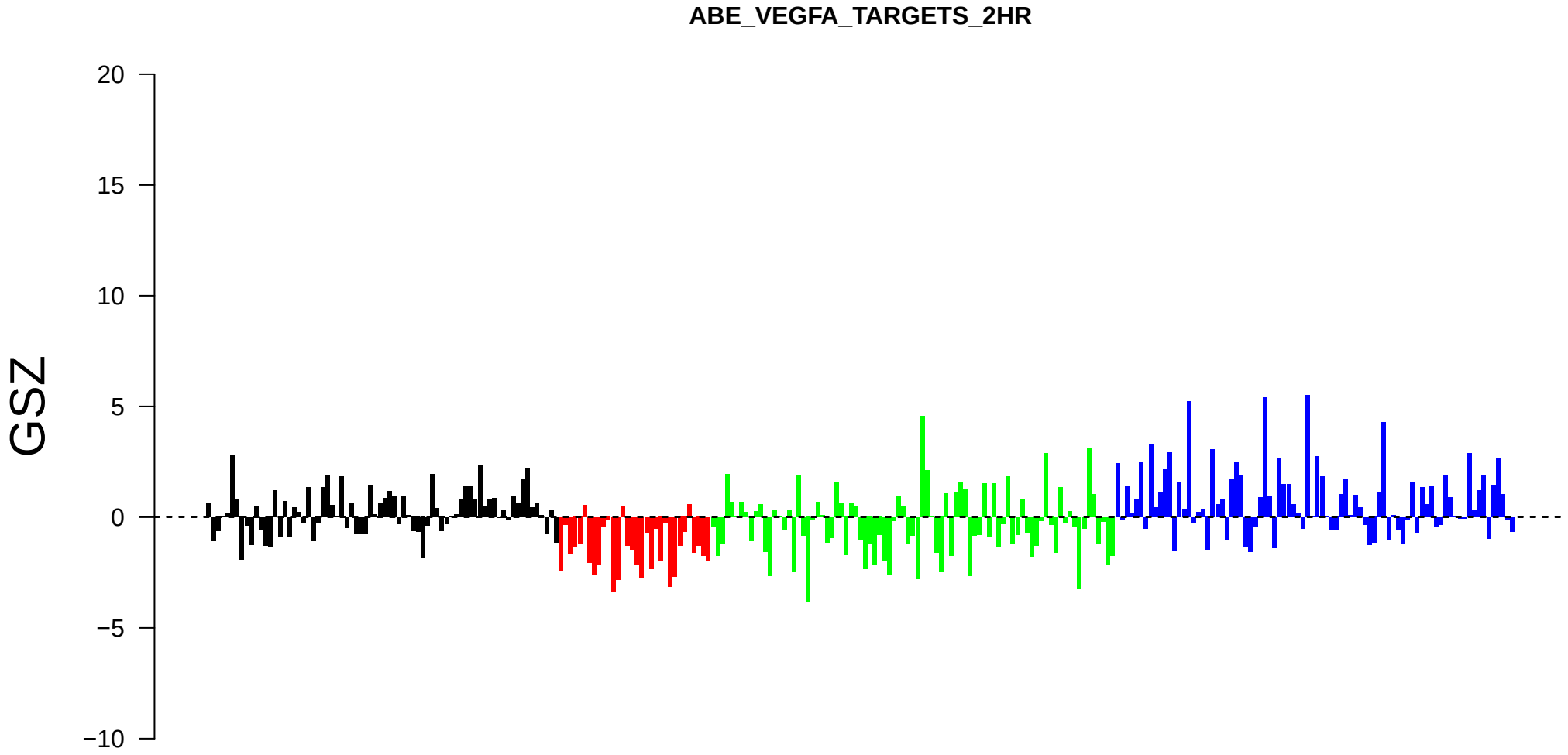
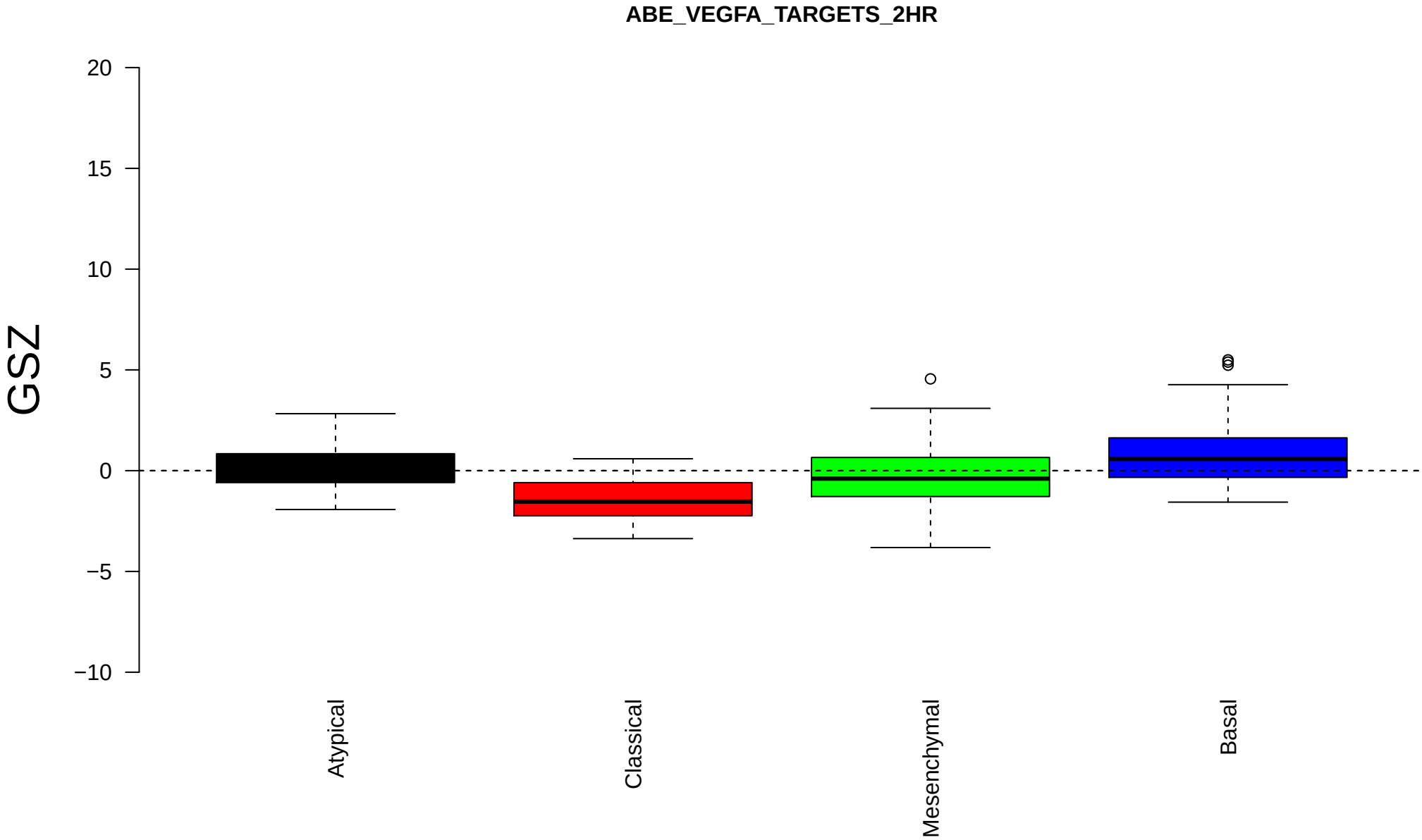
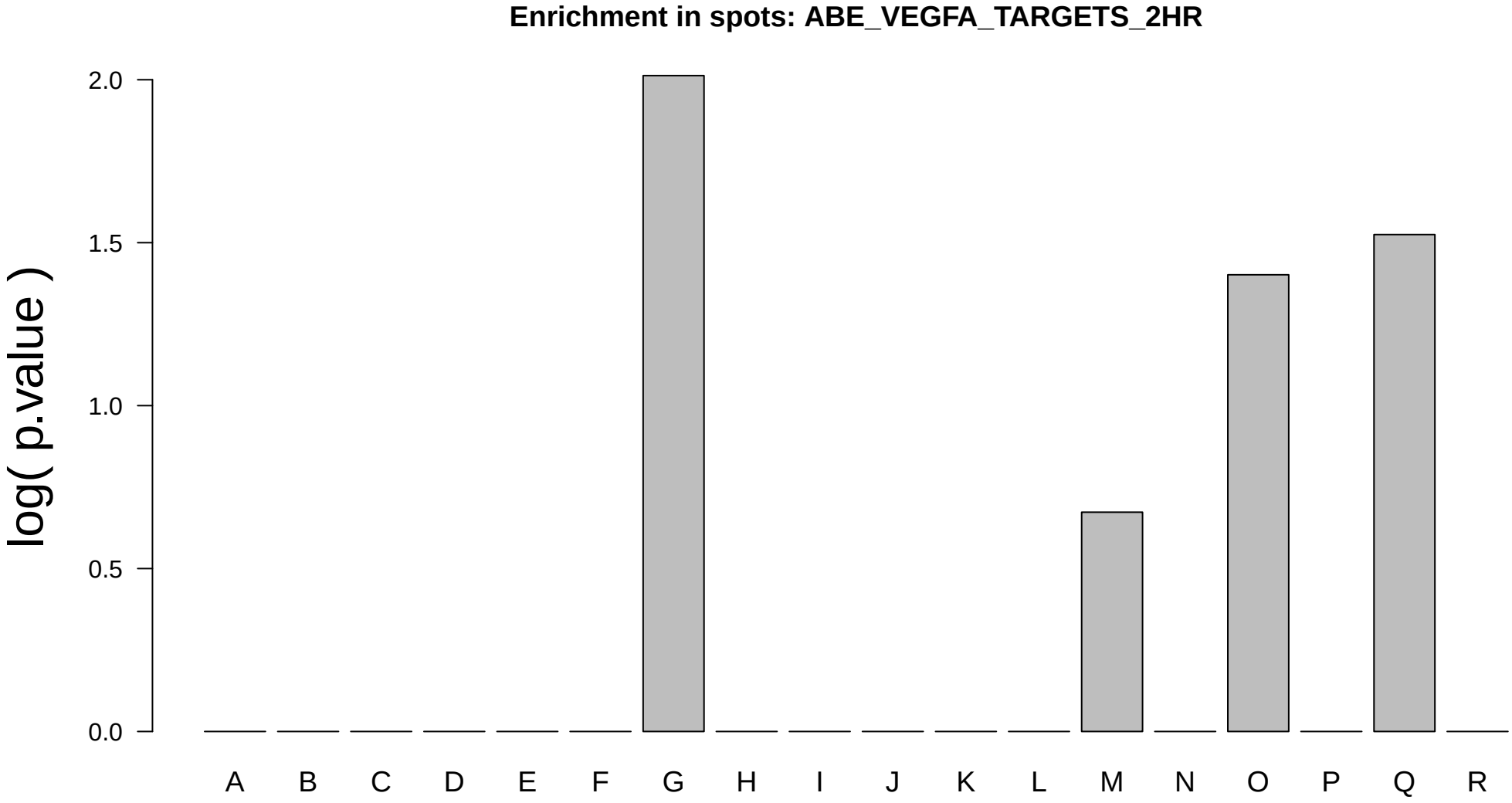
