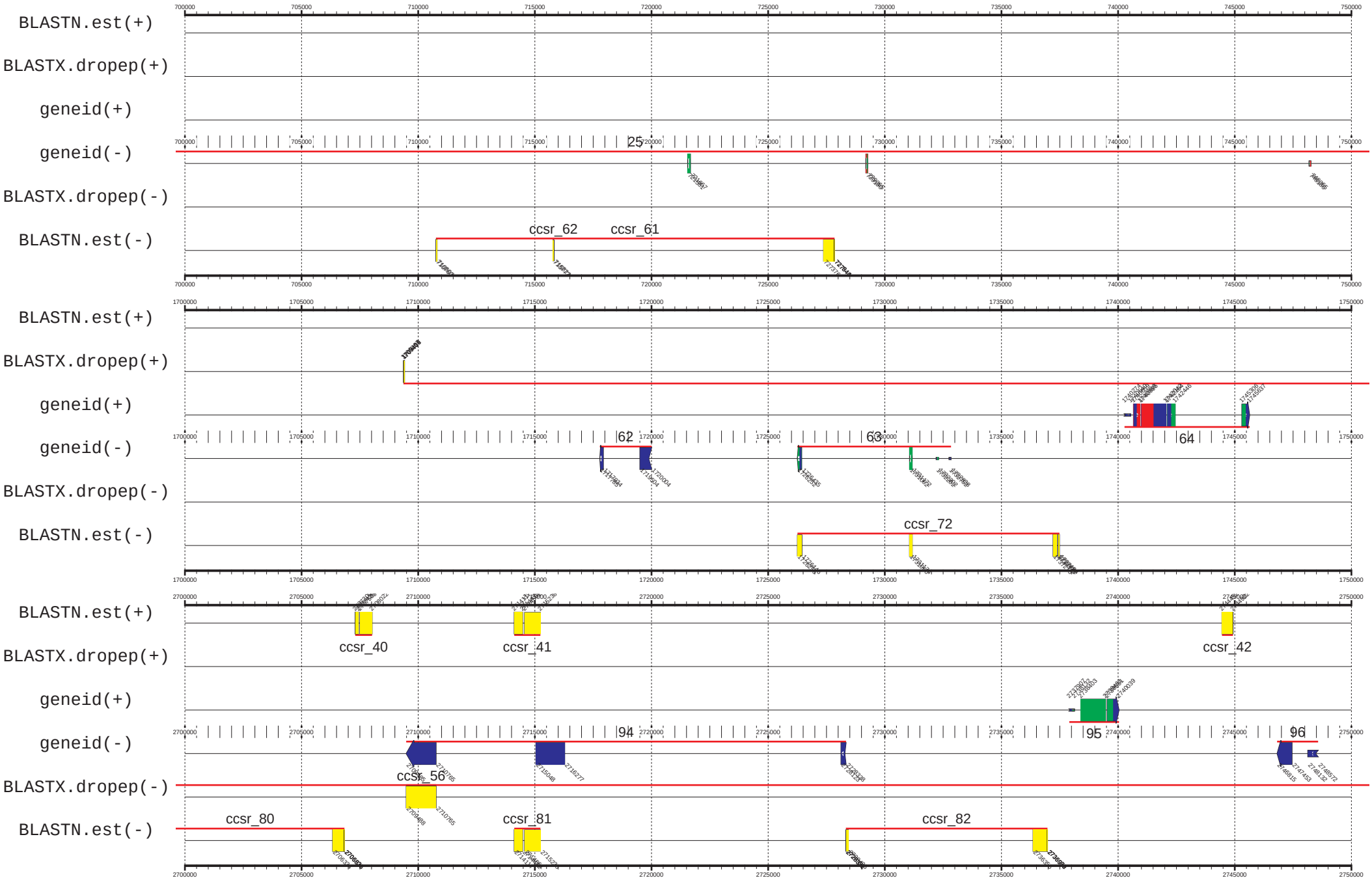
91
92
93
ccsr_55


BLASTN.est(+)
BLASTX.dropep(+)
geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
25
ccsr_62
ccsr_61
62
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ccsr_41
ccsr_42
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