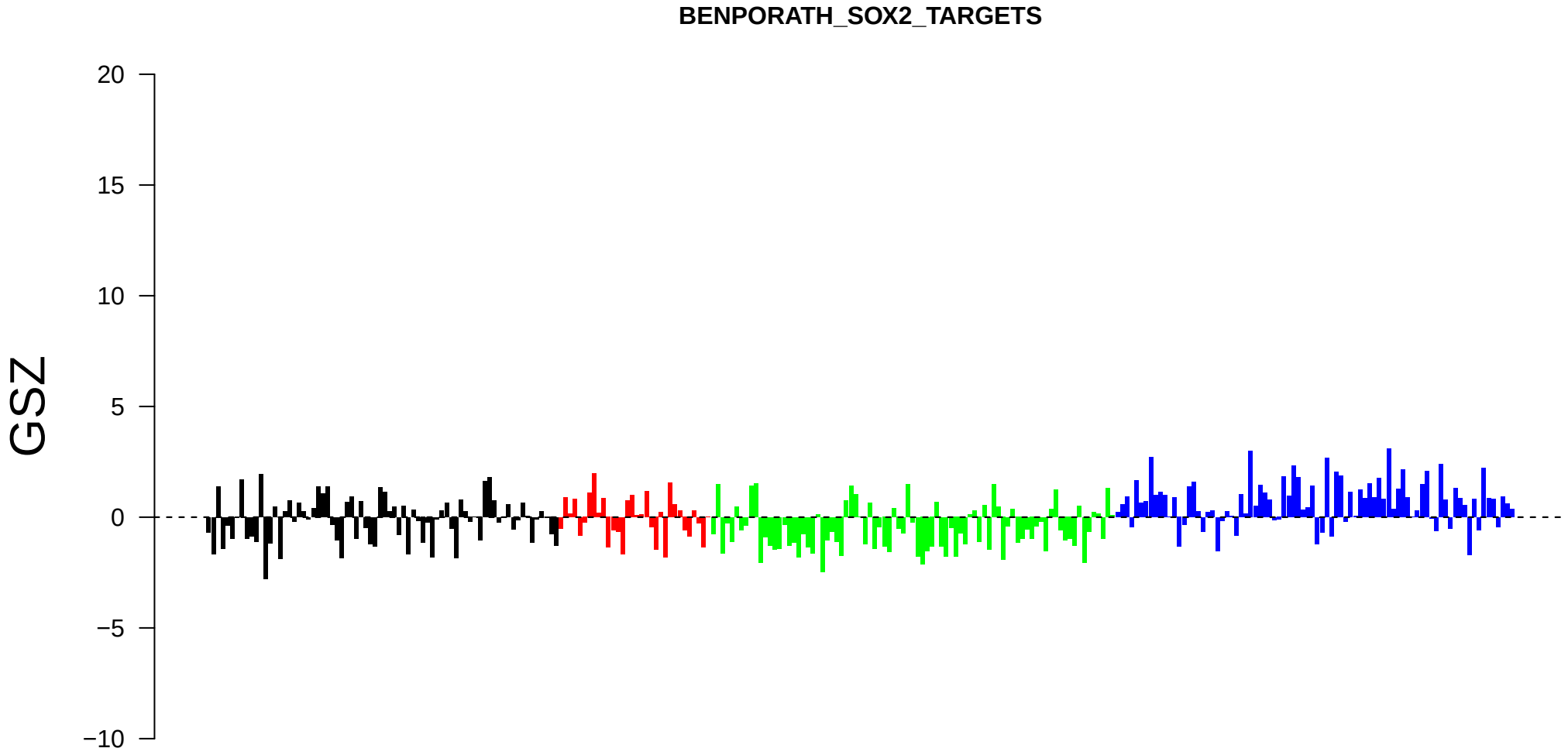
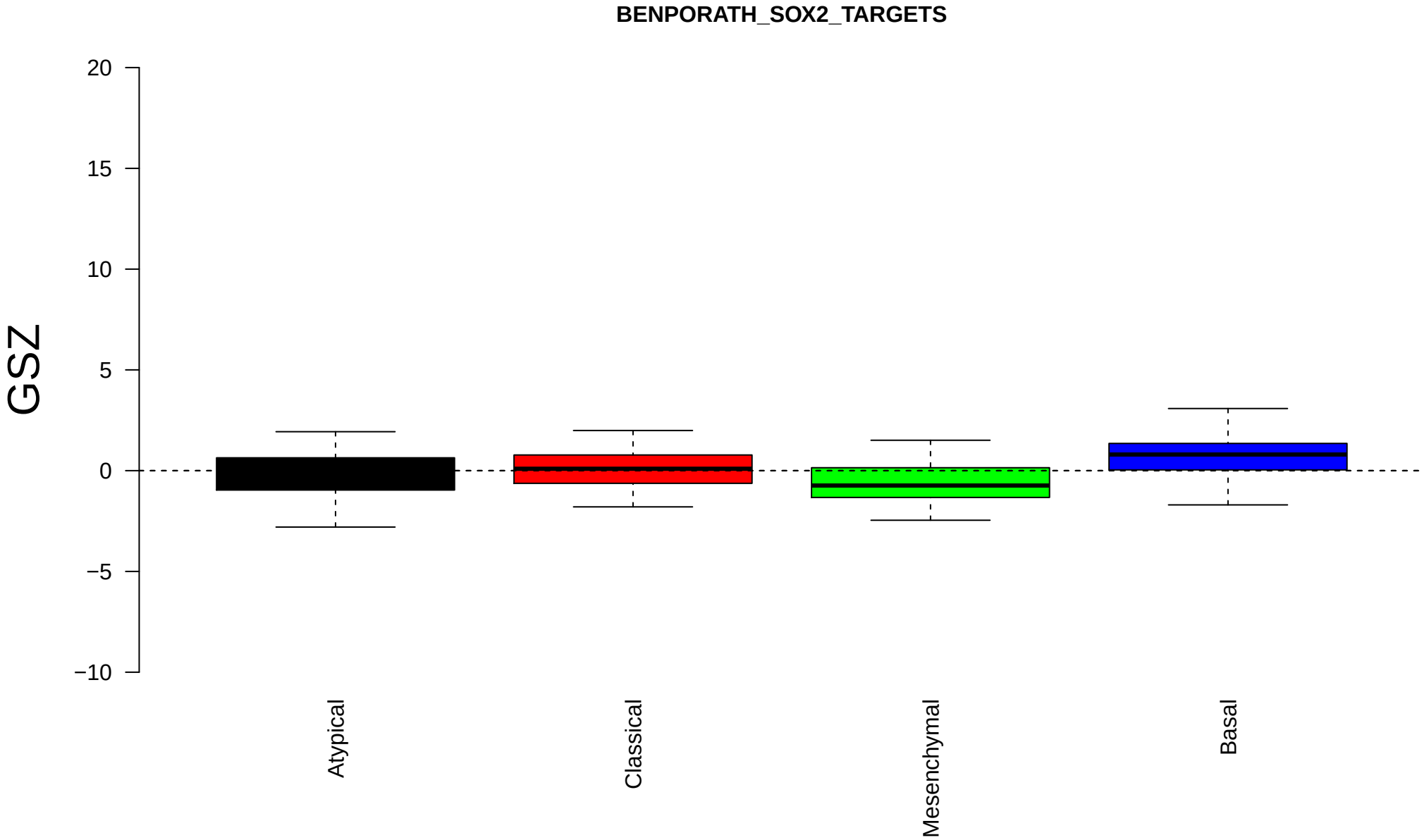
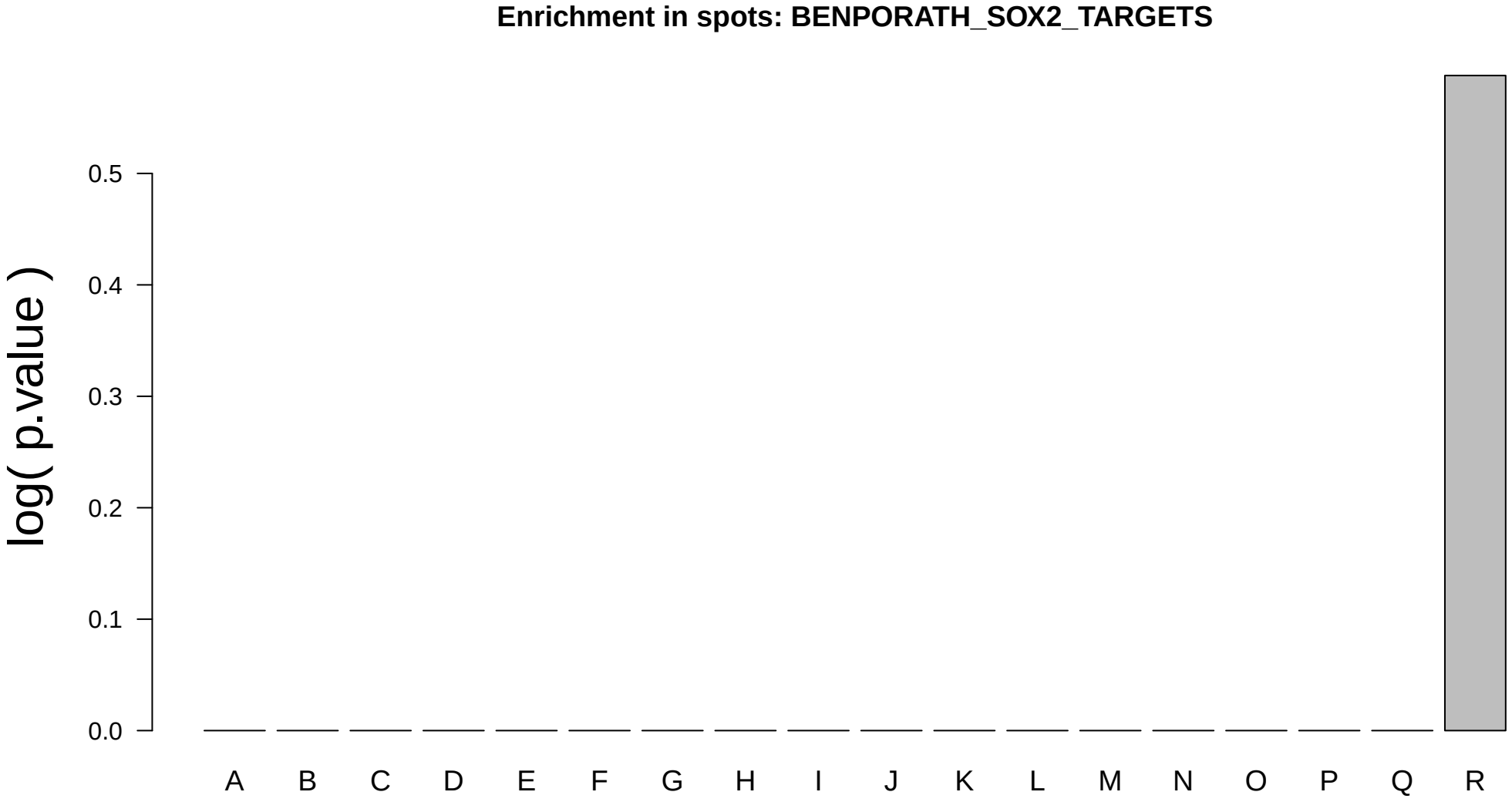