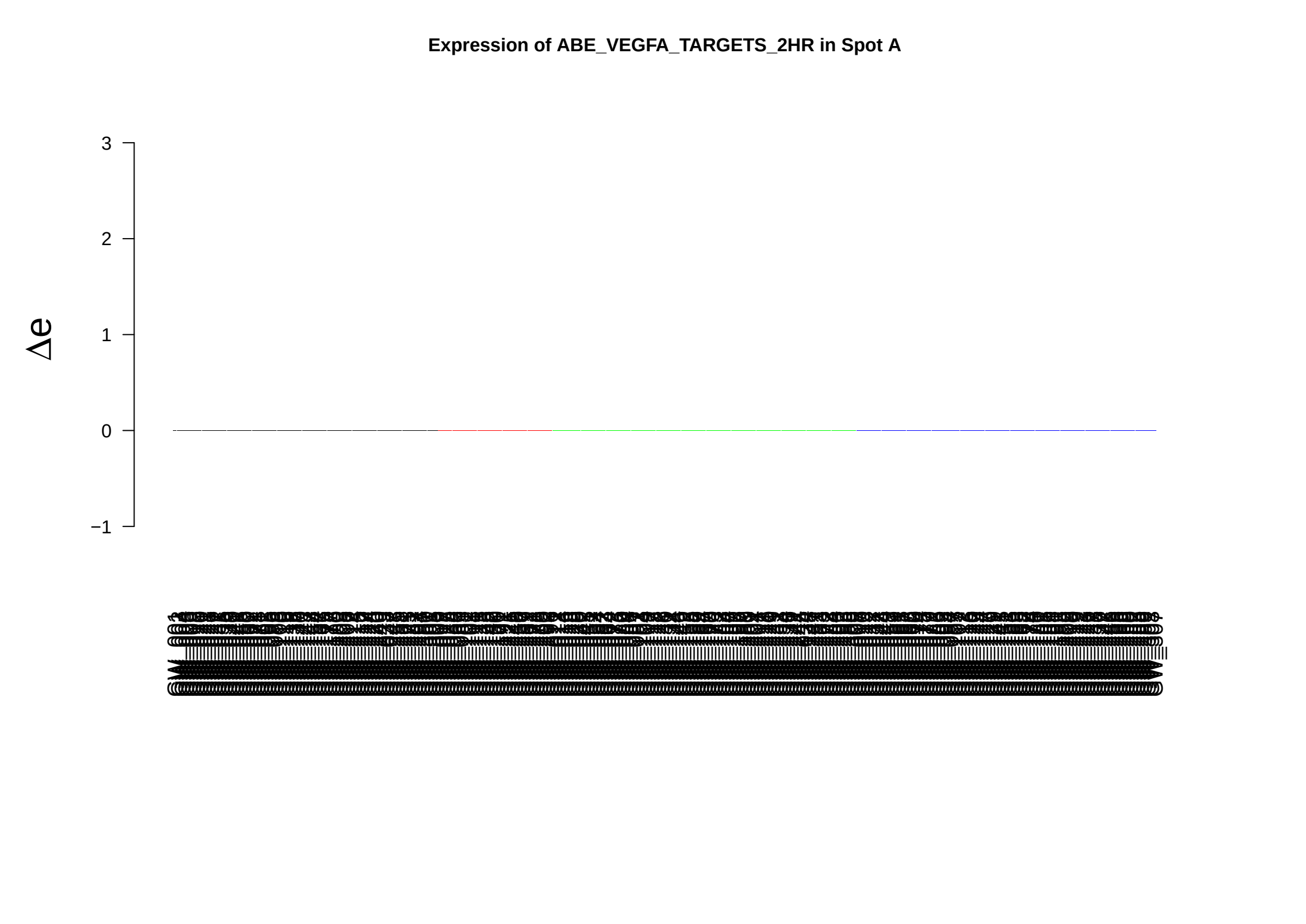
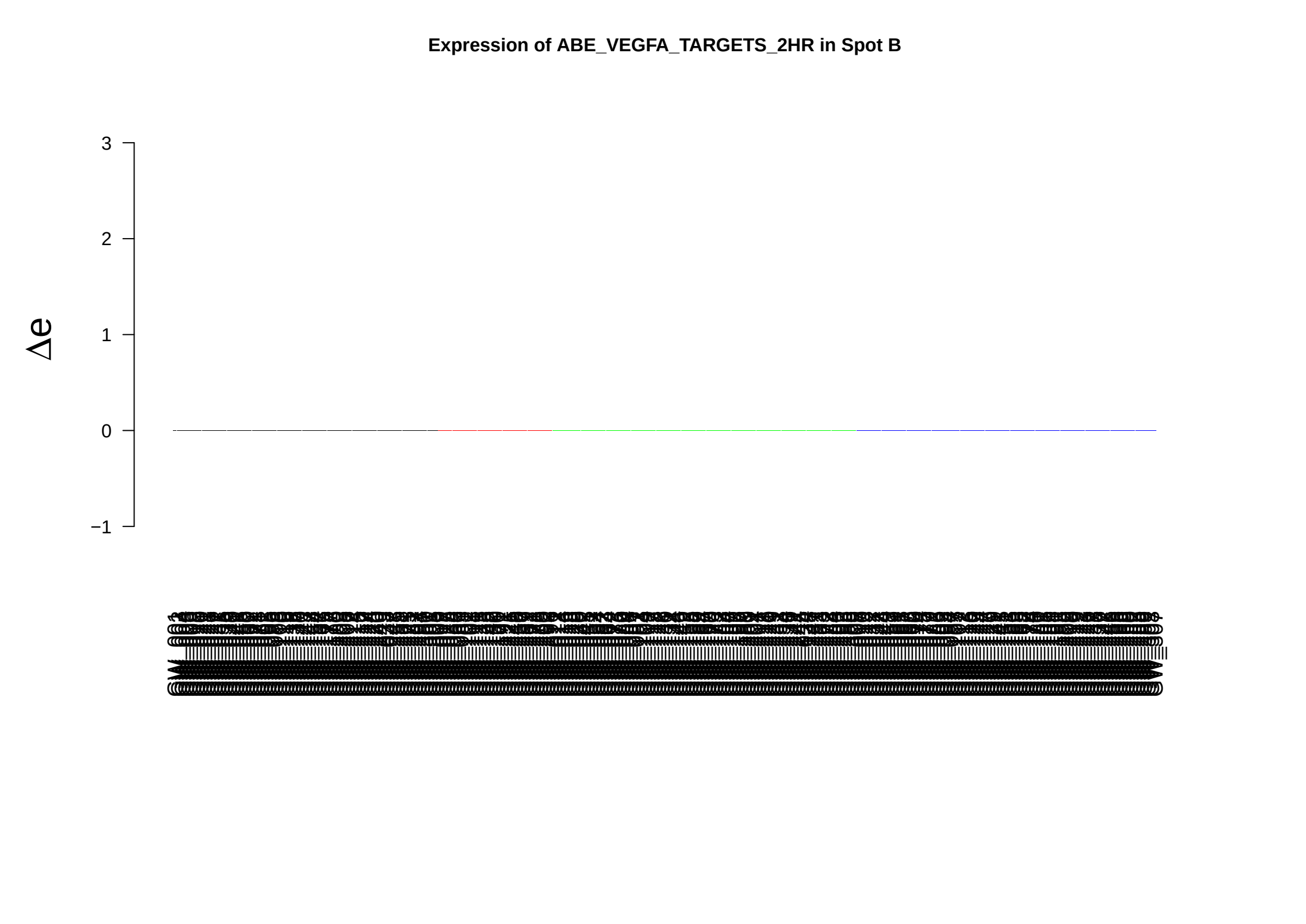
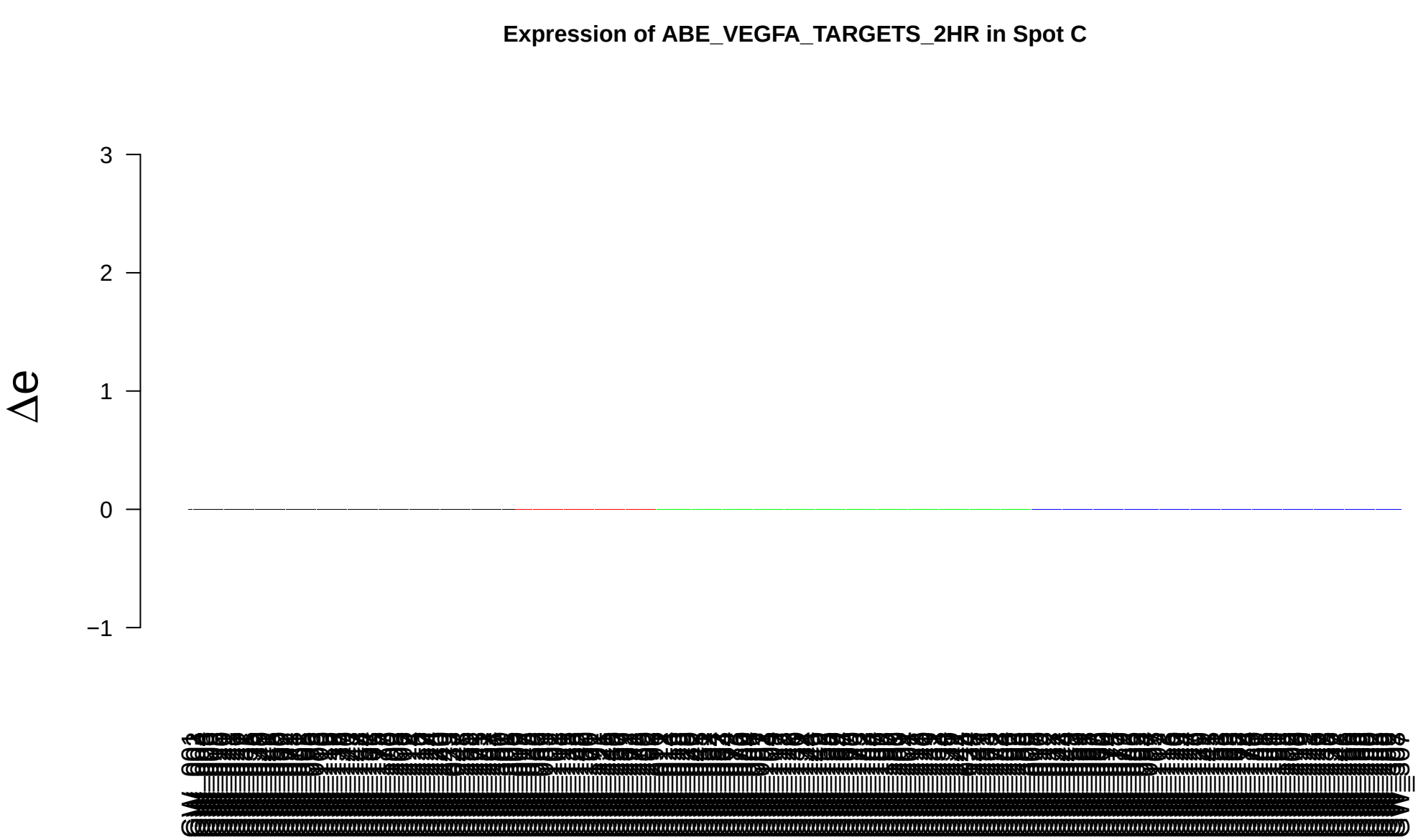

