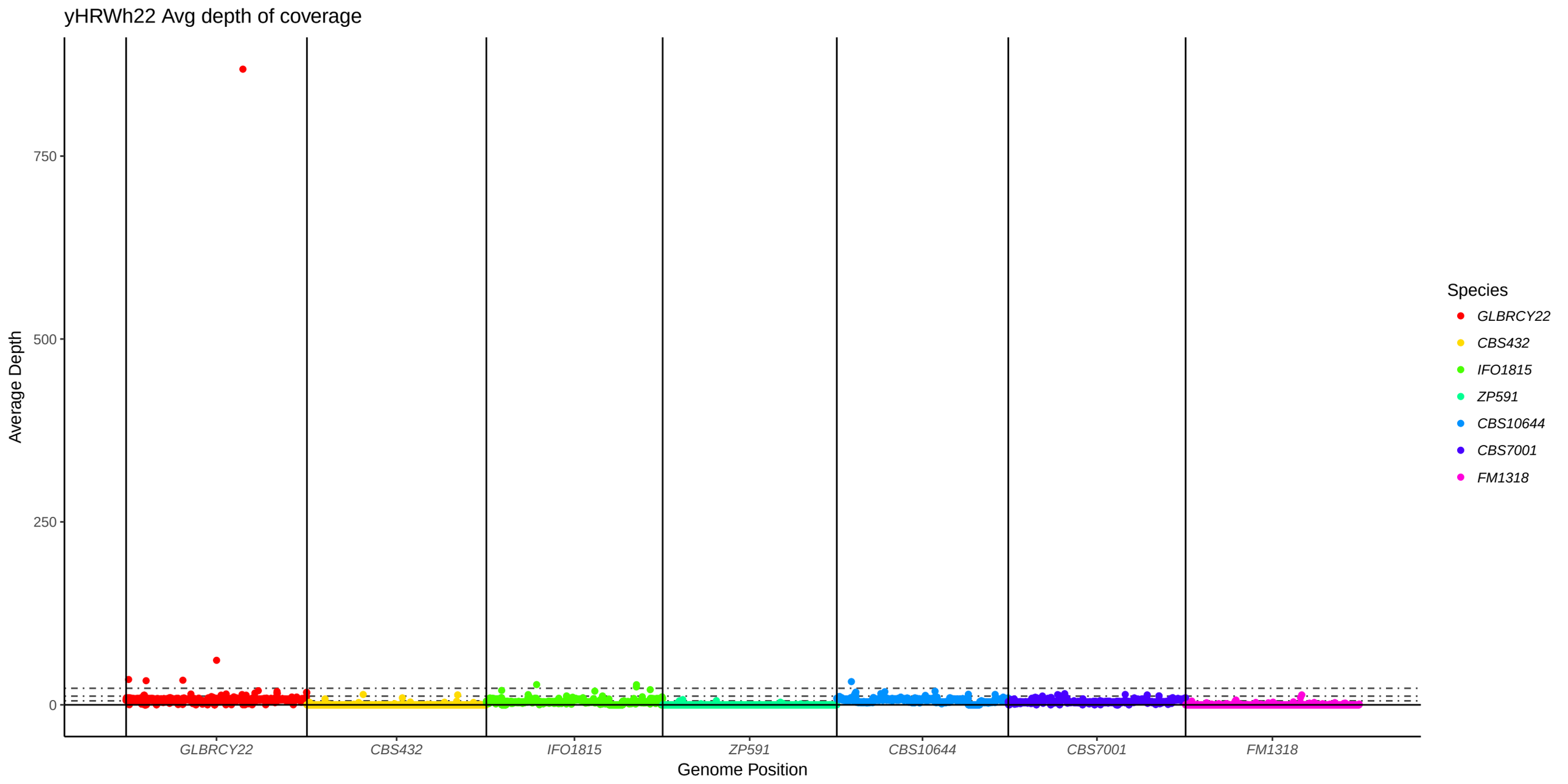
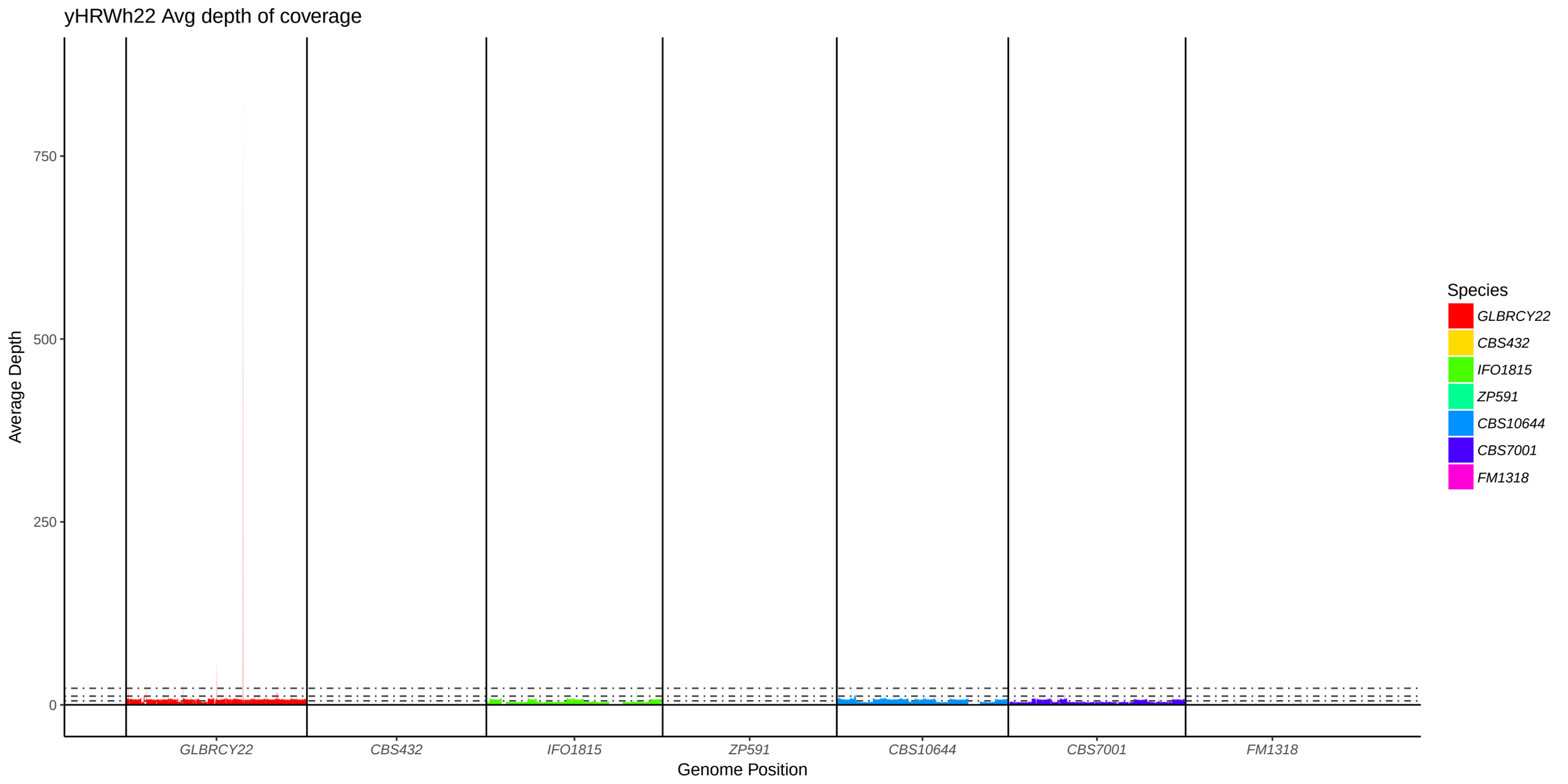
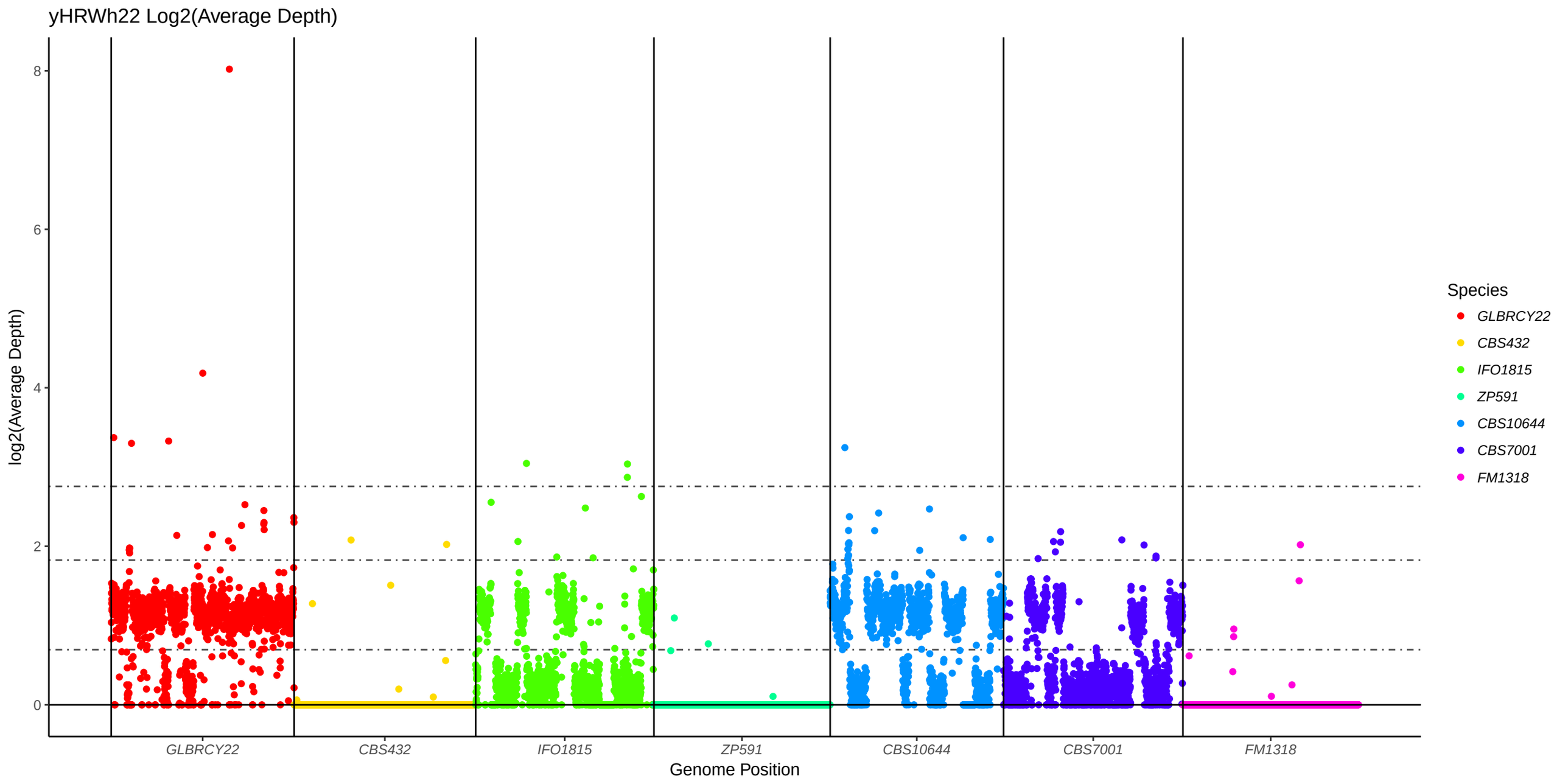
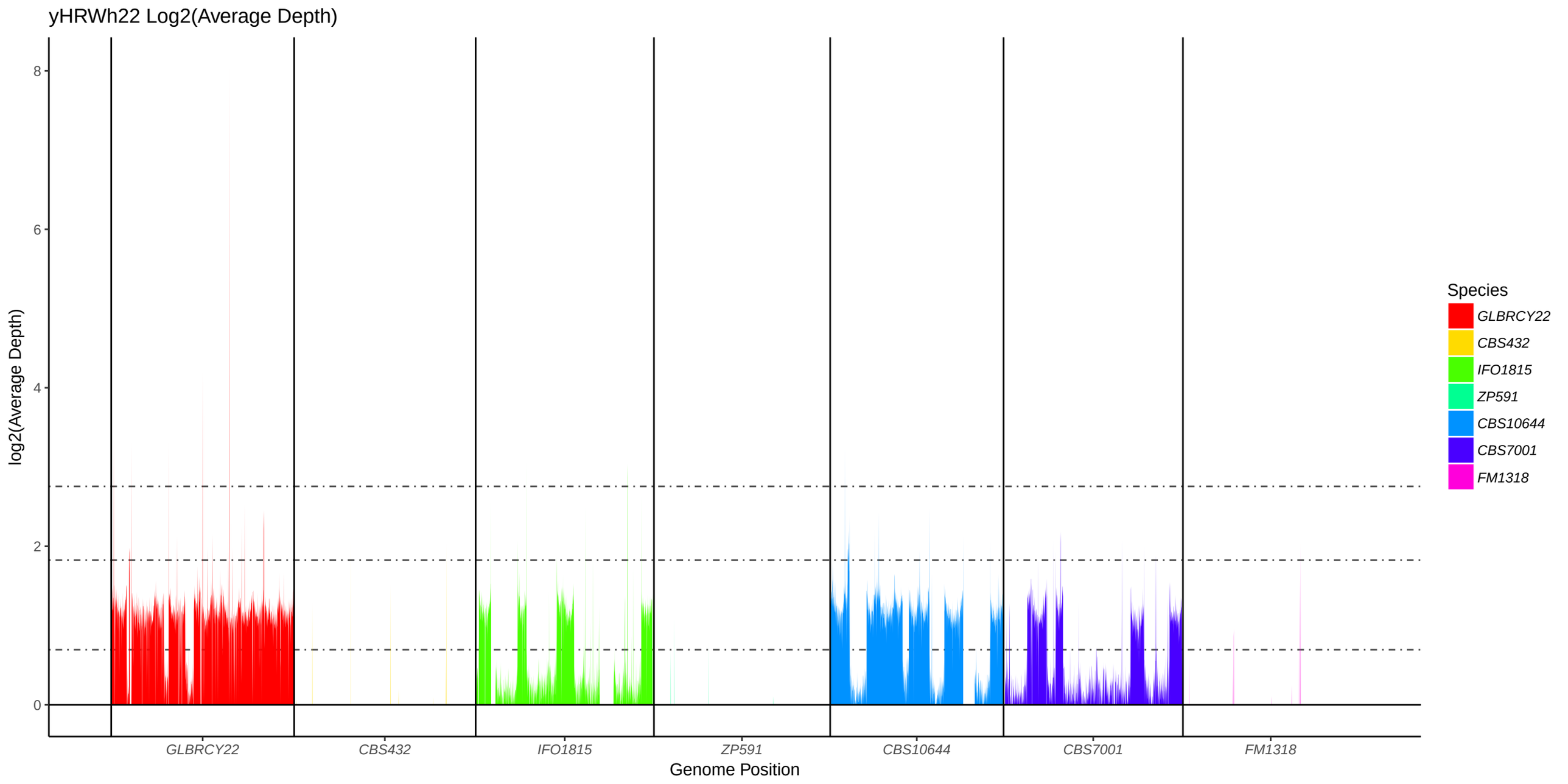

yHRWh22 Avg depth of coverage
Average Depth
750
500
250
0
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
FM1318
Genome Position
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
FM1318


yHRWh22 Avg depth of coverage
Average Depth
750