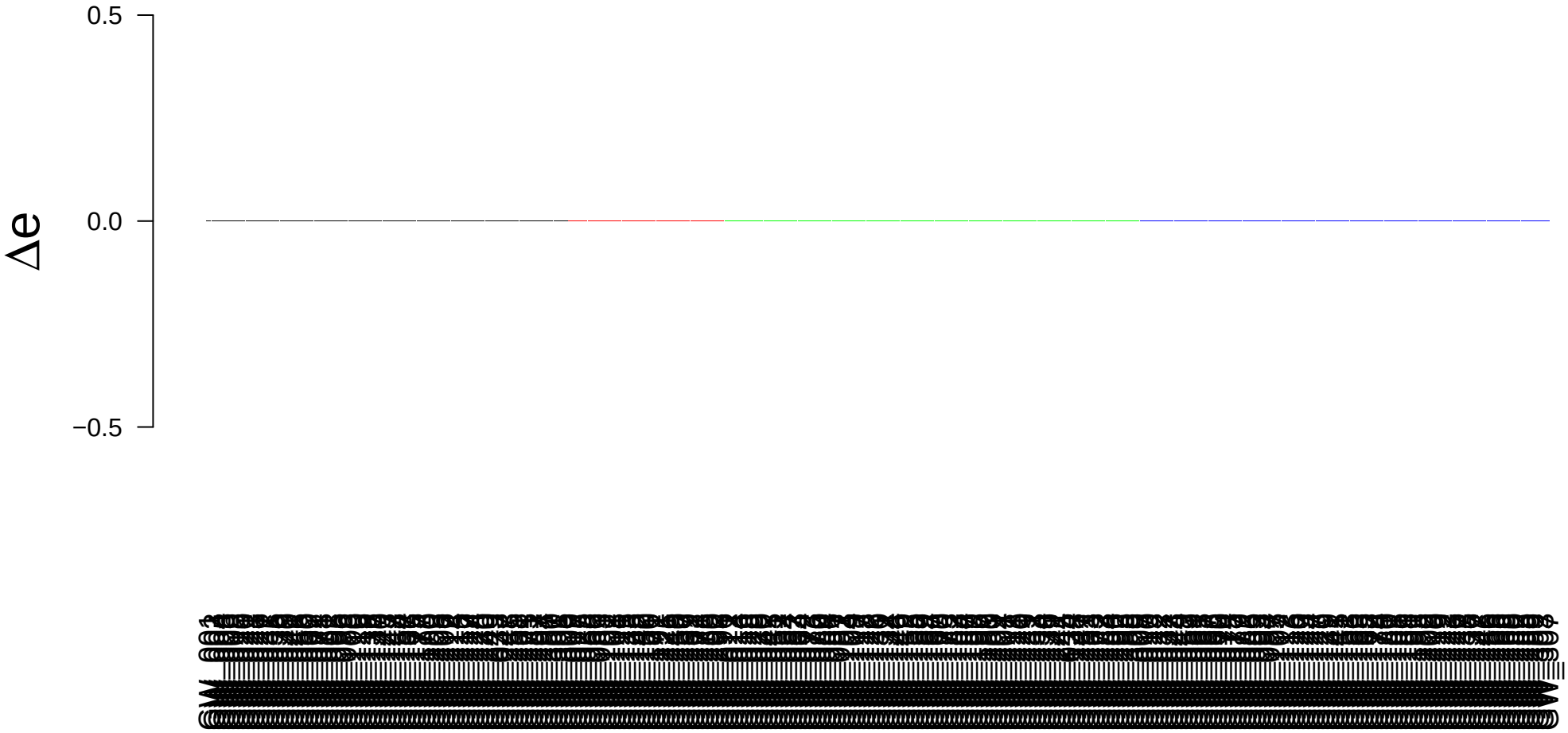
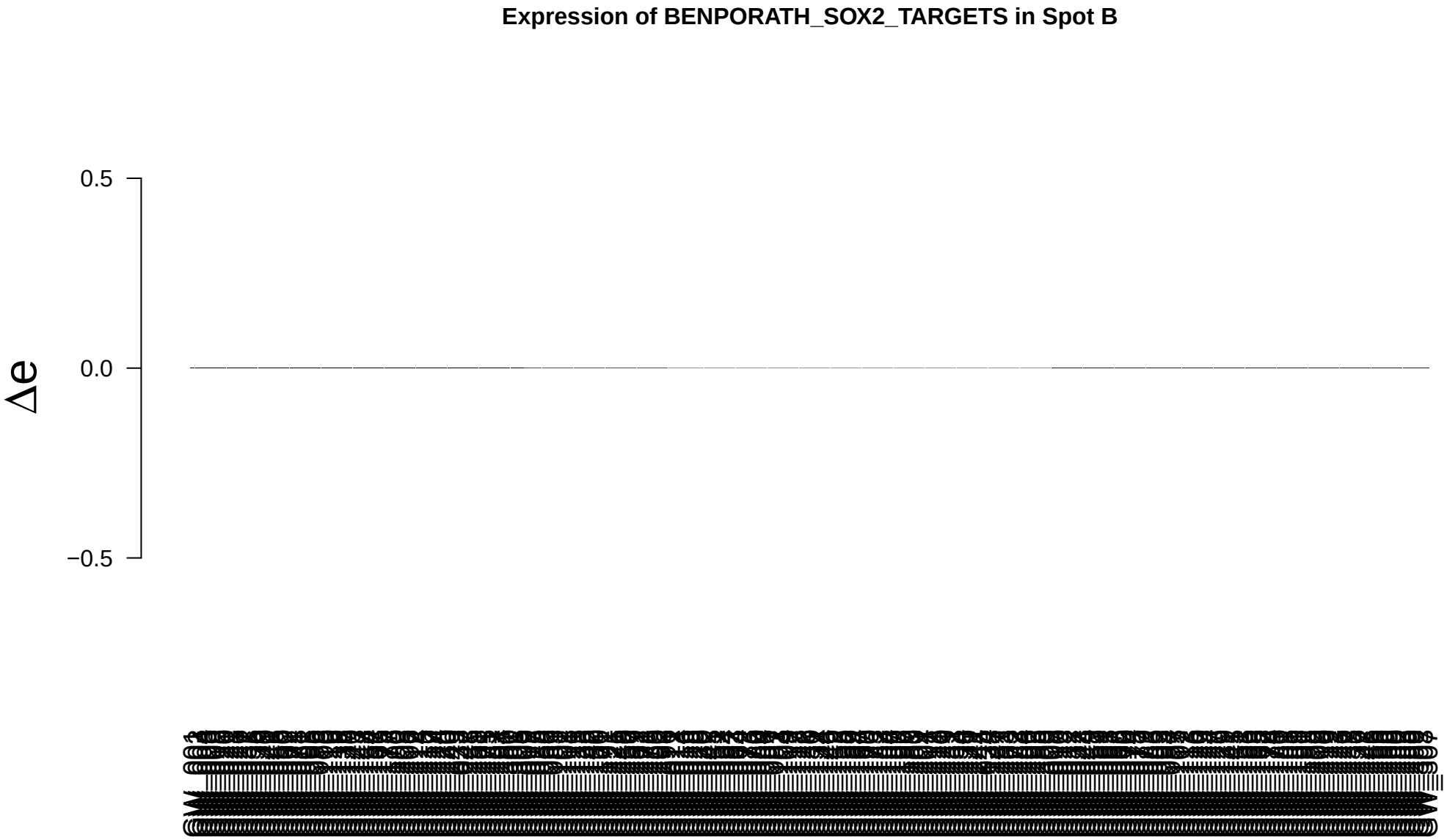
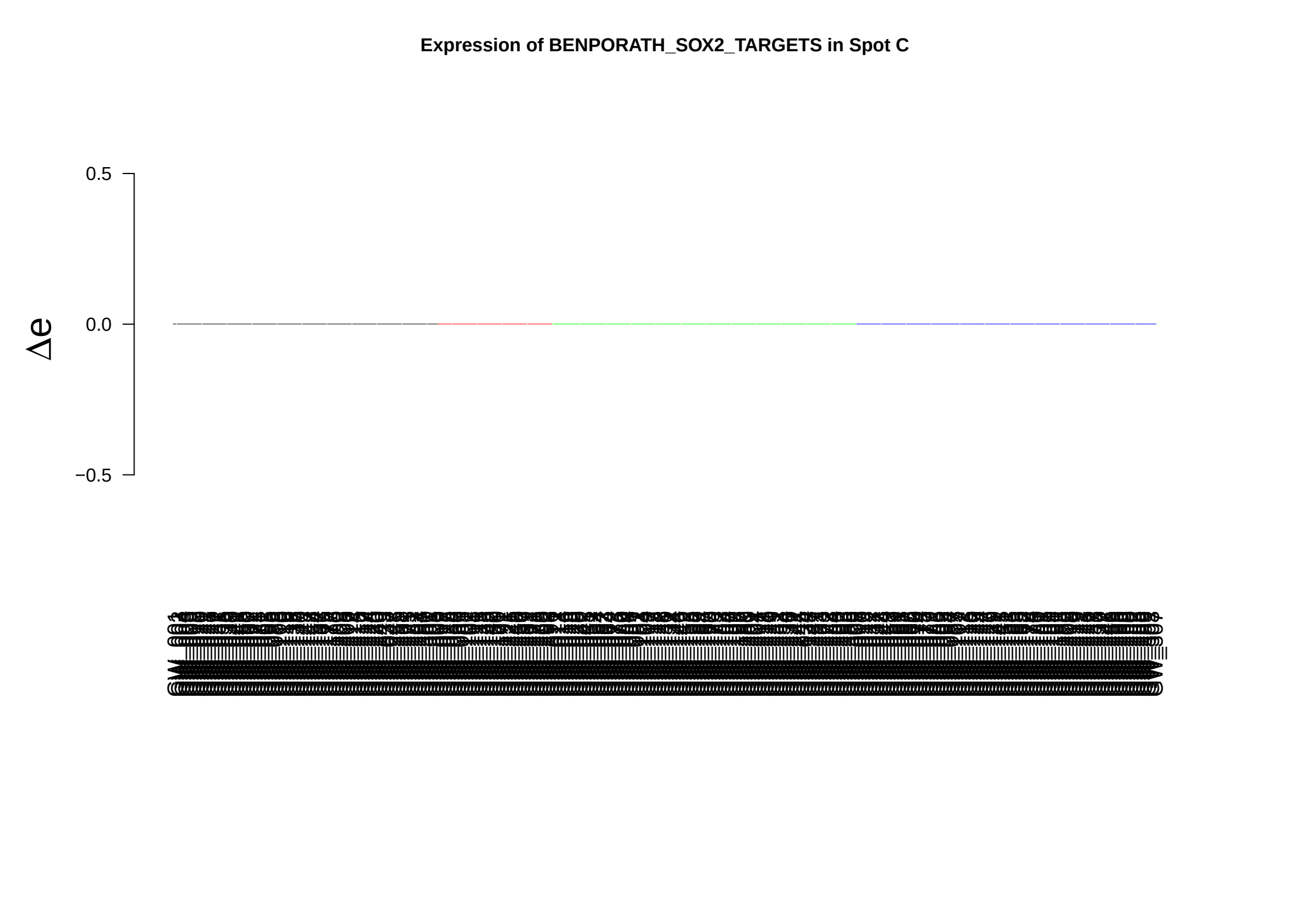

