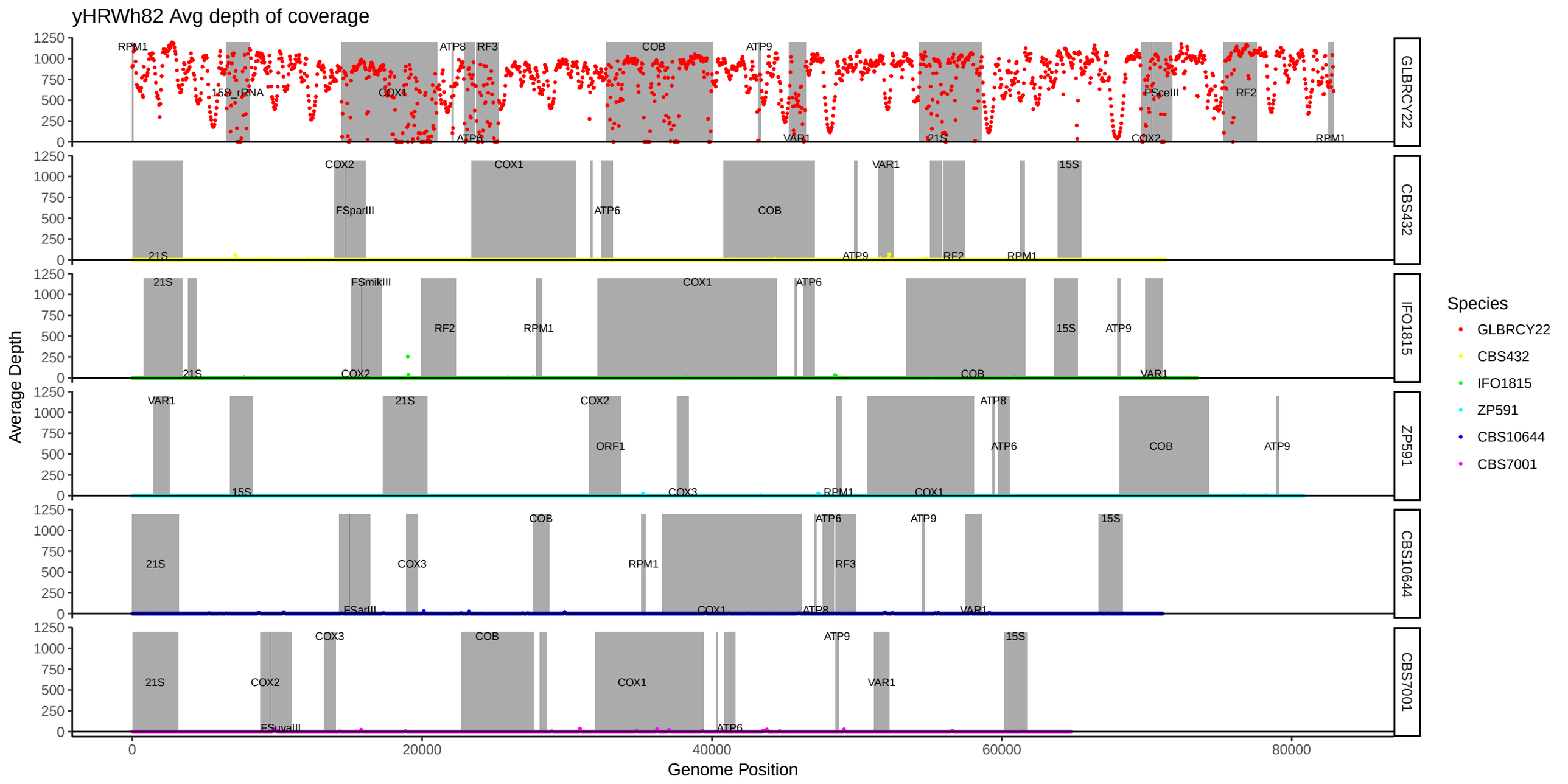
yHRWh82 Avg depth of coverage
Average Depth
Genome Position
0
20000
40000
60000
80000
0
250
500
750
1000
1250
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
FSparIII
FSmikIII
ORF1
COX3
FSarIII
FSuvaIII
Species
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CBS432
IFO1815
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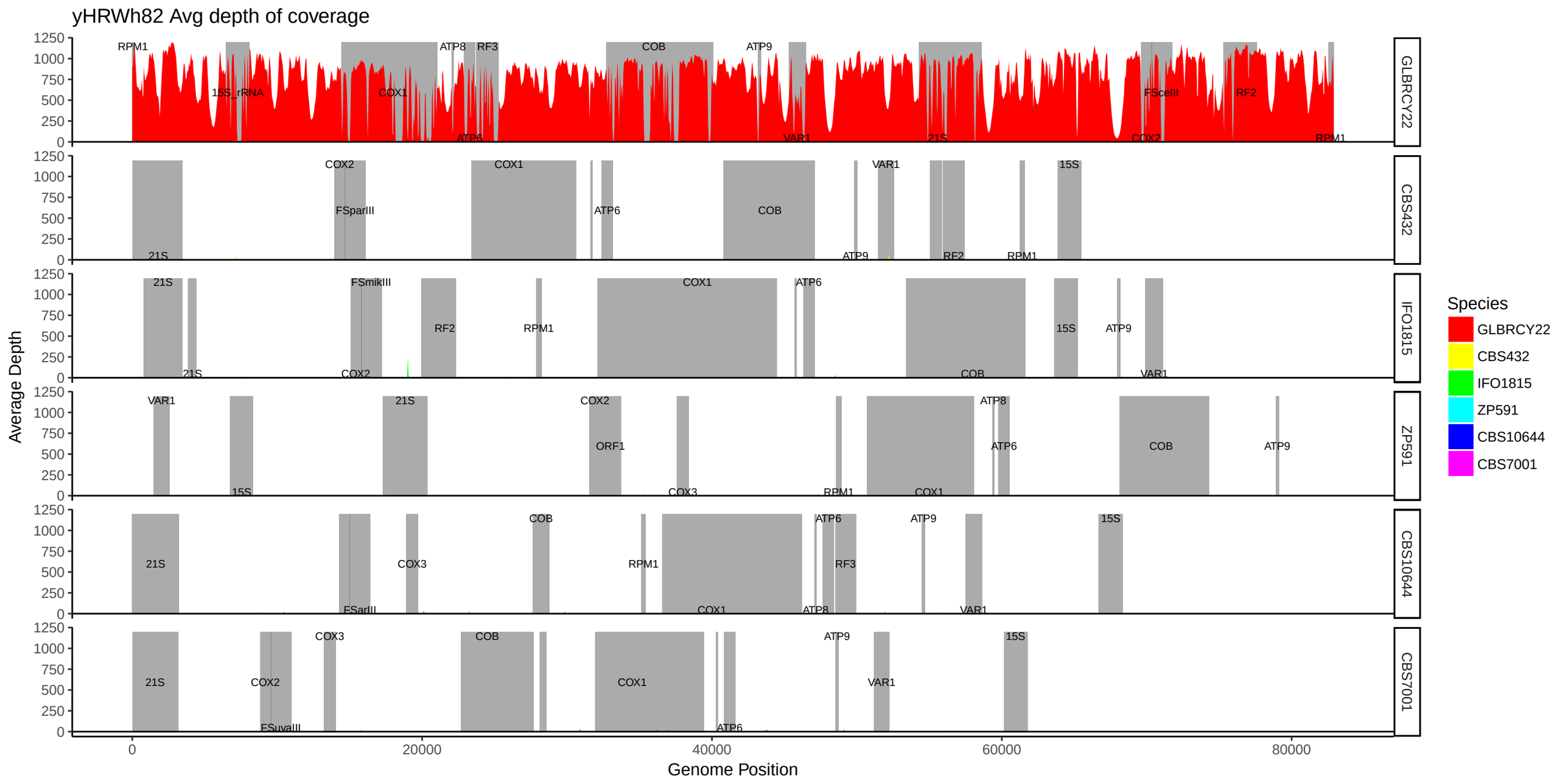
CBS10644
CBS7001


yHRWh82 Avg depth of coverage
Average Depth
Genome Position
0
20000
40000
60000
80000
0
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500
750
1000
1250
