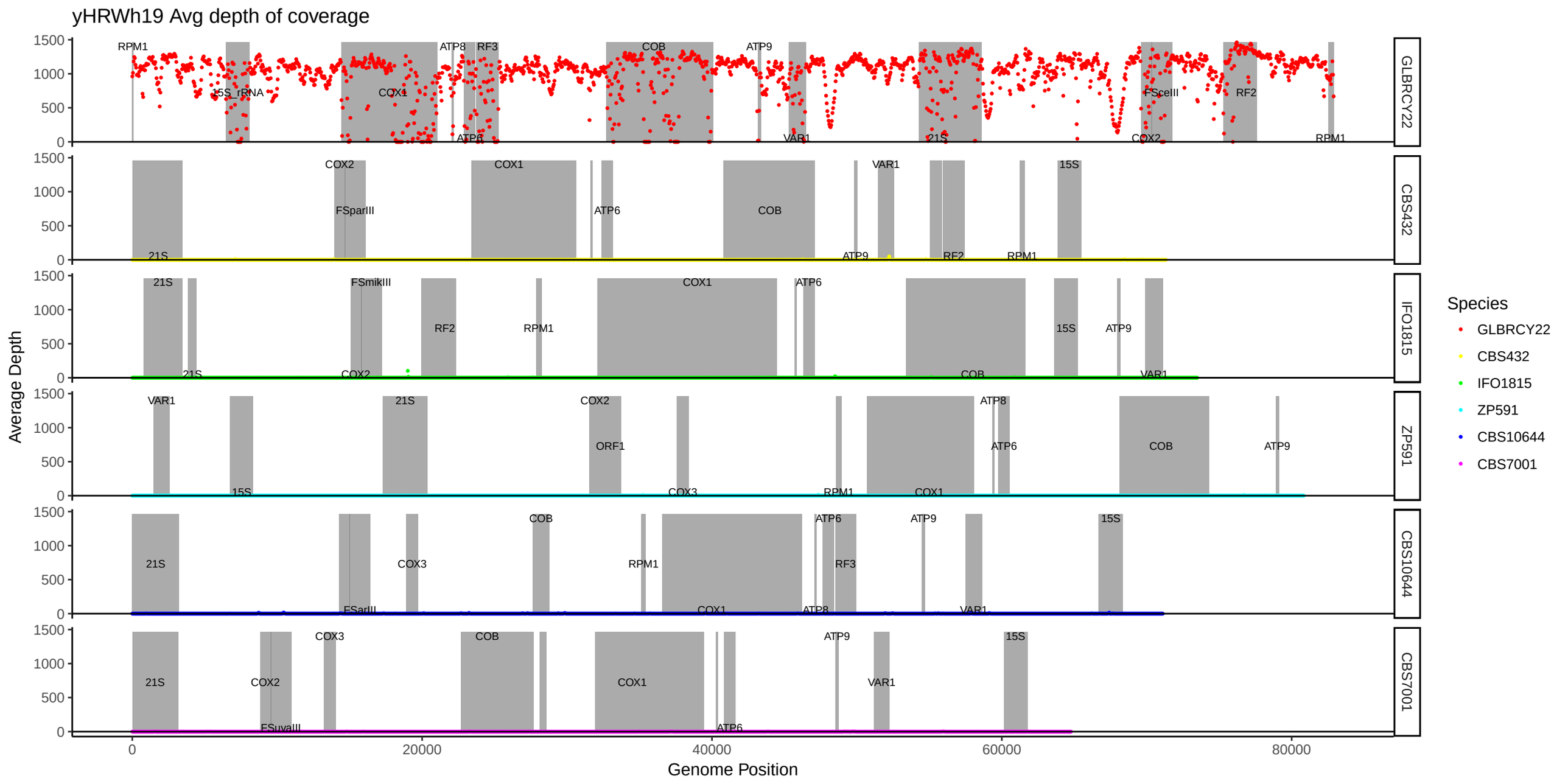
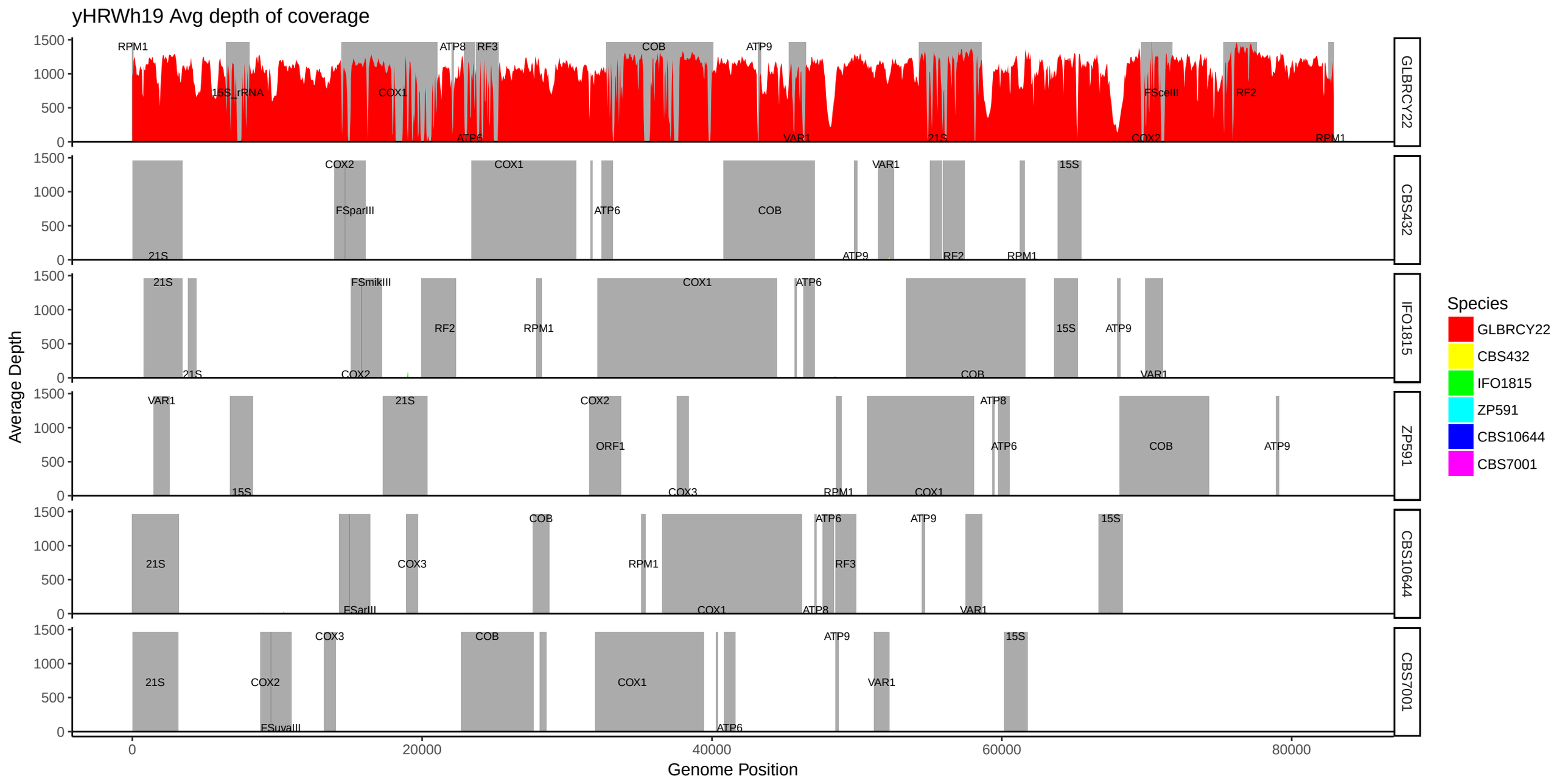