BENPORATH_SOX2_TARGETS
GSZ
20
15
10
5
0
−5
−10

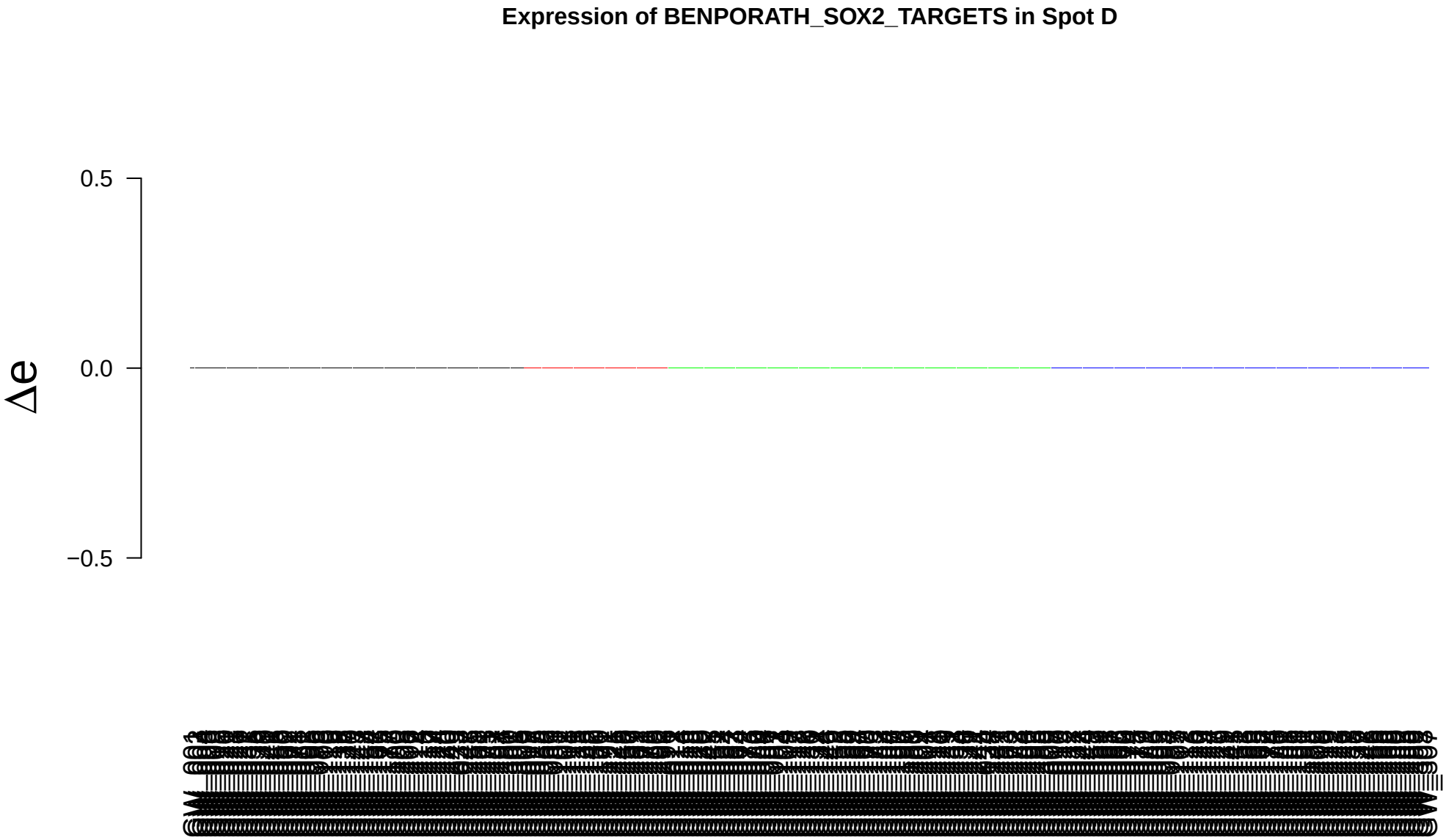

BENPORATH_SOX2_TARGETS
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal

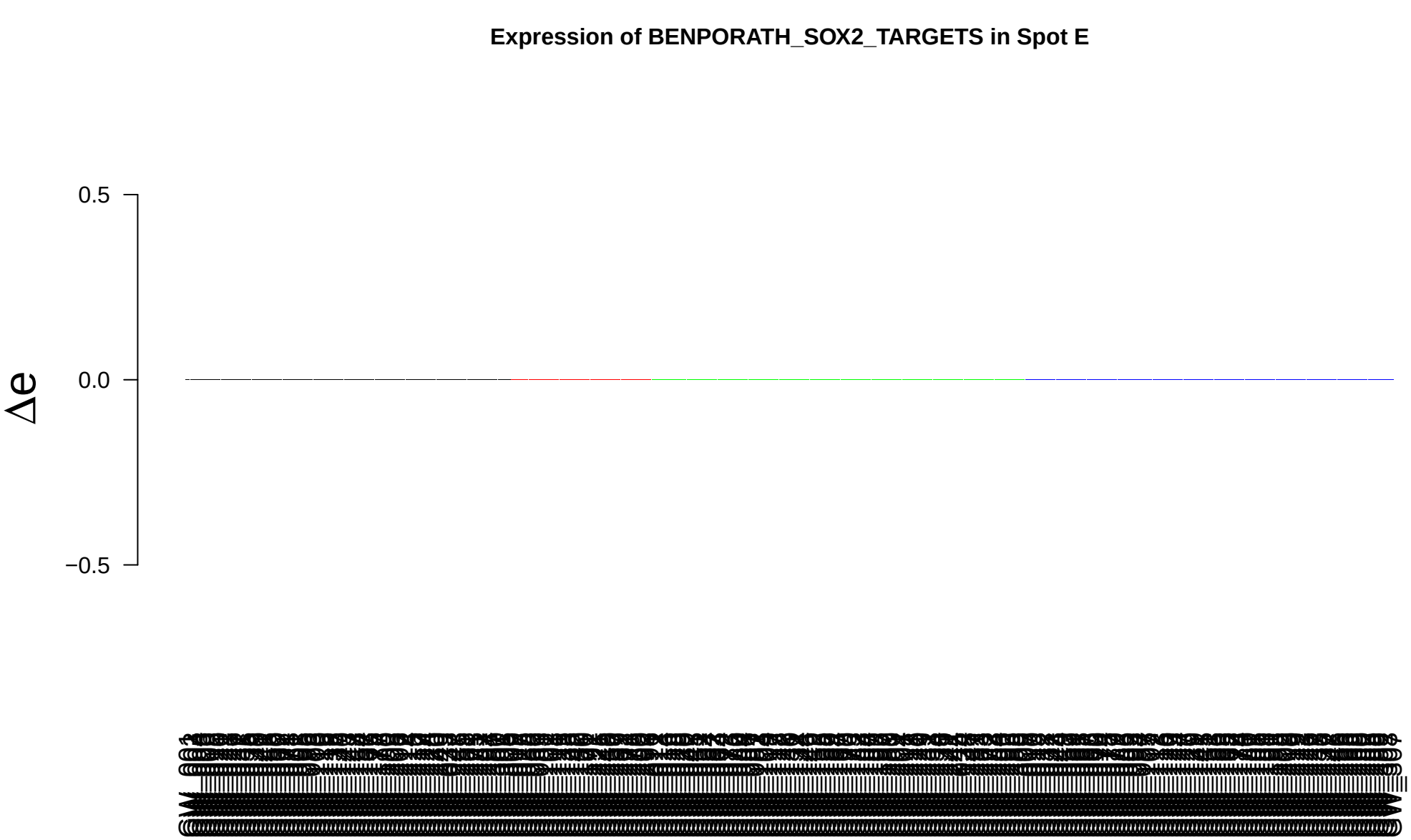

Enrichment in spots: BENPORATH_SOX2_TARGETS
log(p.value)
0.5
0.4
0.3
0.2
0.1
0.0
A
B
C
D
E
F
G
H
I
J
K
L
M
N
O
P
Q
R