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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
COX3
ATP8
ATP6
COB
ATP9
15S
RPM1
COX1
CBS10644
21S
COX3
COB
RPM1
ATP6
RF3
ATP9
15S
FSarIII
COX1
ATP8
VAR1
CBS7001
21S
COX2
COX3
COB
COX1
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VAR1
15S
FSuvaIII
ATP6
Species
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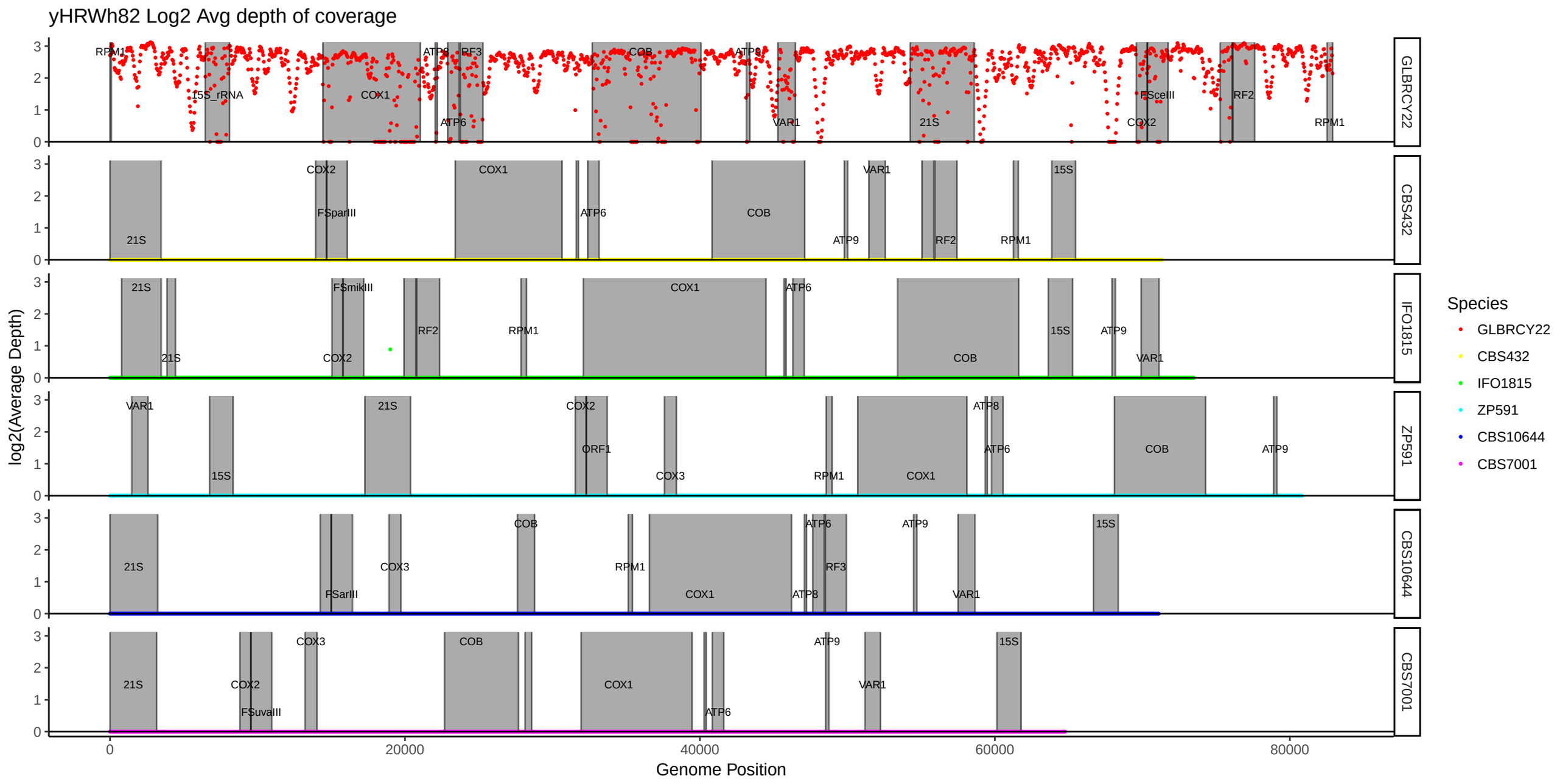
CBS10644
CBS7001


yHRWh82 Log2 Avg depth of coverage
log2(Average Depth)
Genome Position
0
20000
40000
60000
80000
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CBS432
IFO1815
ZP591
CBS10644
CBS7001
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP8
