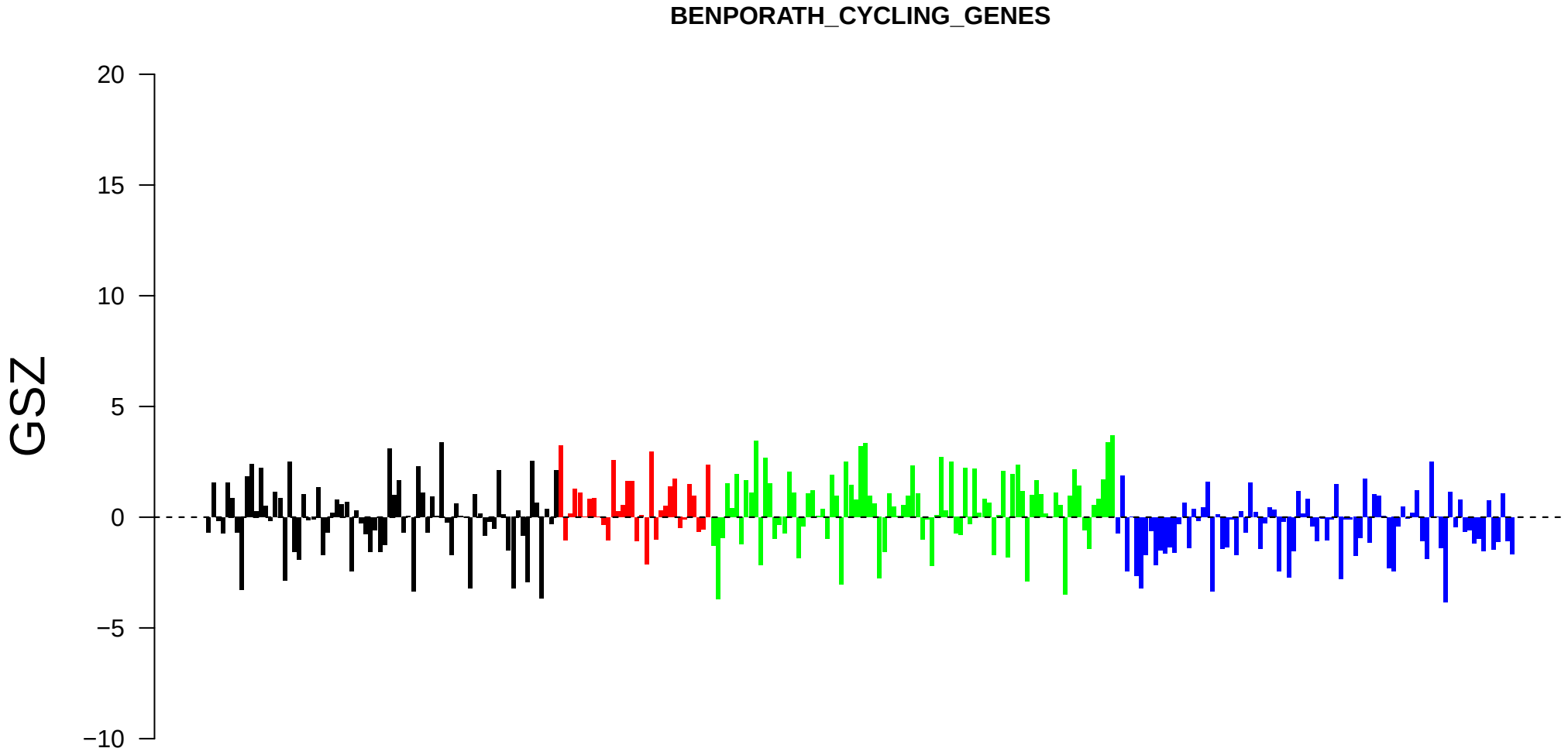
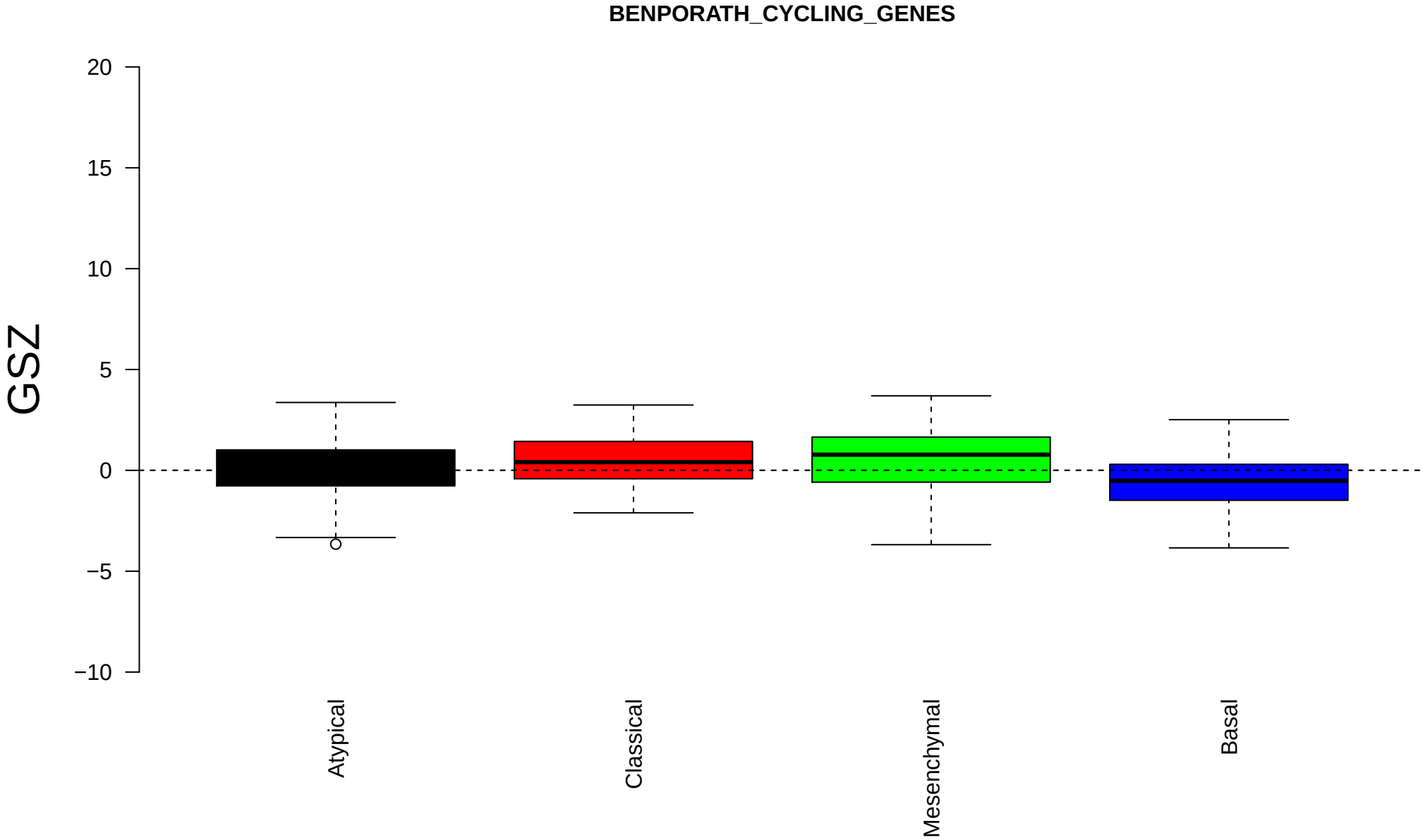
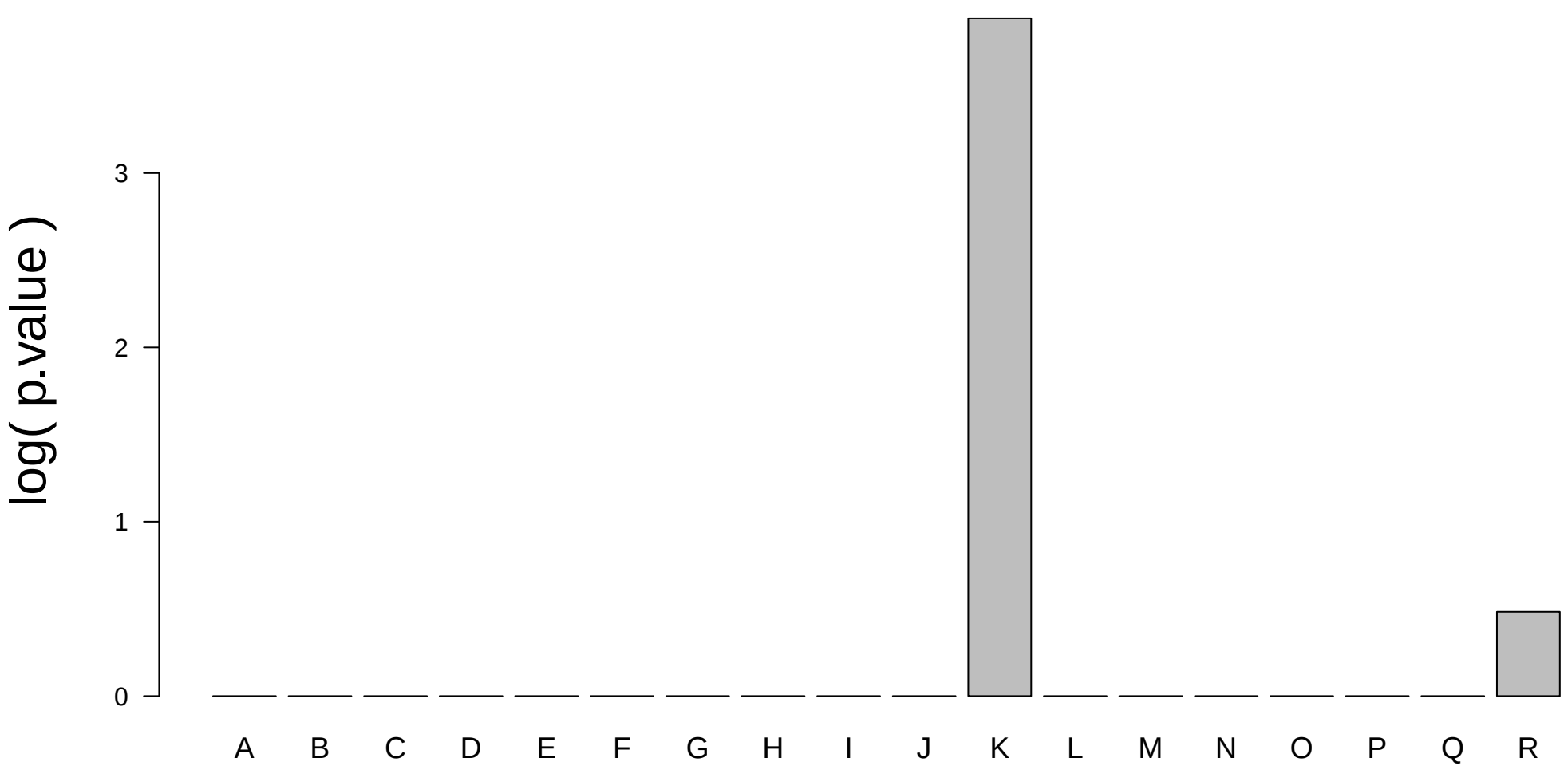