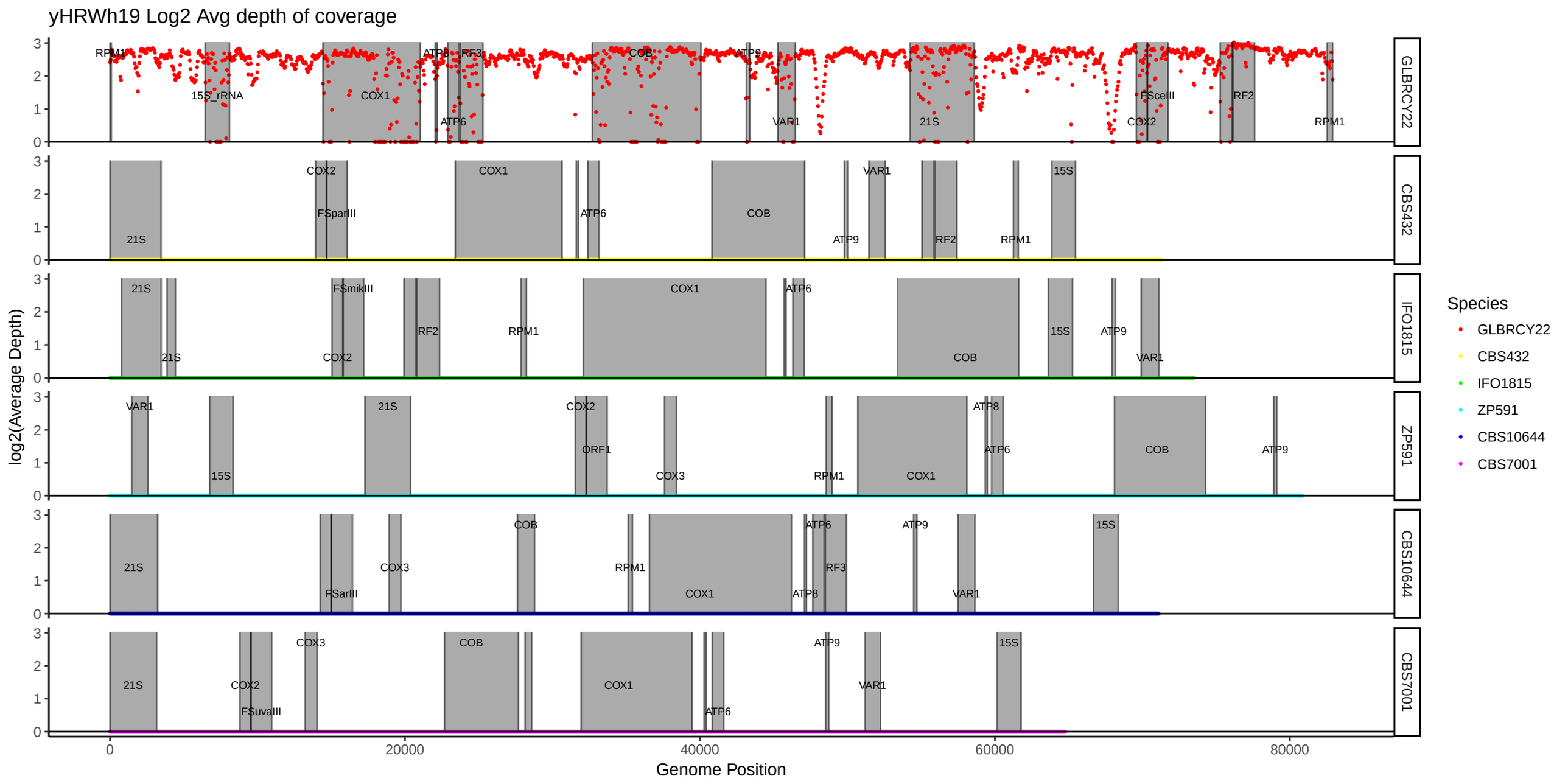
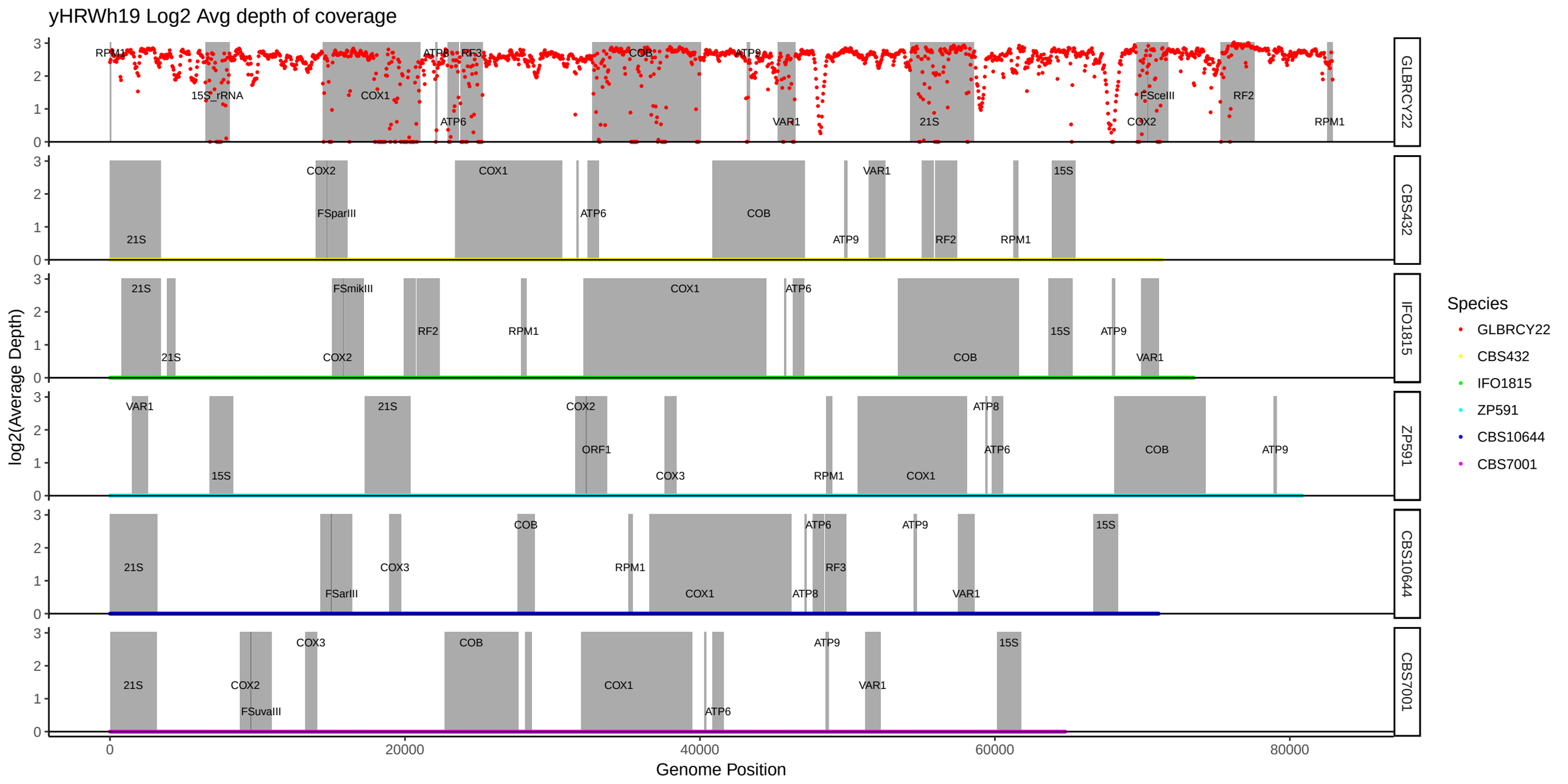