VAR1
21S
FSceII
COX2
RF2
RPM1
COX2
FSparIII
COX1
ATP6
COB
VAR1
15S
21S
ATP9
RF2
RPM1
21S
FSmikIII
COX1
ATP6
RF2
RPM1
15S
ATP9
21S
COX2
COB
VAR1
VAR1
21S
COX2
ATP8
15S
ORF1
COX3
RPM1
COX1
ATP6
COB
ATP9
21S
COB
ATP6
ATP9
15S
COX3
RPM1
RF3
FSarIII
COX1
ATP8
VAR1
21S
COX3
COB
ATP9
15S
COX2
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VAR1
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ATP6
Species
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CBS432
IFO1815
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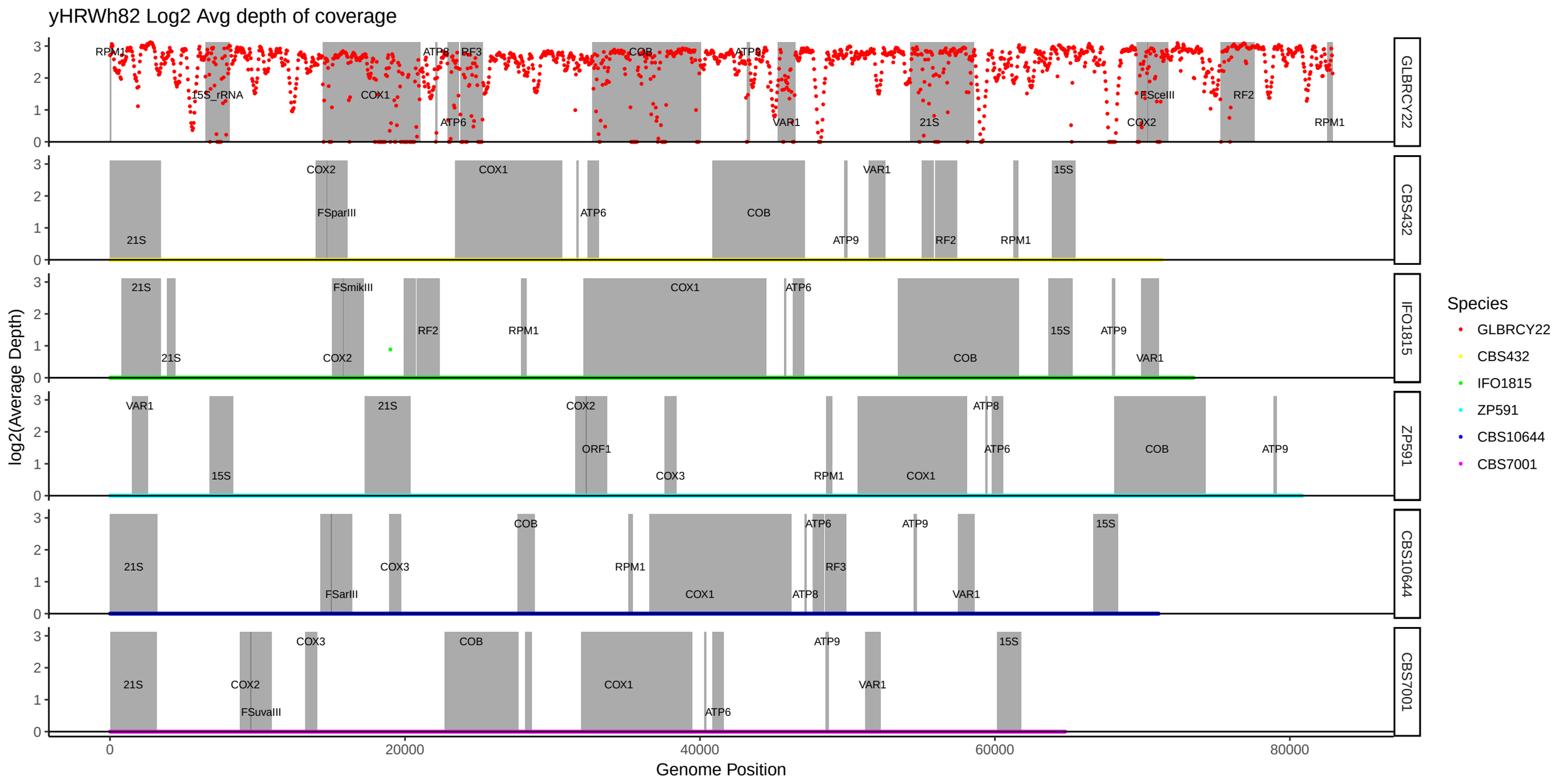
CBS10644
CBS7001