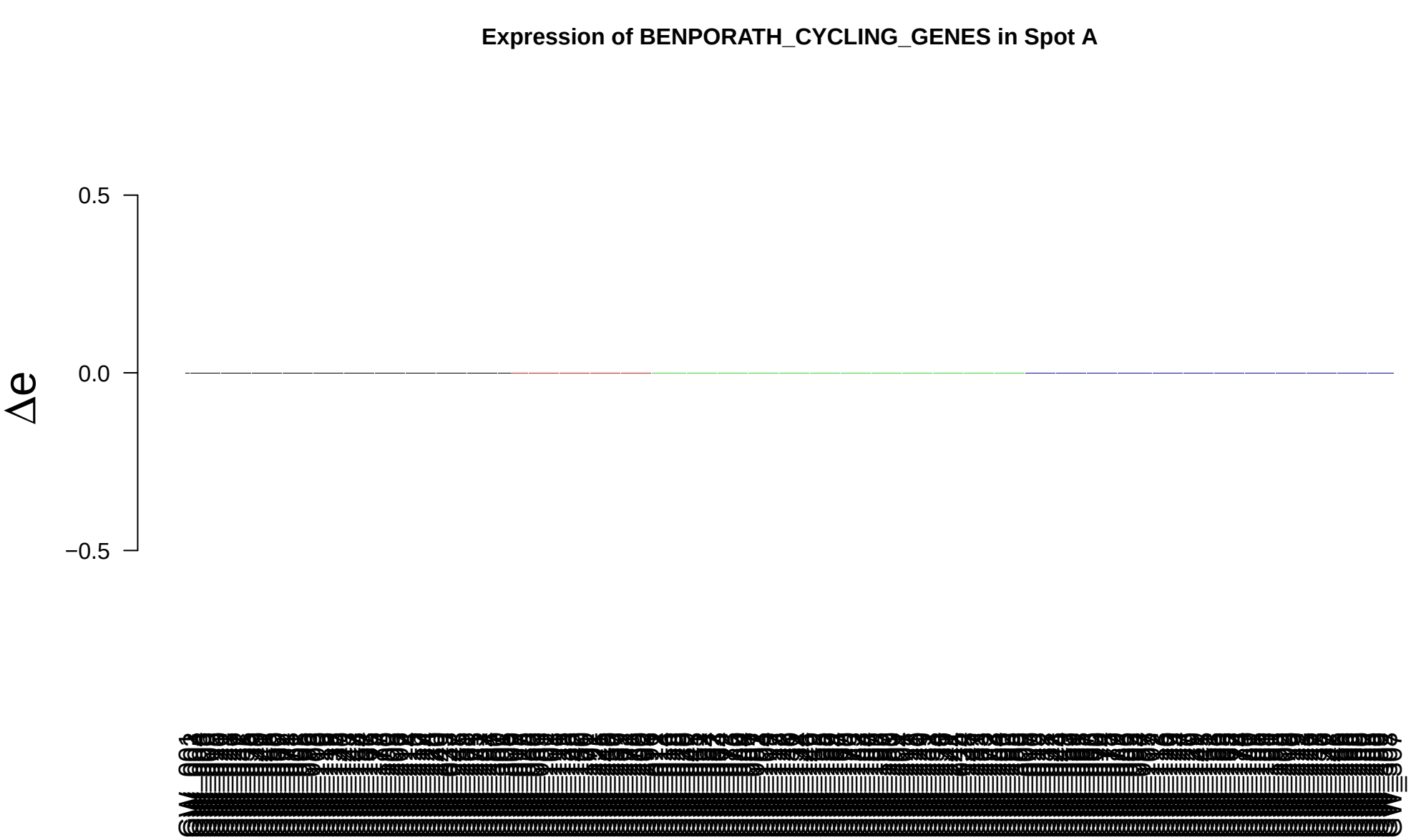
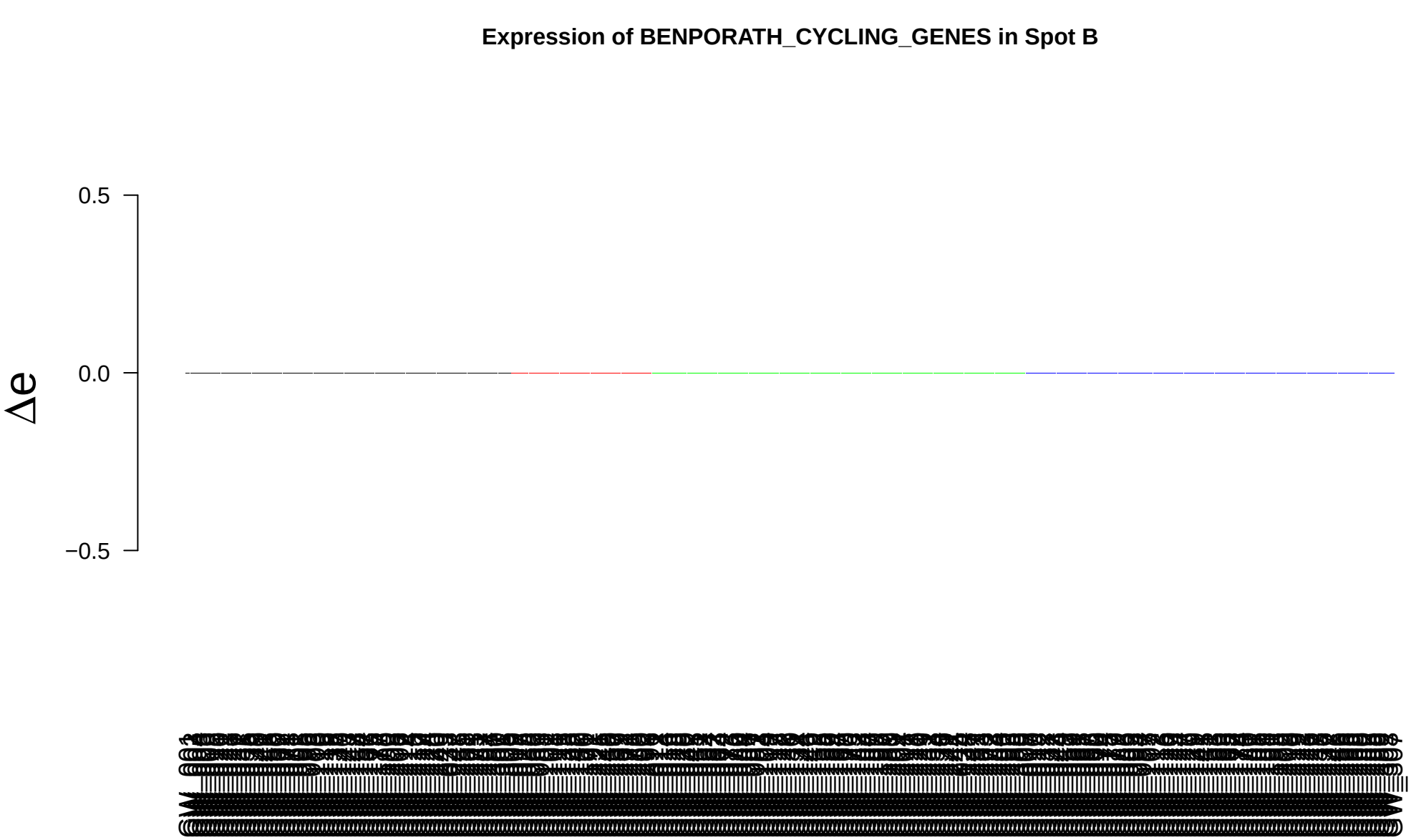
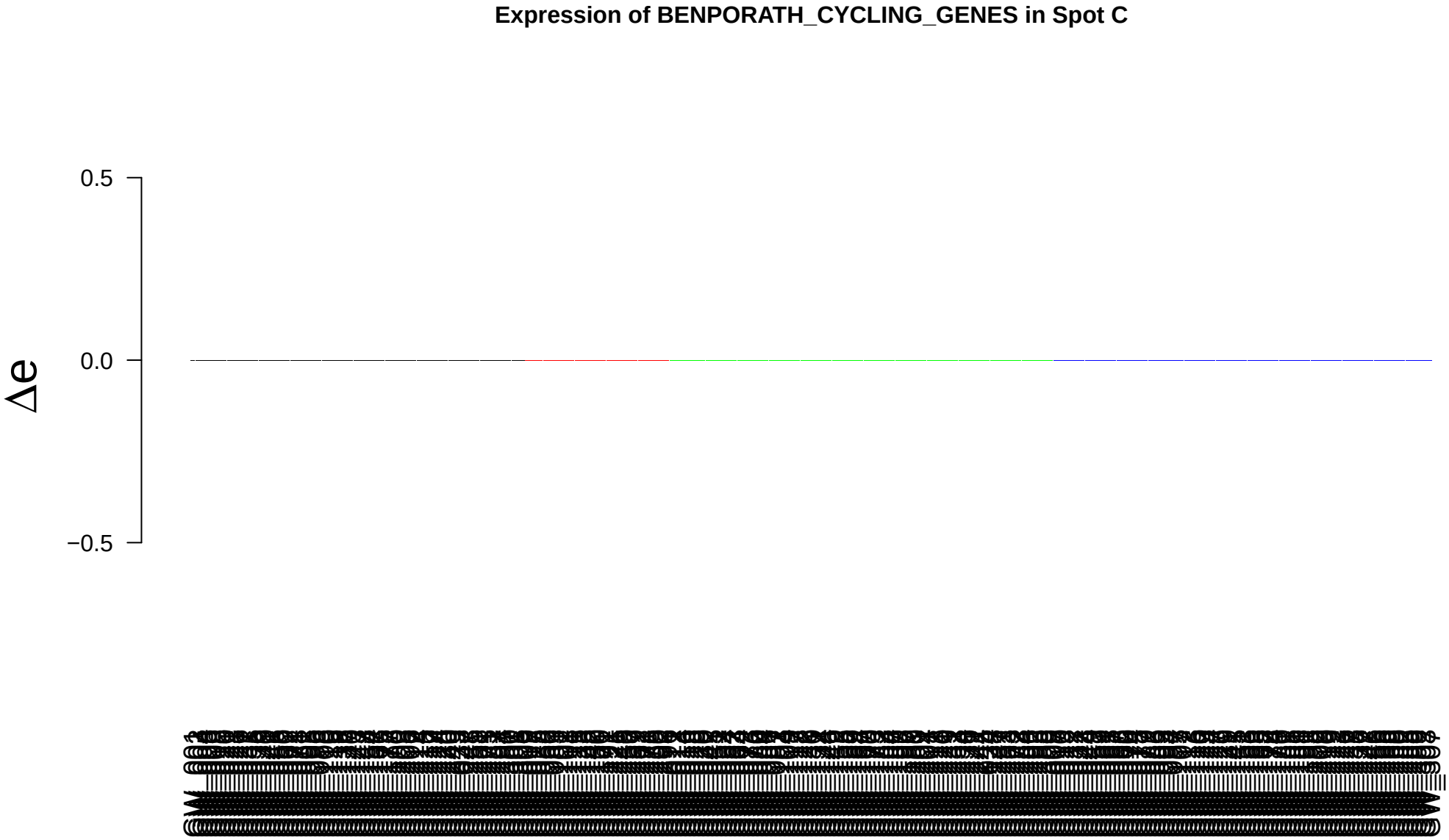