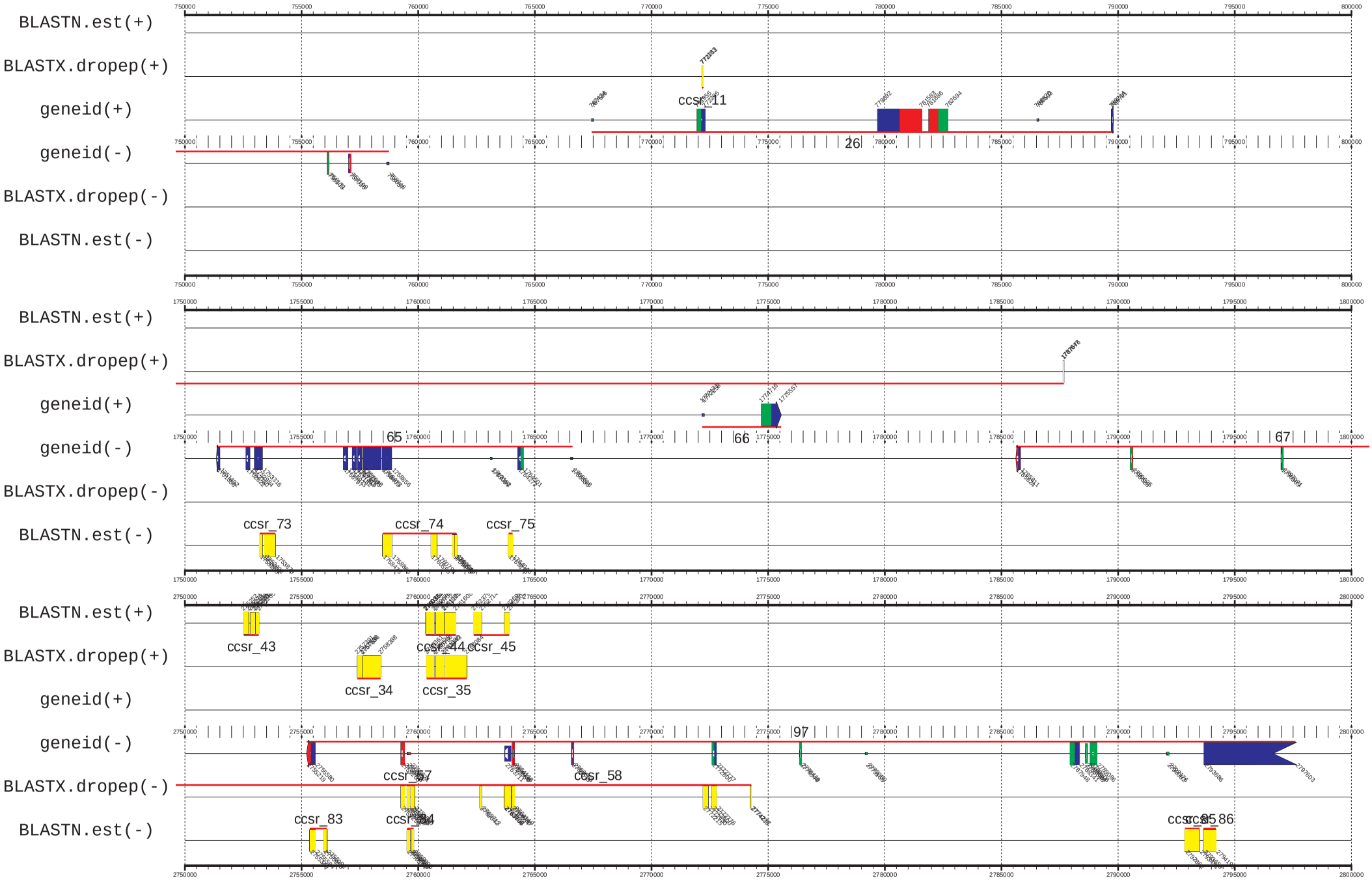
ccsr_80
ccsr_81
ccsr_82


BLASTN.est(+)
BLASTX.dropep(+)
geneid(+)
geneid(-)
BLASTX.dropep(-)
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ccsr_11
26
65
66
67
ccsr_73
ccsr_74
ccsr_75