yHRWh82 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
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FSarIII
FSuvaIII
Species
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CBS432
IFO1815
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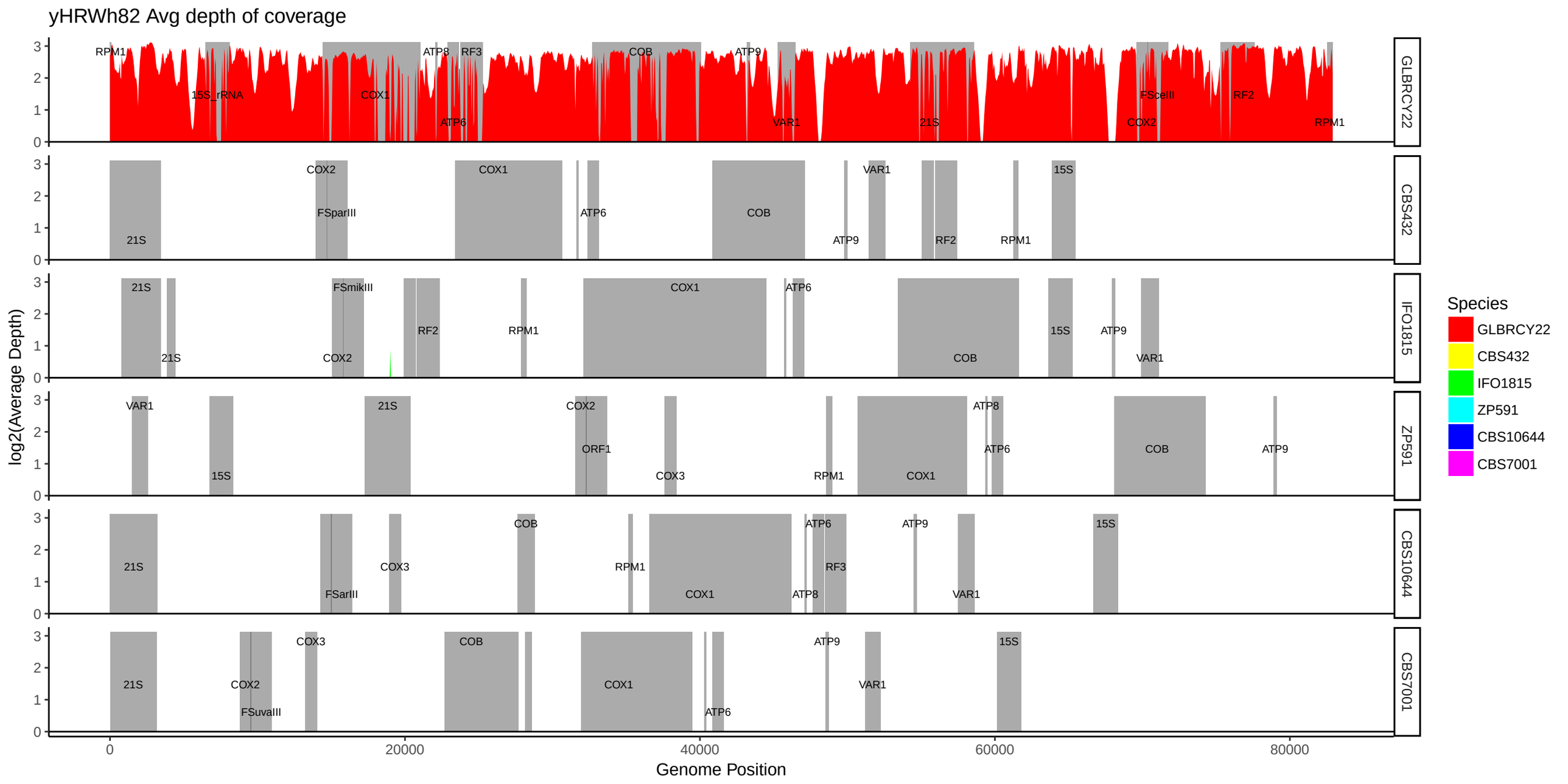
CBS10644
CBS7001


yHRWh82 Avg depth of coverage
log2(Average Depth)
Genome Position
0
20000
40000
60000
80000
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
Species
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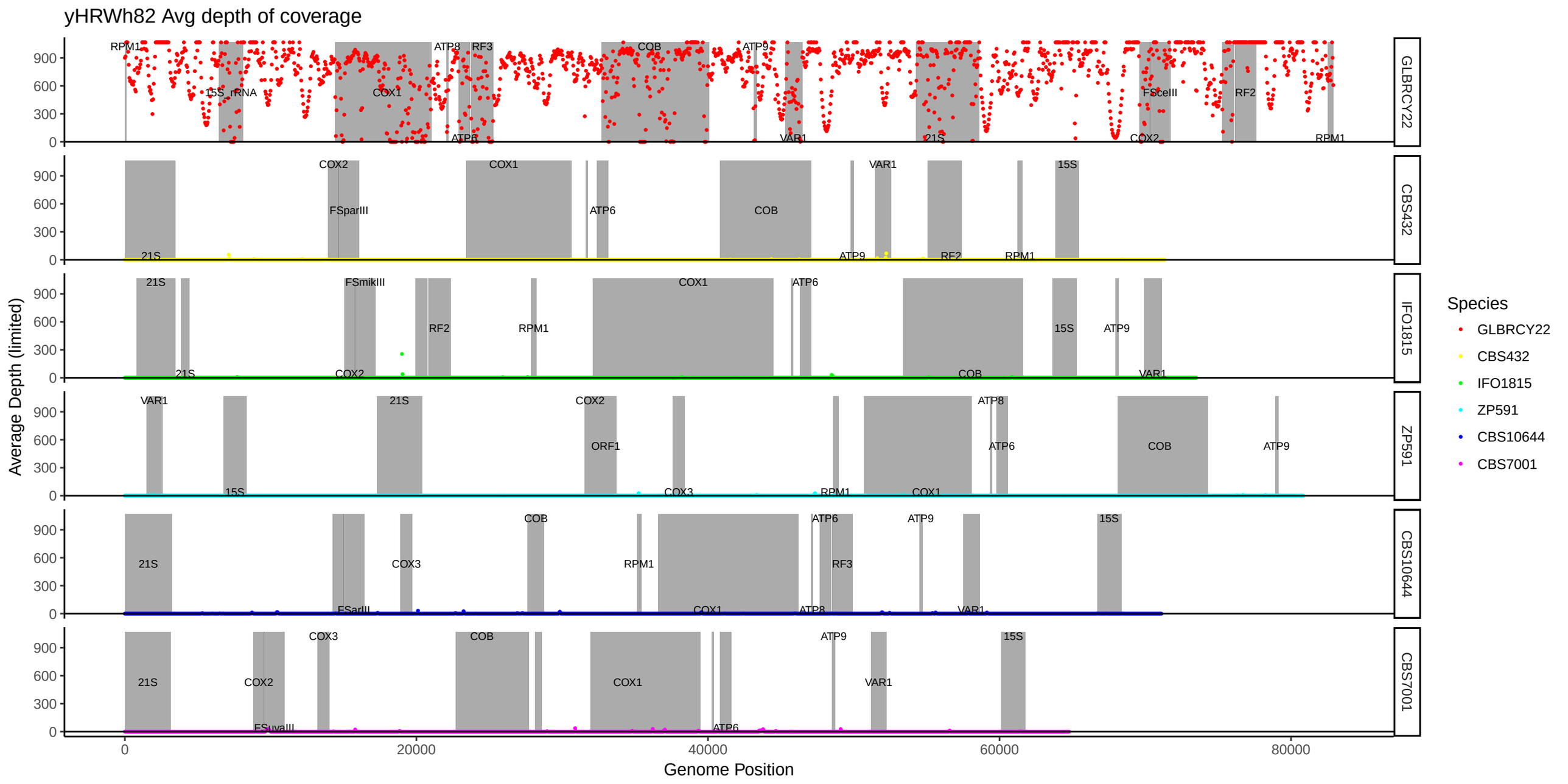
CBS432
IFO1815
ZP591
CBS10644
CBS7001


yHRWh82 Avg depth of coverage
Average Depth (limited)