BENPORATH_CYCLING_GENES
GSZ
20
15
10
5
0
−5
−10

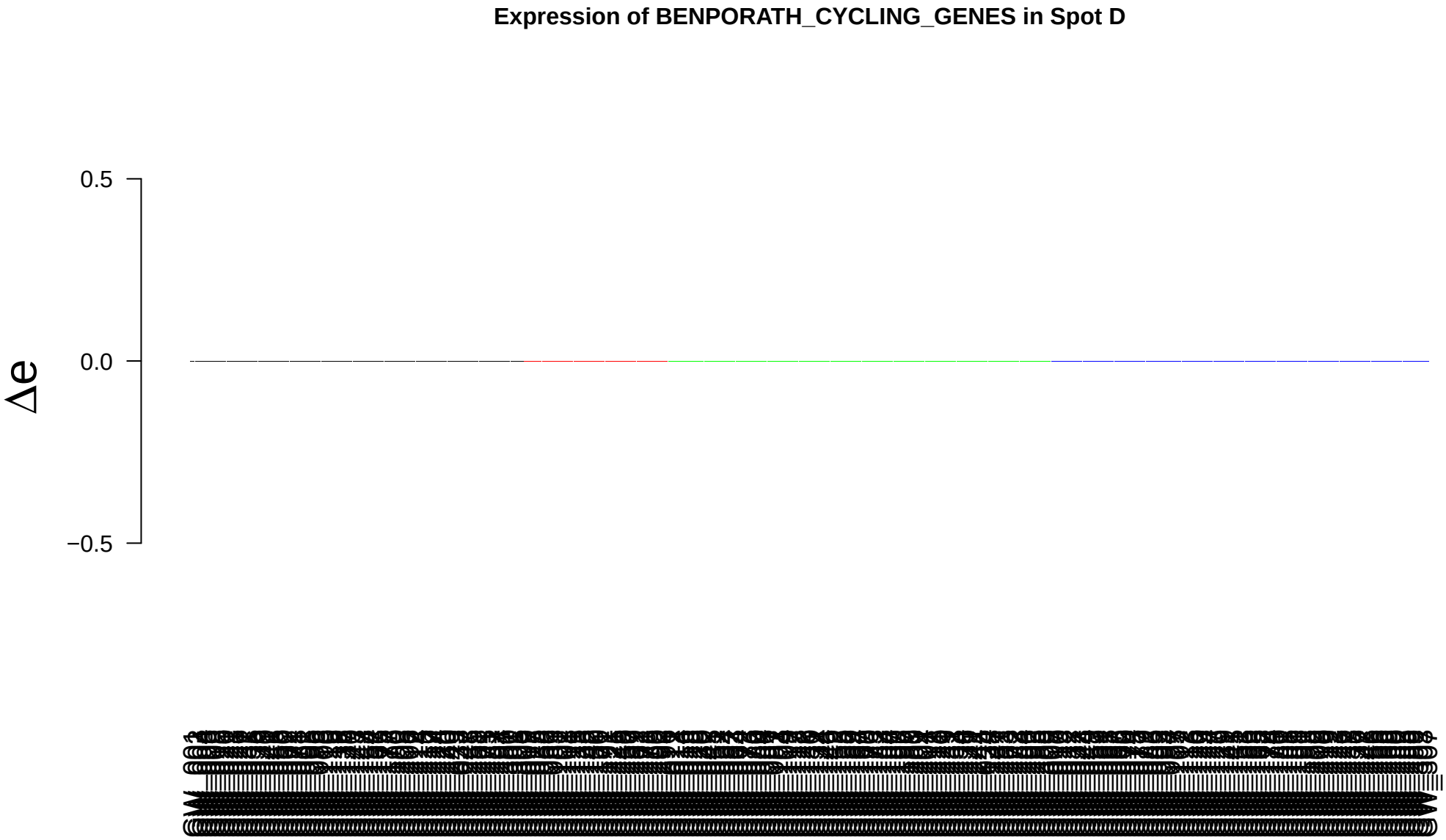

BENPORATH_CYCLING_GENES
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal

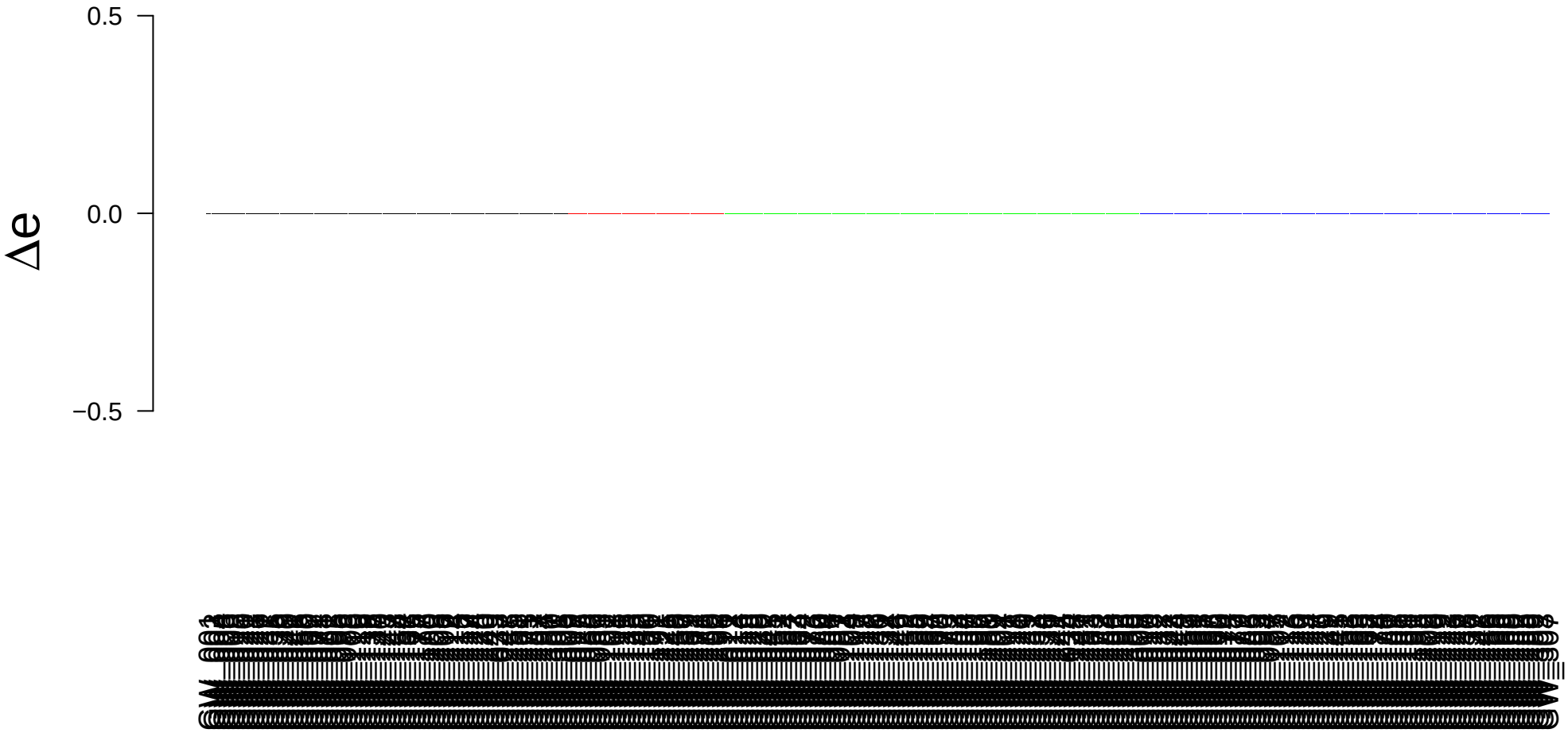

Enrichment in spots: BENPORATH_CYCLING_GENES
log(p.value)
3
2
1
0
A
B
C
D
E
F
G
H
I
J
K
L
M
N
O
P
Q
R