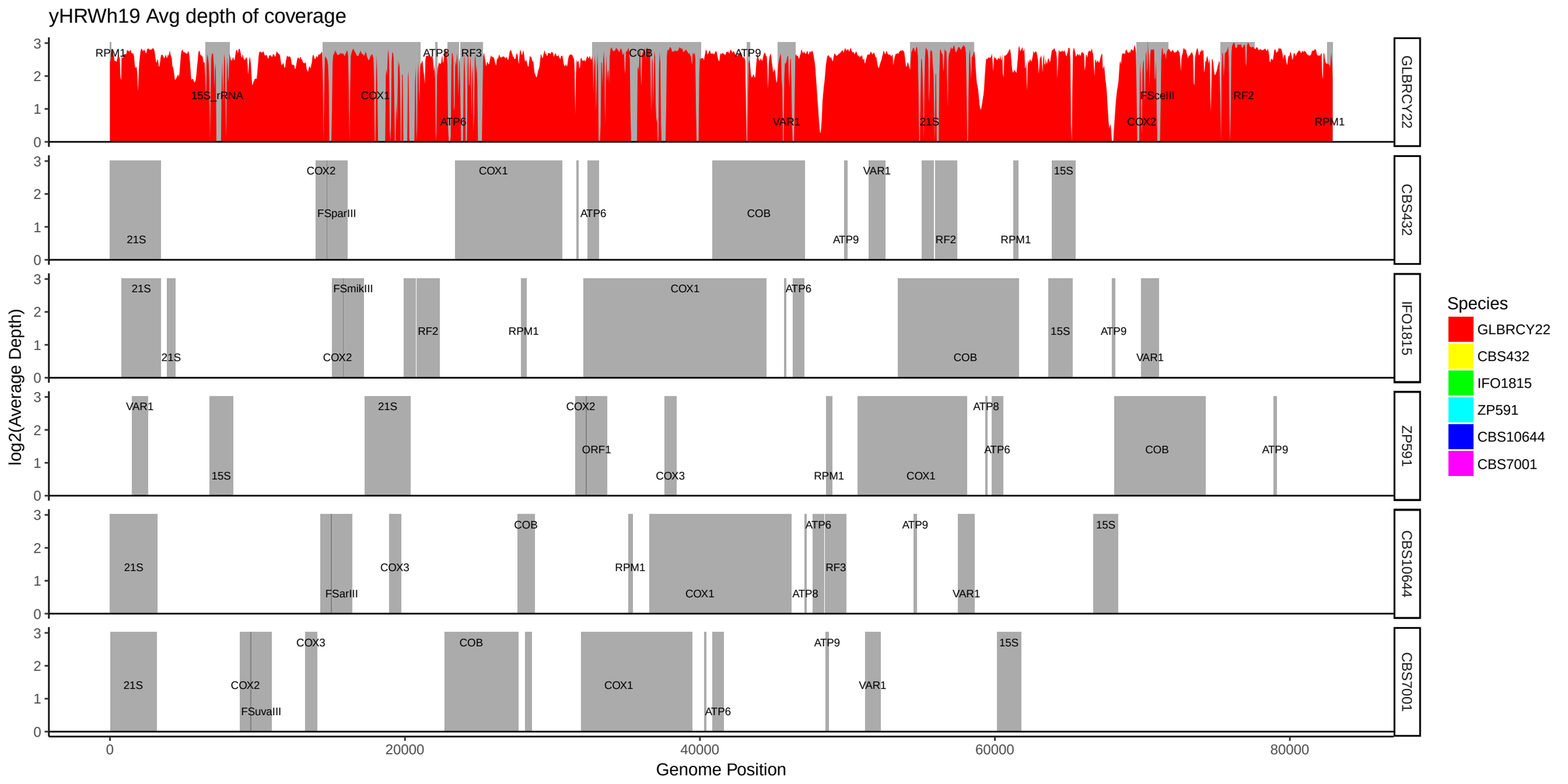
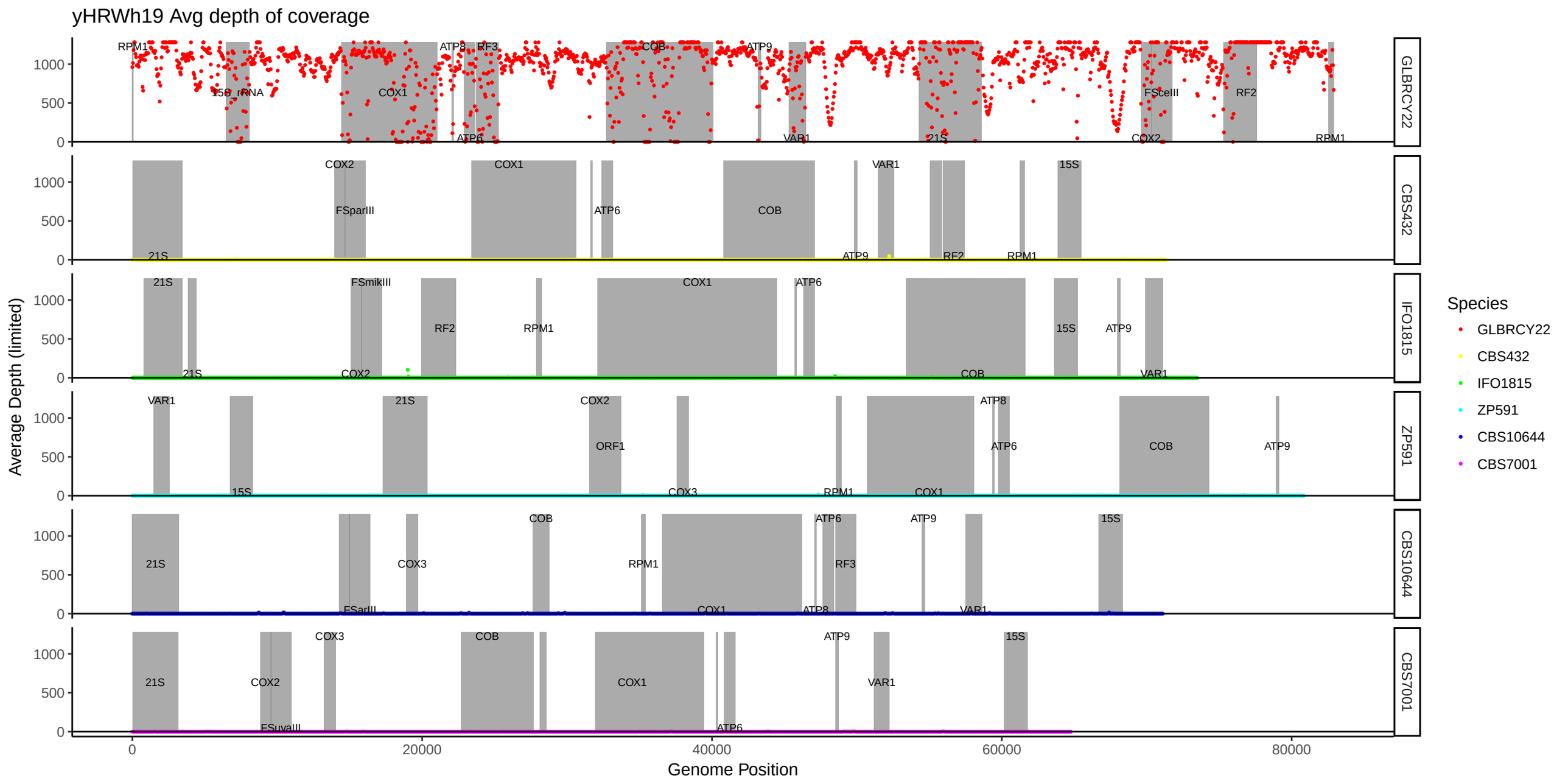