500
250
0
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
FM1318
Genome Position
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
FM1318


yHRWh22 Log2(Average Depth)
log2(Average Depth)
8
6
4
2
0
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
FM1318
Genome Position
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
FM1318


yHRWh22 Log2(Average Depth)
log2(Average Depth)
8
6
4
2
0
GLBRCY22
CBS432
IFO1815
ZP591
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FM1318
Genome Position
Species
GLBRCY22
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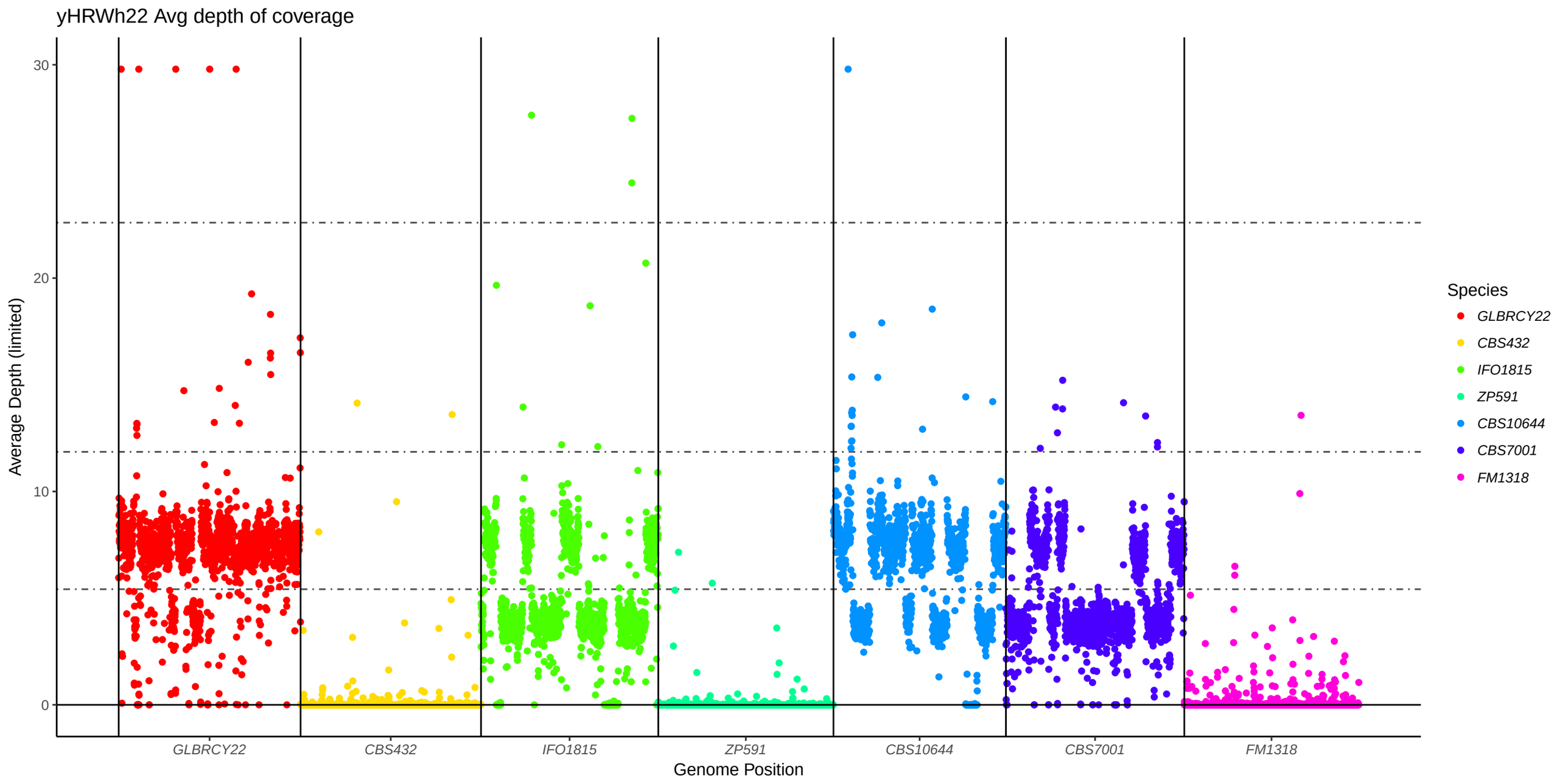
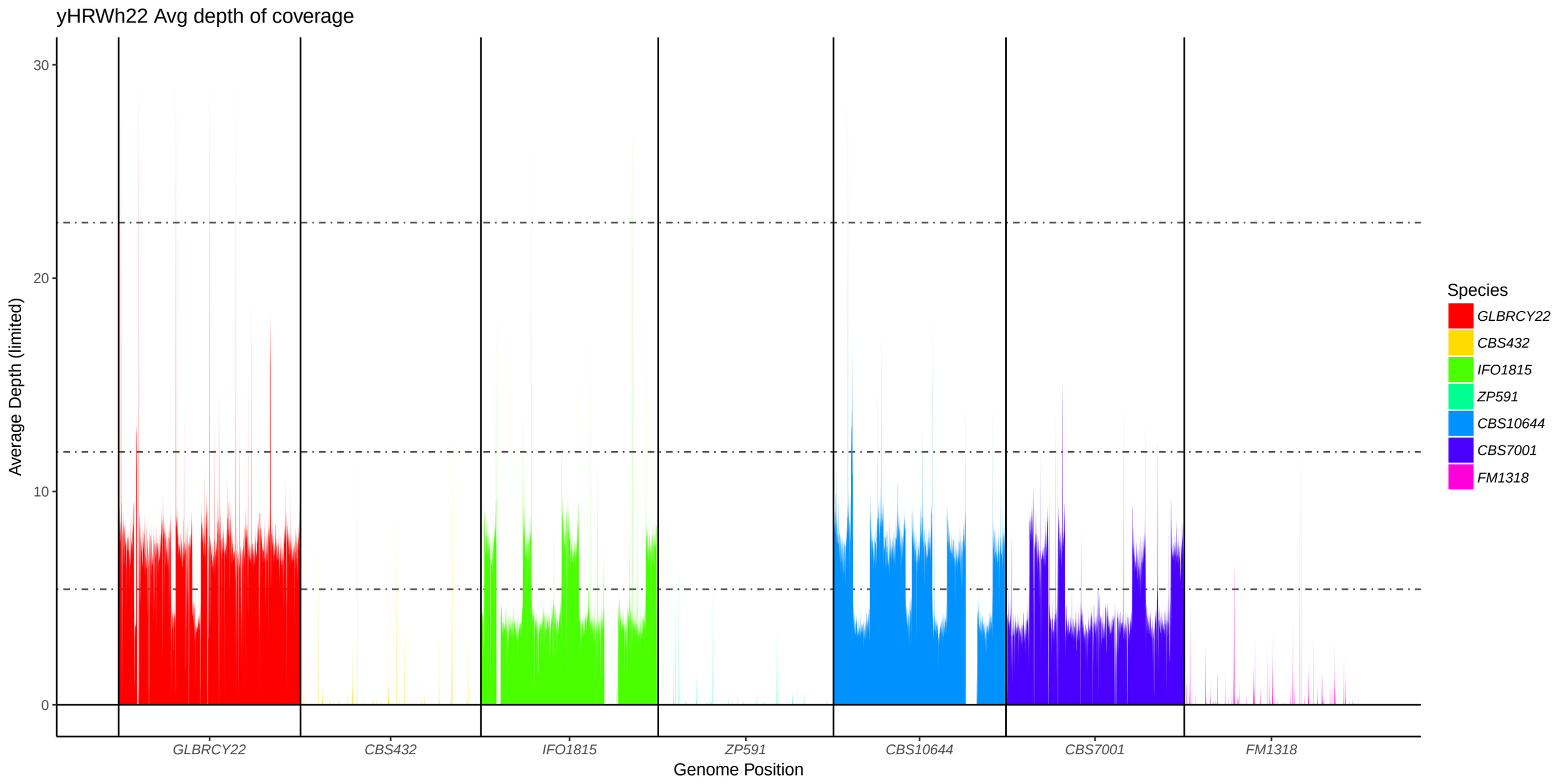
FM1318


yHRWh22 Avg depth of coverage
Average Depth (limited)
30
20
10
0
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
FM1318
Genome Position