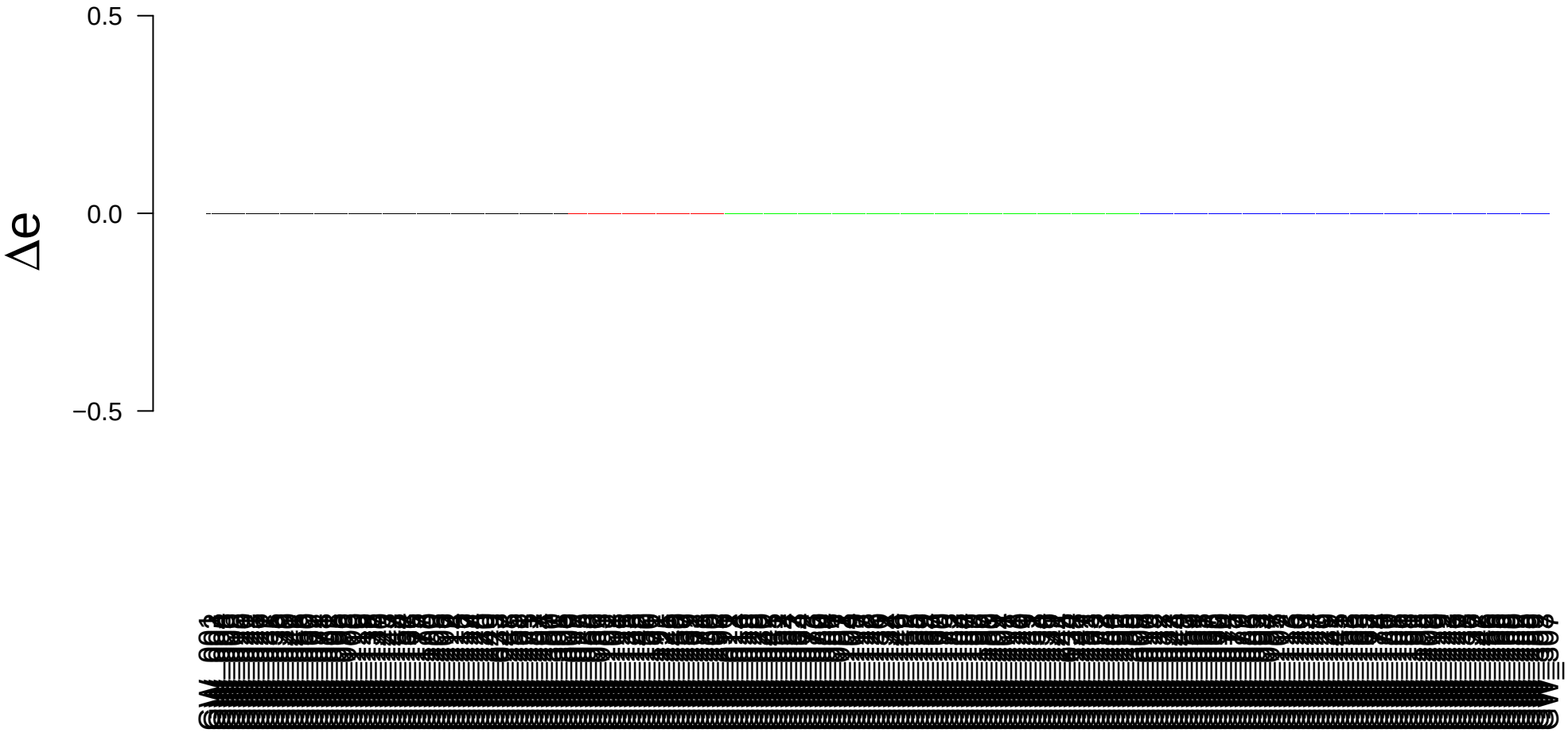

Expression of BENPORATH_CYCLING_GENES in Spot A
0.5
0.0
−0.5
Δe

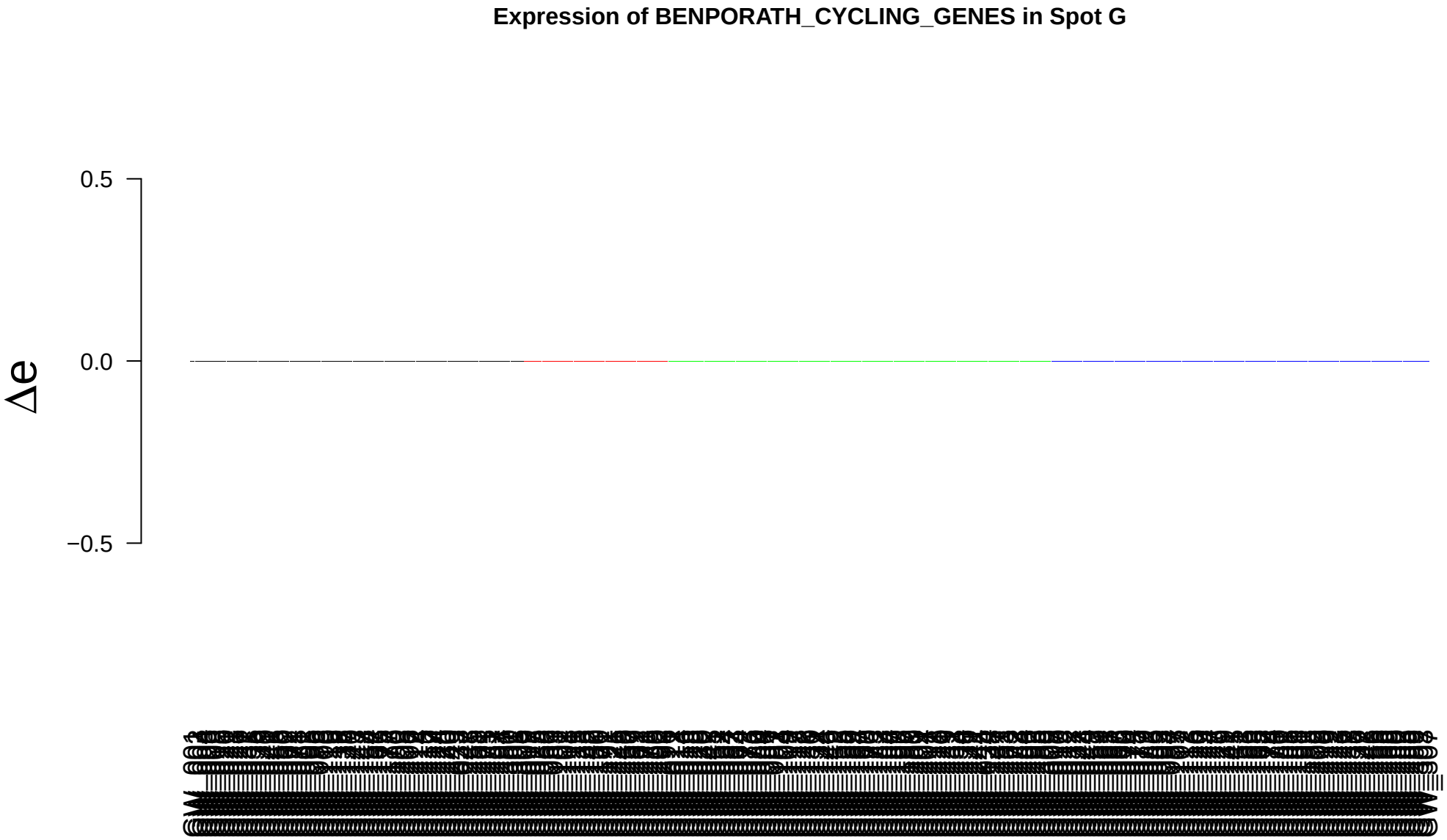

Expression of BENPORATH_CYCLING_GENES in Spot B
0.5
0.0
−0.5
Δe

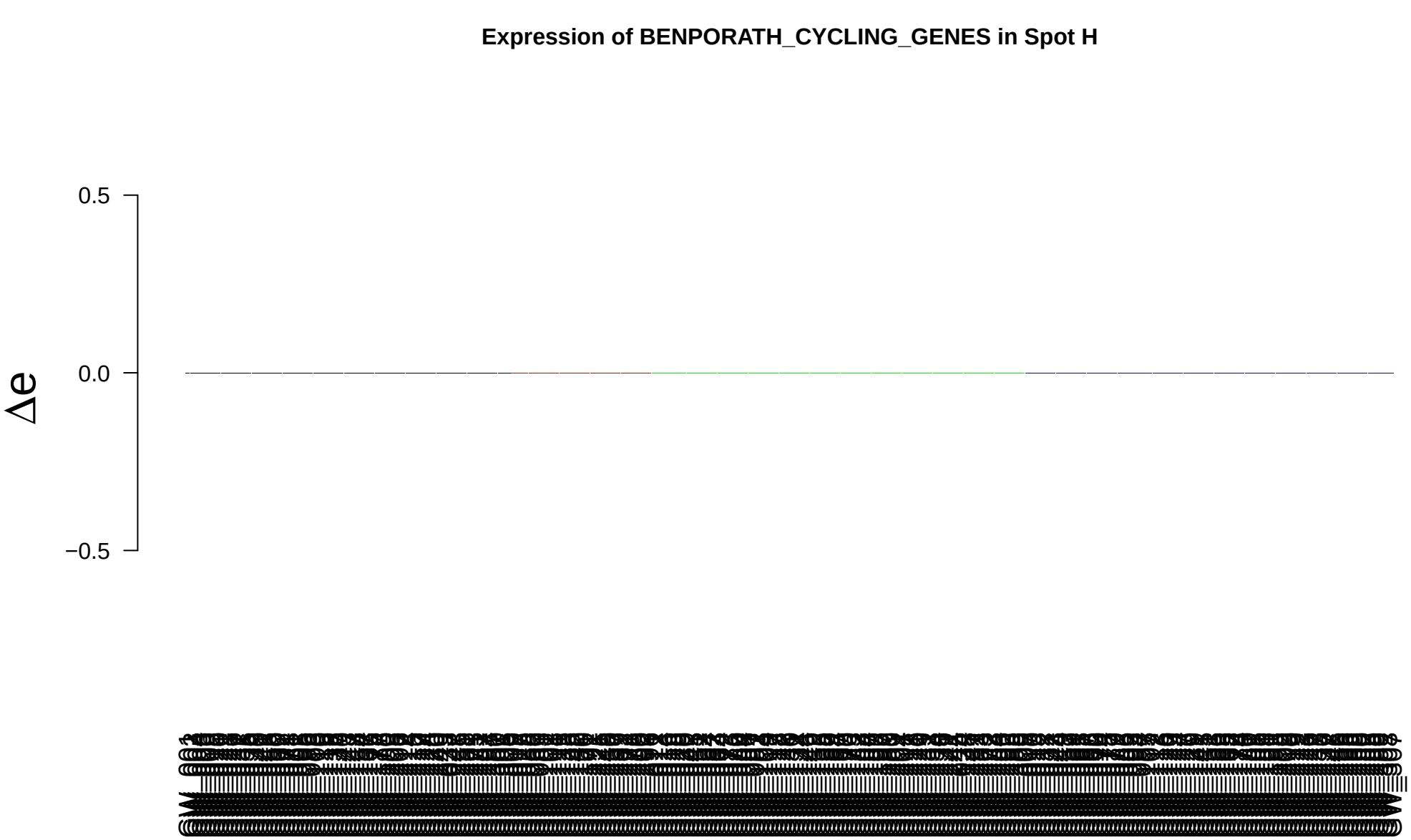

Expression of BENPORATH_CYCLING_GENES in Spot C
0.5
0.0
−0.5
Δe

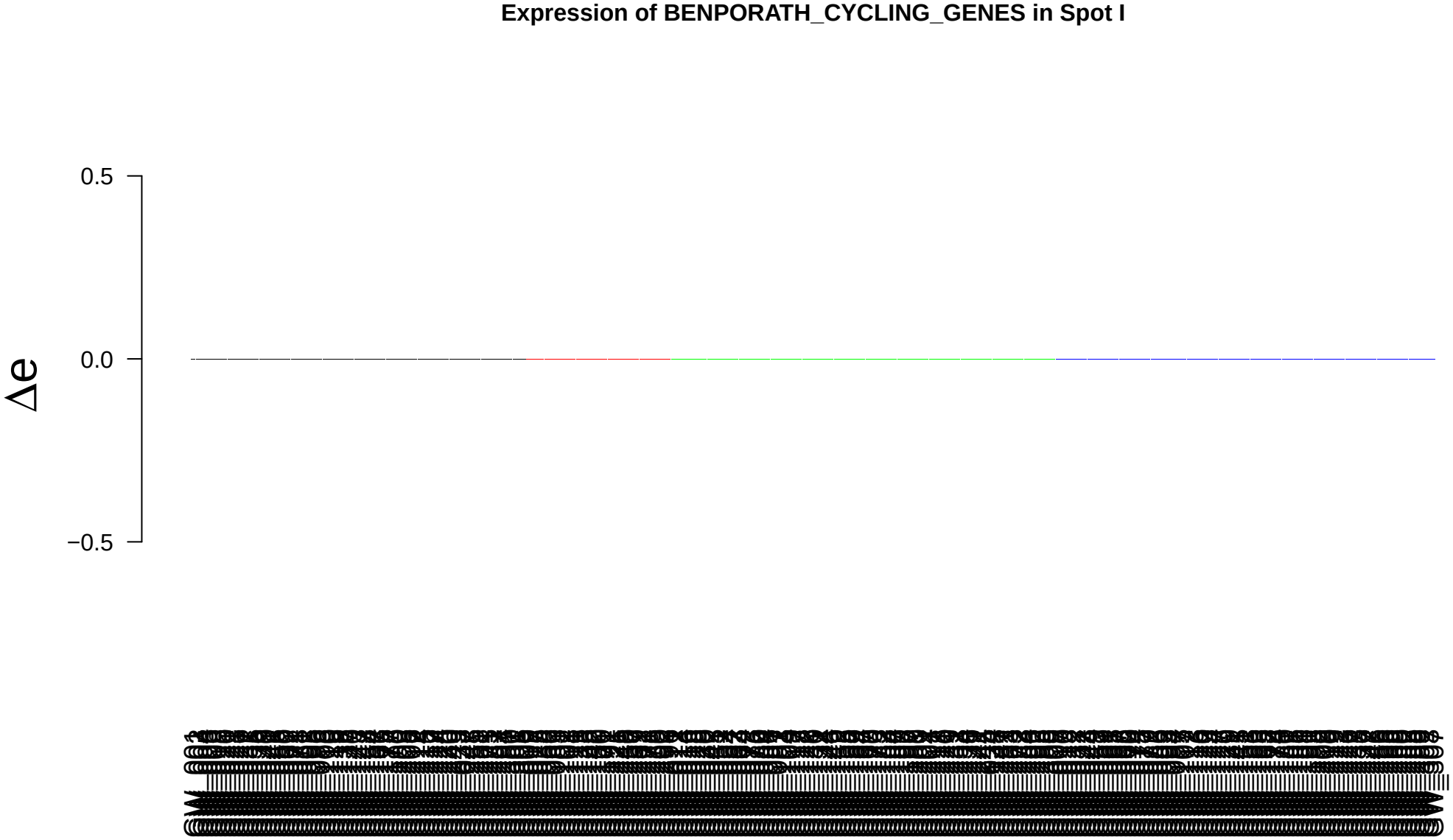