yHRWh19 Avg depth of coverage
Average Depth
Genome Position
0
20000
40000
60000
80000
0
500
1000
1500
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
FSparIII
FSmikIII
ORF1
COX3
FSarIII
FSuvaIII
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001

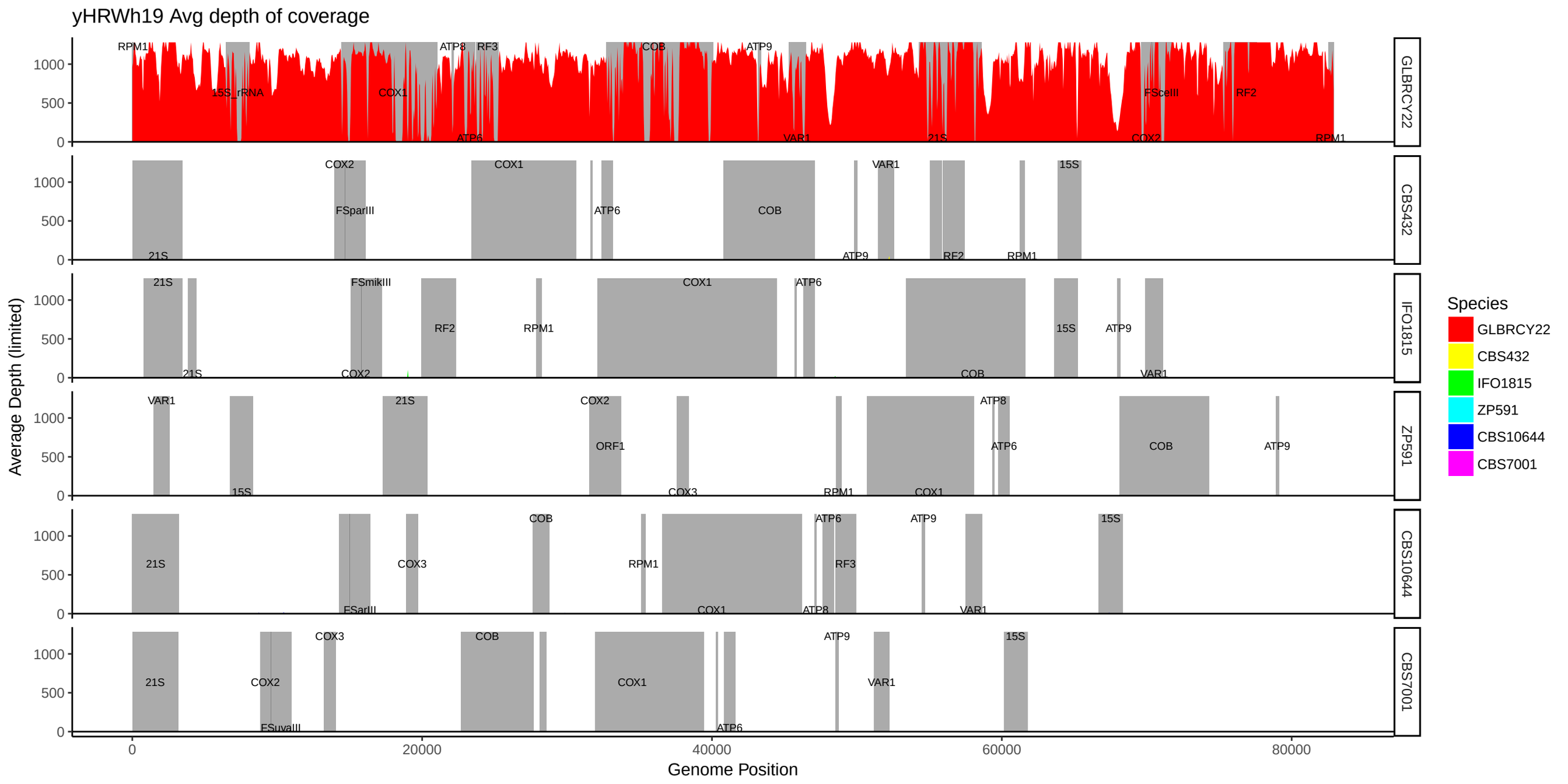

yHRWh19 Avg depth of coverage
Average Depth
Genome Position
0
20000
40000
60000
80000
0
500
1000
1500
GLBRCY22
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
RPM1
CBS432
21S
COX2
FSparIII
COX1
ATP6
COB
ATP9
VAR1
RF2
RPM1
15S
IFO1815
21S
21S
FSmikIII
COX2
RF2
RPM1
COX1
ATP6
COB
15S
ATP9
VAR1
ZP591
VAR1
15S
21S
COX2
ORF1
COX3
RPM1
COX1
ATP8
ATP6
COB
ATP9
CBS10644
21S
FSarIII
COX3
COB
RPM1
COX1
ATP6
ATP8
RF3
ATP9
VAR1
15S
CBS7001
21S
COX2
FSuvaIII
COX3
COB
COX1
ATP6
ATP9
VAR1
15S
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001