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
FM1318


yHRWh22 Avg depth of coverage
Average Depth (limited)
30
20
10
0
GLBRCY22
CBS432
IFO1815
ZP591
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FM1318
Genome Position
Species
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IFO1815
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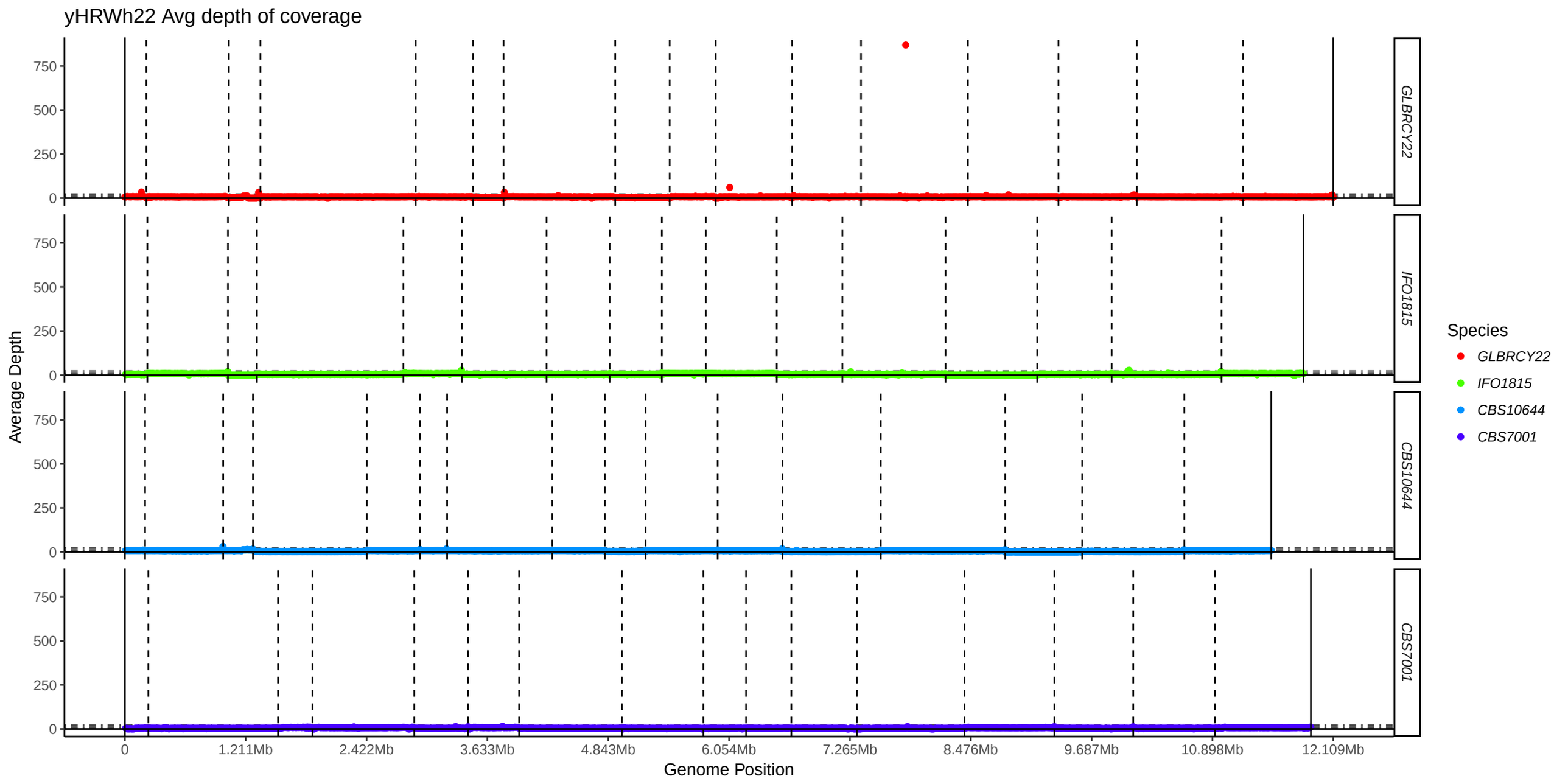
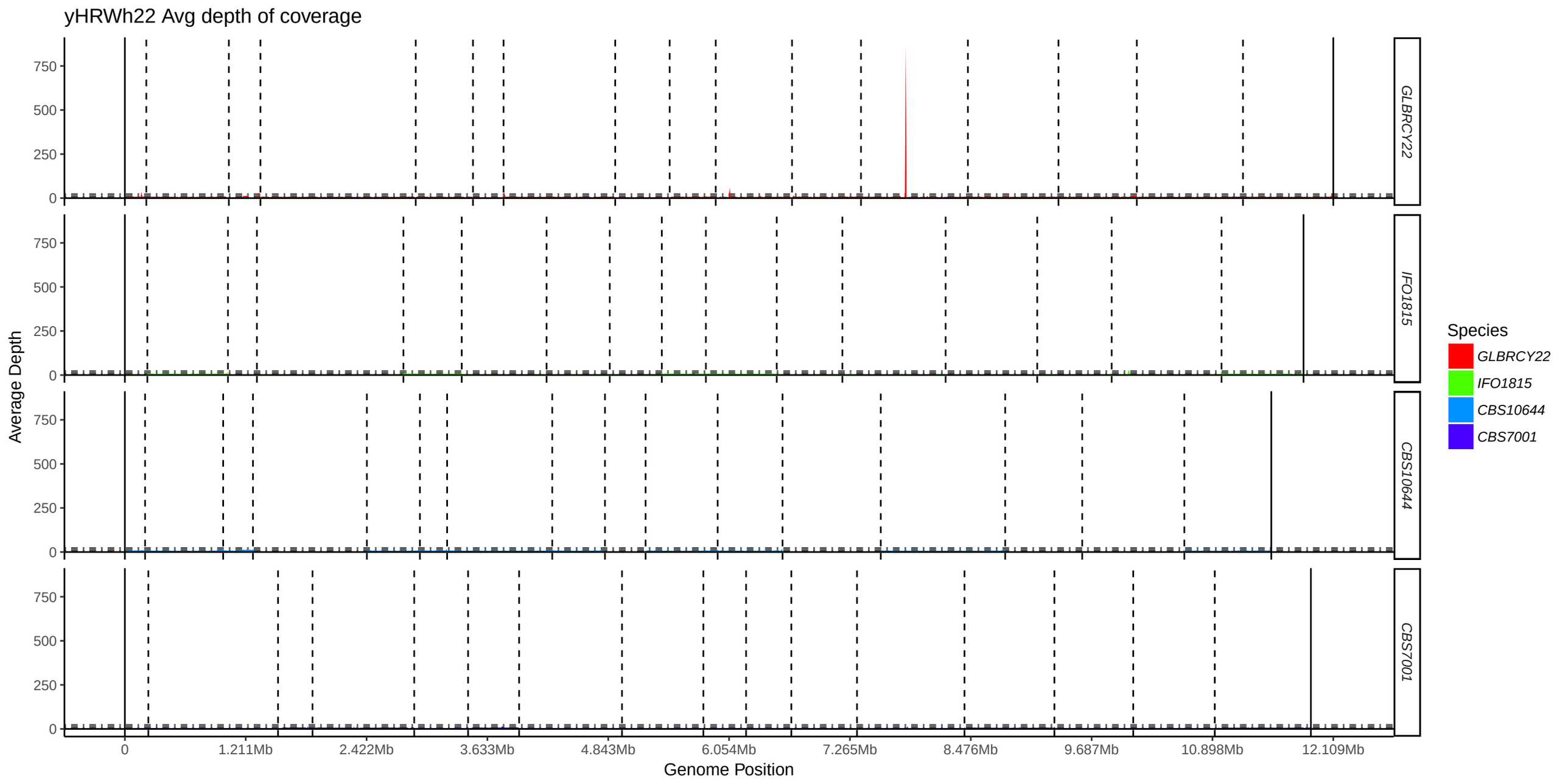