Genome Position
0
20000
40000
60000
80000
0
300
600
900
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
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CBS432
IFO1815
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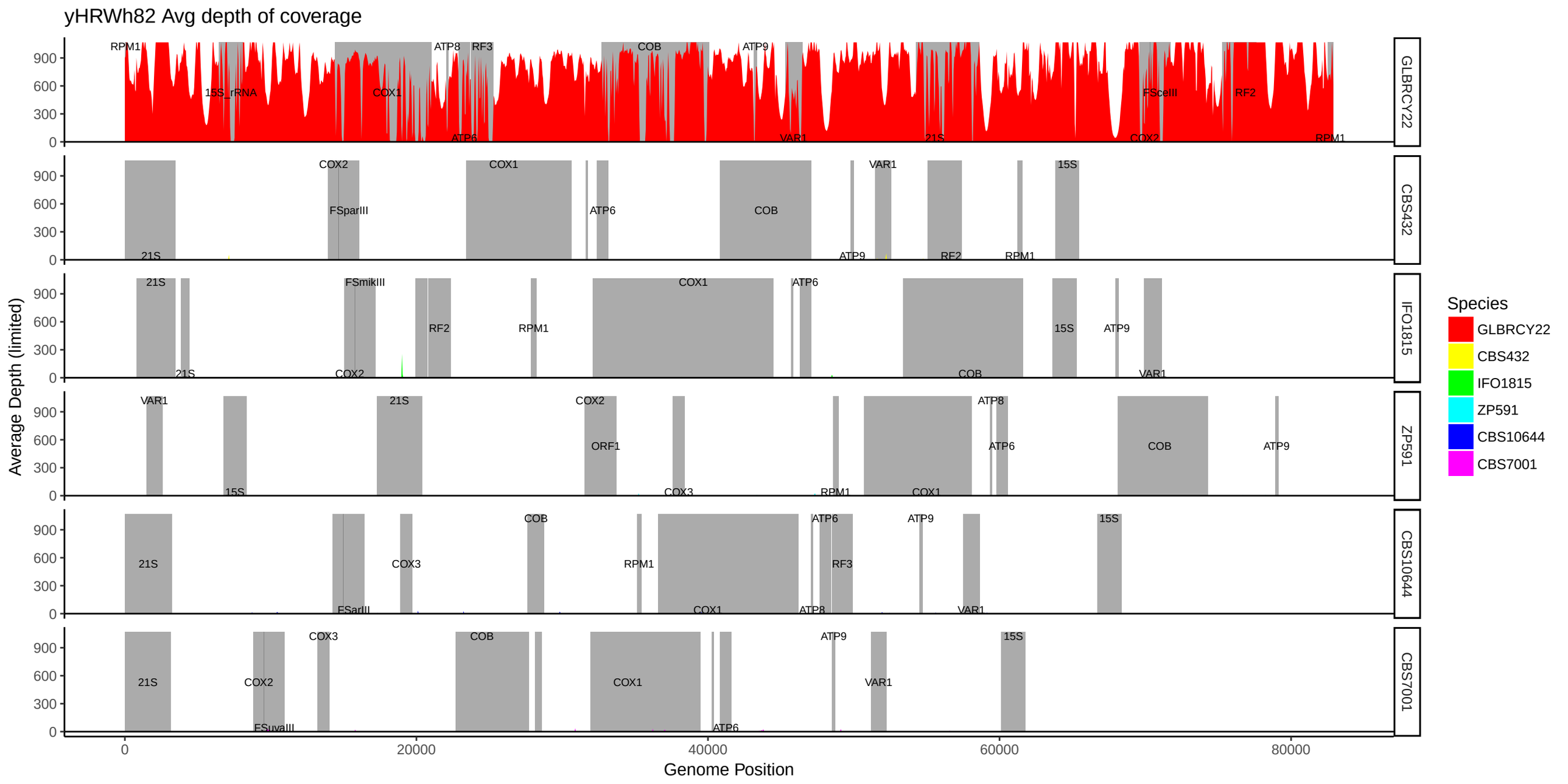
CBS7001


yHRWh82 Avg depth of coverage
Average Depth (limited)
Genome Position
0
20000
40000
60000
80000
0
300
600
900
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001
RPM1
15S_rRNA
ATP8
RF3
COB
ATP9
COX1
ATP6
VAR1
21S
FSceIII
RF2
COX2
FSparIII
15S
RPM1
FSmikIII
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COX3
FSarIII
FSuvaIII
Species