Expression of BENPORATH_CYCLING_GENES in Spot D
0.5
0.0
−0.5
Δe

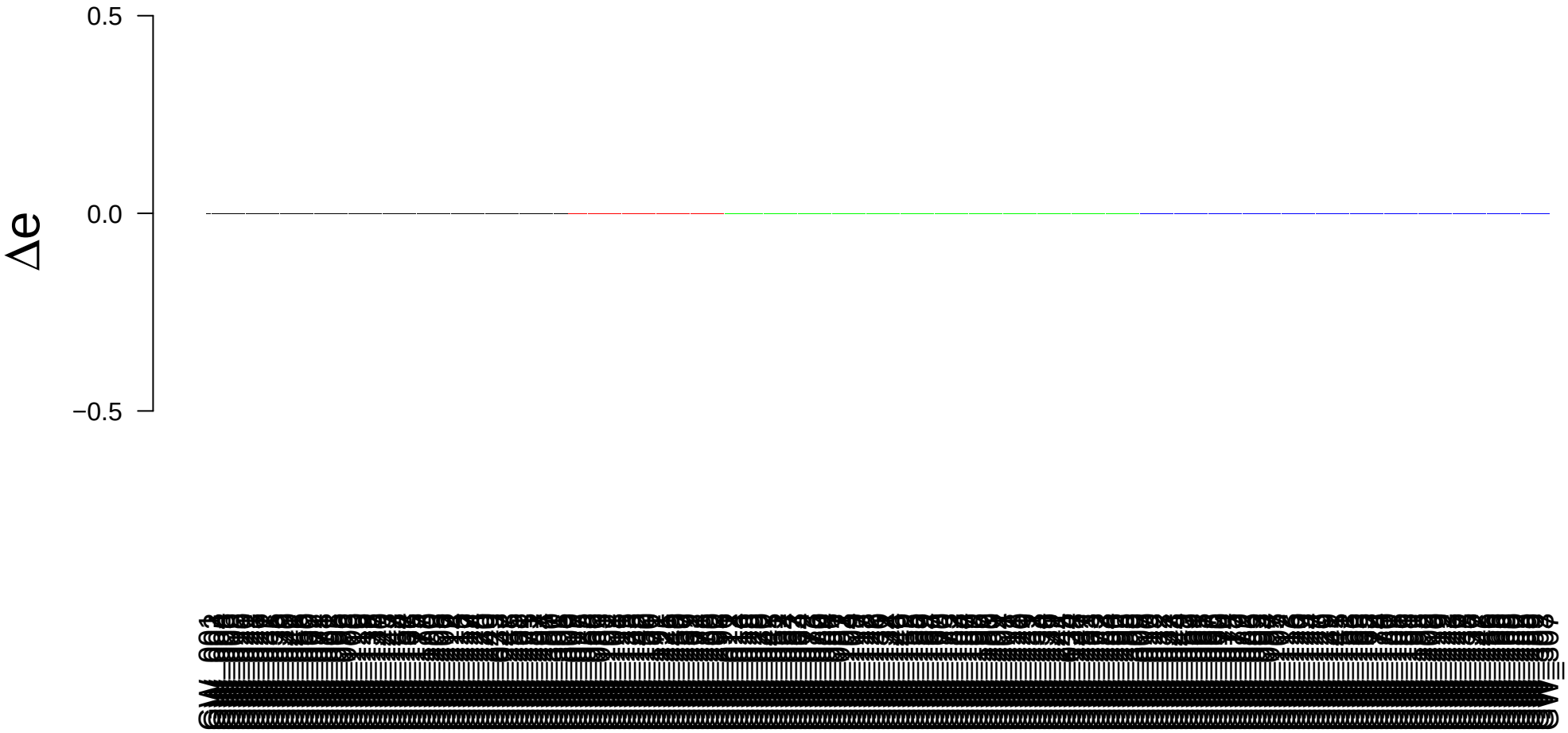

Expression of BENPORATH_CYCLING_GENES in Spot E
0.5
0.0
−0.5
Δe

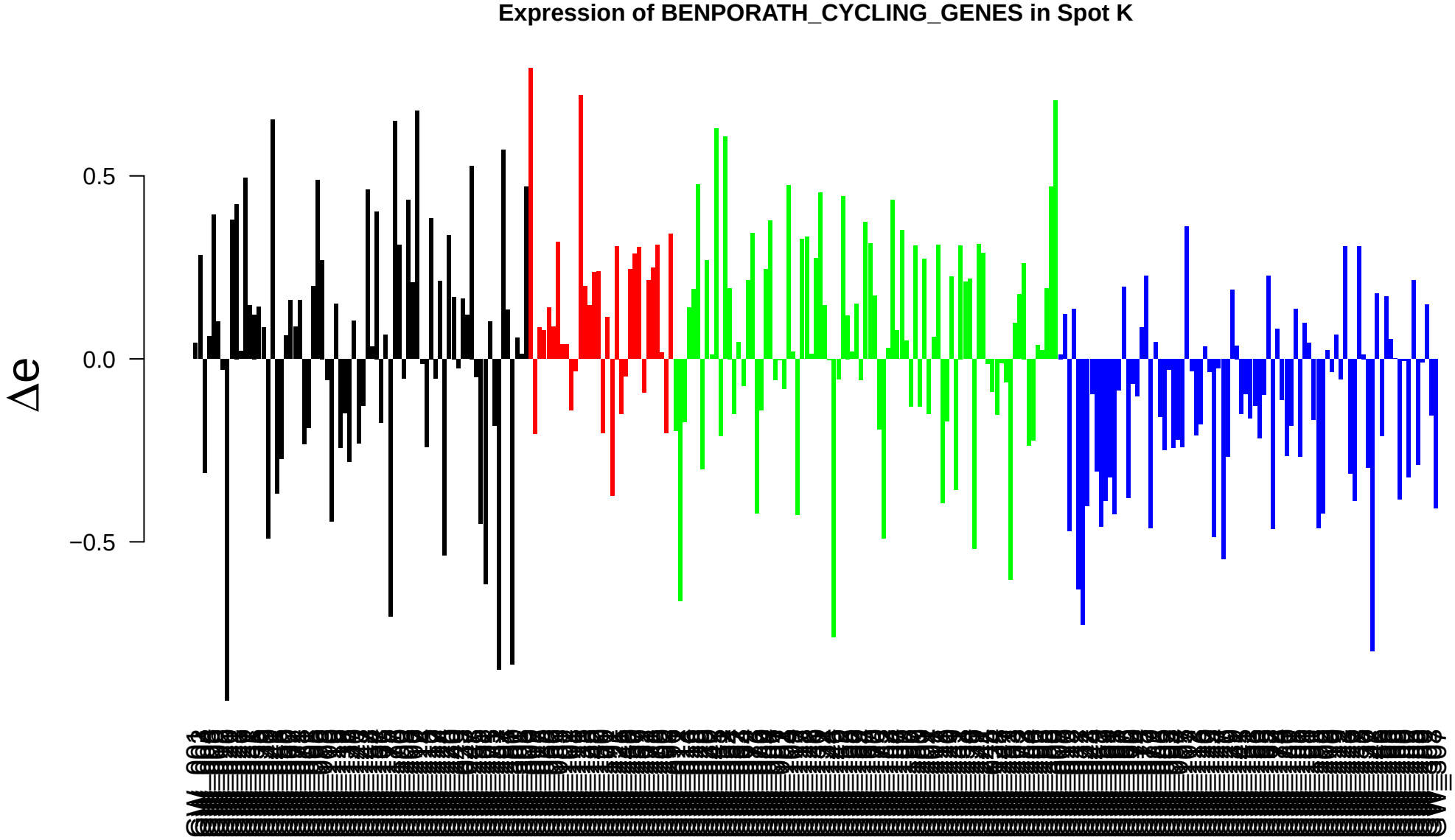

Expression of BENPORATH_CYCLING_GENES in Spot F
0.5
0.0
−0.5
Δe

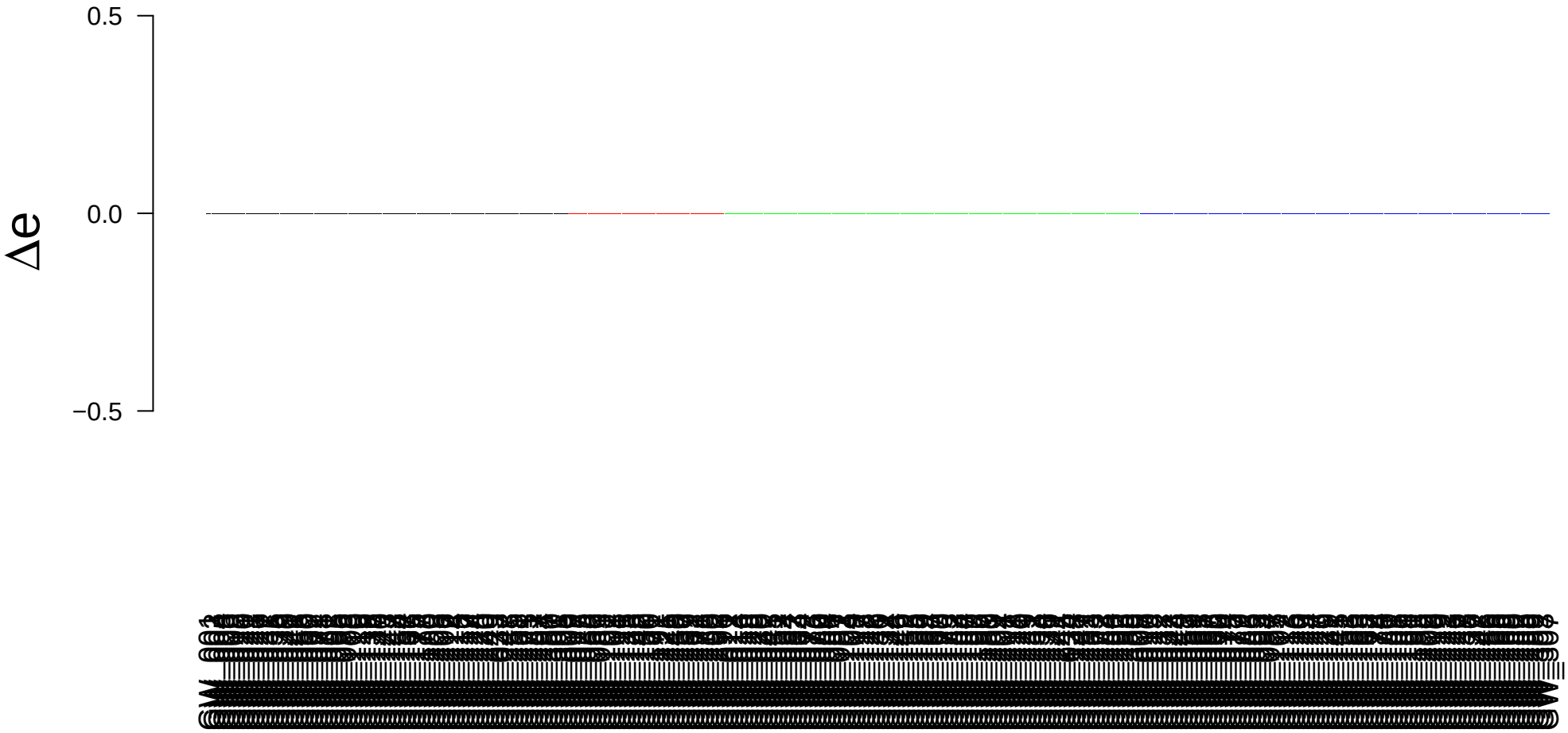