ABE_VEGFA_TARGETS_2HR
GSZ
20
15
10
5
0
−5
−10

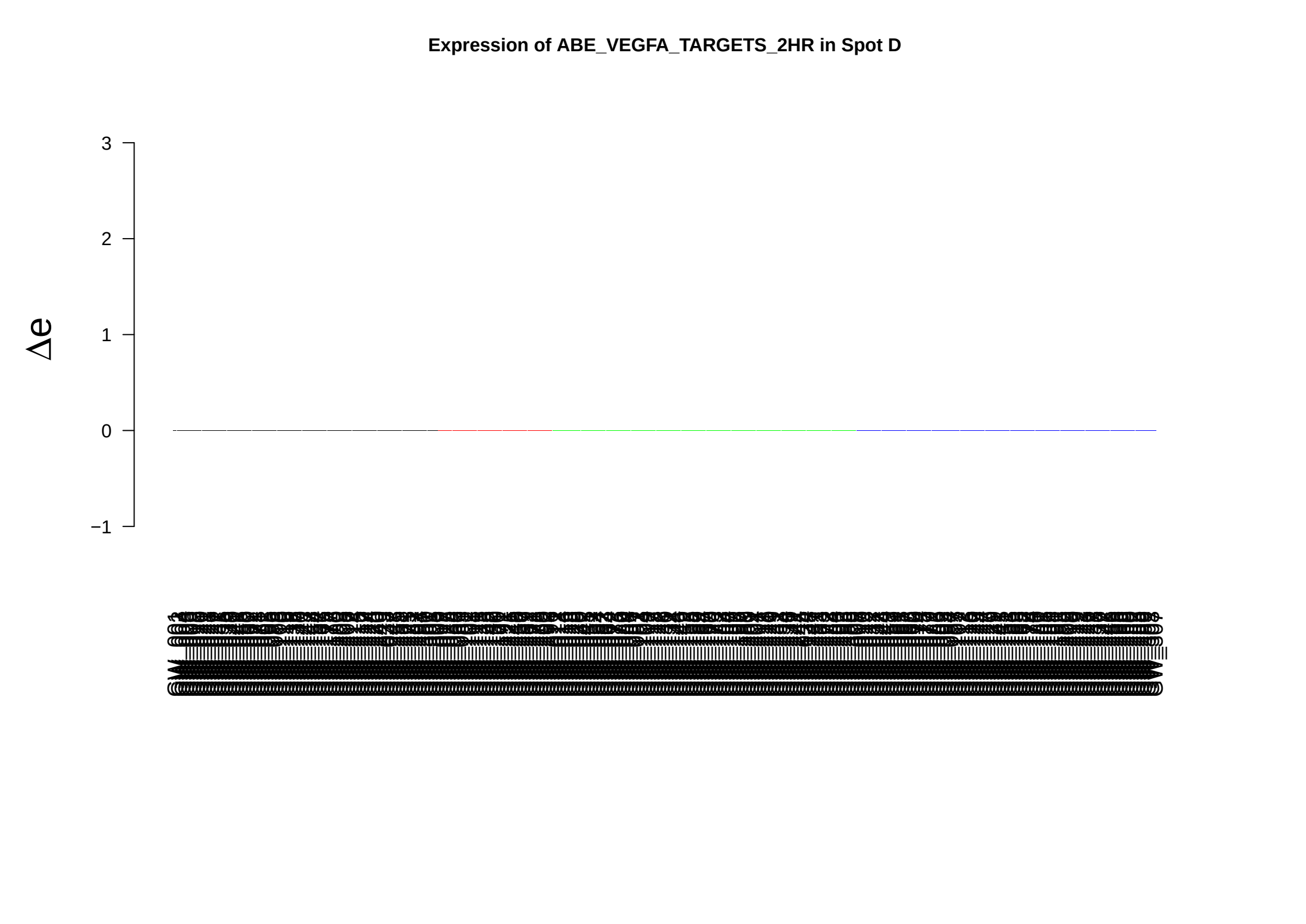

ABE_VEGFA_TARGETS_2HR
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal

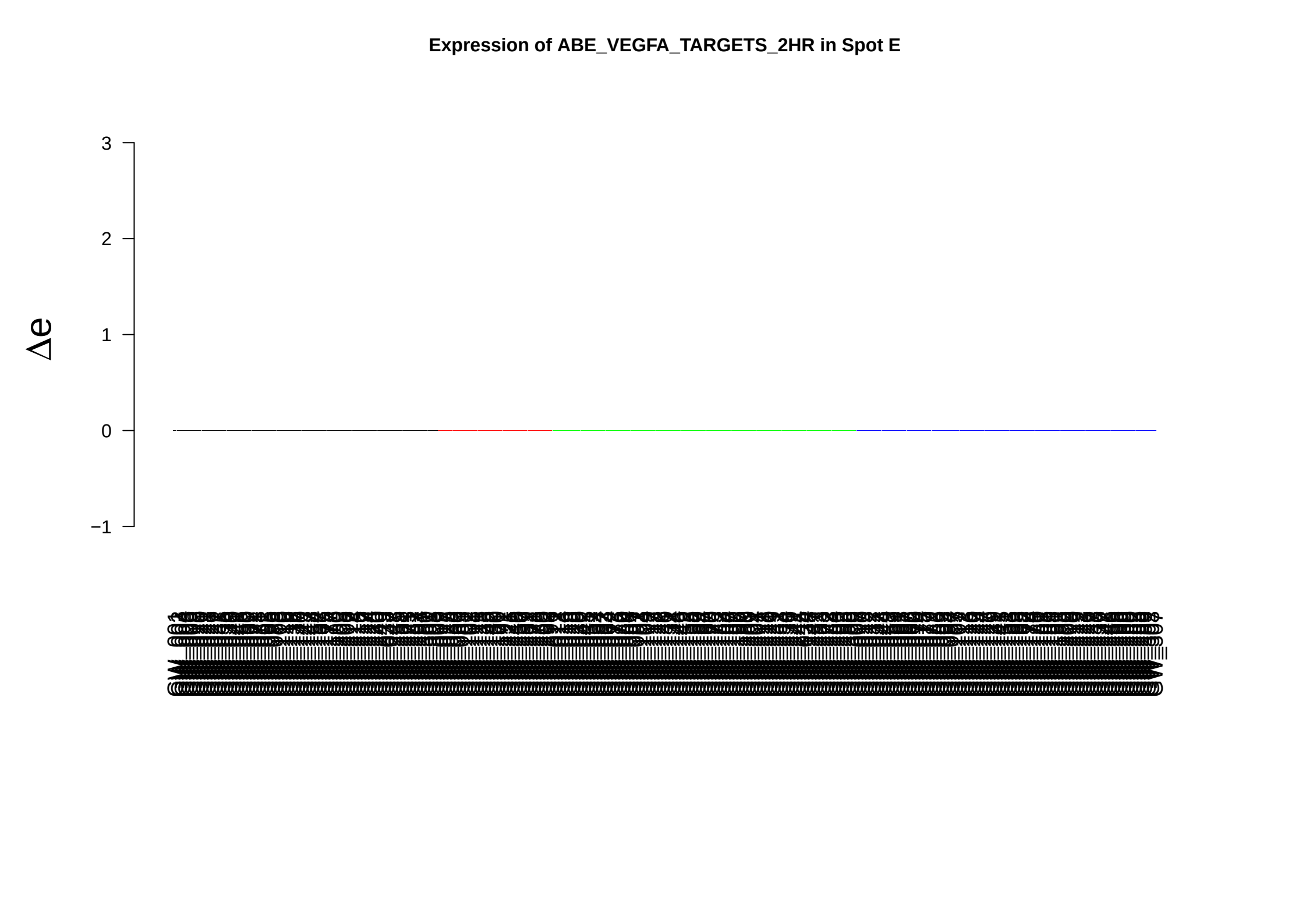

Enrichment in spots: ABE_VEGFA_TARGETS_2HR
log(p.value)
2.0
1.5
1.0
0.5
0.0
A
B
C
D
E
F
G
H
I
J
K
L
M
N
O
P
Q
R