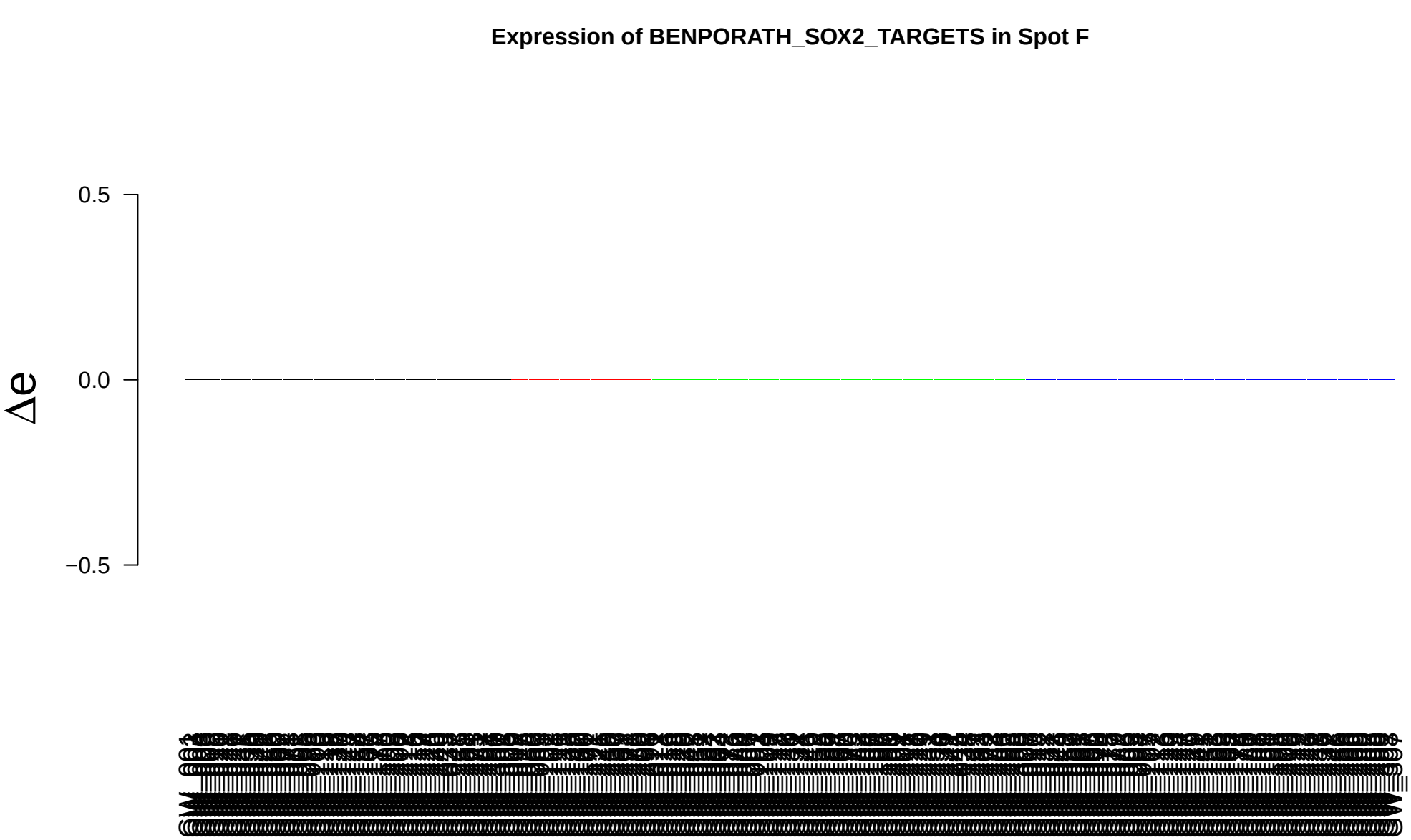

Expression of BENPORATH_SOX2_TARGETS in Spot A
0.5
0.0
−0.5
Δe

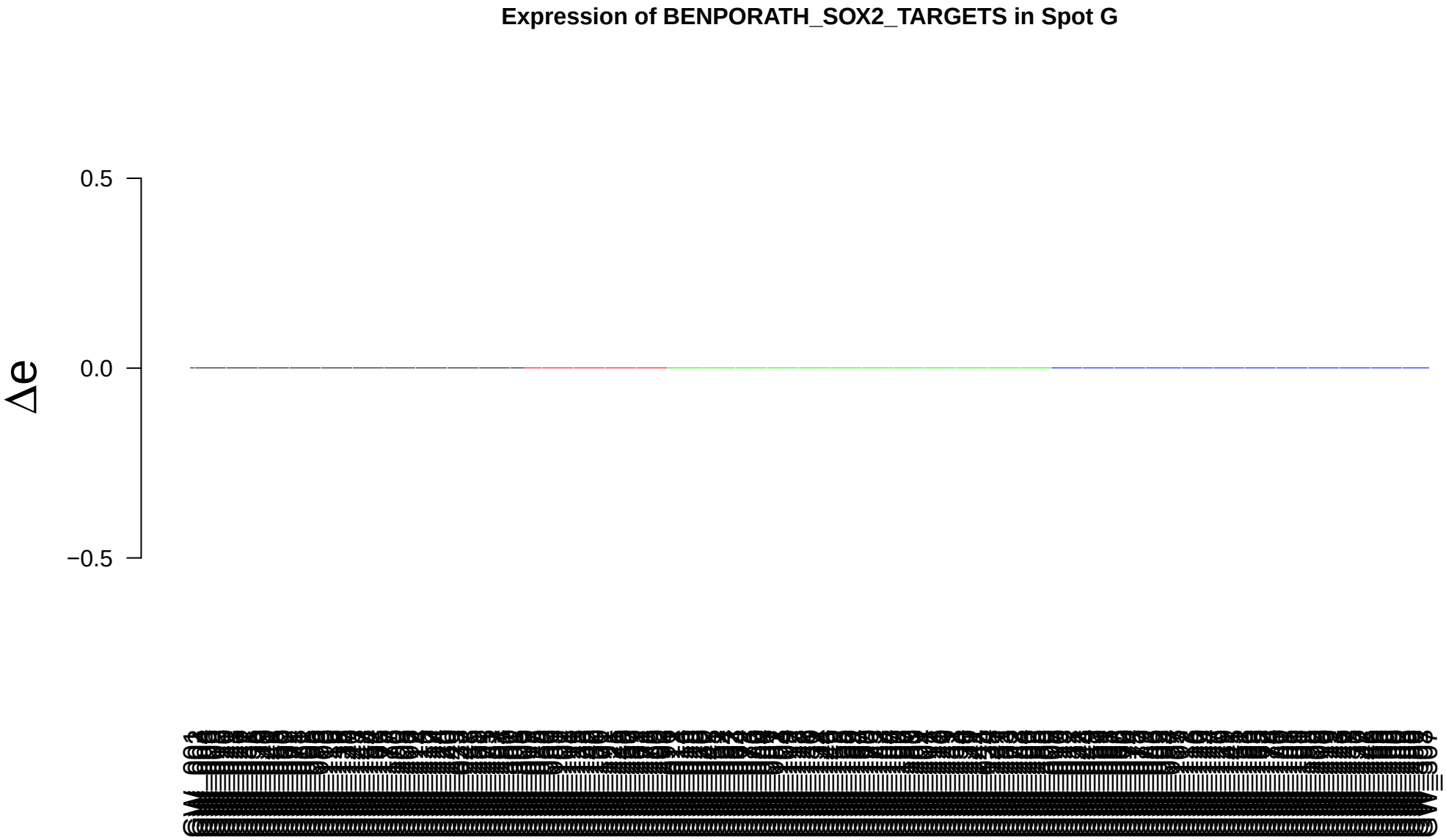

Expression of BENPORATH_SOX2_TARGETS in Spot B
0.5
0.0
−0.5
Δe

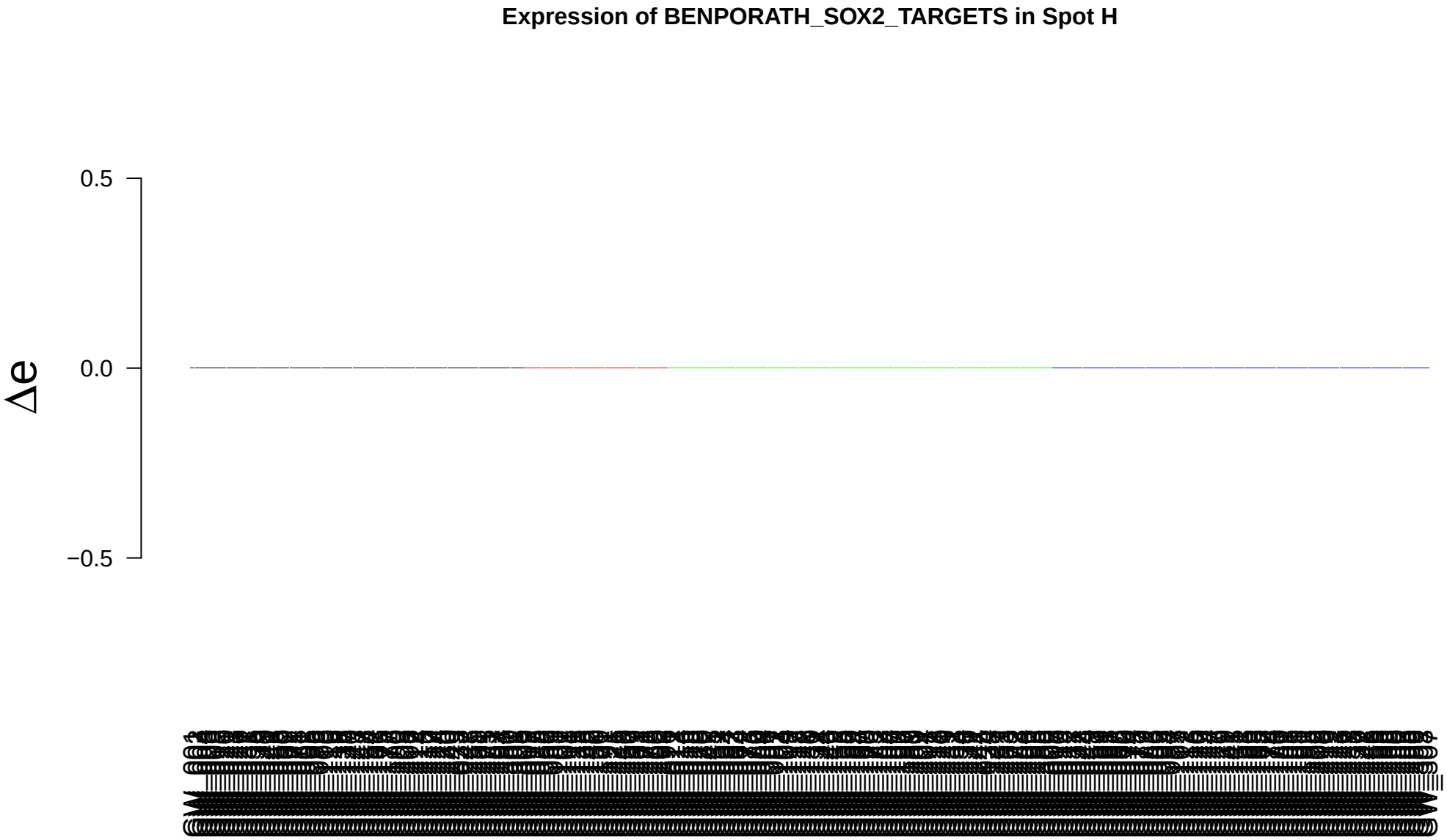

Expression of BENPORATH_SOX2_TARGETS in Spot C
0.5
0.0
−0.5
Δe