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CBS432
IFO1815
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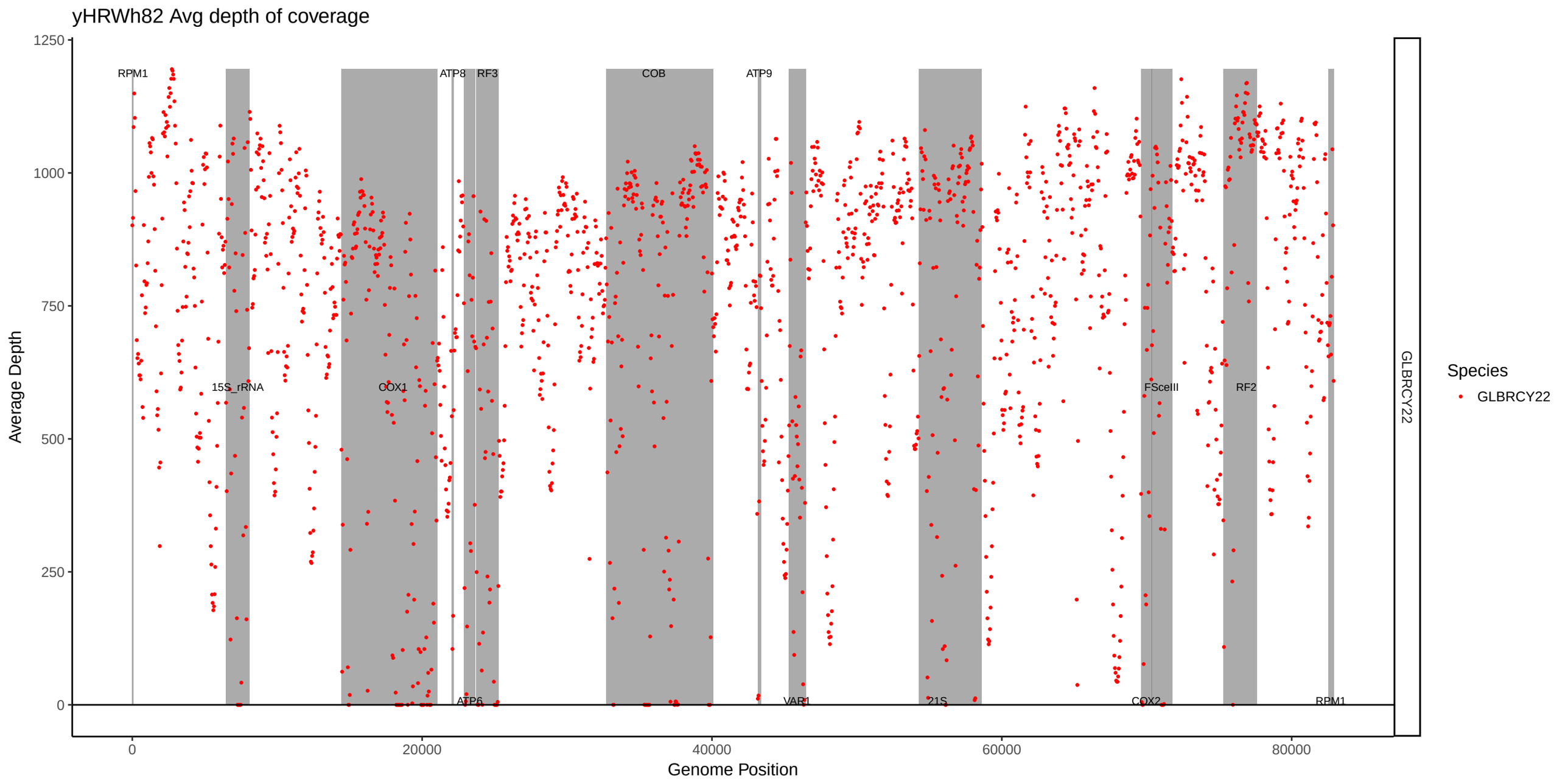
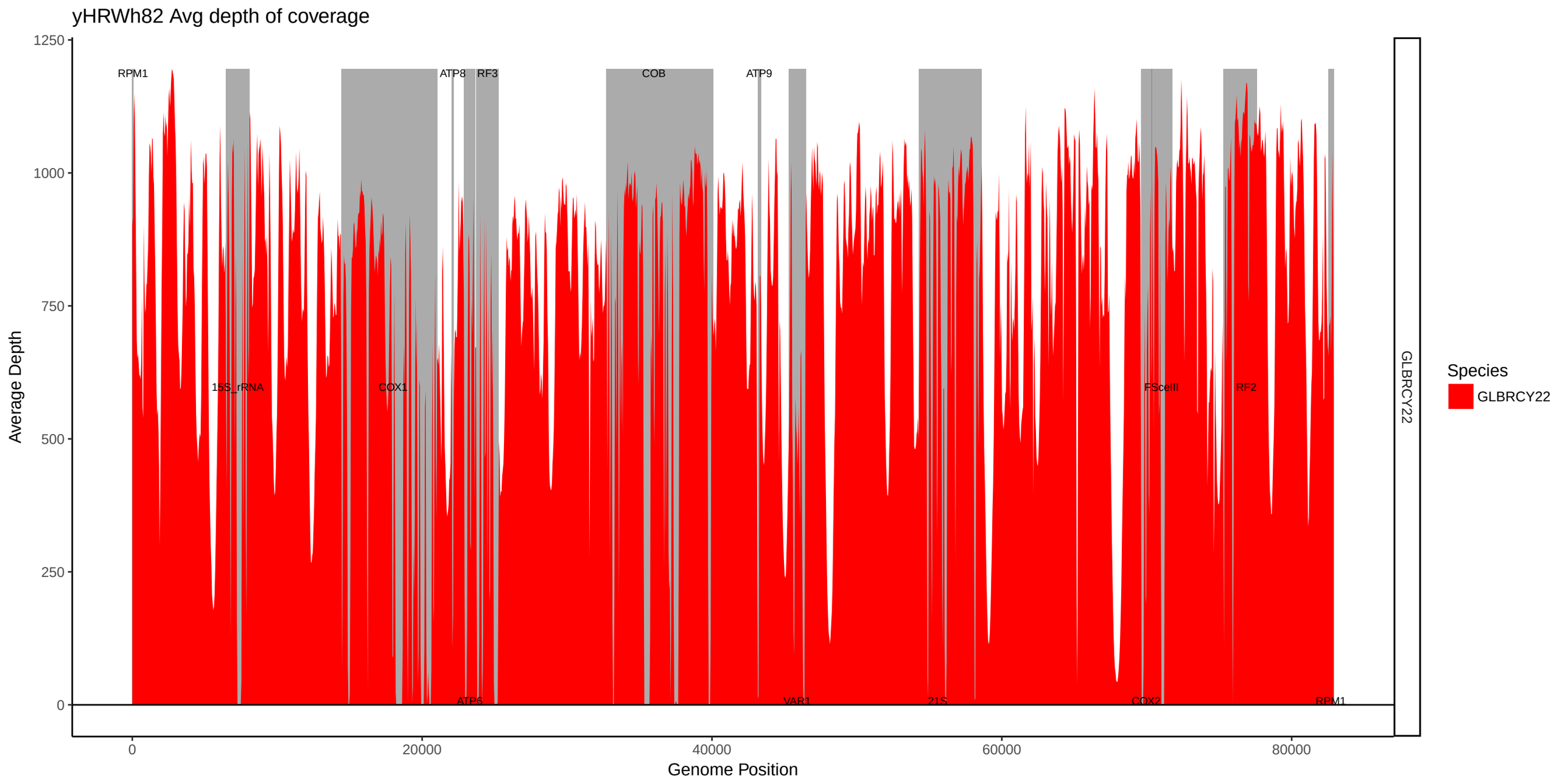
CBS7001


yHRWh82 Avg depth of coverage
Average Depth
Genome Position
1250
1000
750
500
250
0
0
20000
40000
60000
80000
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
RPM1
GLBRCY22
Species
GLBRCY22


yHRWh82 Avg depth of coverage
Average Depth
1250
1000
750
500
250
0
RPM1
ATP8
RF3
COB
ATP9
15S_rRNA
COX1
FSceIII
RF2
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COX2
RPM1
0
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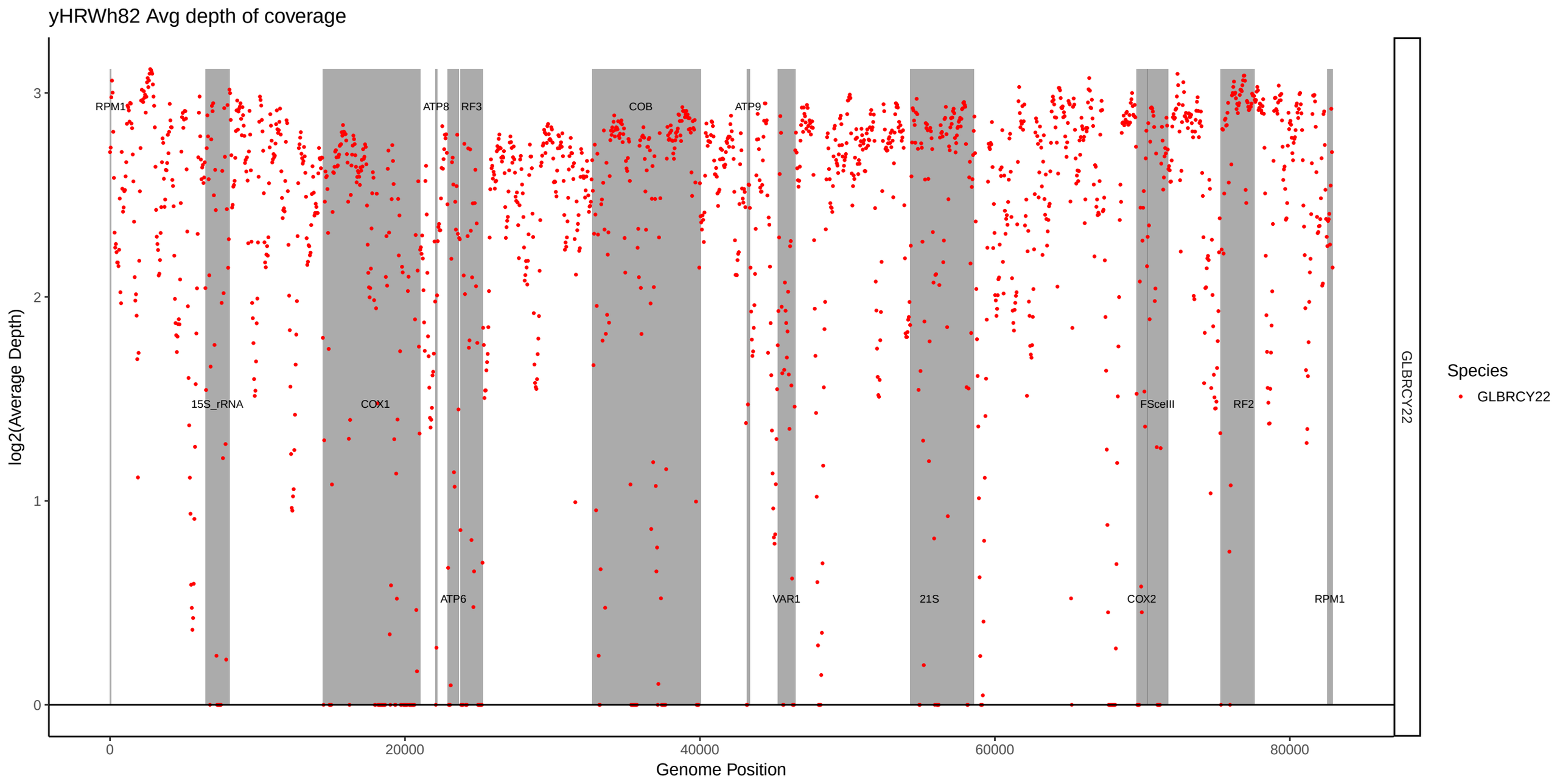