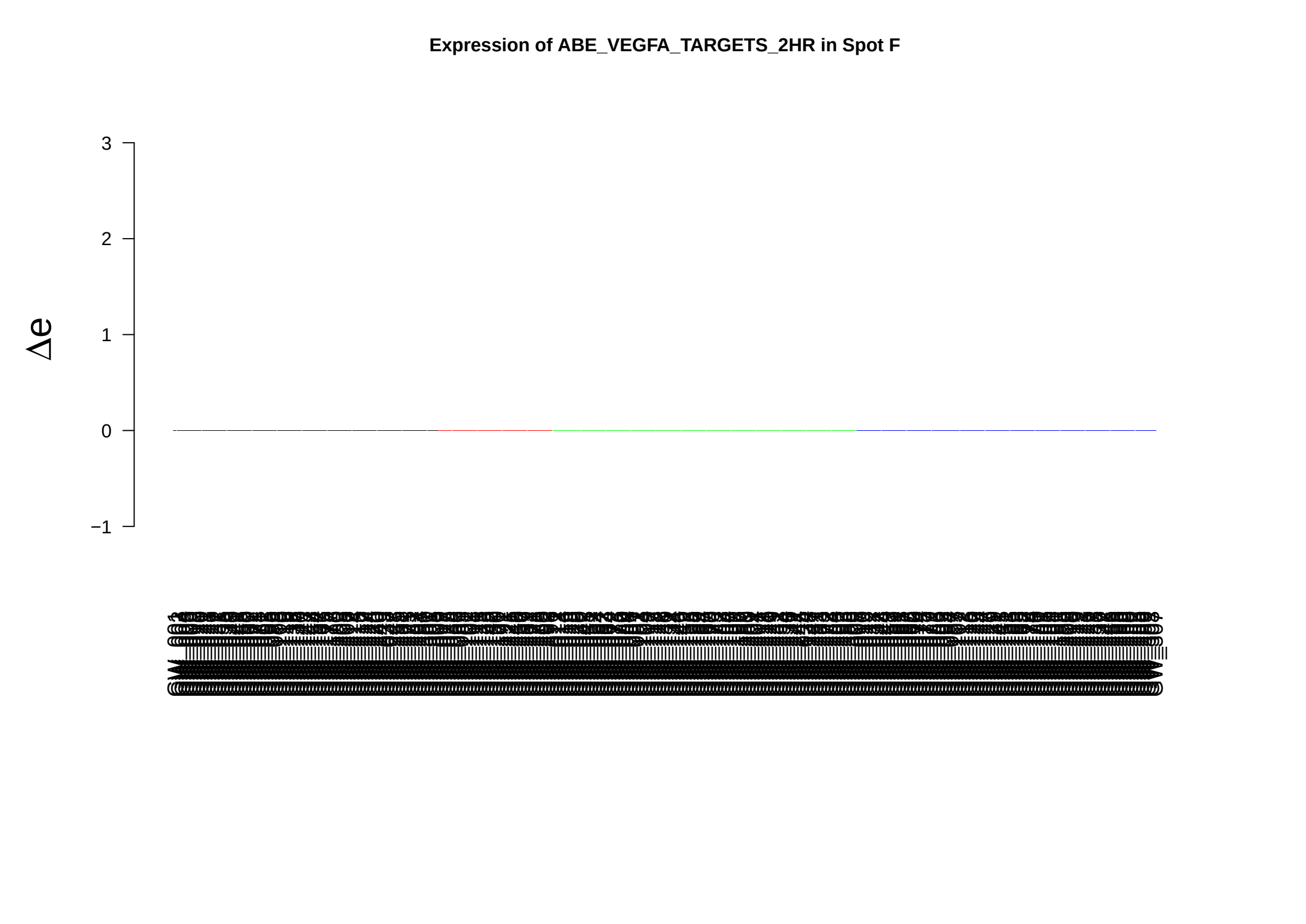

Expression of ABE_VEGFA_TARGETS_2HR in Spot A
Δe
3
2
1
0
−1

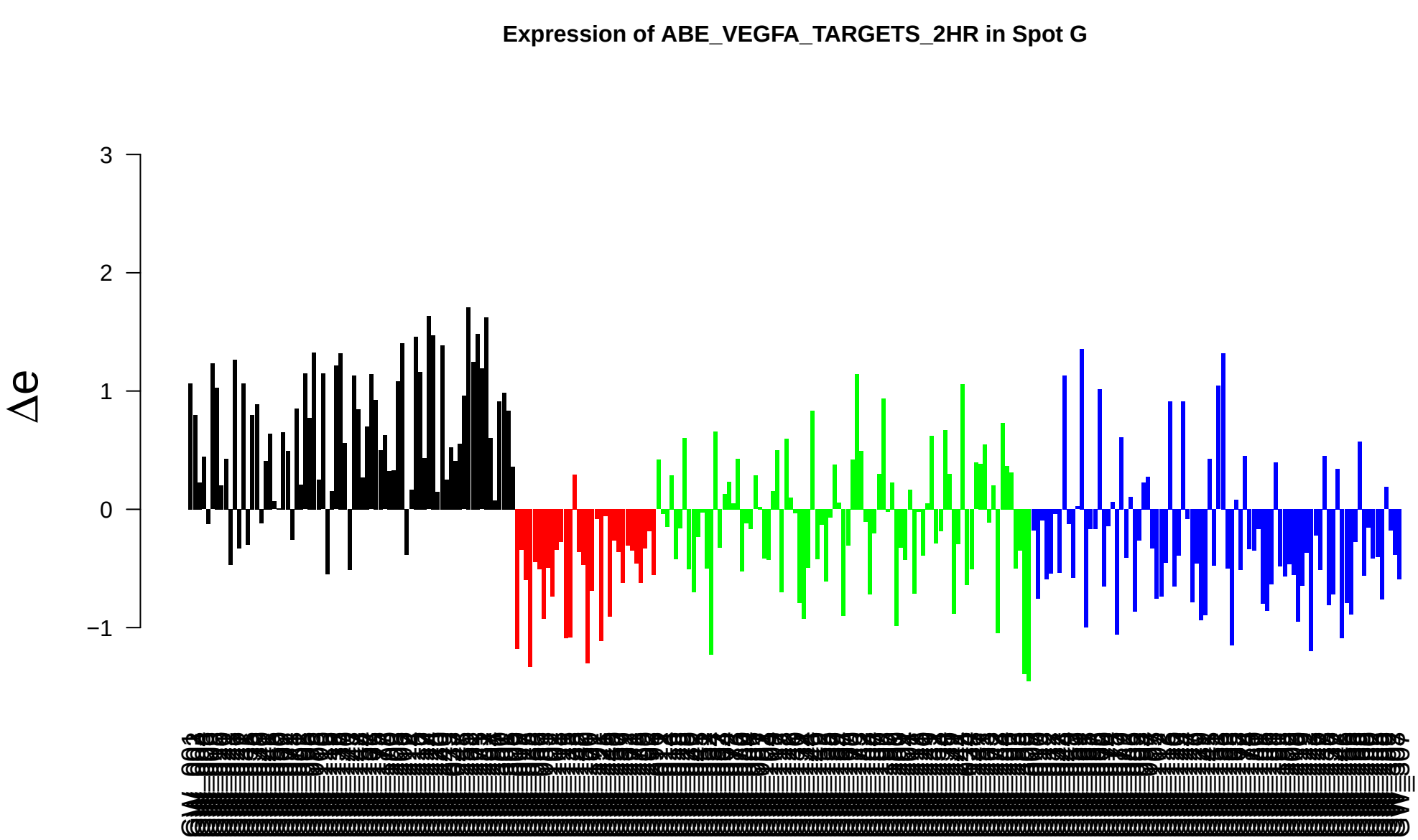

Expression of ABE_VEGFA_TARGETS_2HR in Spot B
Δe
3
2
1
0
−1

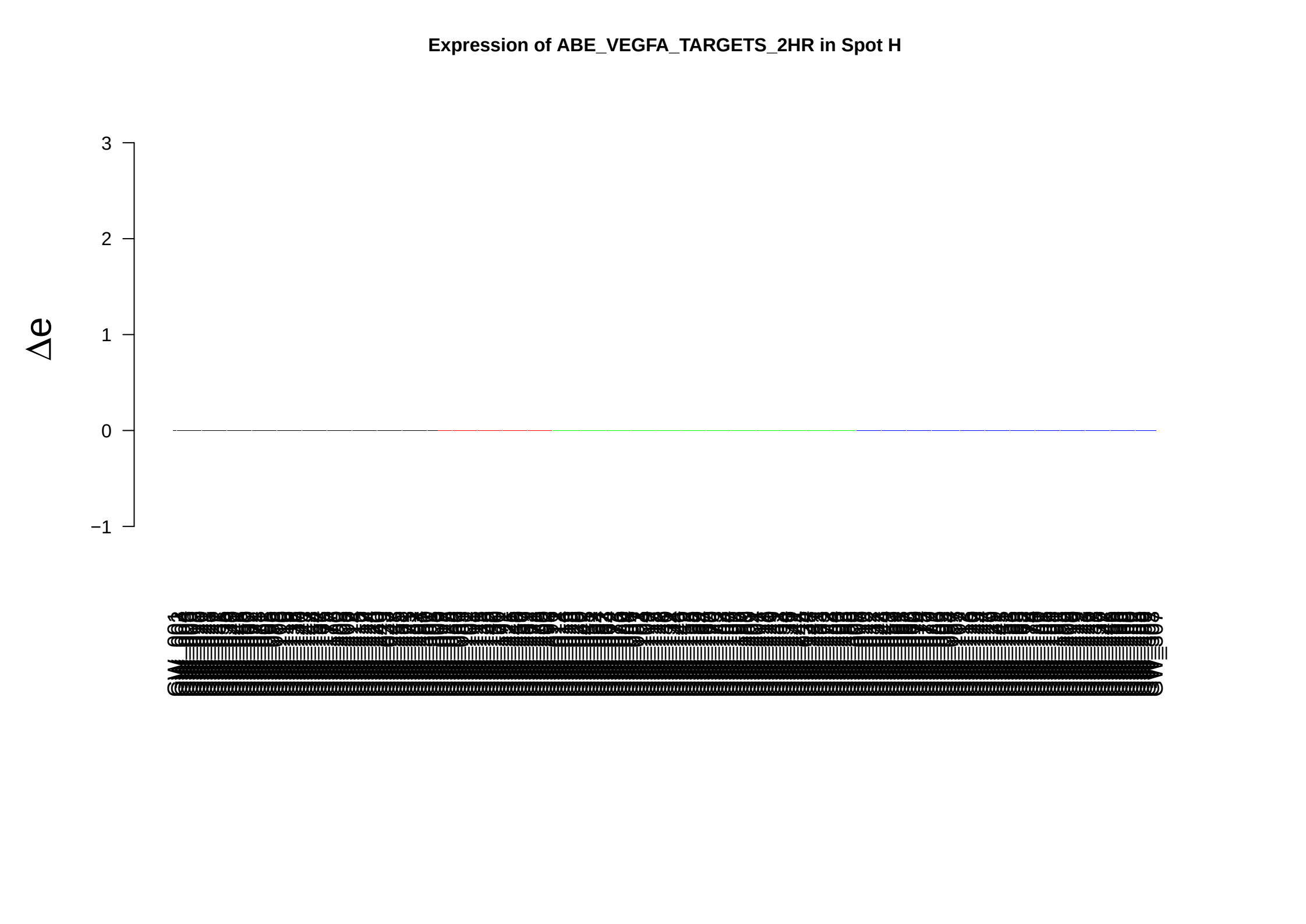

Expression of ABE_VEGFA_TARGETS_2HR in Spot C
Δe
3
2
1
0
−1