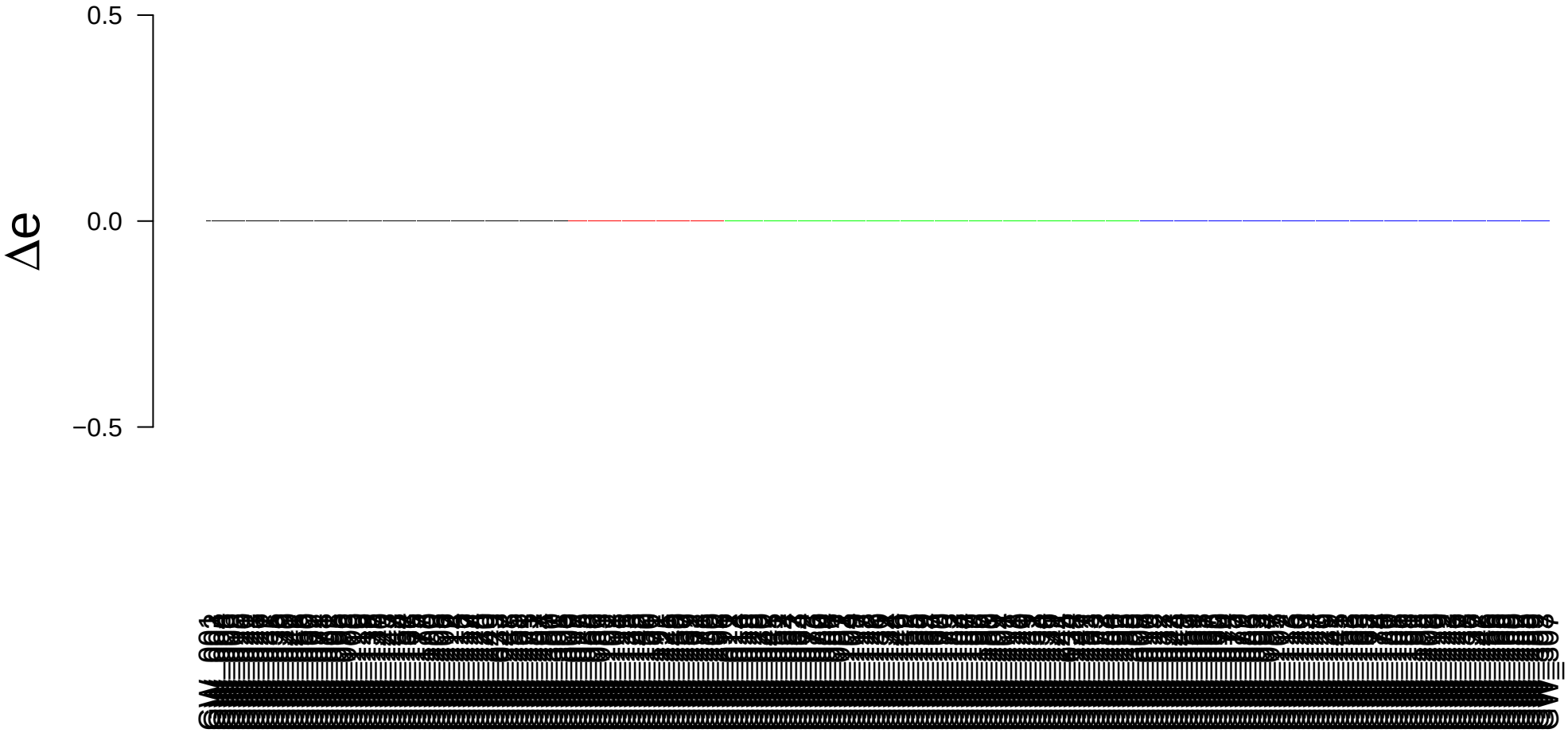

Expression of BENPORATH_SOX2_TARGETS in Spot D
0.5
0.0
−0.5
Δe

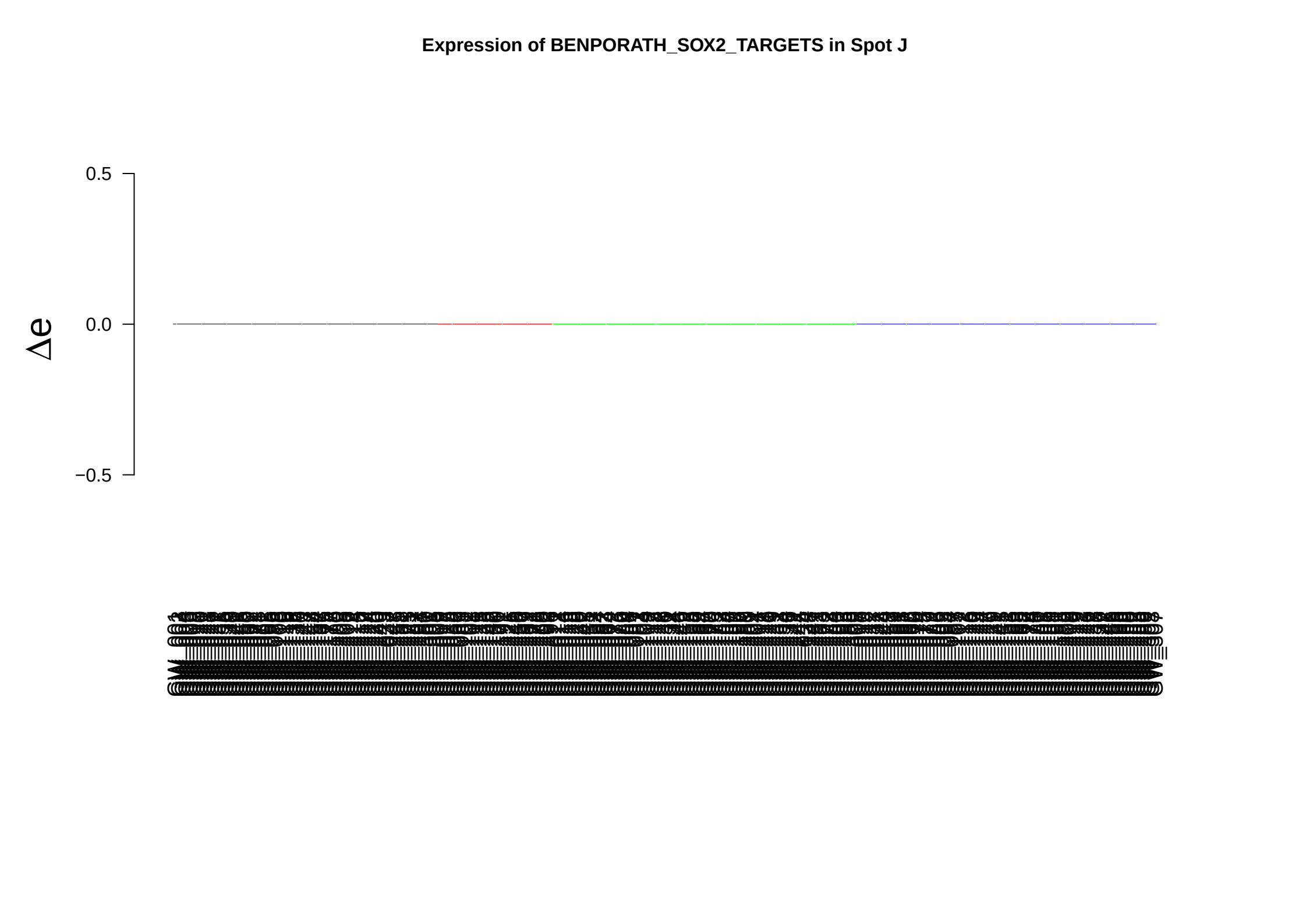

Expression of BENPORATH_SOX2_TARGETS in Spot E
0.5
0.0
−0.5
Δe

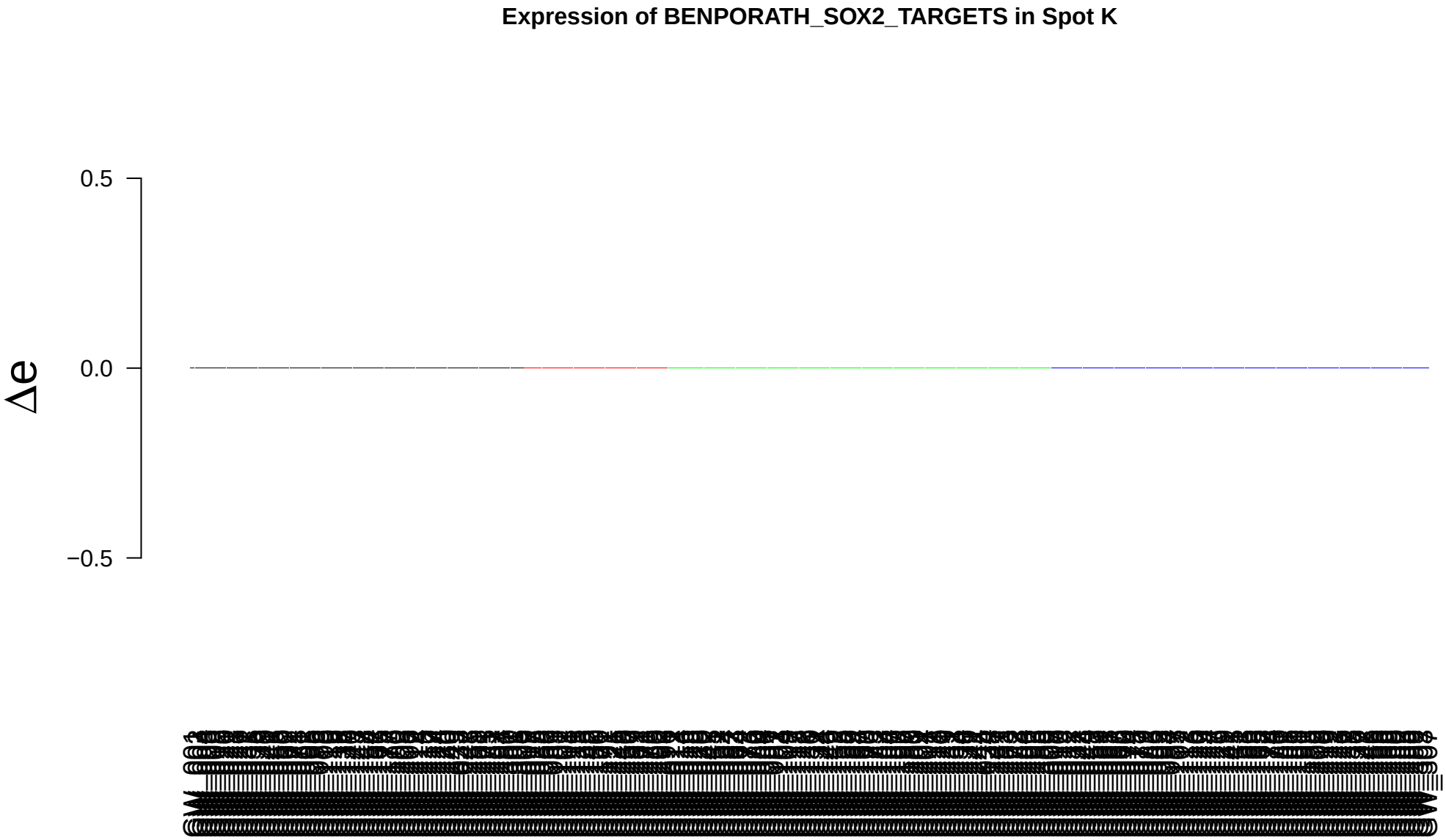

Expression of BENPORATH_SOX2_TARGETS in Spot F
0.5
0.0
−0.5
Δe

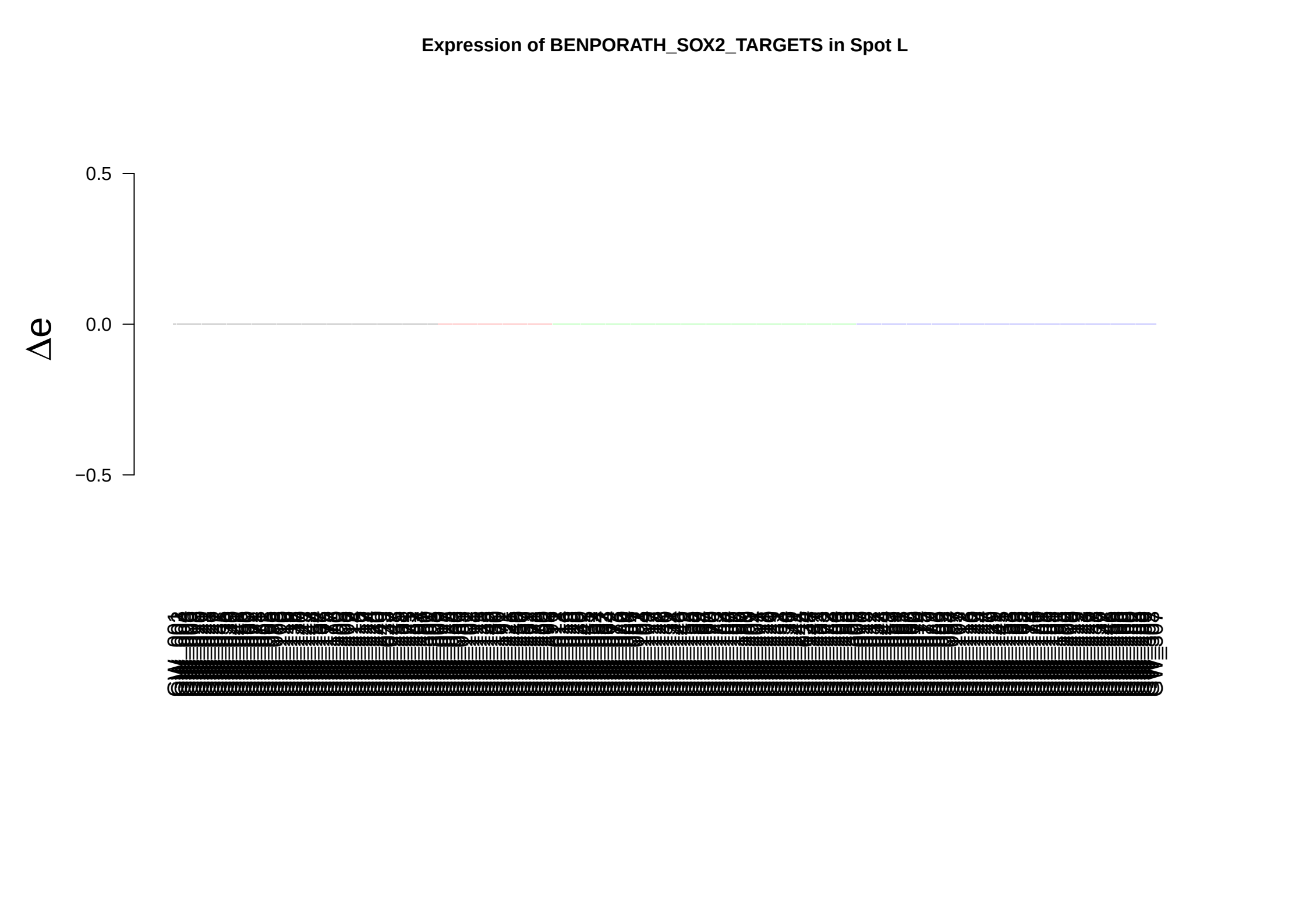