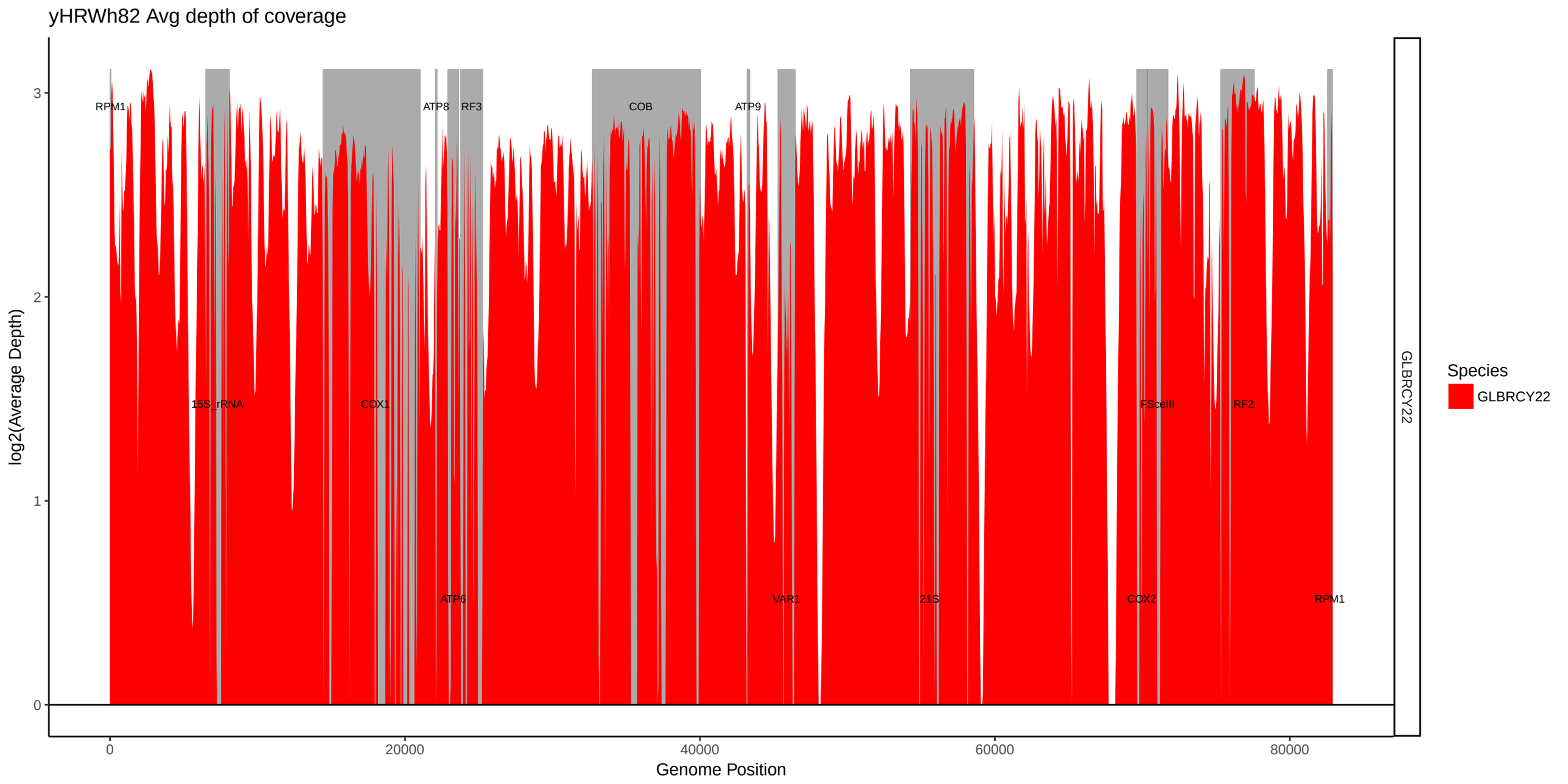
40000
60000
80000
Genome Position
GLBRCY22
Species
GLBRCY22


yHRWh82 Avg depth of coverage
log2(Average Depth)
3
2
1
0
RPM1
15S_rRNA
COX1
ATP8
ATP6
RF3
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
RPM1
0
20000
40000
60000
80000
Genome Position
GLBRCY22
Species
GLBRCY22


yHRWh82 Avg depth of coverage
log2(Average Depth)
3
2
1
0
RPM1
15S_rRNA
COX1
ATP8
RF3
ATP6
COB
ATP9
VAR1
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RF2
RPM1
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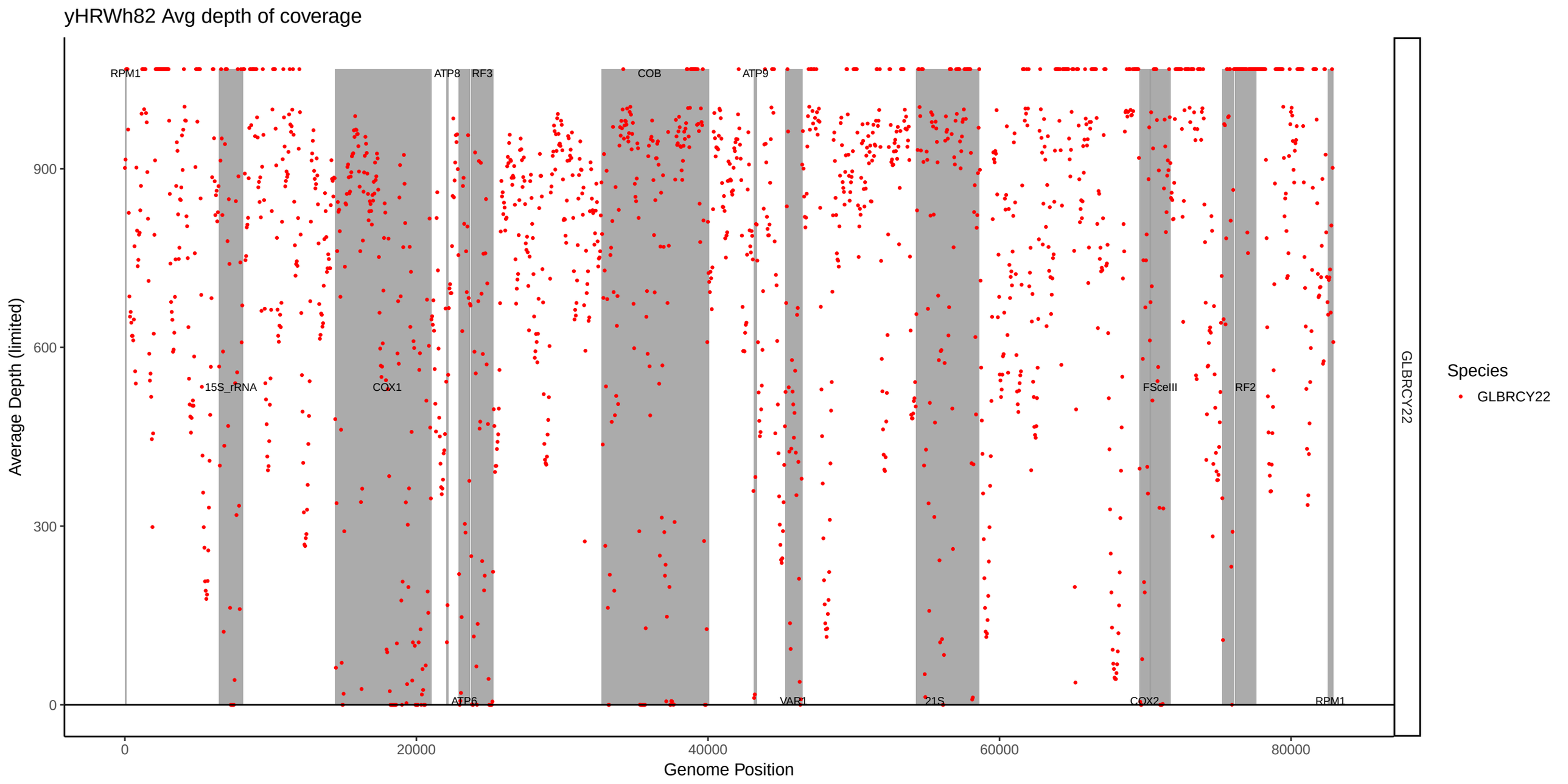
Genome Position
