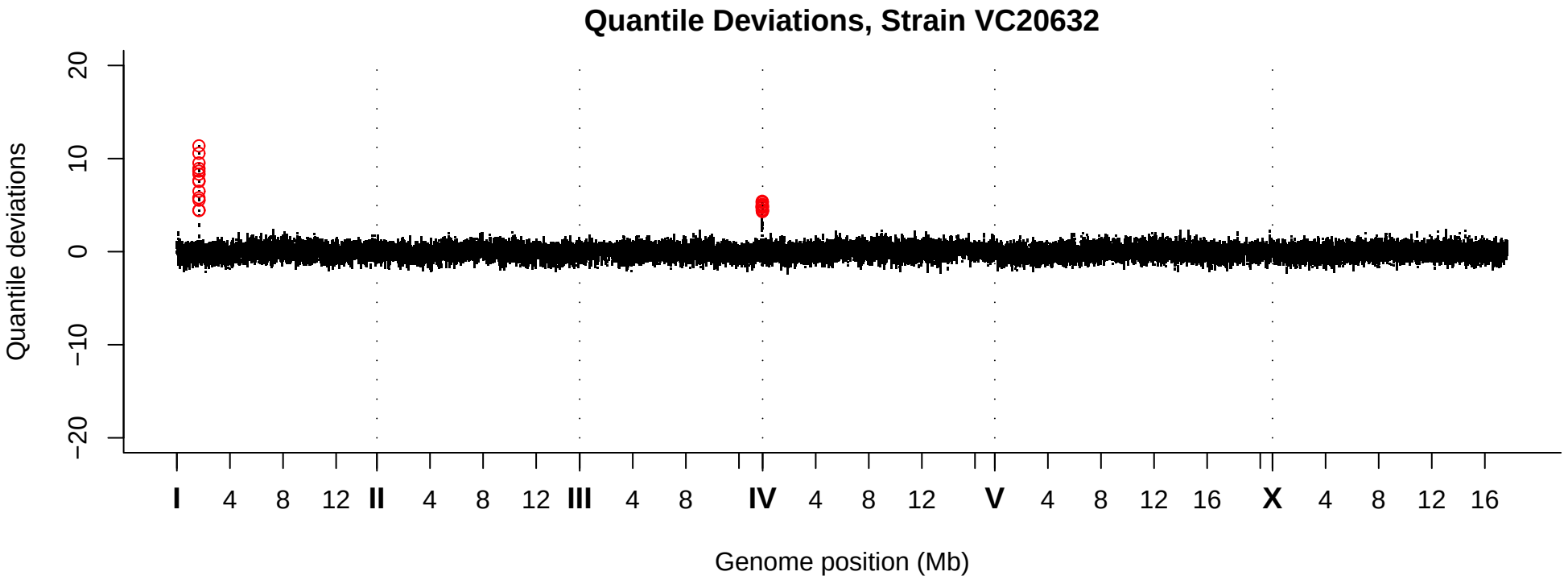
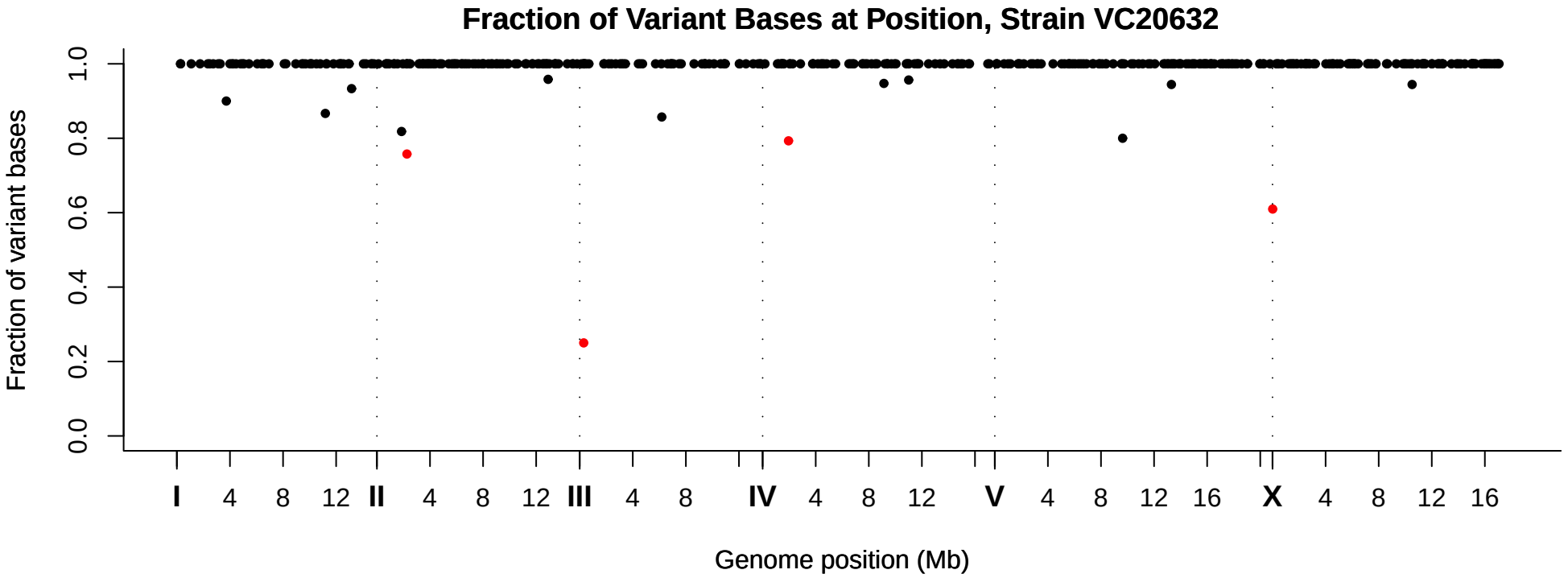

Quantile Deviations, Strain VC20632
Quantile deviations
20
10
0
−10
−20
I
4
8
12
II
4
8
12
III
4
8
IV
4
8
12
V
4
8
12
16
X
4
8
12
16
Genome position (Mb)


Fraction of Variant Bases at Position, Strain VC20632
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)