Expression of BENPORATH_CYCLING_GENES in Spot G
0.5
0.0
−0.5
Δe

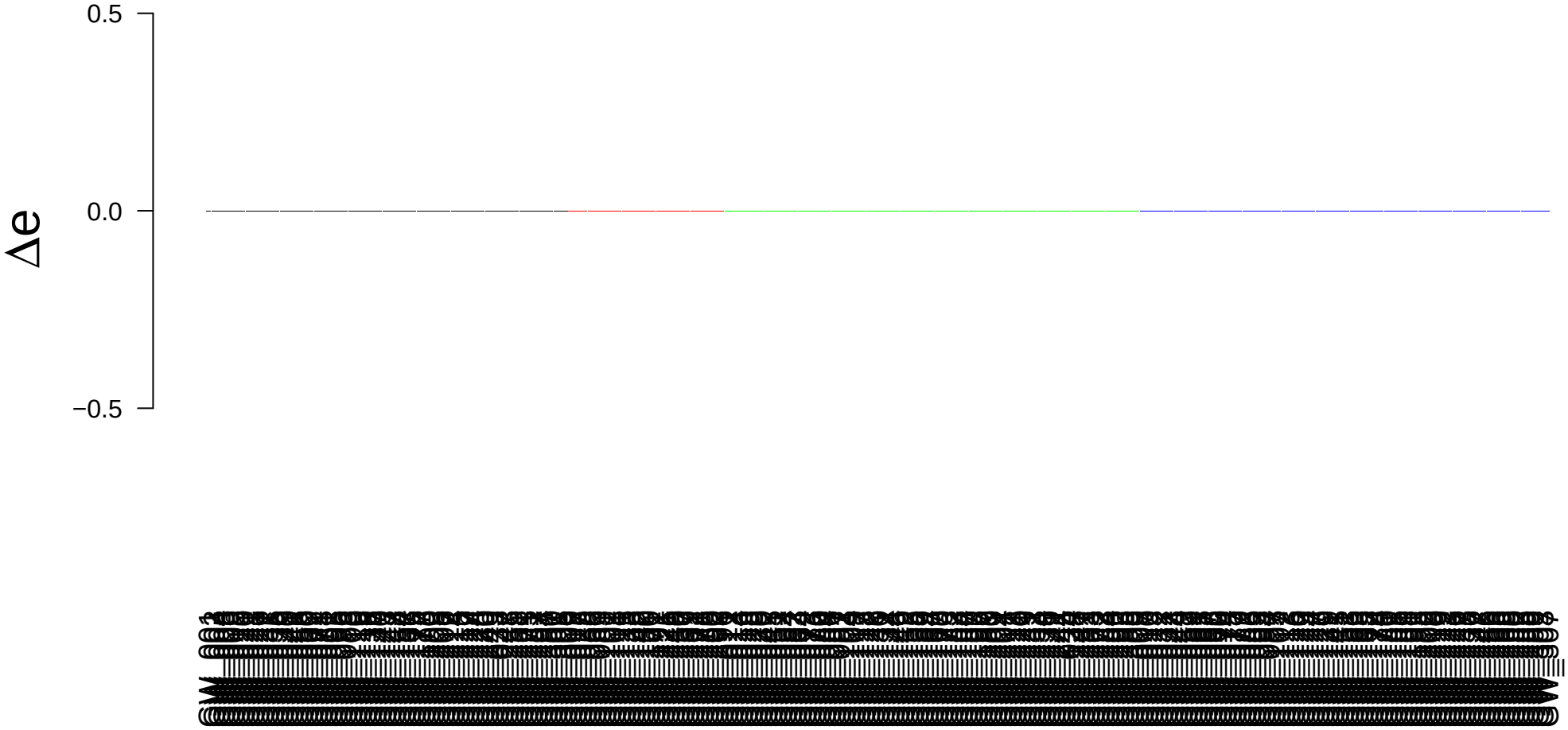

Expression of BENPORATH_CYCLING_GENES in Spot H
0.5
0.0
−0.5
Δe

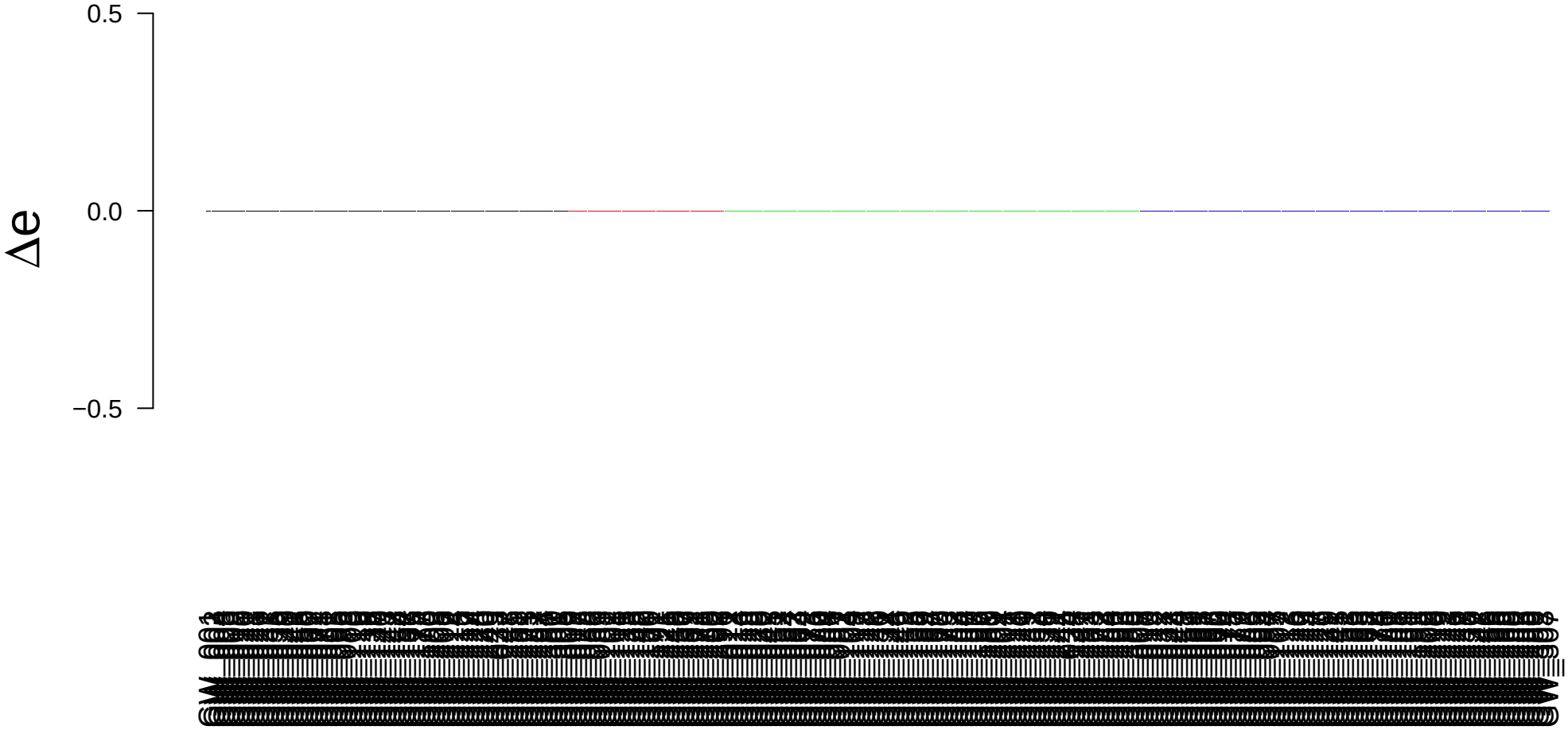

Expression of BENPORATH_CYCLING_GENES in Spot I
0.5
0.0
−0.5
Δe

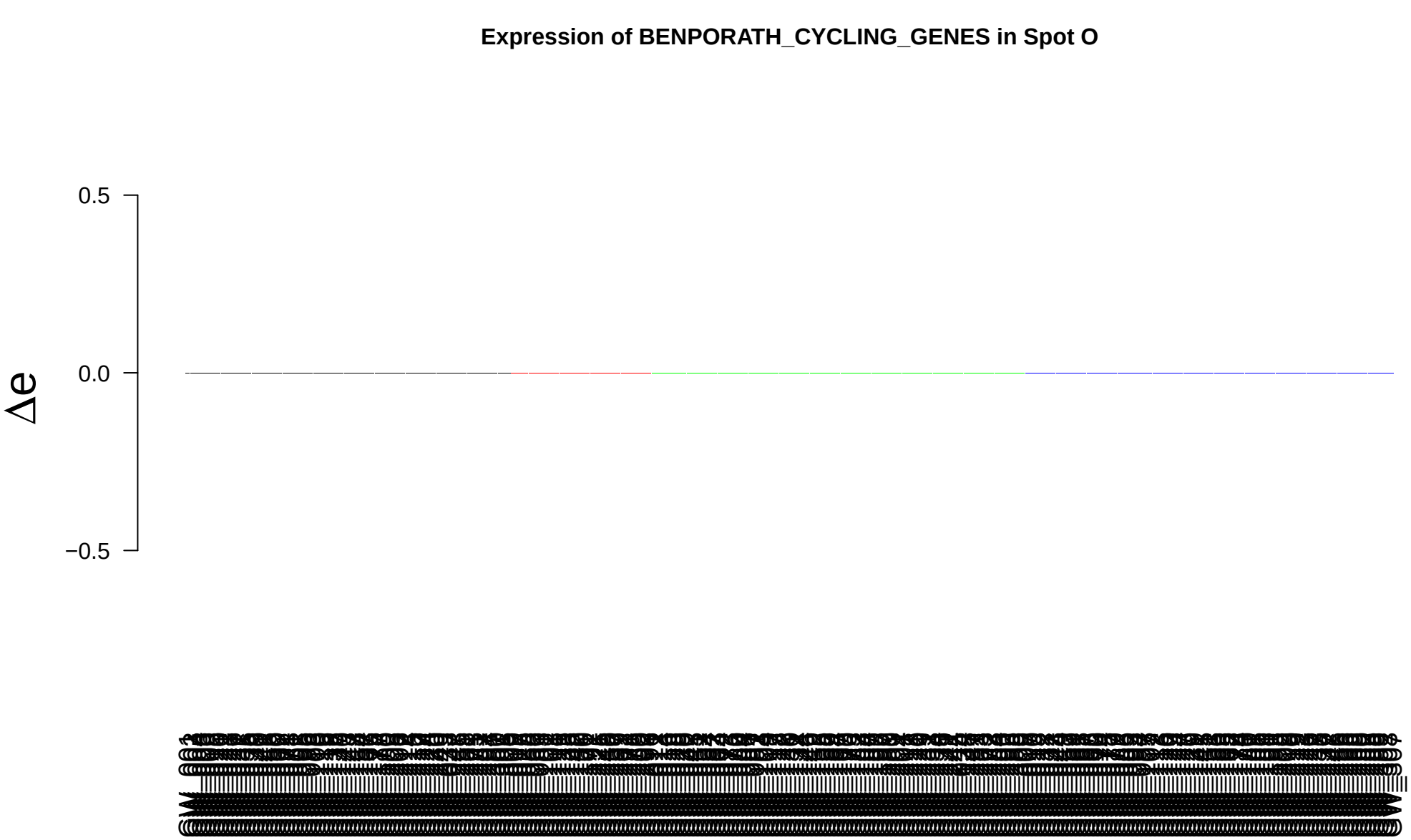

Expression of BENPORATH_CYCLING_GENES in Spot J