GLBRCY22
Species
GLBRCY22


yHRWh82 Avg depth of coverage
RPM1
15S_rRNA
COX1
ATP8
ATP6
RF3
COB
ATP9
VAR1
21S
FSceIII
COX2
RF2
RPM1
Average Depth (limited)
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300
600
900
0
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40000
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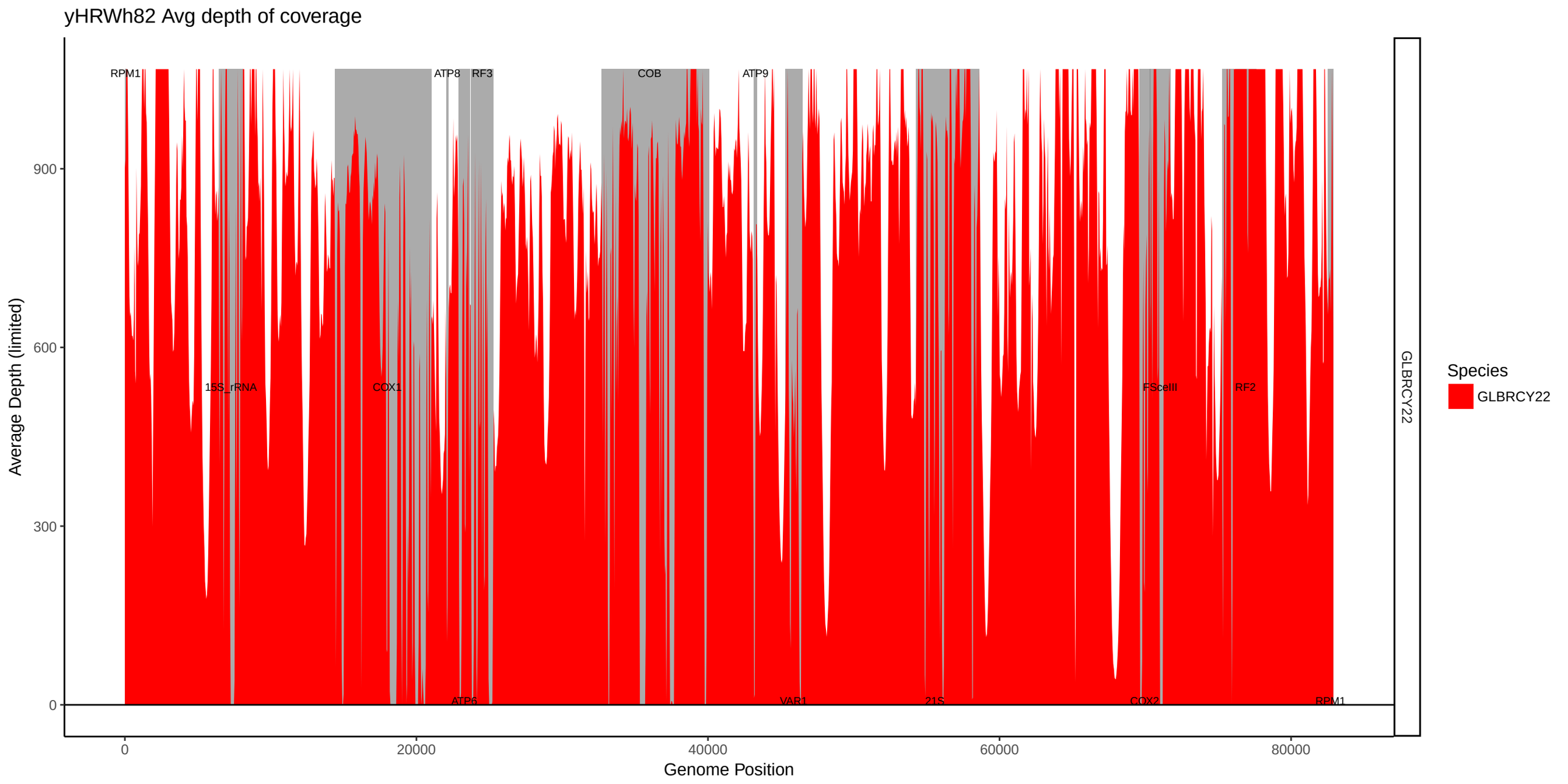
Genome Position
GLBRCY22
Species
GLBRCY22


yHRWh82 Avg depth of coverage
RPM1
ATP8
RF3
COB
ATP9
15S_rRNA
COX1
FSceIII
RF2
ATP6
VAR1
21S
COX2
RPM1
900
600
300
0
Average Depth (limited)
0
20000
40000
60000
80000
Genome Position
GLBRCY22
Species
GLBRCY22