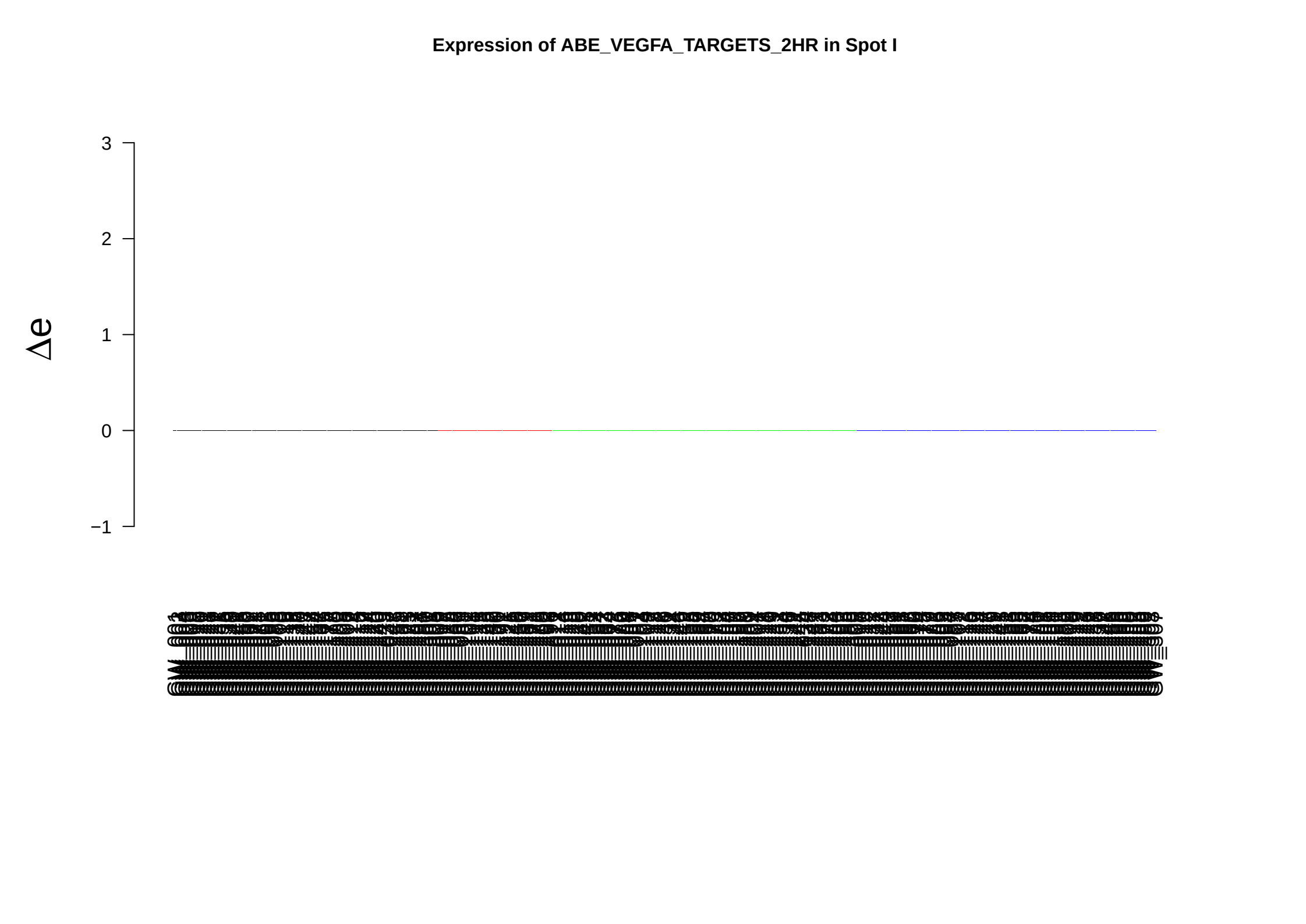

Expression of ABE_VEGFA_TARGETS_2HR in Spot D
Δe
3
2
1
0
−1

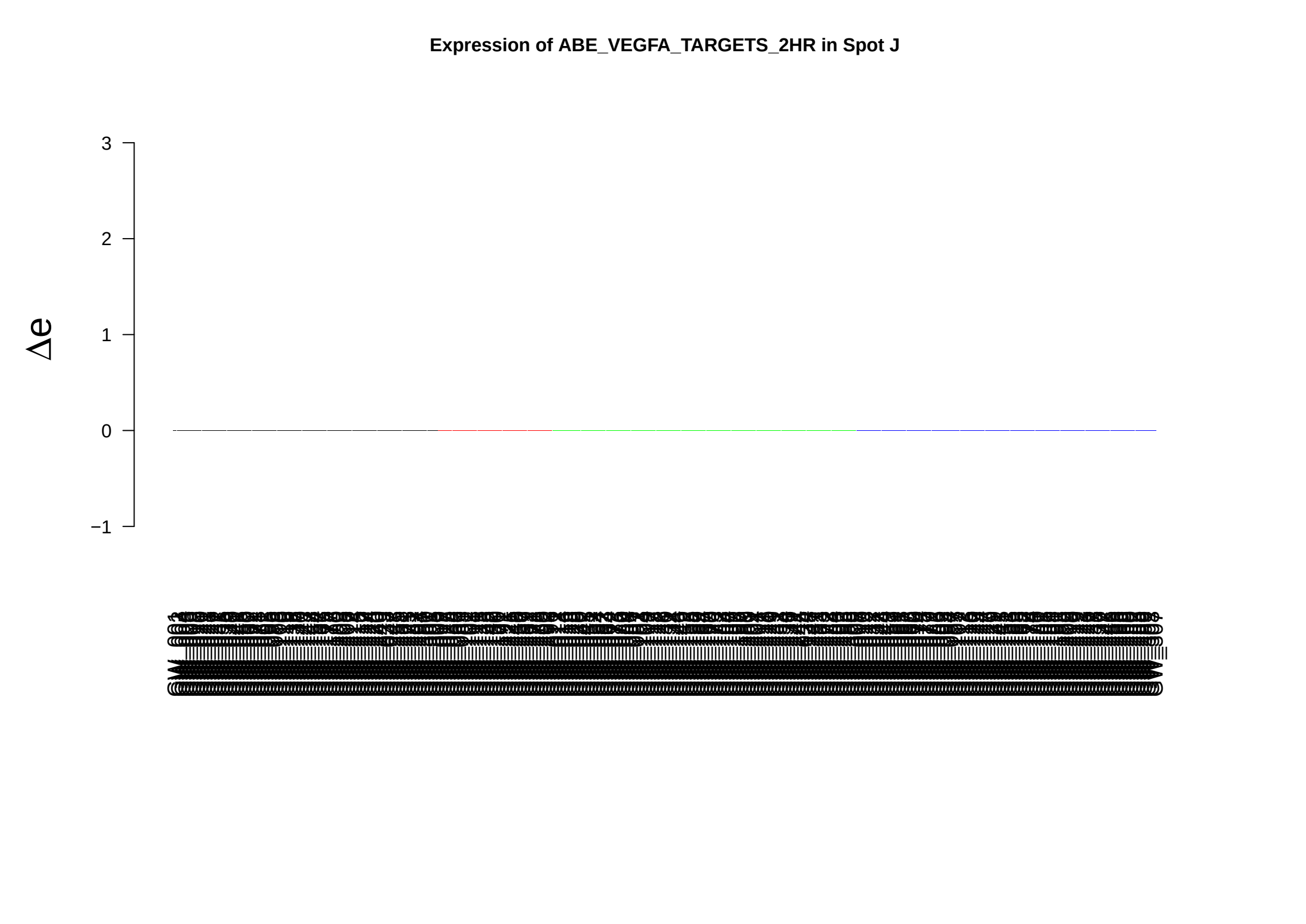

Expression of ABE_VEGFA_TARGETS_2HR in Spot E
Δe
3
2
1
0
−1

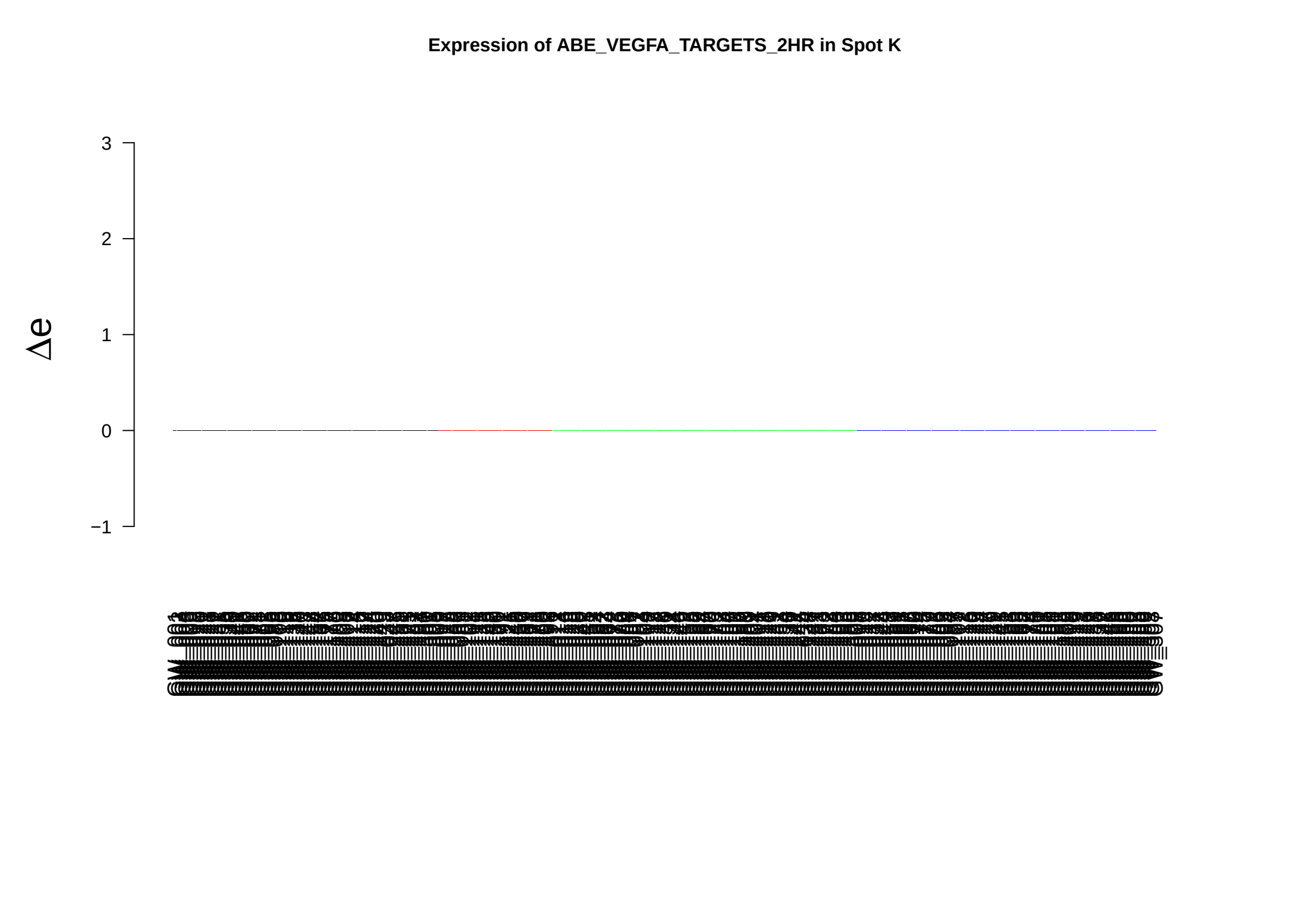

Expression of ABE_VEGFA_TARGETS_2HR in Spot F
Δe
3
2
1
0
−1

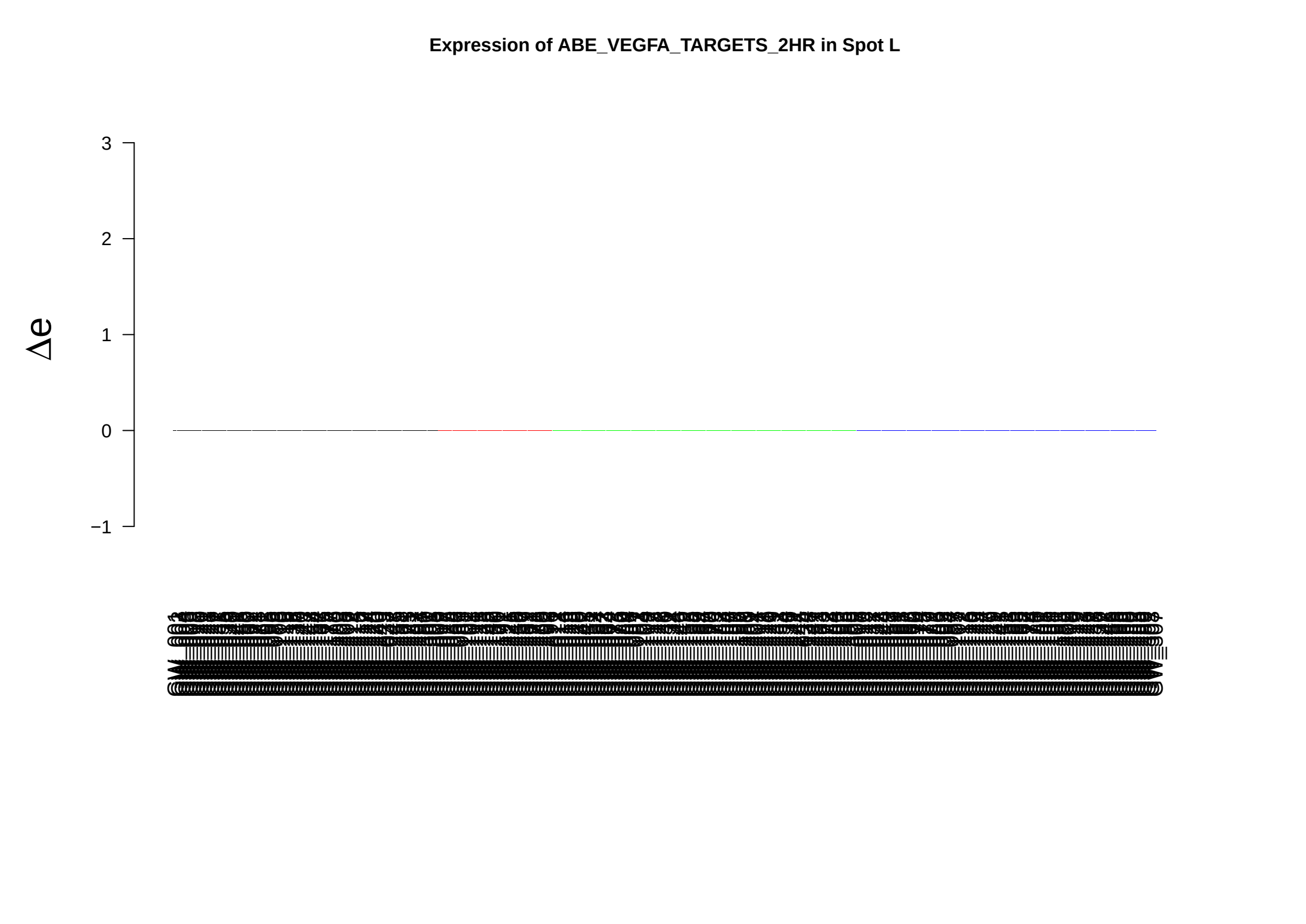