Expression of BENPORATH_SOX2_TARGETS in Spot G
0.5
0.0
−0.5
Δe


Expression of BENPORATH_SOX2_TARGETS in Spot H
0.5
0.0
−0.5
Δe

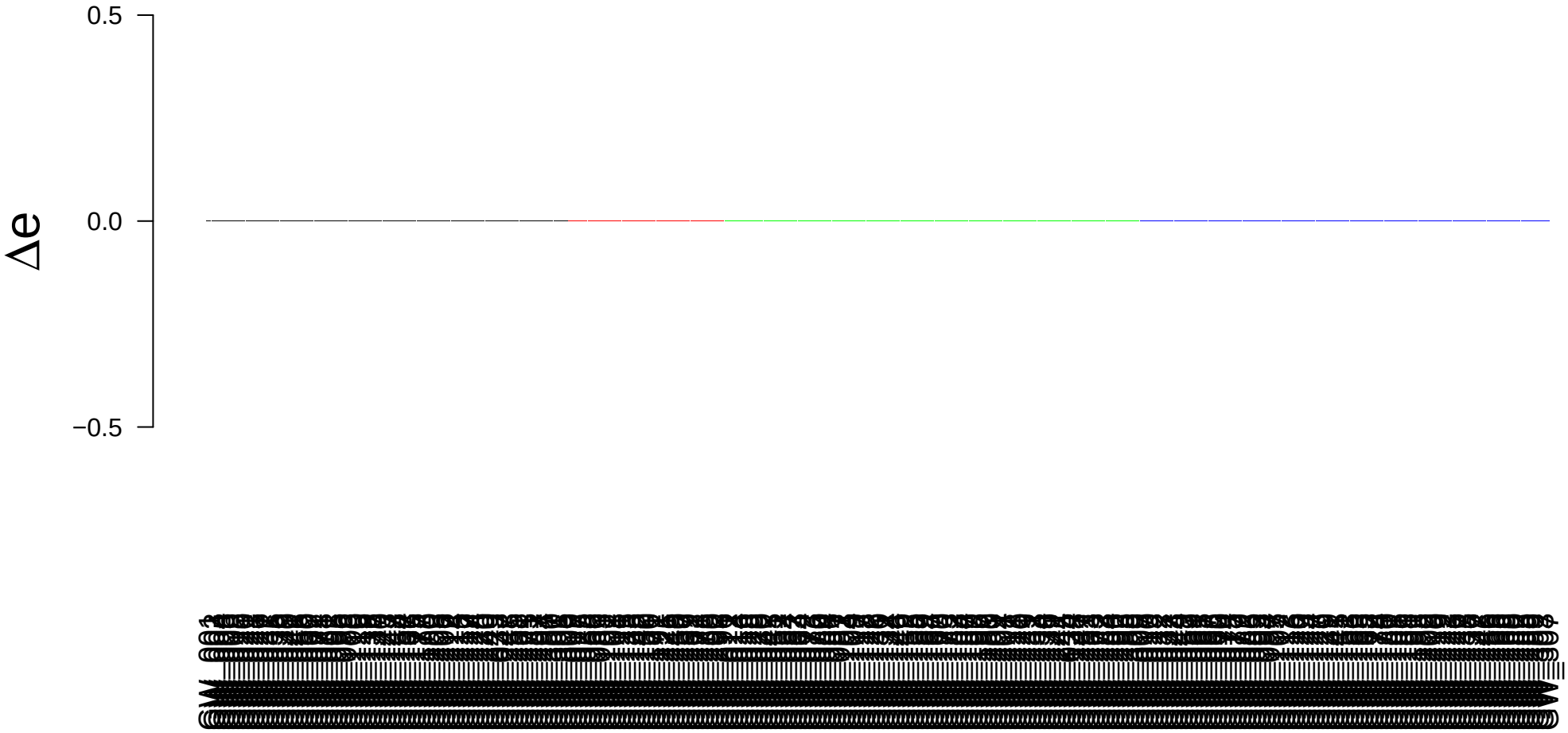

Expression of BENPORATH_SOX2_TARGETS in Spot I
0.5
0.0
−0.5
Δe

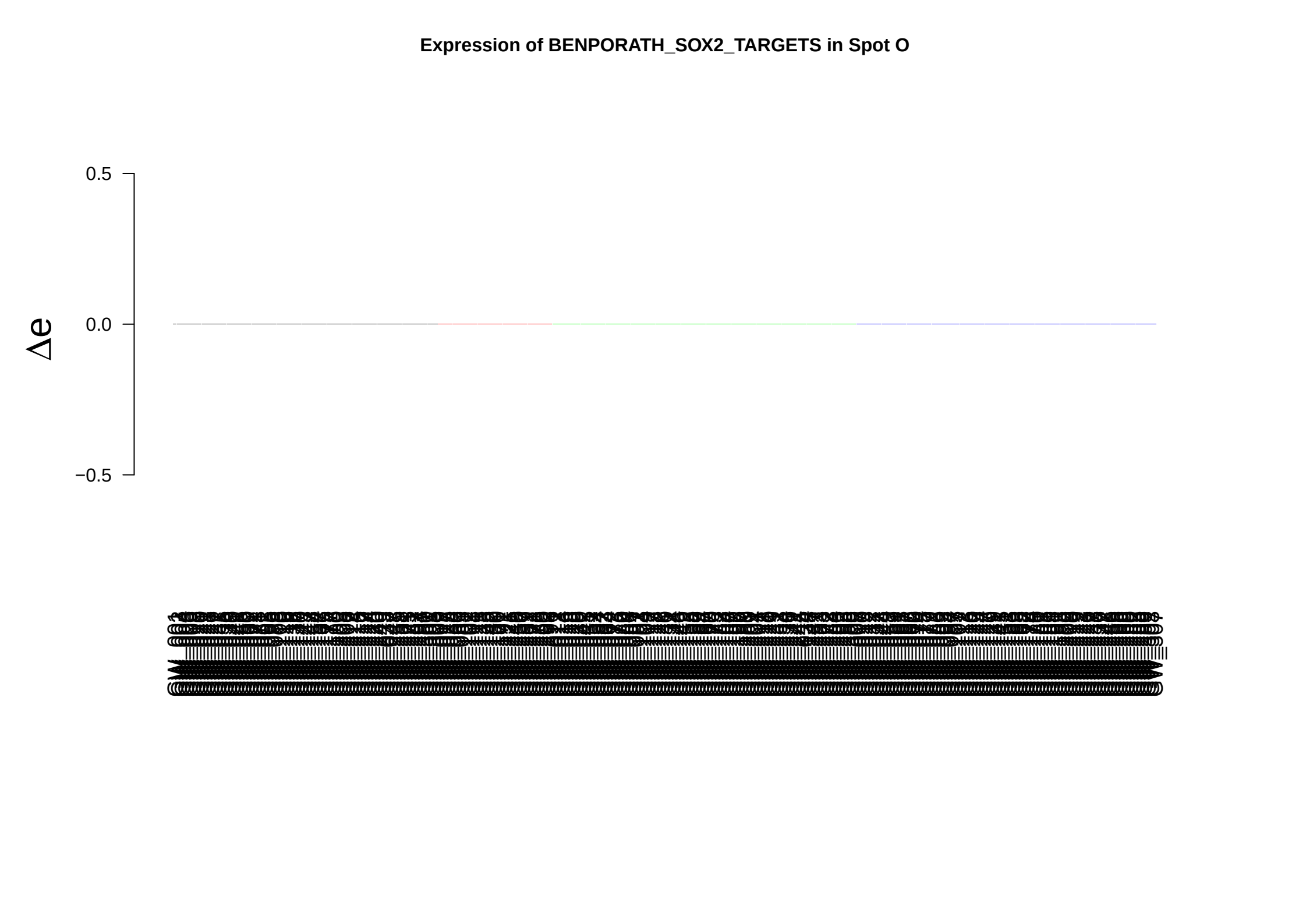

Expression of BENPORATH_SOX2_TARGETS in Spot J
