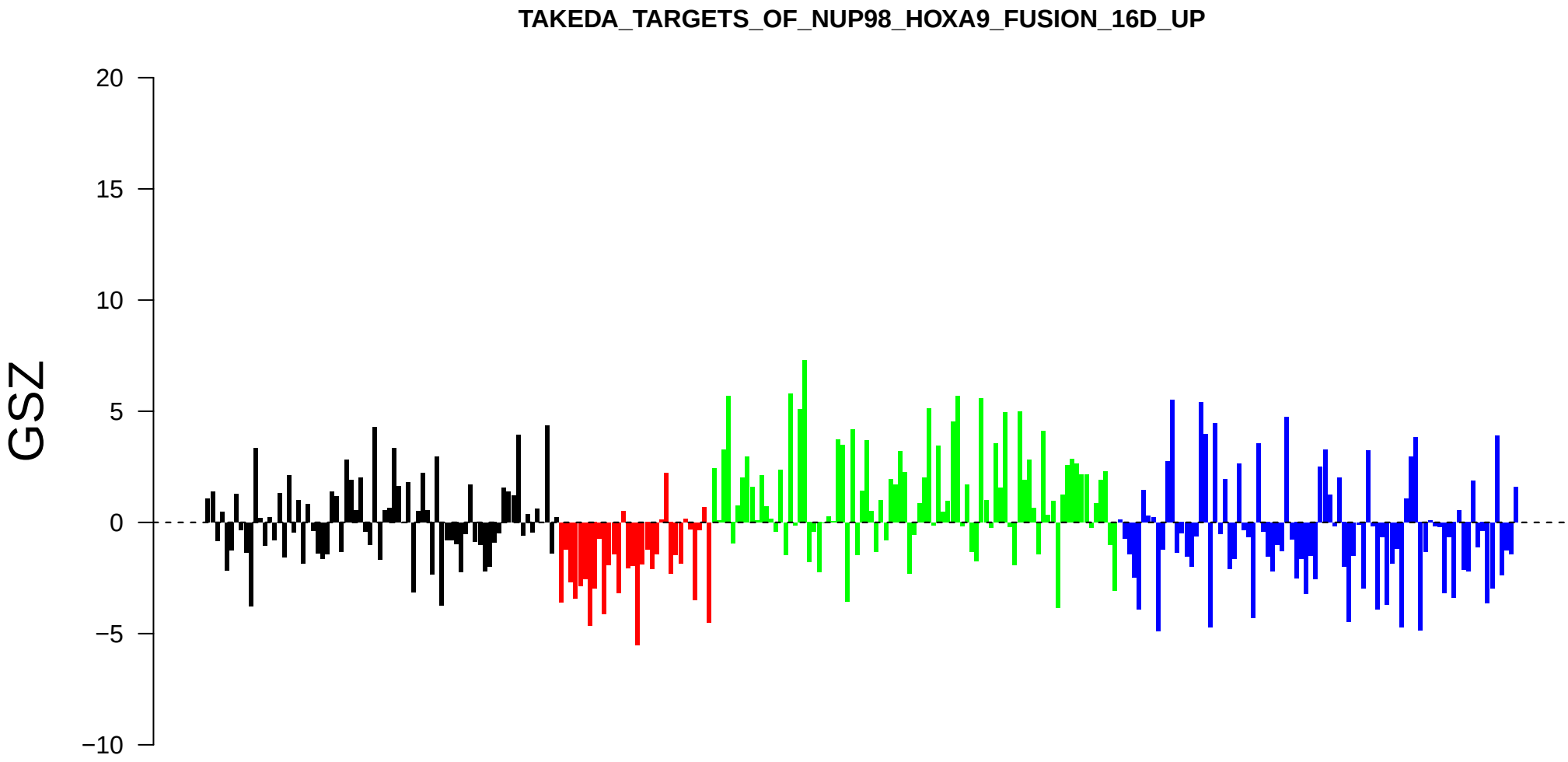
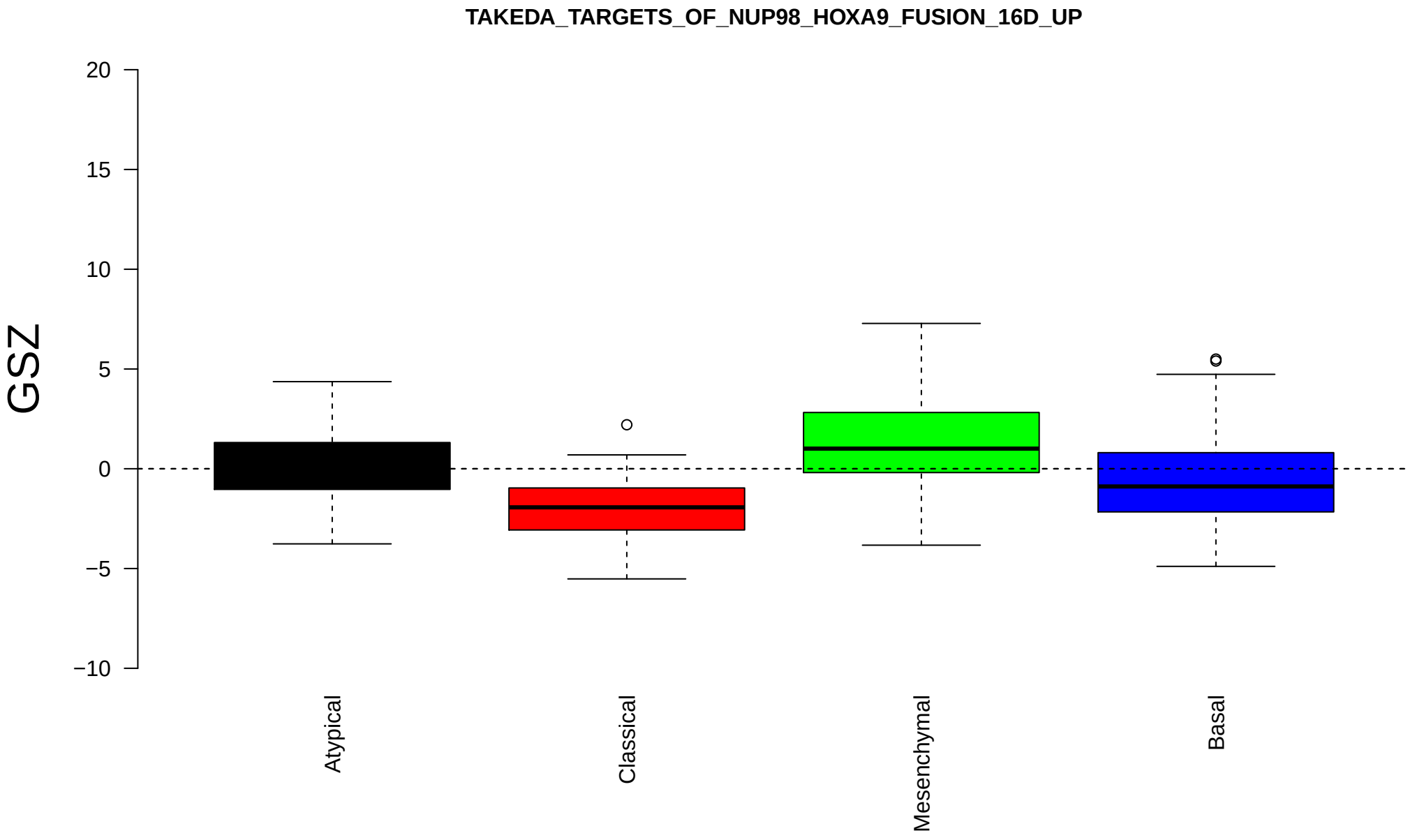