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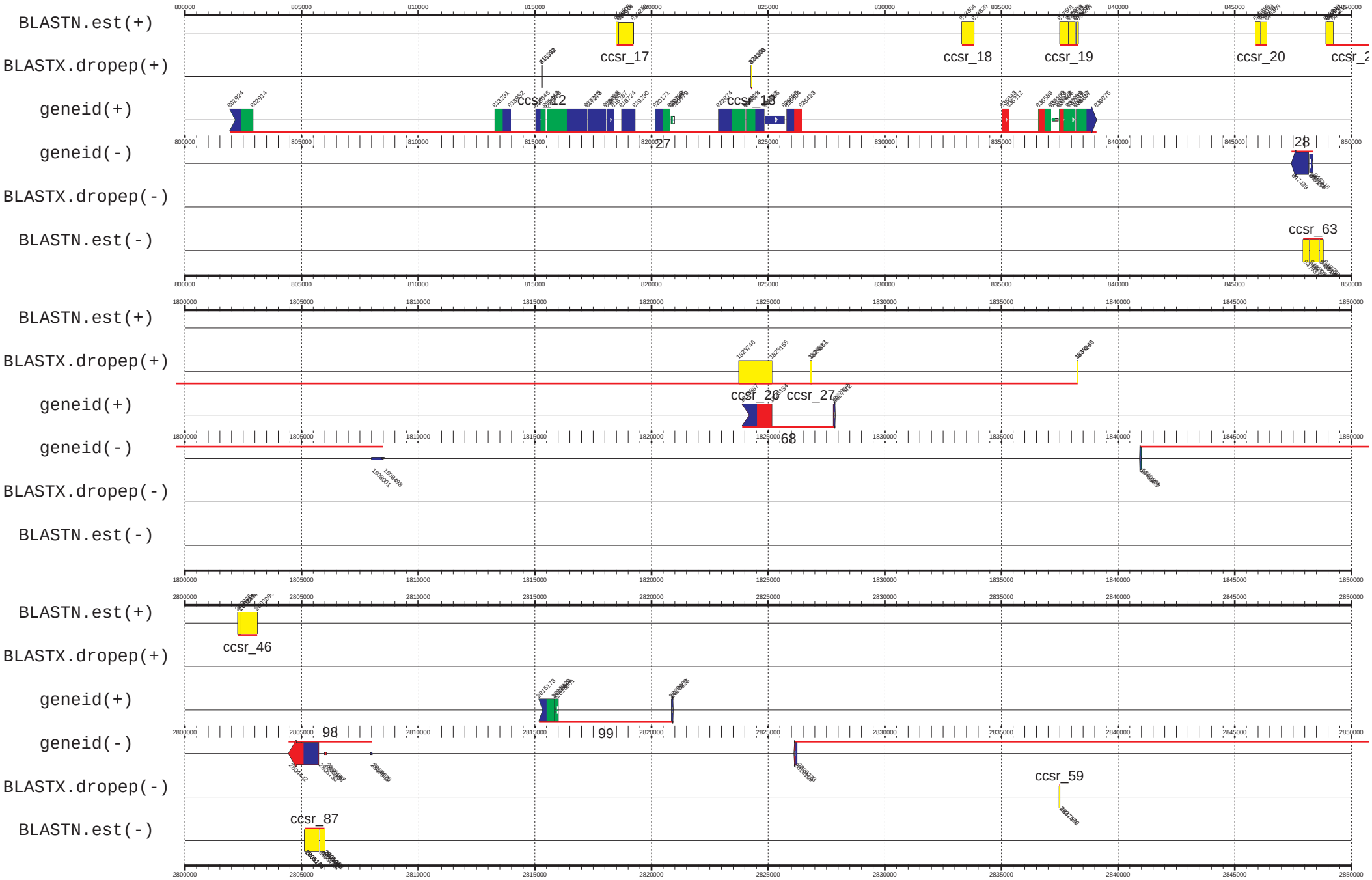
ccsr_85
ccsr_86


BLASTN.est(+)
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geneid(+)
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BLASTX.dropep(-)
BLASTN.est(-)
ccsr_17
ccsr_18
ccsr_19
ccsr_20