Expression of ABE_VEGFA_TARGETS_2HR in Spot G
Δe
3
2
1
0
−1

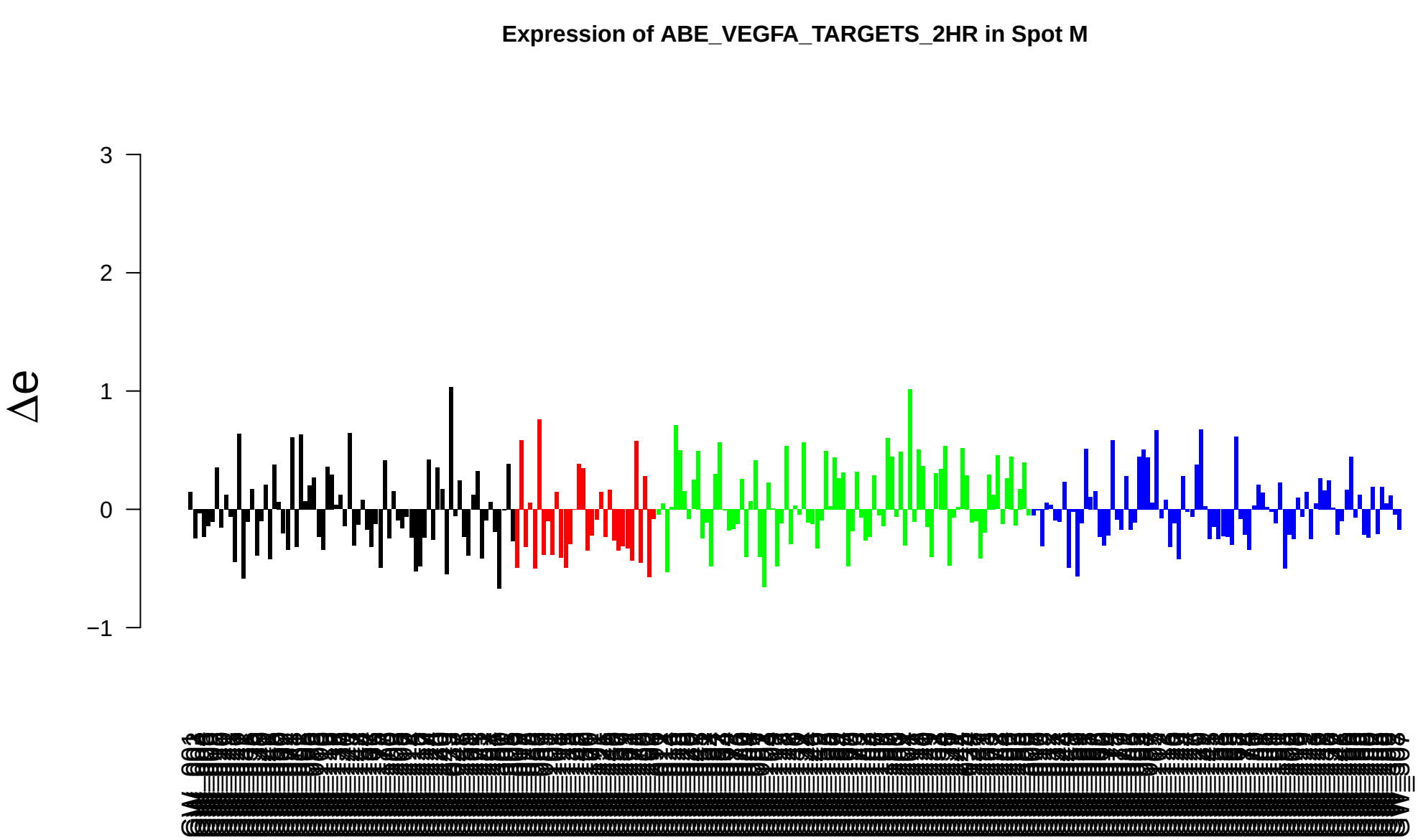

Expression of ABE_VEGFA_TARGETS_2HR in Spot H
Δe
3
2
1
0
−1

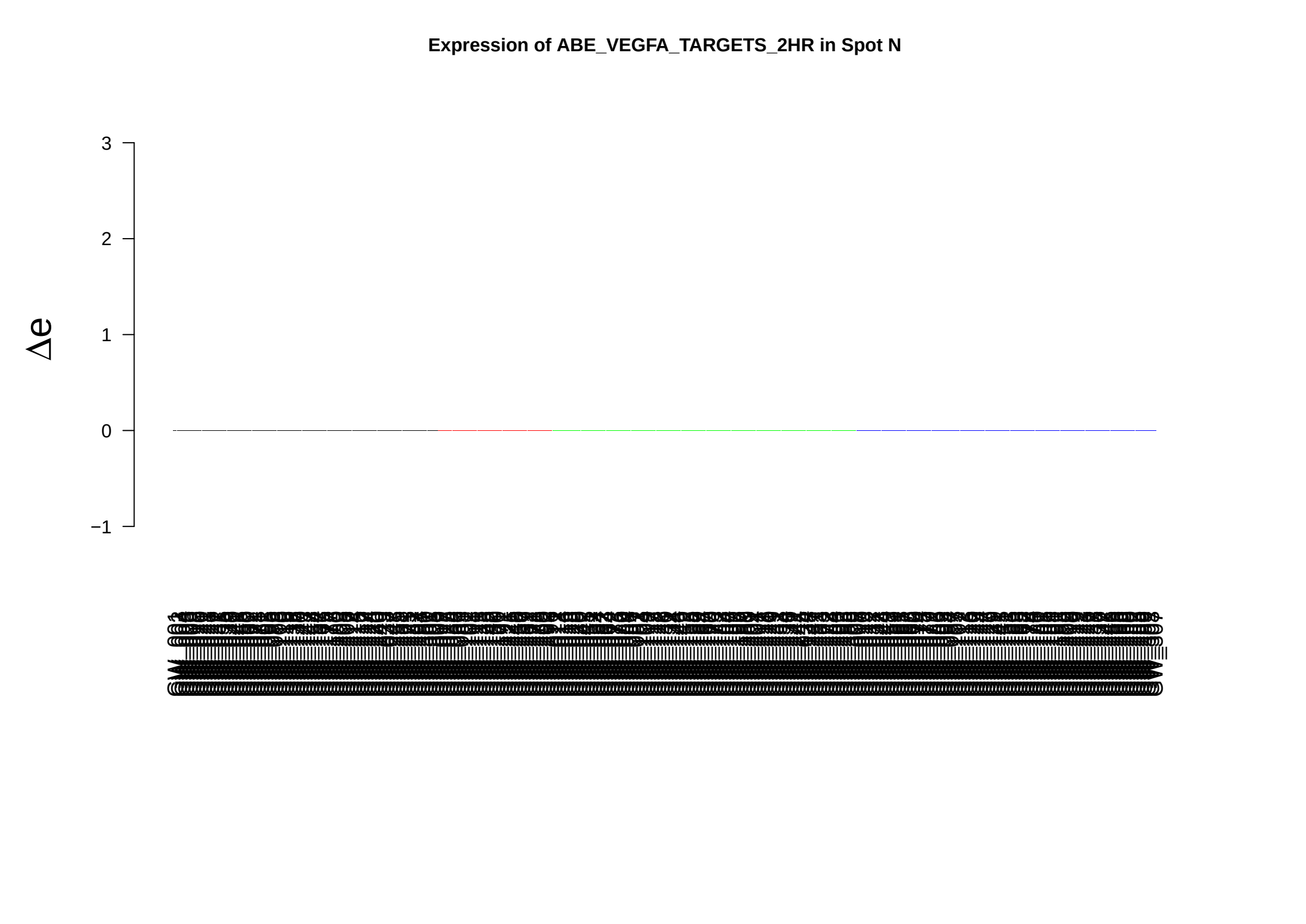

Expression of ABE_VEGFA_TARGETS_2HR in Spot I
Δe
3
2
1
0
−1

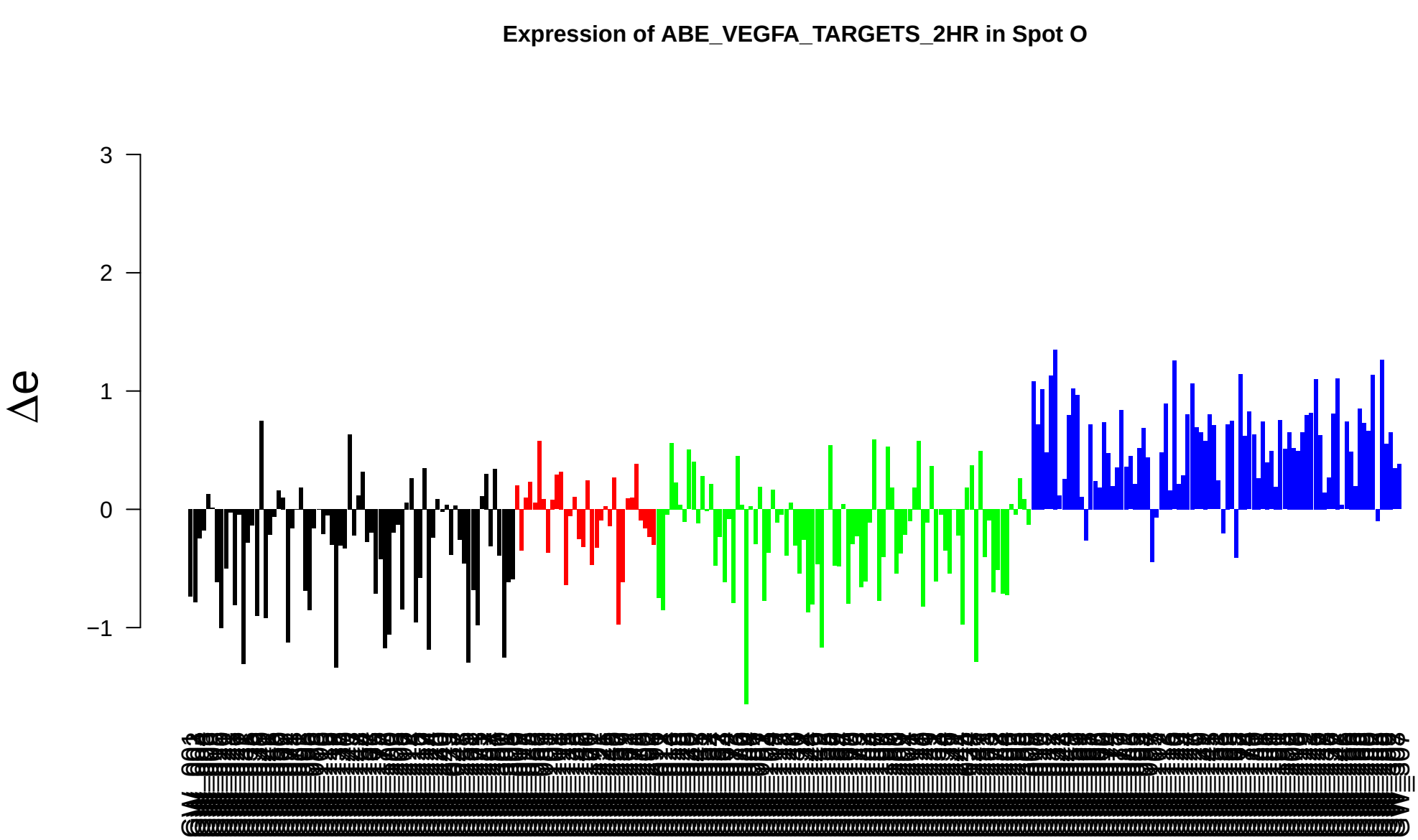