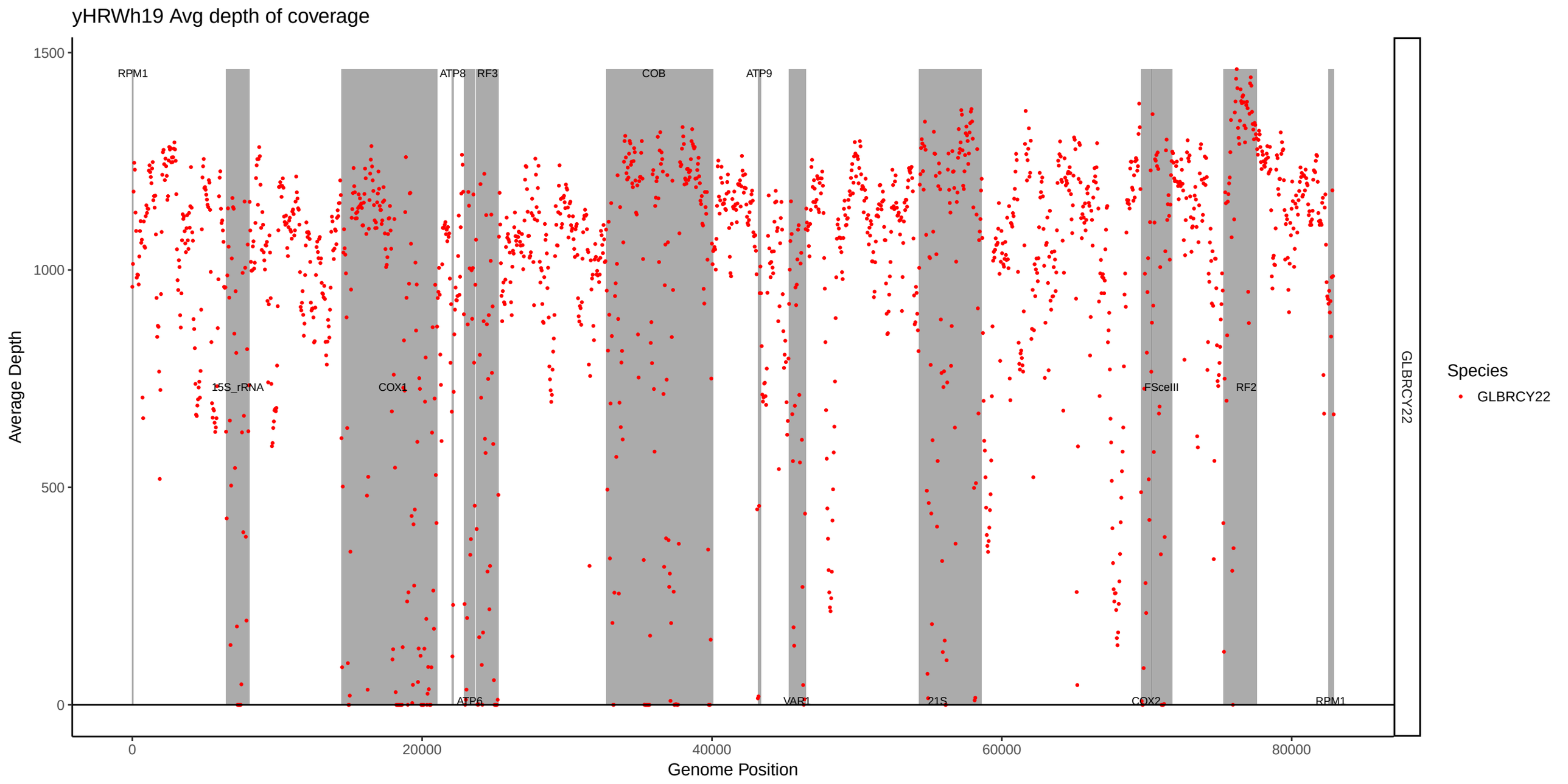

yHRWh19 Log2 Avg depth of coverage
log2(Average Depth)
Genome Position
0
20000
40000
60000
80000
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
FSparIII
FSmikIII
ORF1
COX3
FSarIII
FSuvaIII
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001

