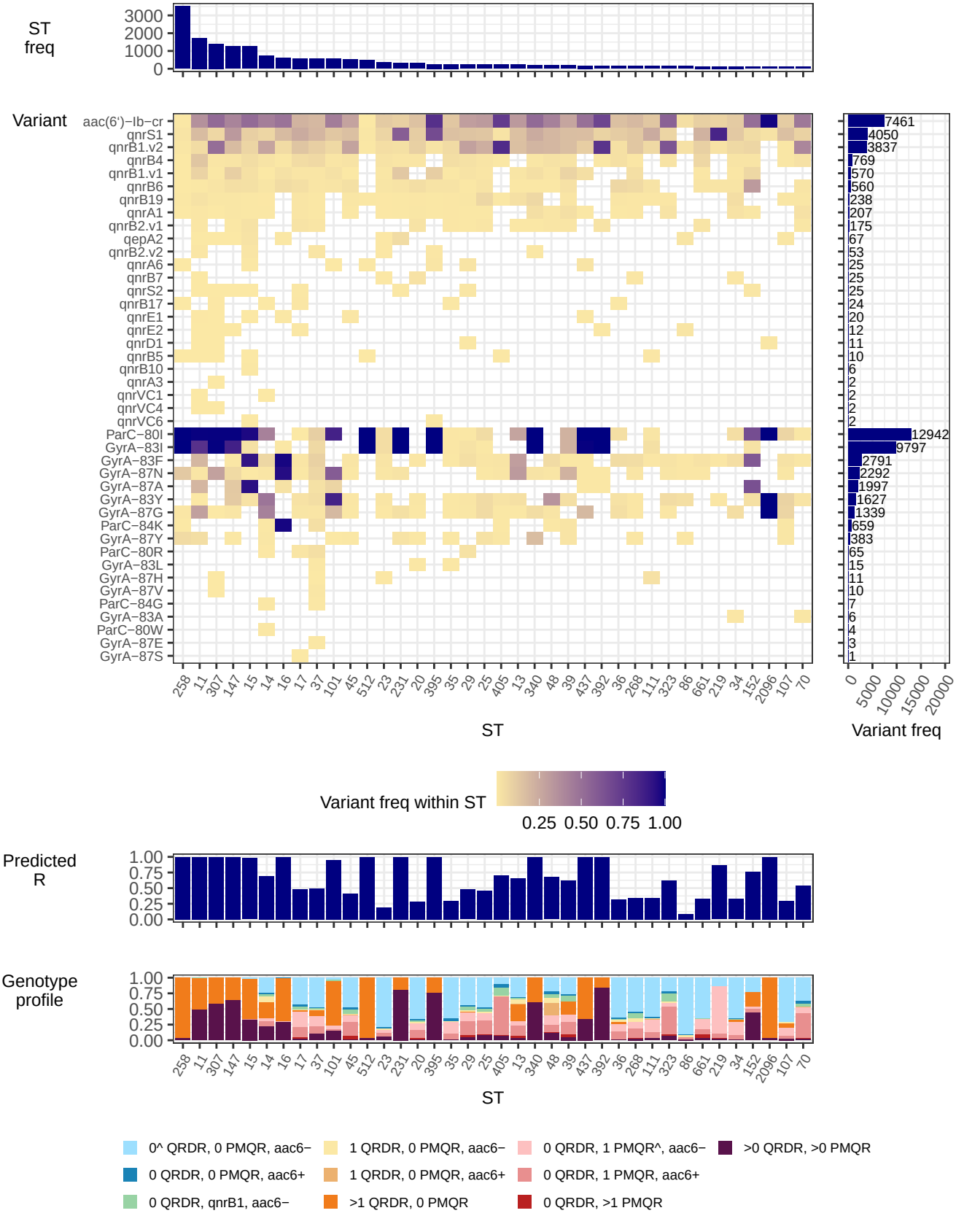
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Variant
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qnrA1
qnrB2.v1
qepA2
qnrB2.v2
qnrA6
qnrB7
qnrS2
qnrB17
qnrE1
qnrE2
qnrD1
qnrB5
qnrB10
qnrA3
qnrVC1
qnrVC4
qnrVC6
ParC−80I
GyrA−83I
GyrA−83F
GyrA−87N
GyrA−87A
GyrA−83Y
GyrA−87G
ParC−84K
GyrA−87Y
ParC−80R
GyrA−83L
GyrA−87H
GyrA−87V
ParC−84G
GyrA−83A
ParC−80W
GyrA−87E
GyrA−87S
7461
4050
3837
769
570
560
238
207
175
67
53
25
25
25
24
20
12
11
10
6
2
2
2
2
12942
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2791
2292
1997
1627
1339
659
383
65
15
11
10
7
6
4
3
1
258 11 307 147 15 14 16 17 37 101 45 512 23 231 20 395 35 29 25 405 13 340 48 39 437 392 36 268 111 323 86 661 219 34 152 2096 107 70
ST
0 5000 10000 15000 20000
Variant freq
Variant freq within ST
0.25 0.50 0.75 1.00
Predicted R
1.00
0.75
0.50
0.25
0.00
Genotype profile
1.00
0.75
0.50
0.25
0.00
258 11 307 147 15 14 16 17 37 101 45 512 23 231 20 395 35 29 25 405 13 340 48 39 437 392 36 268 111 323 86 661 219 34 152 2096 107 70
ST
0^ QRDR, 0 PMQR, aac6−
1 QRDR, 0 PMQR, aac6−
0 QRDR, 1 PMQR^, aac6−
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0 QRDR, qnrB1, aac6−
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0 QRDR, >1 PMQR