Expression of ABE_VEGFA_TARGETS_2HR in Spot J
Δe
3
2
1
0
−1

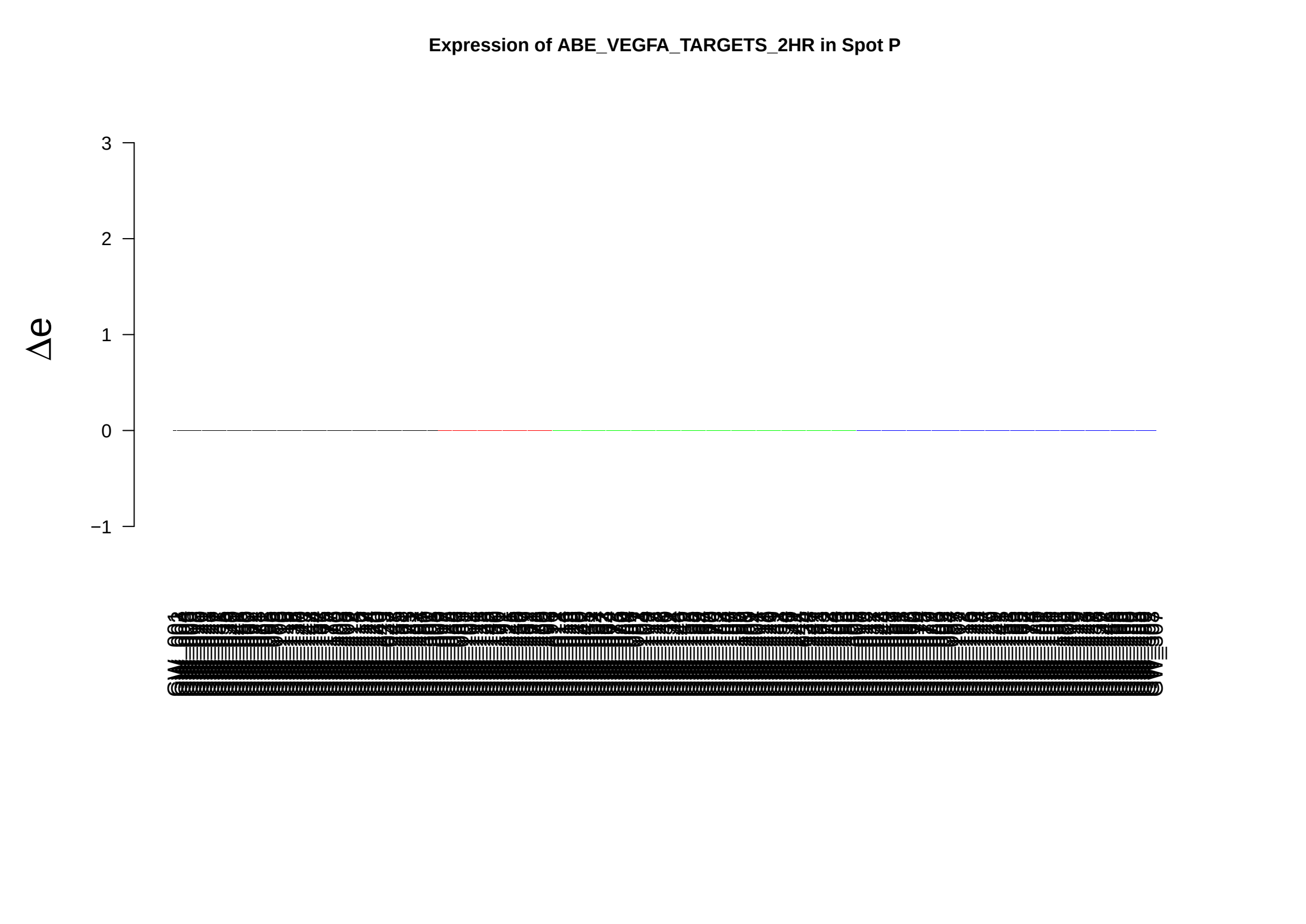

Expression of ABE_VEGFA_TARGETS_2HR in Spot K
Δe
3
2
1
0
−1

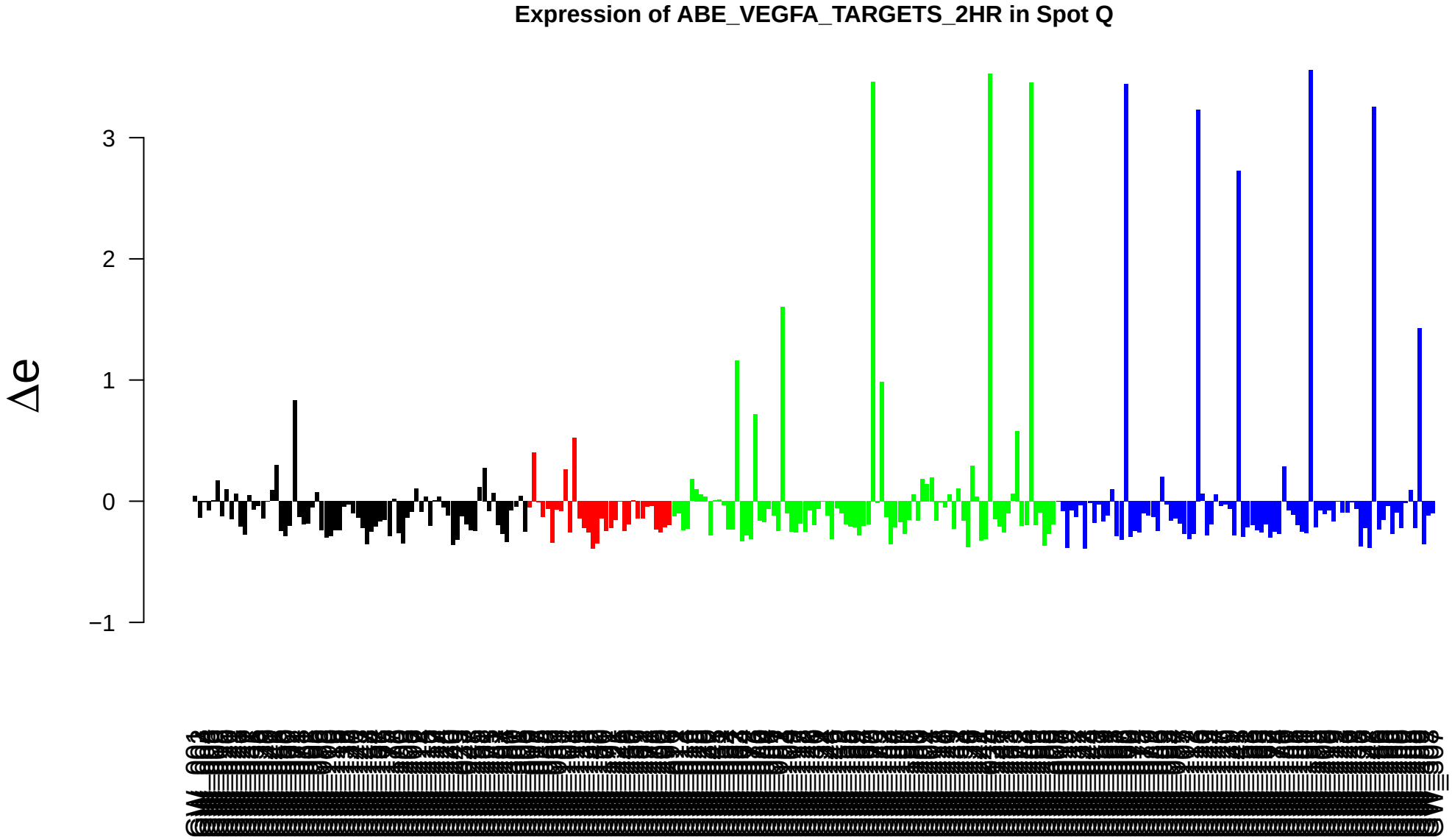

Expression of ABE_VEGFA_TARGETS_2HR in Spot L
Δe
3
2
1
0
−1

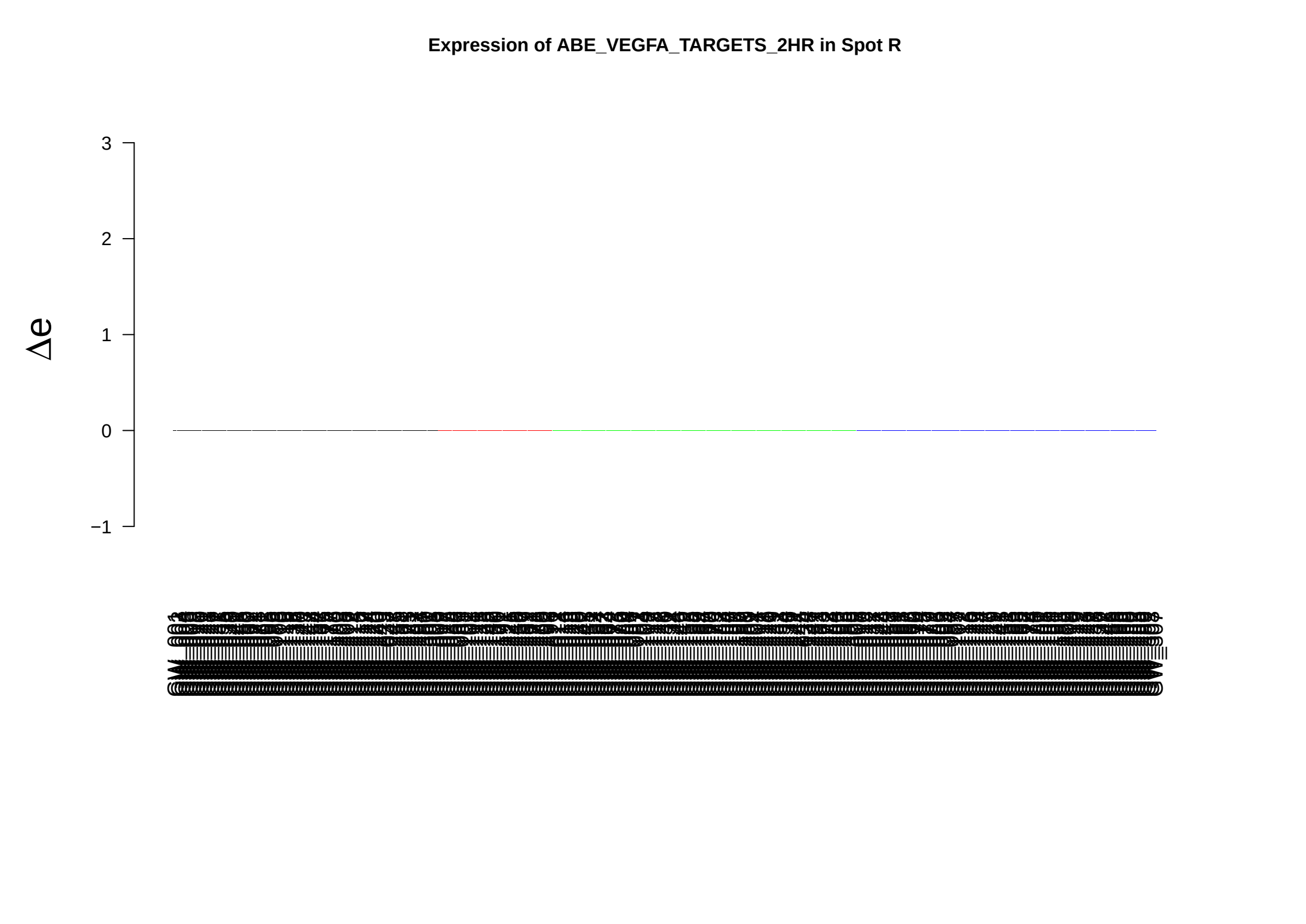

Expression of ABE_VEGFA_TARGETS_2HR in Spot M
Δe
3
2
1
0
−1


Expression of ABE_VEGFA_TARGETS_2HR in Spot N
Δe
3
2
1
0
−1


Expression of ABE_VEGFA_TARGETS_2HR in Spot O
Δe
3
2
1
0
−1


Expression of ABE_VEGFA_TARGETS_2HR in Spot P
Δe
3
2
1
0
−1


Expression of ABE_VEGFA_TARGETS_2HR in Spot Q
Δe
3
2
1
0
−1


Expression of ABE_VEGFA_TARGETS_2HR in Spot R
Δe
3
2
1
0
−1