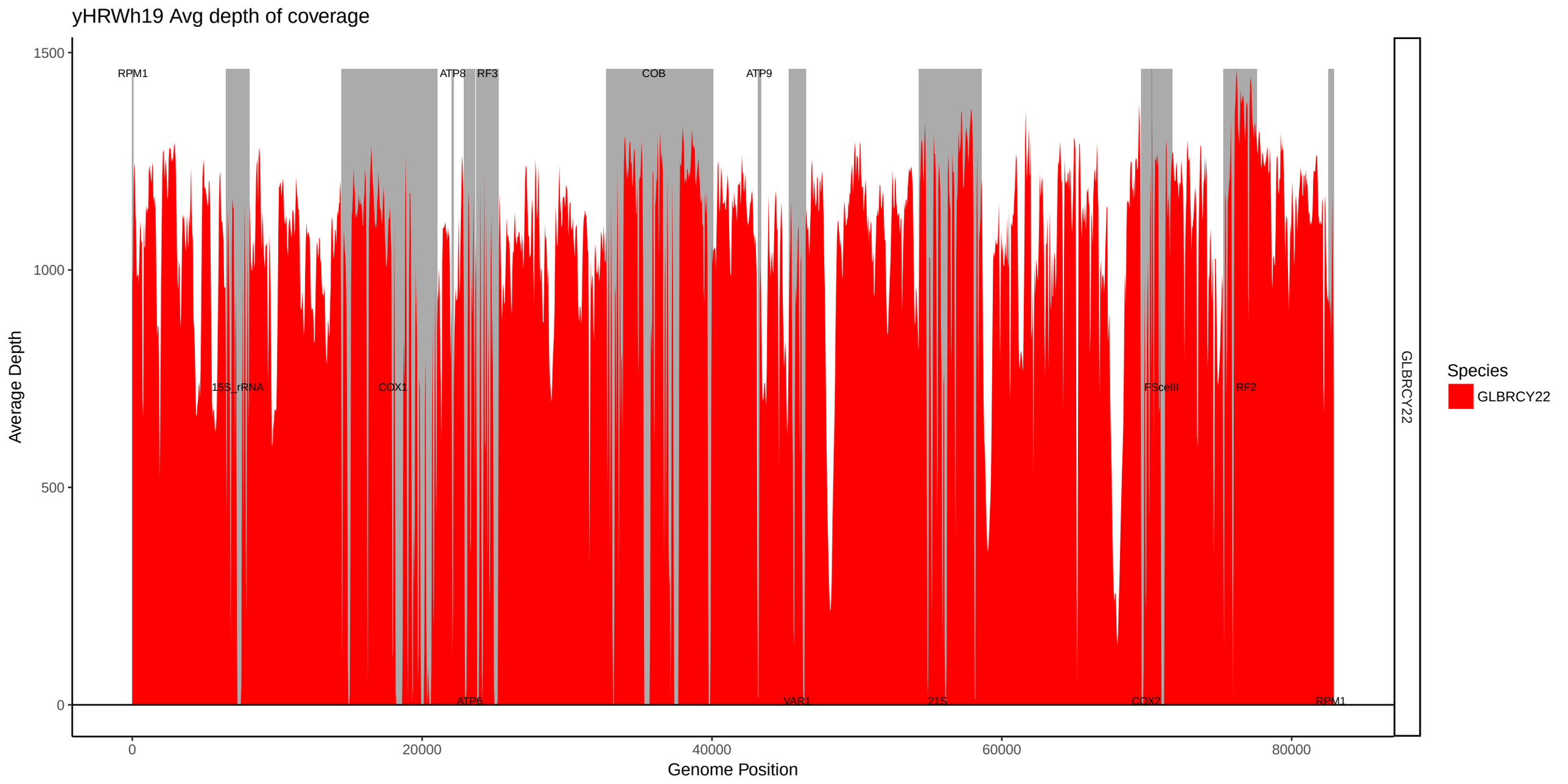

yHRWh19 Log2 Avg depth of coverage
log2(Average Depth)
Genome Position
0
20000
40000
60000
80000
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
FSparIII
15S
FSmikIII
ORF1
COX3
FSarIII
FSuvaIII
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001

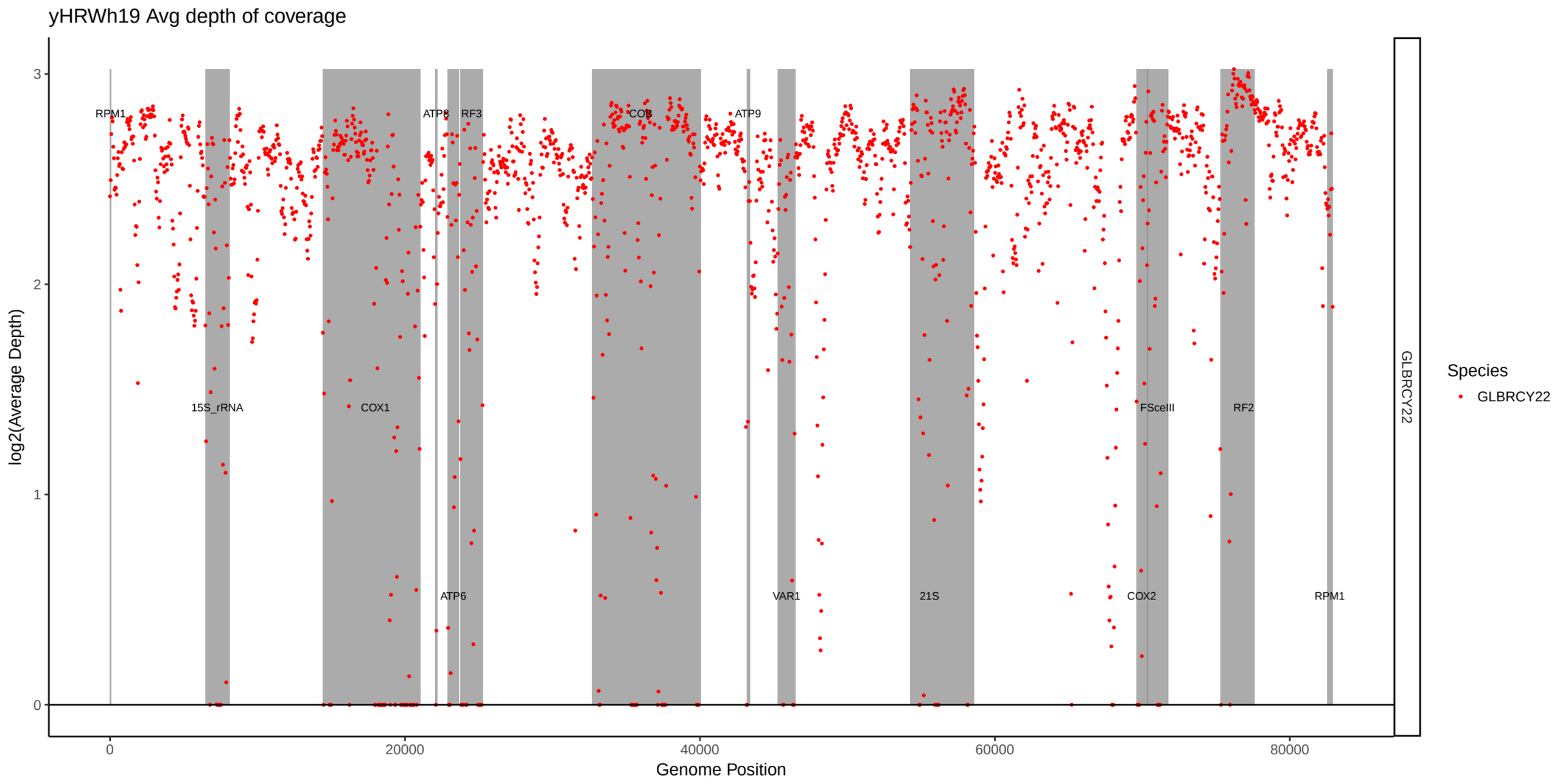

yHRWh19 Avg depth of coverage
log2(Average Depth)
Genome Position
0
20000
40000
60000
80000
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
FSparIII
ORF1
COX3
FSmikIII
FSarIII
FSuvaIII
15S
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001