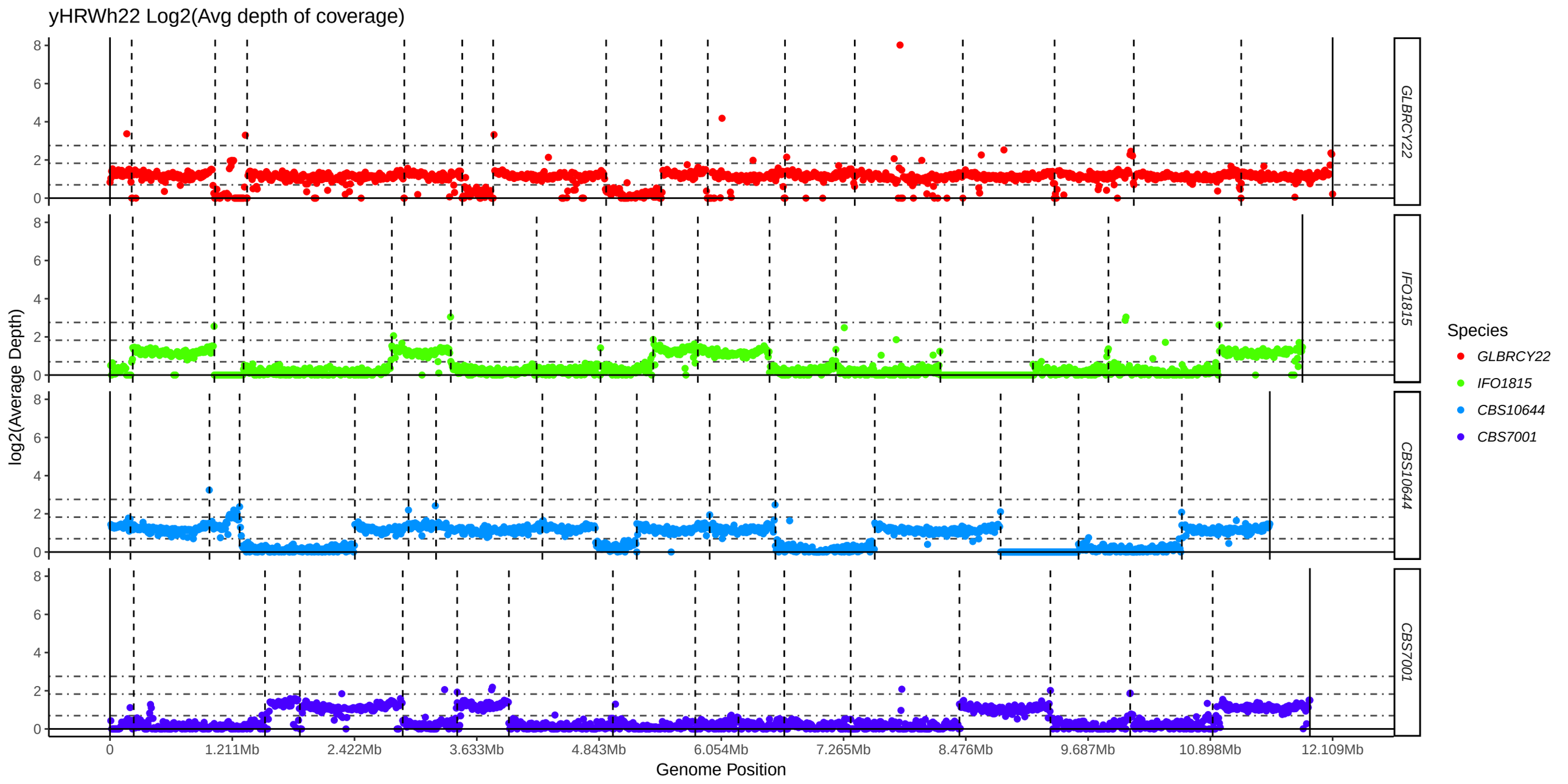
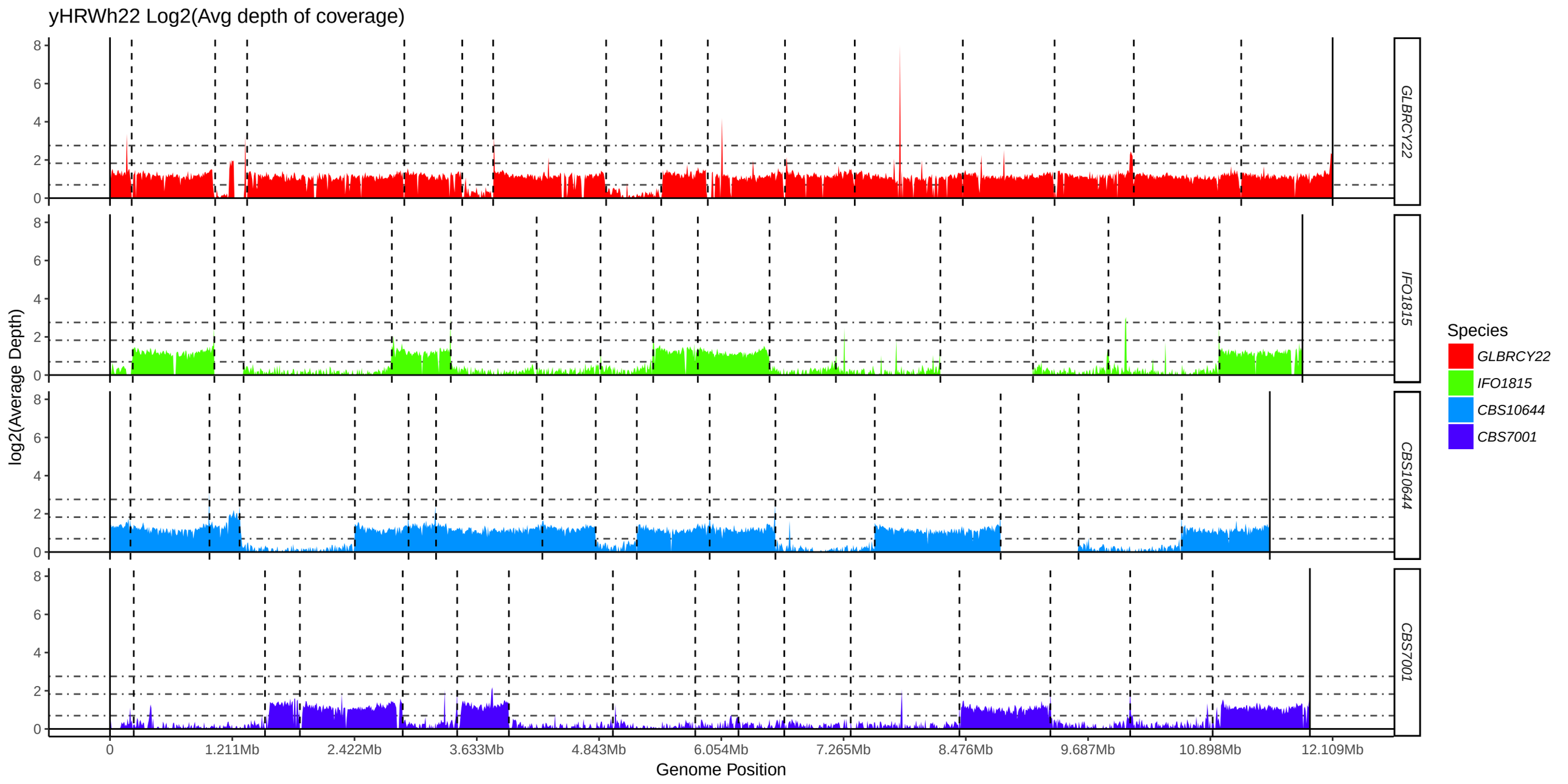
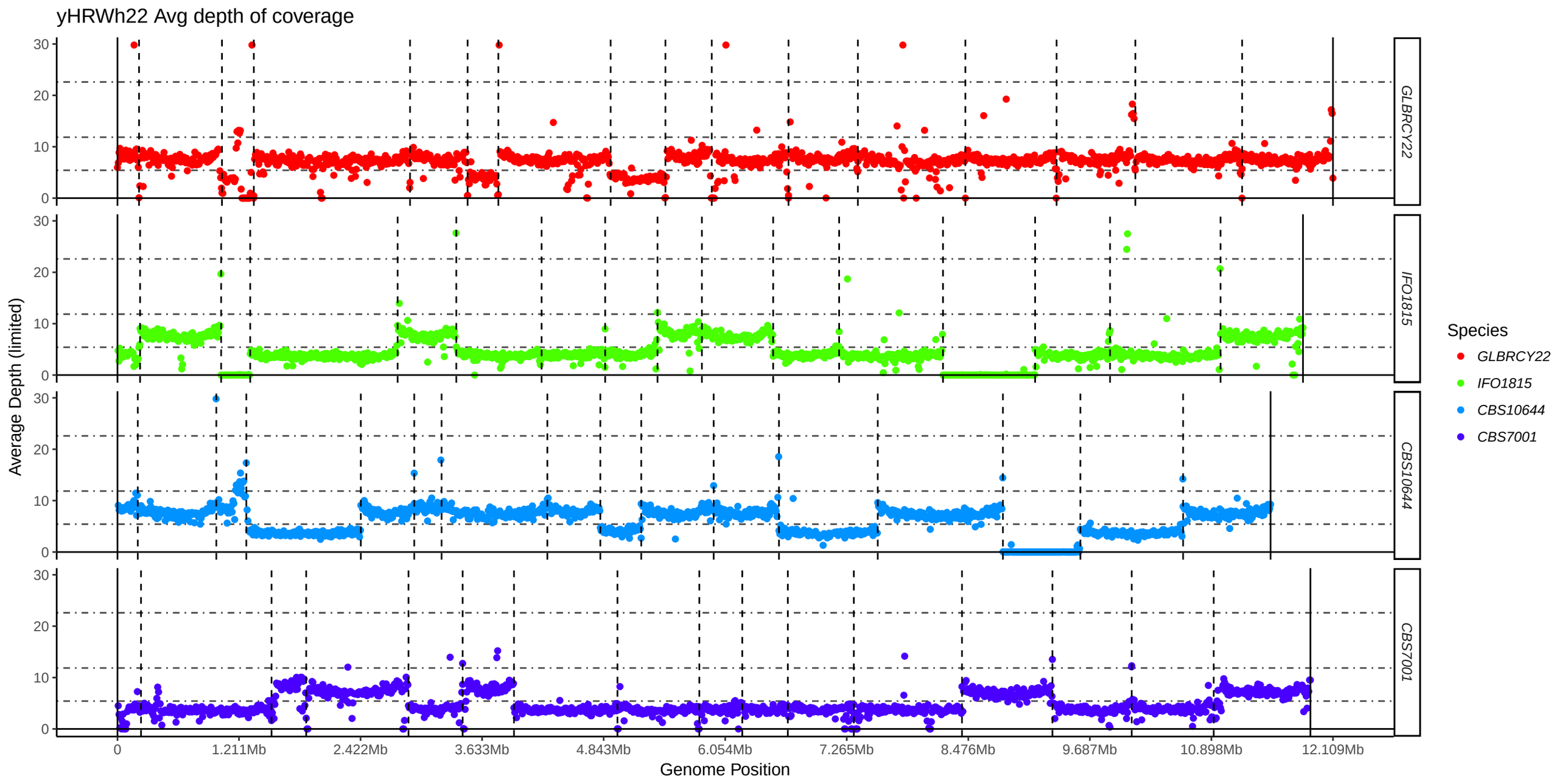
CBS7001
FM1318

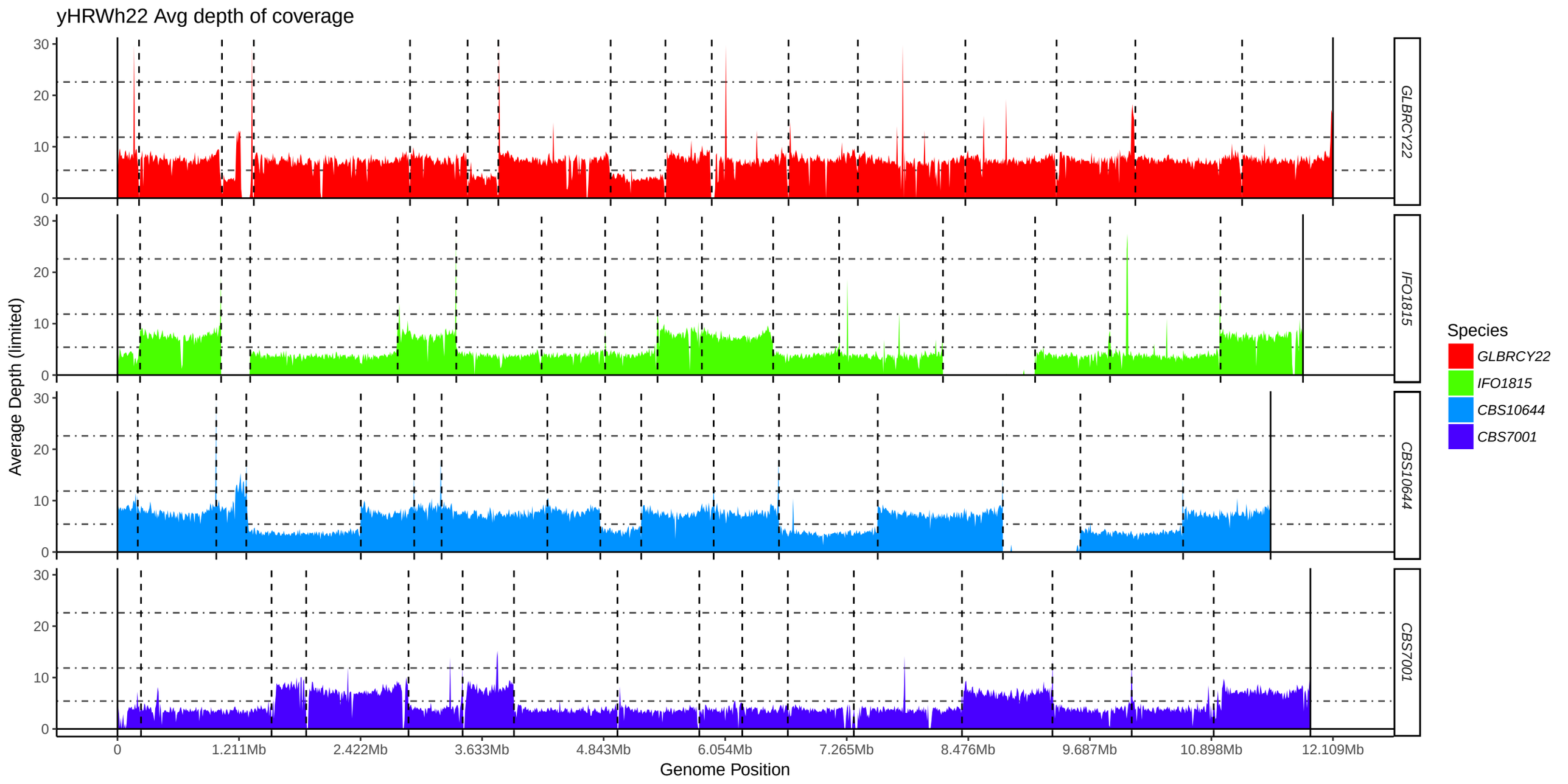

yHRWh22 Avg depth of coverage
Average Depth
750
500
250
0
GLBRCY22
IFO1815
CBS10644
CBS7001
0
1.211Mb
2.422Mb
3.633Mb
4.843Mb
6.054Mb
7.265Mb
8.476Mb
9.687Mb
10.898Mb
12.109Mb
Genome Position
Species
GLBRCY22
IFO1815
