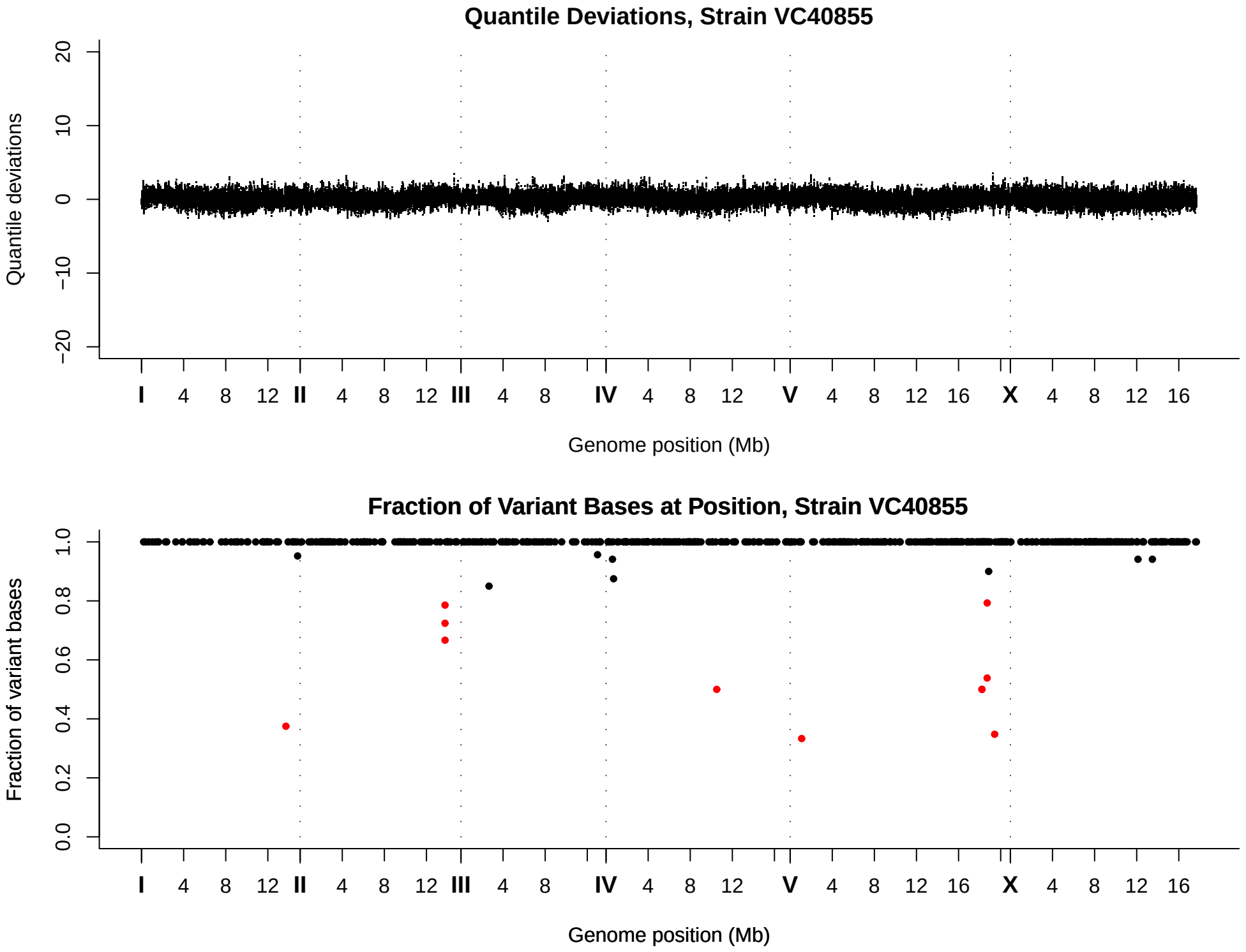
Quantile Deviations, Strain VC40855
Quantile deviations
20
10
0
−10
−20
I 4 8 12 II 4 8 12 III 4 8 IV 4 8 12 V 4 8 12 16 X 4 8 12 16
Genome position (Mb)
Fraction of Variant Bases at Position, Strain VC40855
Fraction of variant bases
1.0
0.8
0.6
0.4
0.2
0.0
I 4 8 12 II 4 8 12 III 4 8 IV 4 8 12 V 4 8 12 16 X 4 8 12 16
Genome position (Mb)