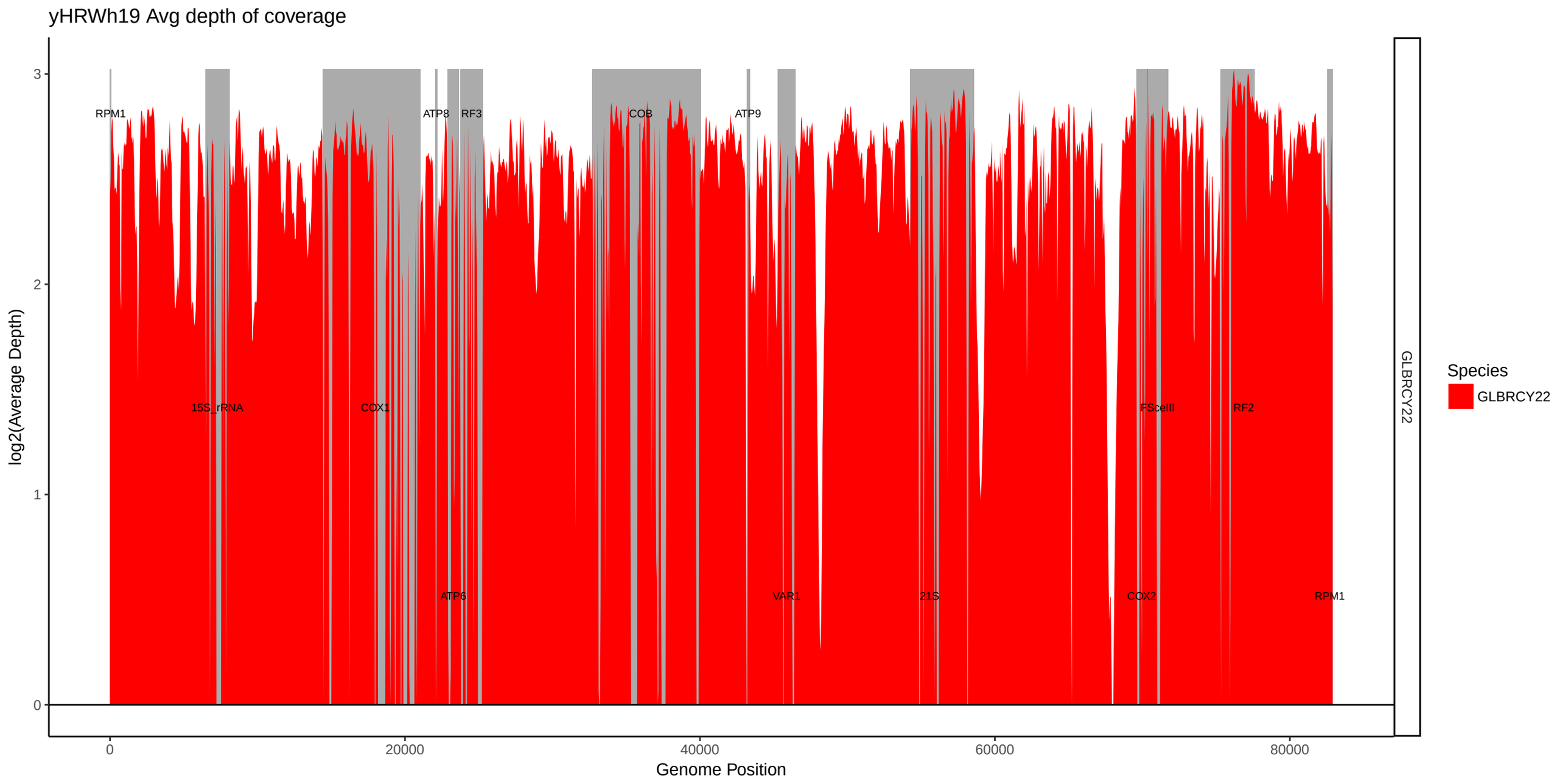

yHRWh19 Avg depth of coverage
Average Depth (limited)
Genome Position
0
20000
40000
60000
80000
0
500
1000
GLBRCY22
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
RPM1
CBS432
COX2
FSparIII
COX1
ATP6
COB
VAR1
15S
21S
ATP9
RF2
RPM1
IFO1815
21S
FSmikIII
RF2
RPM1
COX1
ATP6
15S
ATP9
21S
COX2
COB
VAR1
ZP591
VAR1
21S
COX2
ORF1
ATP8
ATP6
COB
ATP9
15S
COX3
RPM1
COX1
CBS10644
COB
ATP6
ATP9
15S
21S
COX3
RPM1
RF3
FSarIII
COX1
ATP8
VAR1
CBS7001
COX3
COB
ATP9
15S
21S
COX2
COX1
VAR1
FSuvaIII
ATP6
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001