CBS10644
CBS7001


yHRWh22 Avg depth of coverage
Average Depth
750
500
250
0
GLBRCY22
IFO1815
CBS10644
CBS7001
0
1.211Mb
2.422Mb
3.633Mb
4.843Mb
6.054Mb
7.265Mb
8.476Mb
9.687Mb
10.898Mb
12.109Mb
Genome Position
Species
GLBRCY22
IFO1815
CBS10644
CBS7001


yHRWh22 Log2(Avg depth of coverage)
log2(Average Depth)
GLBRCY22
IFO1815
CBS10644
CBS7001
0
2
4
6
8
0
1.211Mb
2.422Mb
3.633Mb
4.843Mb
6.054Mb
7.265Mb
8.476Mb
9.687Mb
10.898Mb
12.109Mb
Genome Position
Species
GLBRCY22
IFO1815
CBS10644
CBS7001


yHRWh22 Log2(Avg depth of coverage)
log2(Average Depth)
GLBRCY22
IFO1815
CBS10644
CBS7001
0
2
4
6
8
0
1.211Mb
2.422Mb
3.633Mb
4.843Mb
6.054Mb
7.265Mb
8.476Mb
9.687Mb
10.898Mb
12.109Mb
Genome Position
Species
GLBRCY22
IFO1815
CBS10644
CBS7001


yHRWh22 Avg depth of coverage
Average Depth (limited)
Genome Position
0
1.211Mb
2.422Mb
3.633Mb
4.843Mb
6.054Mb
7.265Mb
8.476Mb
9.687Mb
10.898Mb
12.109Mb
GLBRCY22
IFO1815
CBS10644
CBS7001
Species
GLBRCY22
IFO1815
CBS10644
CBS7001


yHRWh22 Avg depth of coverage
Average Depth (limited)
30
20
10
0
GLBRCY22
IFO1815
CBS10644
CBS7001
0
1.211Mb
2.422Mb
3.633Mb
4.843Mb
6.054Mb
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Genome Position
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