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ccsr_63
ccsr_26
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ccsr_46
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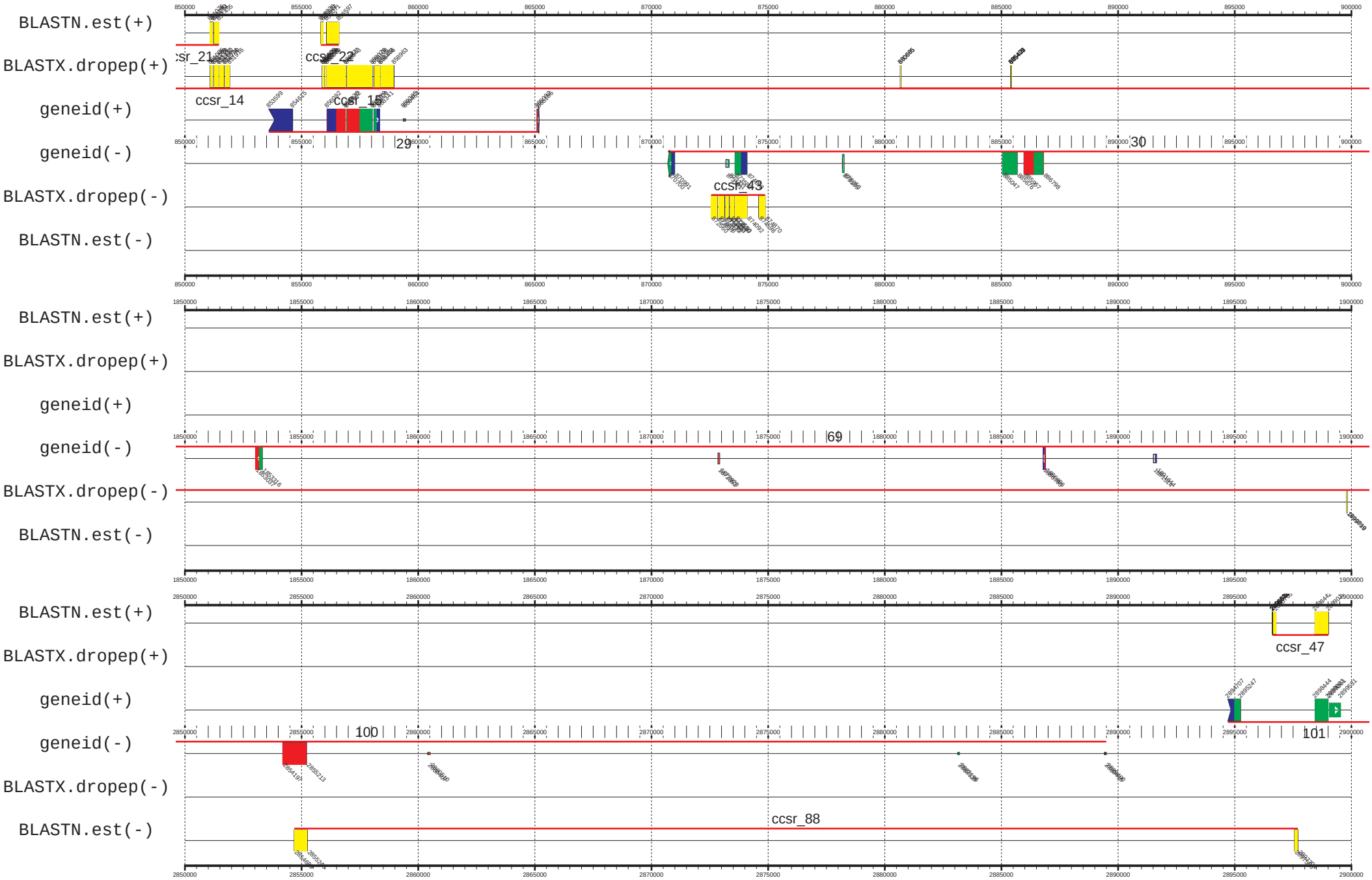
ccsr_87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geneid(+)