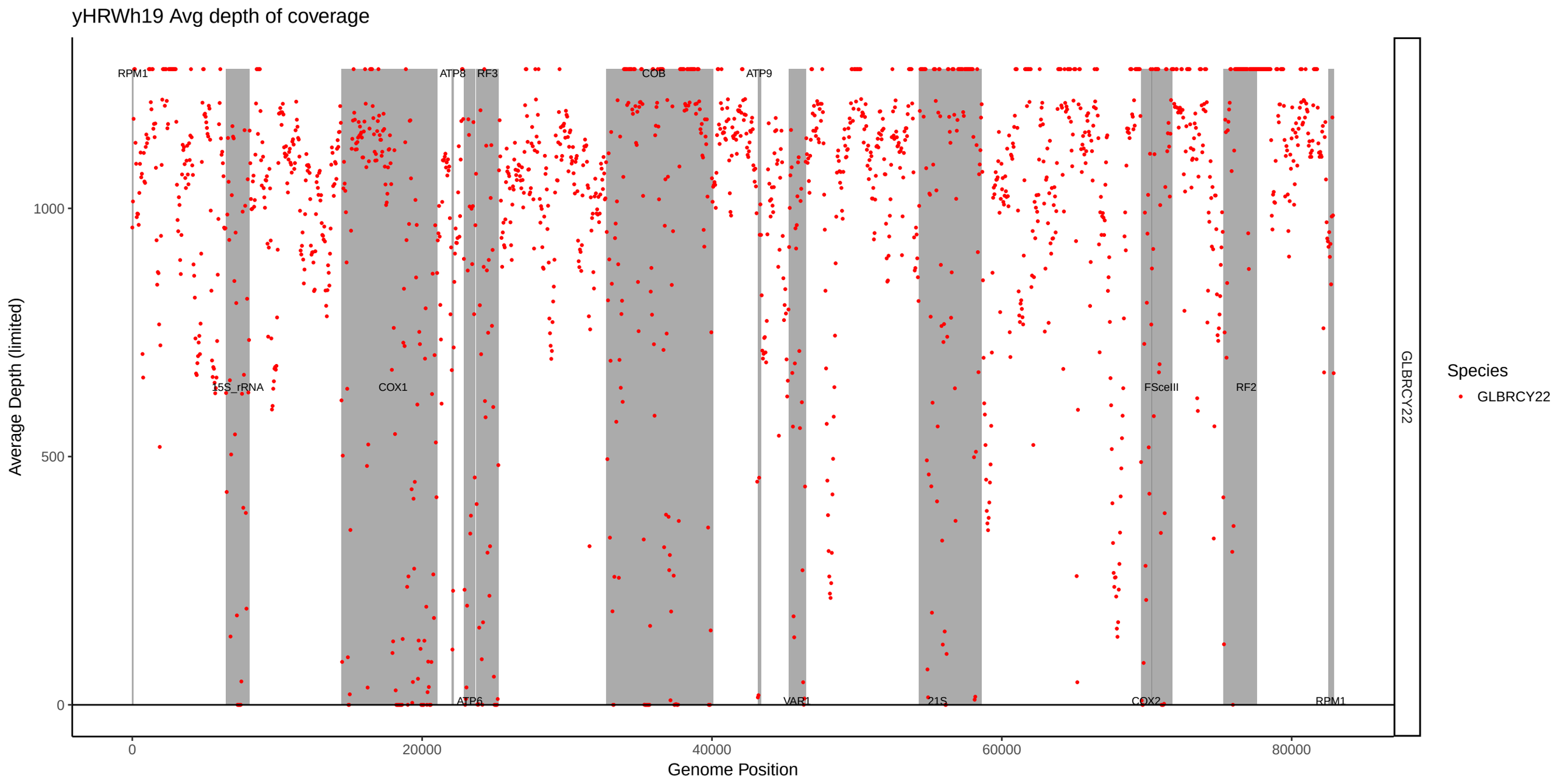

yHRWh19 Avg depth of coverage
Average Depth (limited)
Genome Position
0
20000
40000
60000
80000
0
500
1000
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
FSparIII
FSmikIII
ORF1
COX3
FSarIII
FSuvaIII
Species
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001

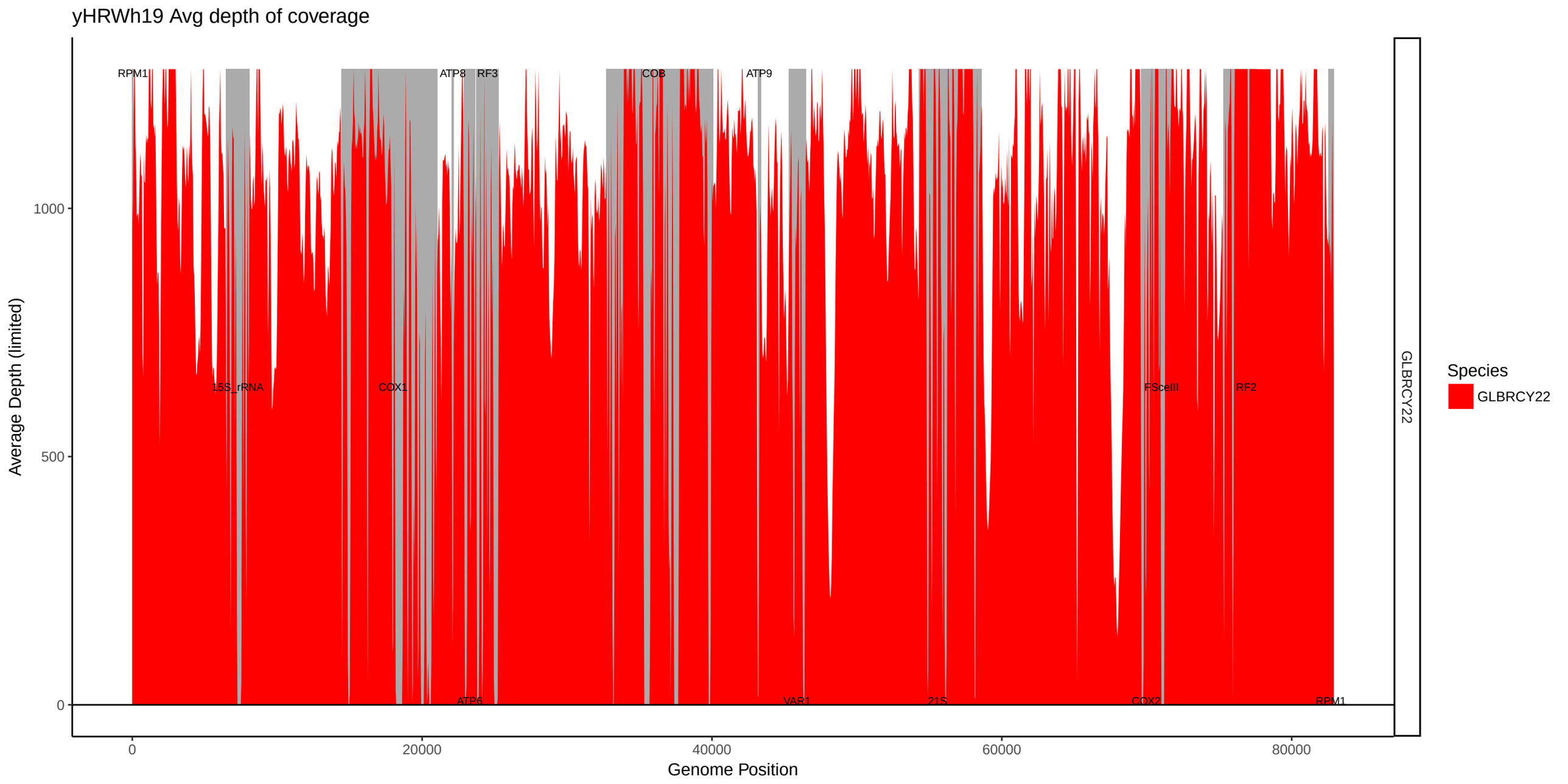

yHRWh19 Avg depth of coverage
Average Depth
1500
1000
500
0
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
RPM1
0
20000
40000
60000
80000
Genome Position
GLBRCY22
Species
GLBRCY22


yHRWh19 Avg depth of coverage
Average Depth
1500
1000
500
0
RPM1
15S_rRNA
ATP8
RF3
COX1
COB
ATP9
ATP6
VAR1
21S
FSceIII
COX2
RF2
RPM1
0
20000
40000
60000
80000
Genome Position
GLBRCY22
Species
GLBRCY22


yHRWh19 Avg depth of coverage
log2(Average Depth)
3
2
1
0
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
RPM1
0
20000
40000
60000
80000
Genome Position
GLBRCY22
Species
GLBRCY22


yHRWh19 Avg depth of coverage
log2(Average Depth)
3
2
1
0
RPM1
15S_rRNA
ATP8
RF3
ATP6
COX1
COB
VAR1
ATP9
21S
FSceIII
COX2
RF2
RPM1
0
20000
40000
60000
80000
Genome Position
GLBRCY22
Species
GLBRCY22


yHRWh19 Avg depth of coverage
RPM1
ATP8
RF3
COB
ATP9
15S_rRNA
COX1
FSceIII
RF2
ATP6
VAR1
21S
COX2
RPM1
Average Depth (limited)
1000
500
0
0
20000
40000
60000
80000
Genome Position
GLBRCY22
Species
GLBRCY22


yHRWh19 Avg depth of coverage
RPM1
ATP8
RF3
COB
ATP9
15S_rRNA
COX1
FSceIII
RF2
ATP6
VAR1
21S
COX2
RPM1
1000
500
0
Average Depth (limited)
0
20000
40000
60000
80000
Genome Position
GLBRCY22
Species
GLBRCY22