0.5
0.0
−0.5
Δe

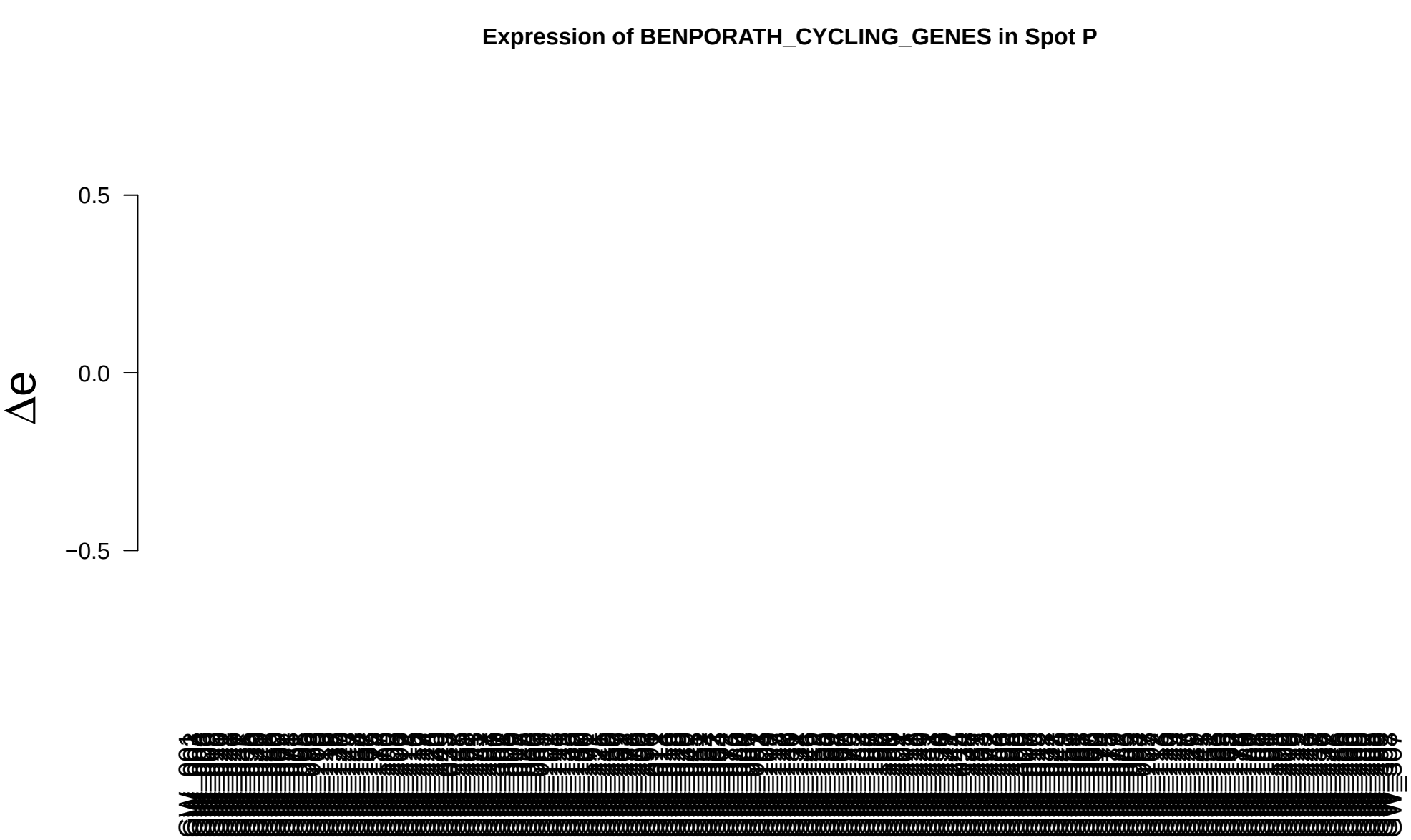

Expression of BENPORATH_CYCLING_GENES in Spot K
Δe
0.5
0.0
−0.5

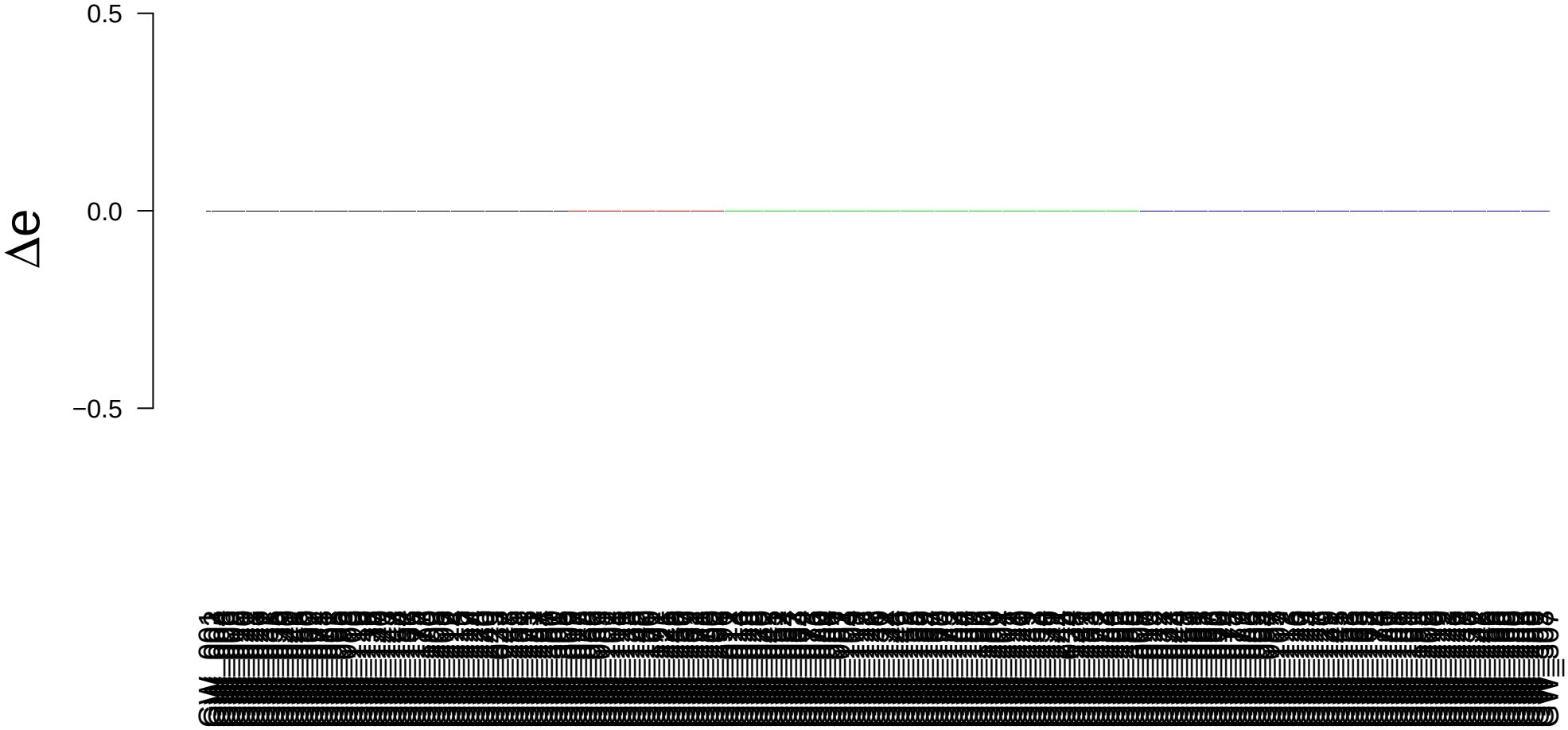

Expression of BENPORATH_CYCLING_GENES in Spot L
0.5
0.0
−0.5
Δe

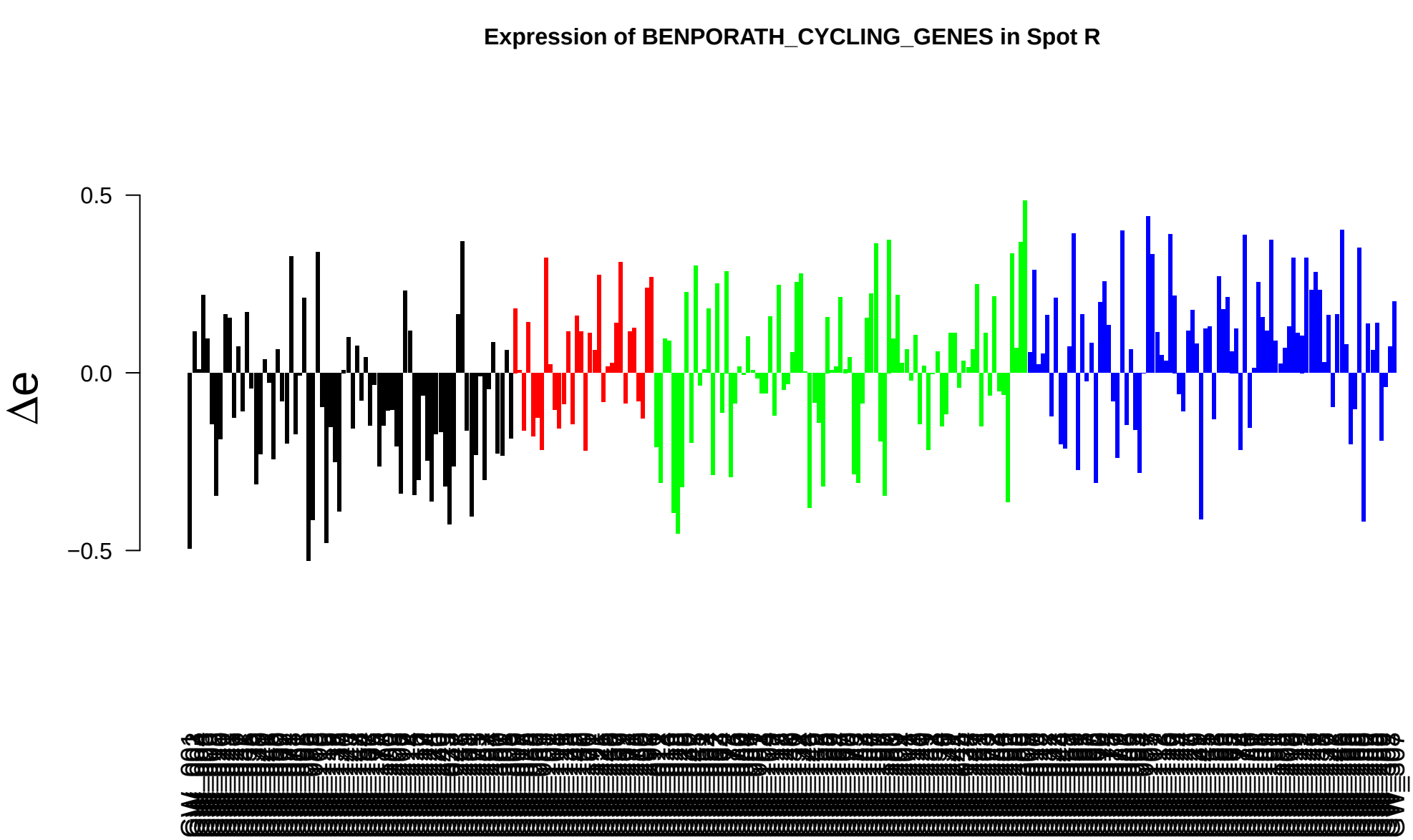

Expression of BENPORATH_CYCLING_GENES in Spot M
0.5
0.0
−0.5
Δe


Expression of BENPORATH_CYCLING_GENES in Spot N
0.5
0.0
−0.5
Δe


Expression of BENPORATH_CYCLING_GENES in Spot O
0.5
0.0
−0.5
Δe


Expression of BENPORATH_CYCLING_GENES in Spot P
0.5
0.0
−0.5
Δe


Expression of BENPORATH_CYCLING_GENES in Spot Q
0.5
0.0
−0.5
Δe


Expression of BENPORATH_CYCLING_GENES in Spot R
0.5
0.0
−0.5
Δe