geneid(-)
BLASTX.dropep(-)
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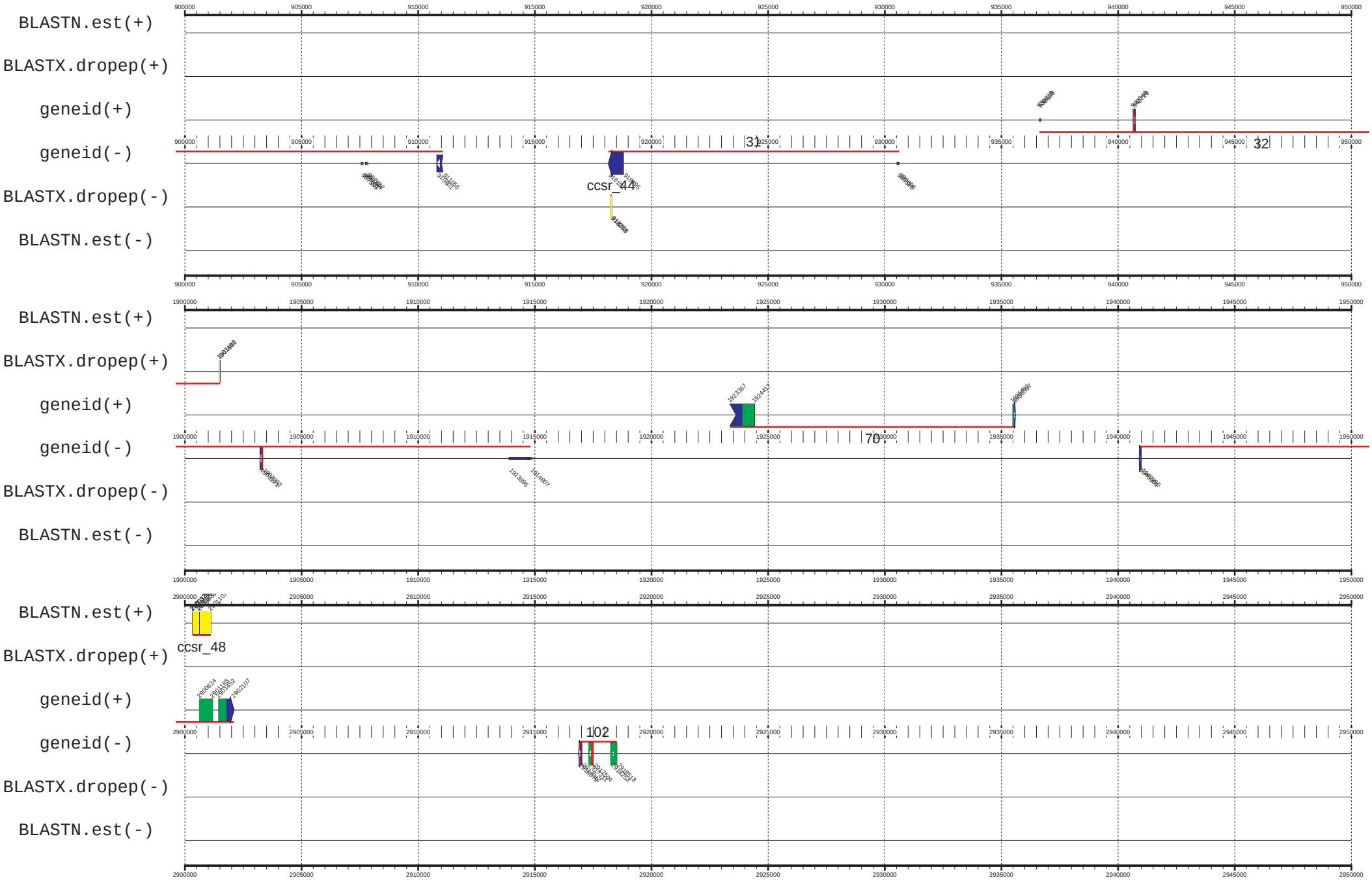
ccsr_14
ccsr_15
ccsr_43
ccsr_47
ccsr_88
29
30
69
100
101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BLASTX.dropep(+)
geneid(+)
geneid(-)
BLASTX.dropep(-)