0.5
0.0
−0.5
Δe

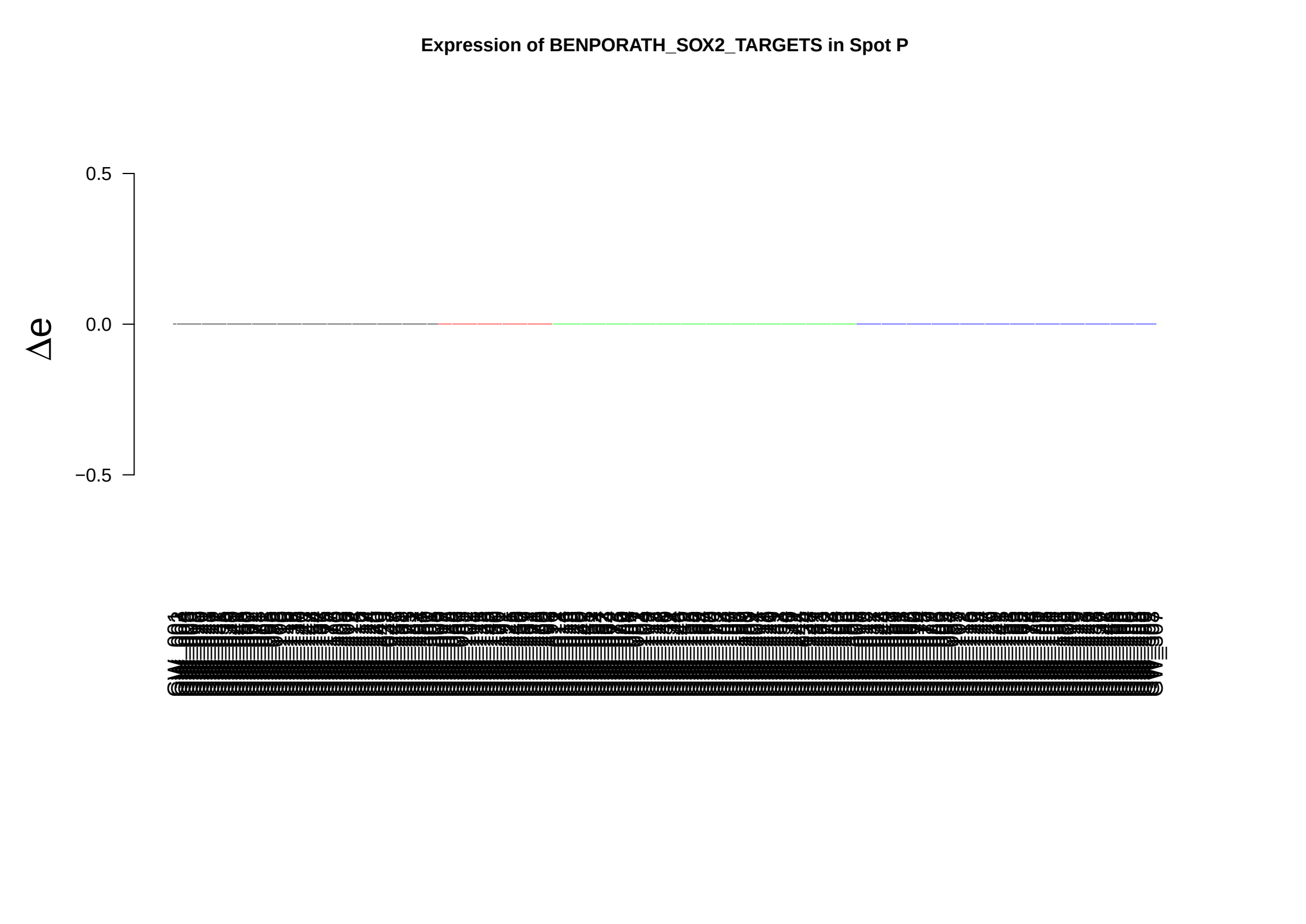

Expression of BENPORATH_SOX2_TARGETS in Spot K
0.5
0.0
−0.5
Δe

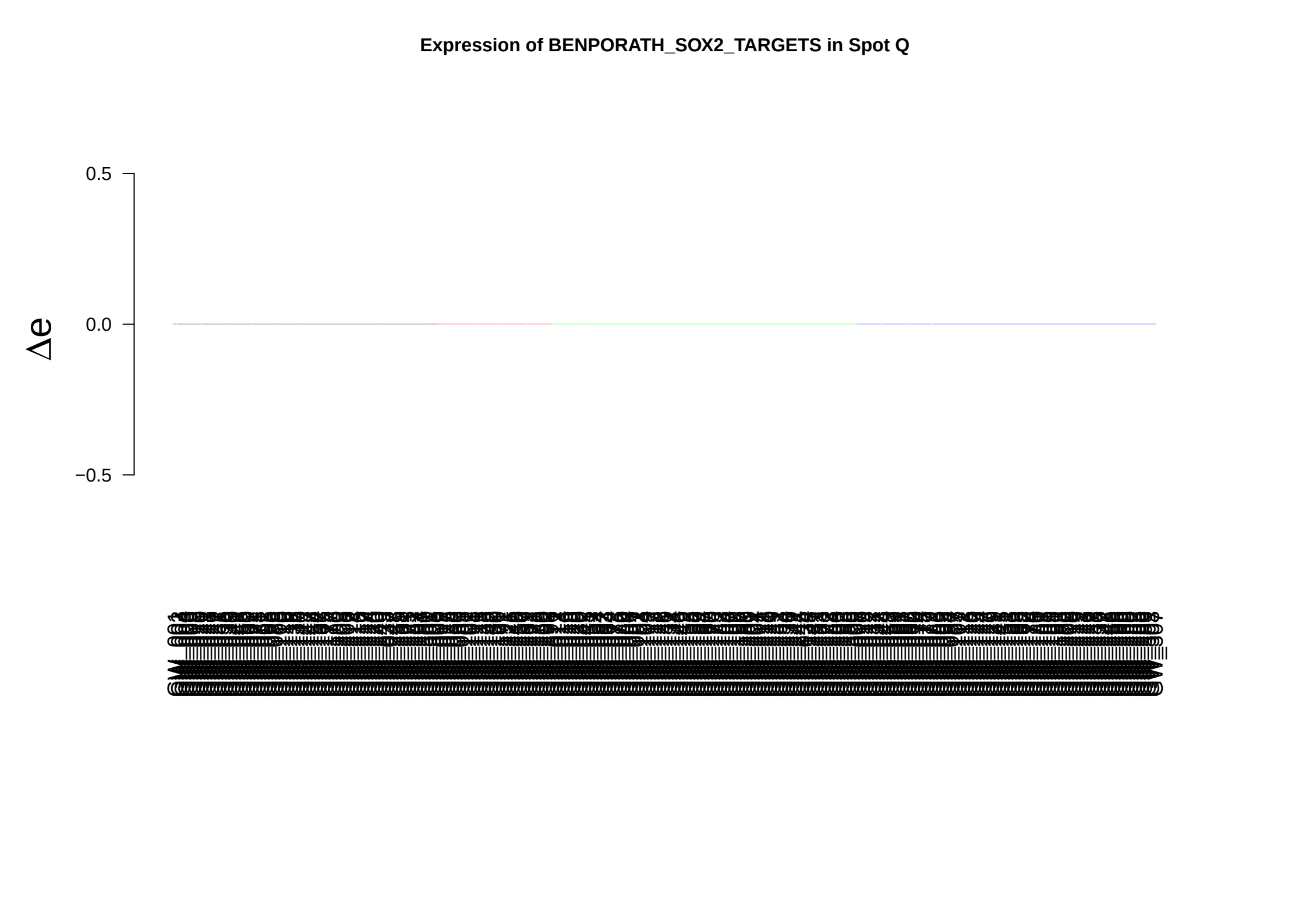

Expression of BENPORATH_SOX2_TARGETS in Spot L
0.5
0.0
−0.5
Δe

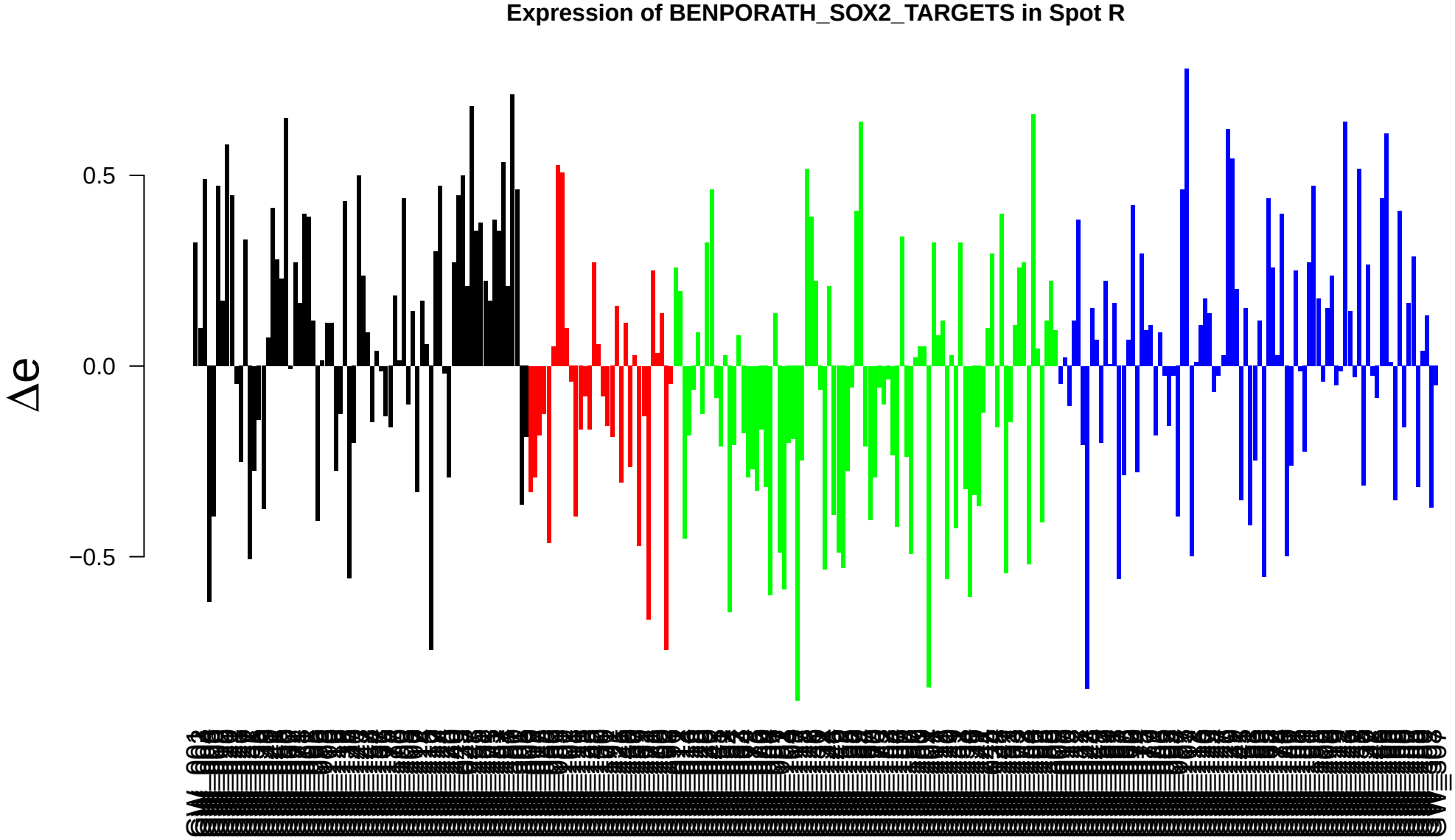
Expression of BENPORATH_SOX2_TARGETS in Spot M
0.5
0.0
−0.5
Δe


Expression of BENPORATH_SOX2_TARGETS in Spot N
0.5
0.0
−0.5
Δe


Expression of BENPORATH_SOX2_TARGETS in Spot O
0.5
0.0
−0.5
Δe


Expression of BENPORATH_SOX2_TARGETS in Spot P
0.5
0.0
−0.5
Δe


Expression of BENPORATH_SOX2_TARGETS in Spot Q
0.5
0.0
−0.5
Δe


Expression of BENPORATH_SOX2_TARGETS in Spot R
Δe
0.5
0.0
−0.5