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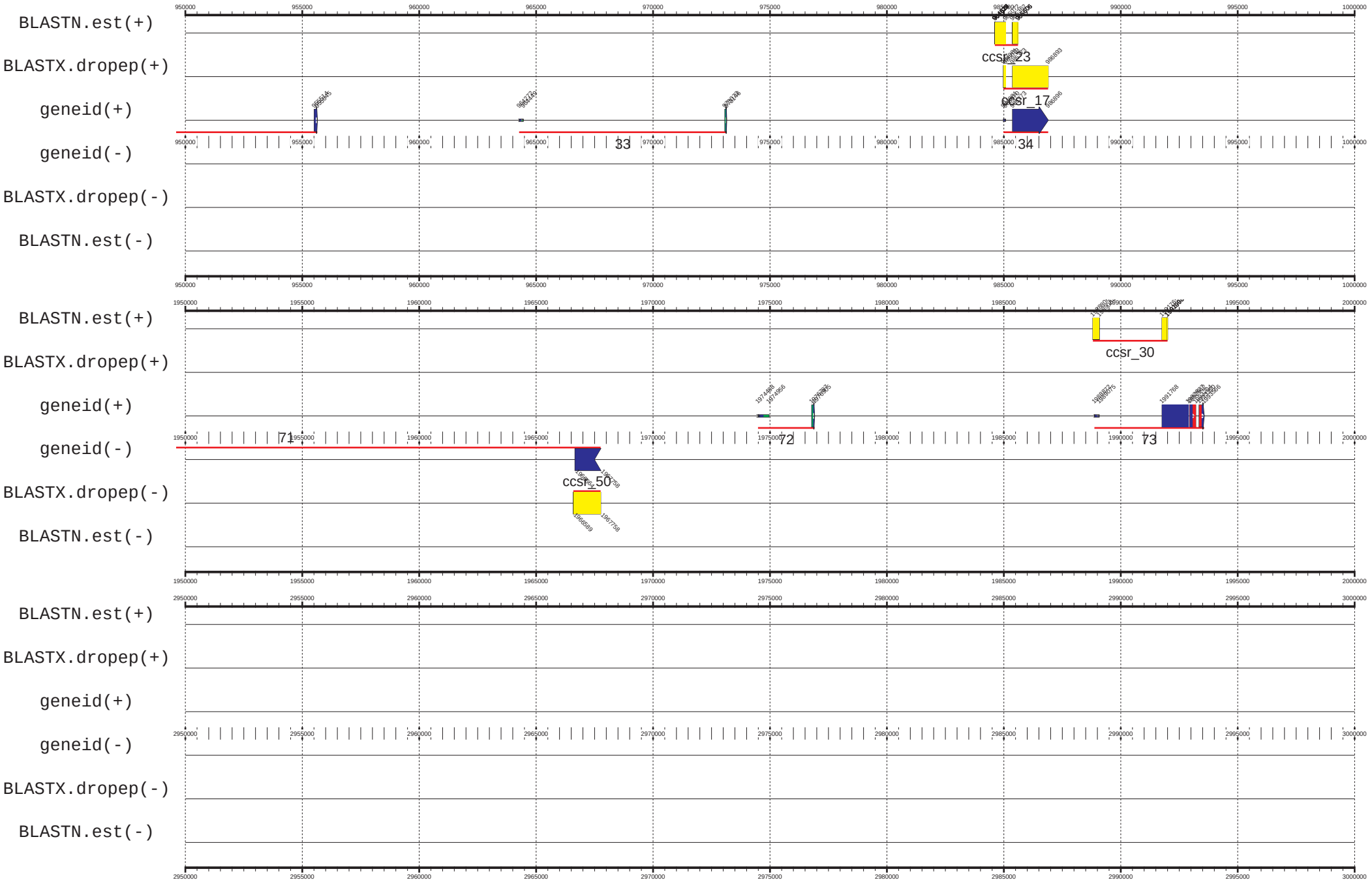
31
32
70
ccsr_48
102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BLASTX.dropep(+)
geneid(+)
geneid(-)
BLASTX.dropep(-)
BLASTN.est(-)
ccsr_23
ccsr_17
33
34
ccsr_30
71
72
73
ccsr_50