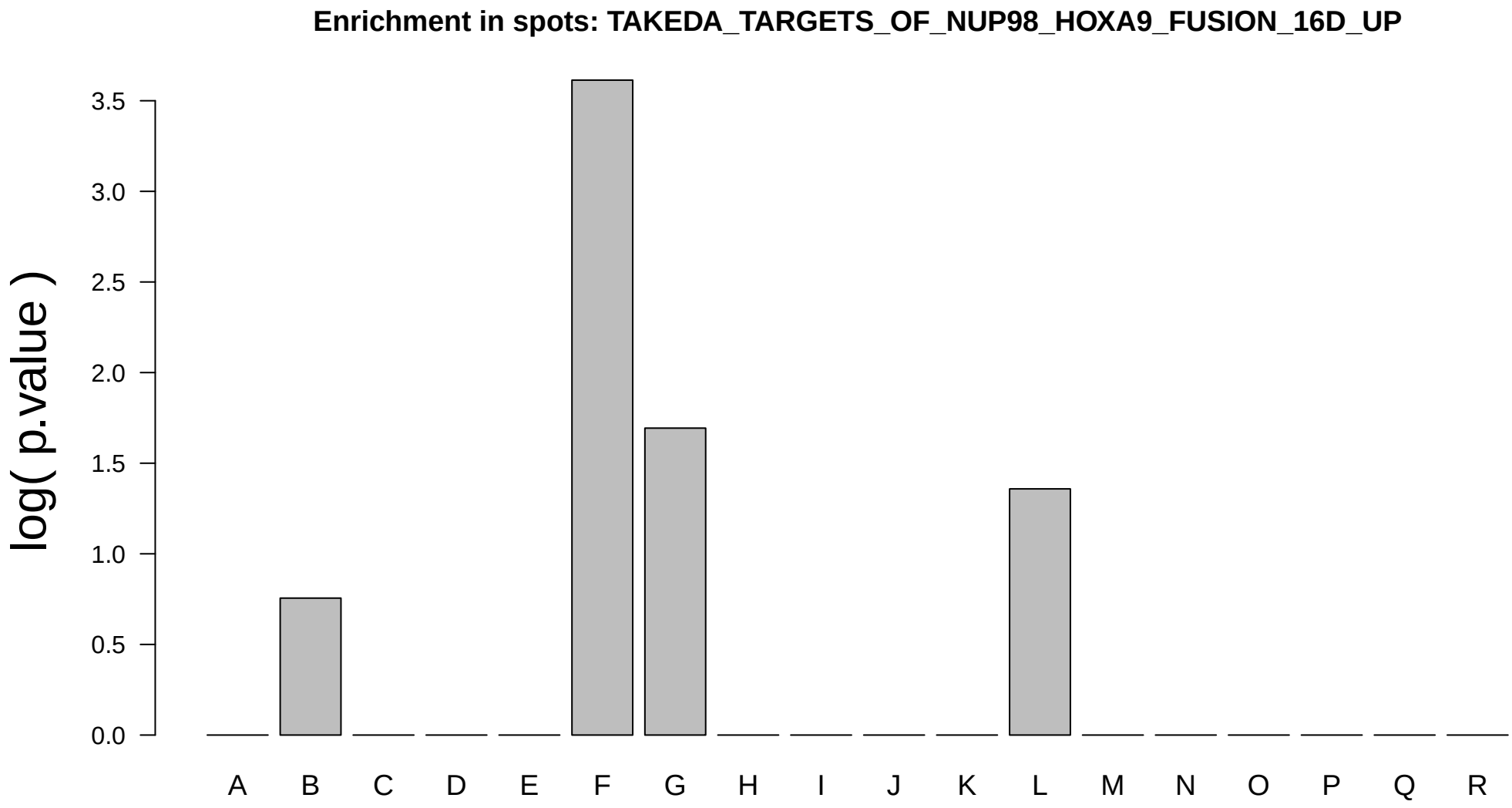
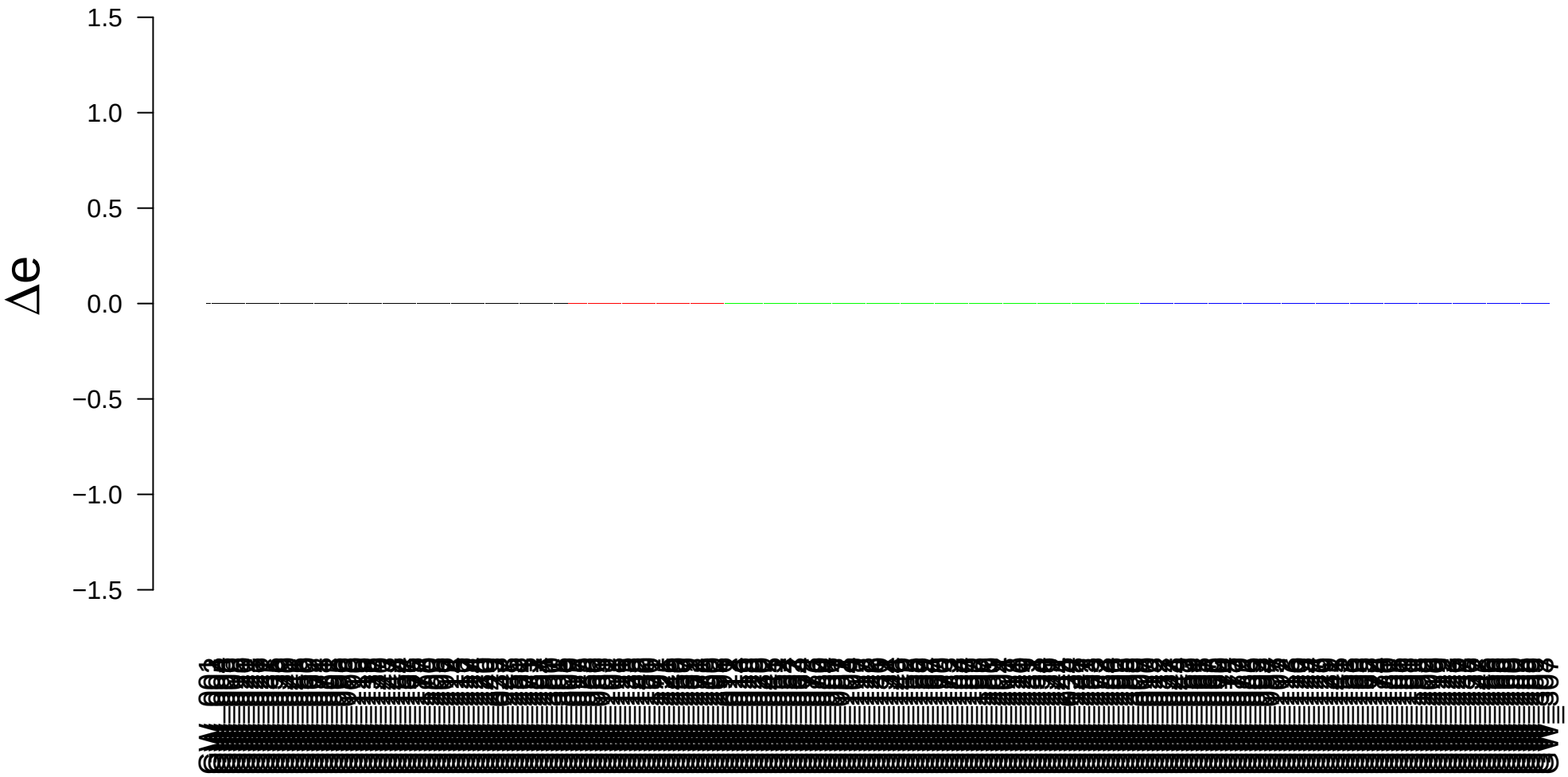
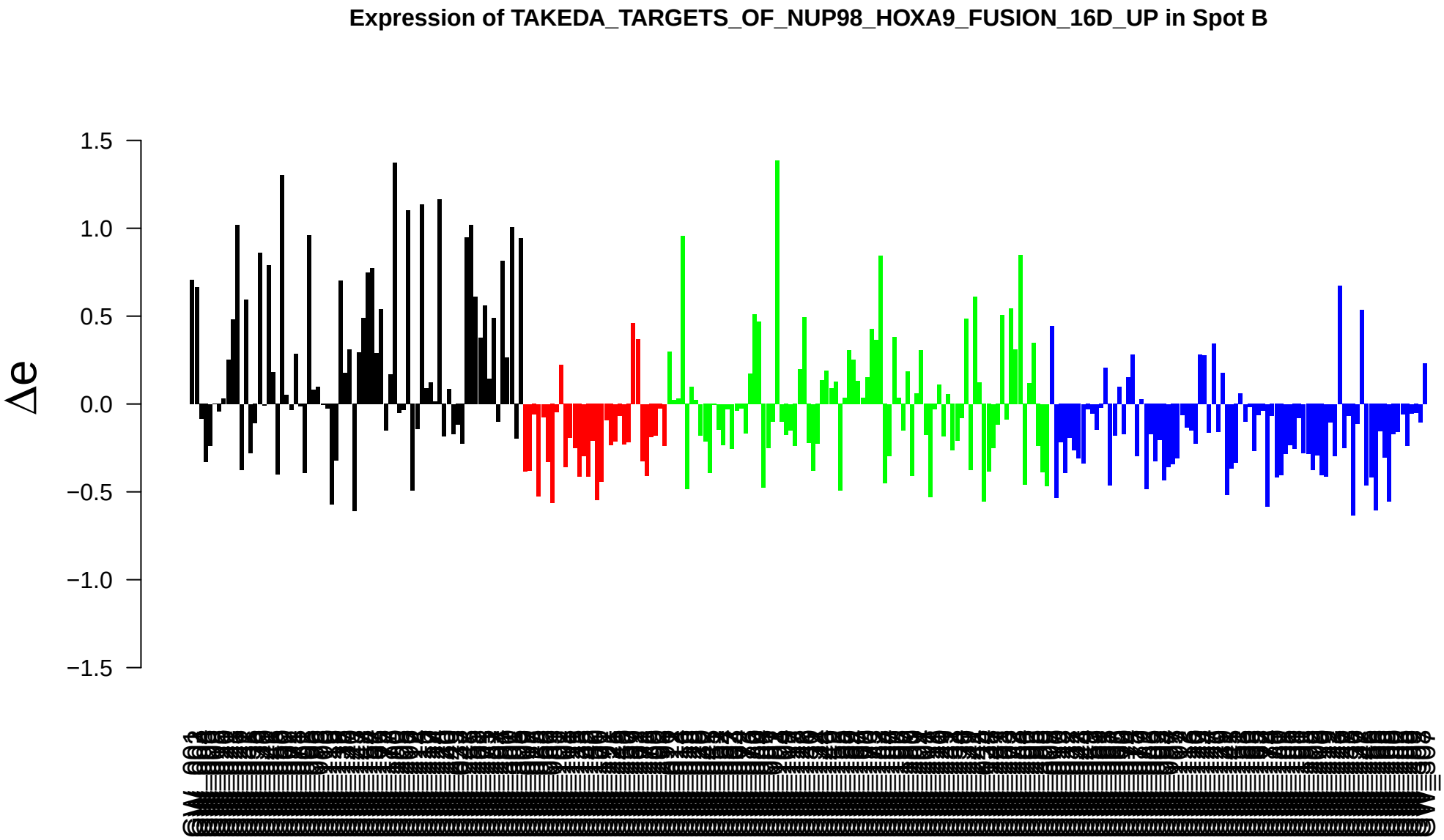

TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP
GSZ
20
15
10
5
0
−5
−10

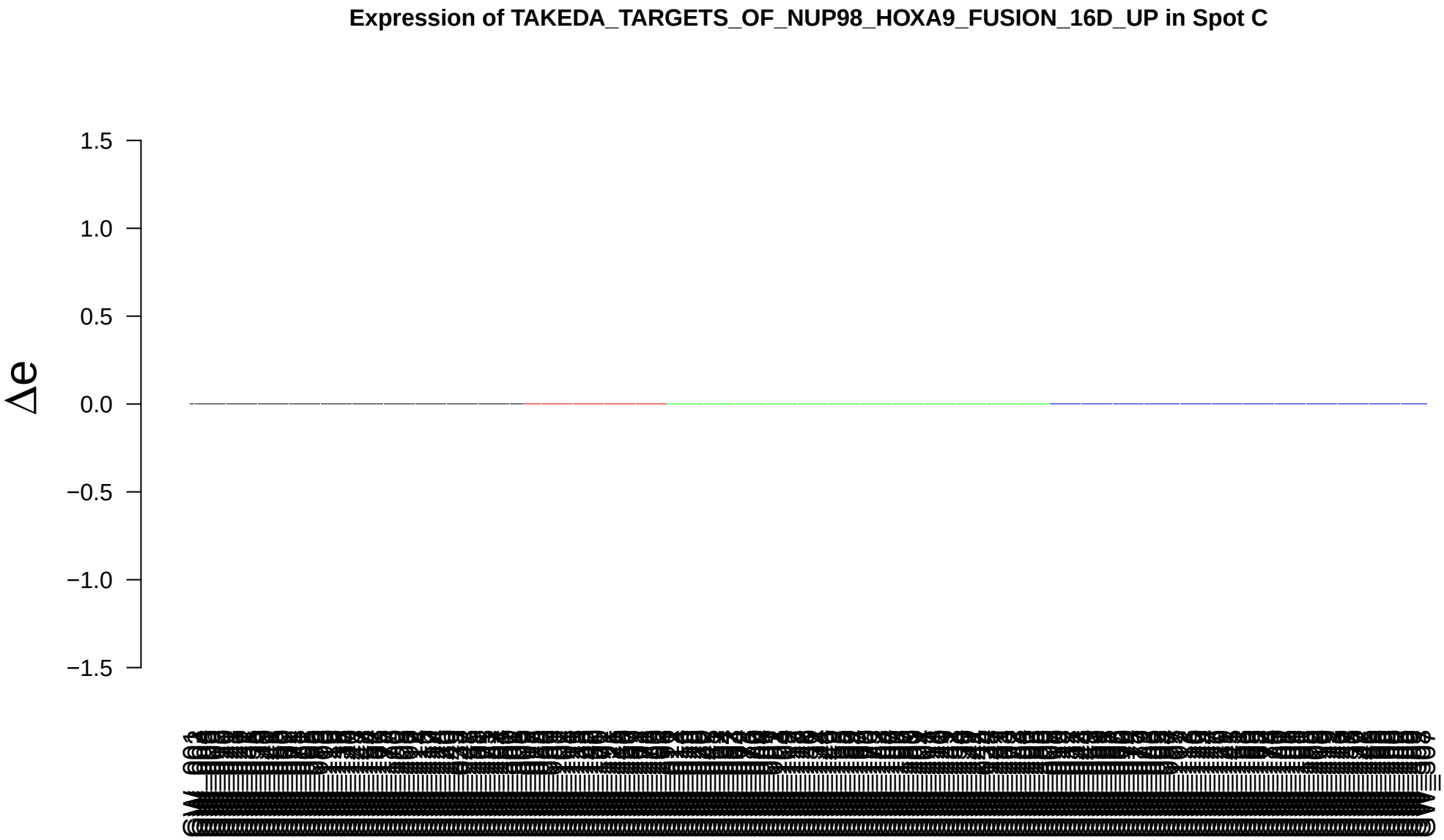

TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal


Enrichment in spots: TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP
log(p.value)
3.5
3.0
2.5
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

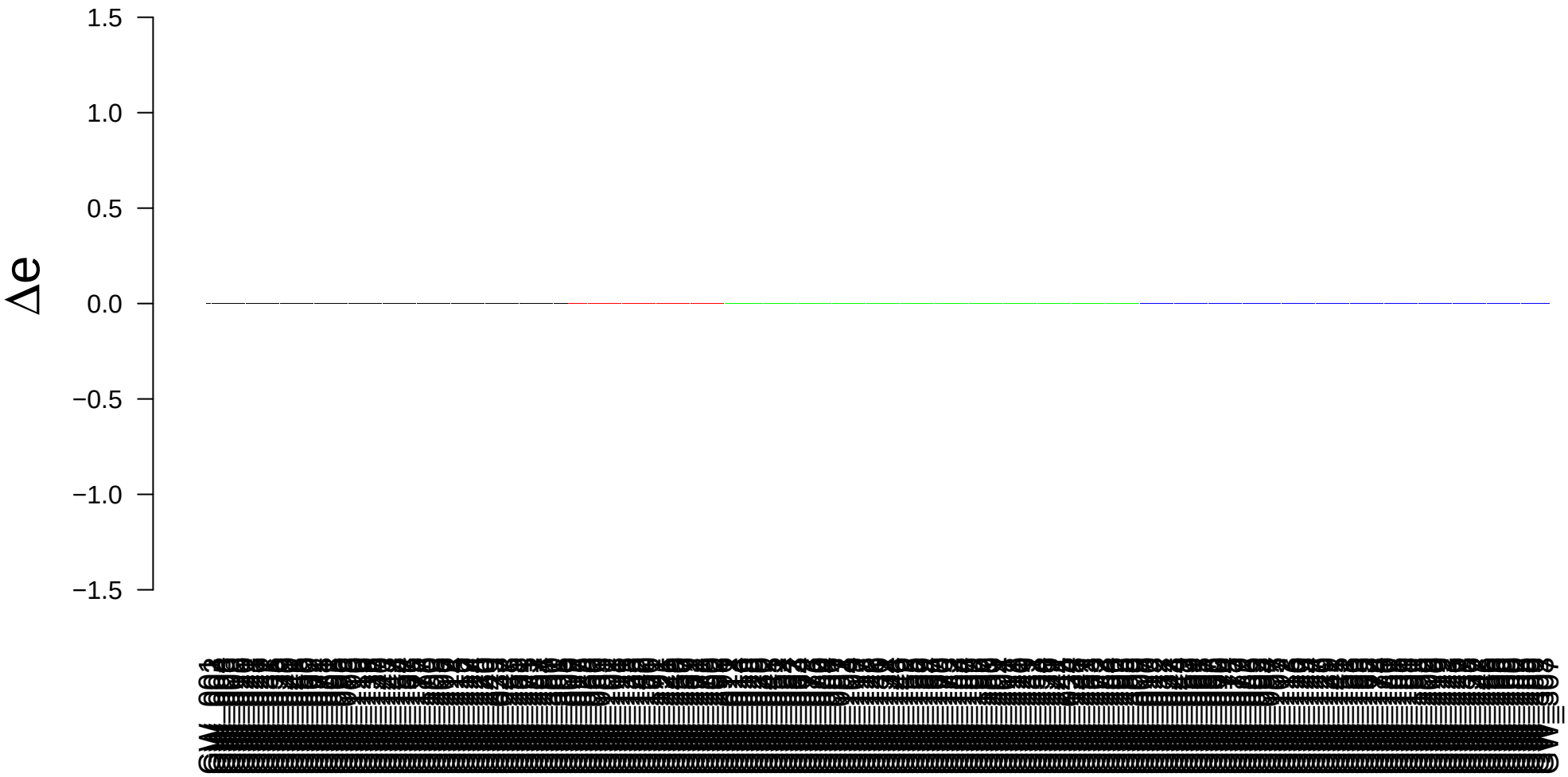
Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot A



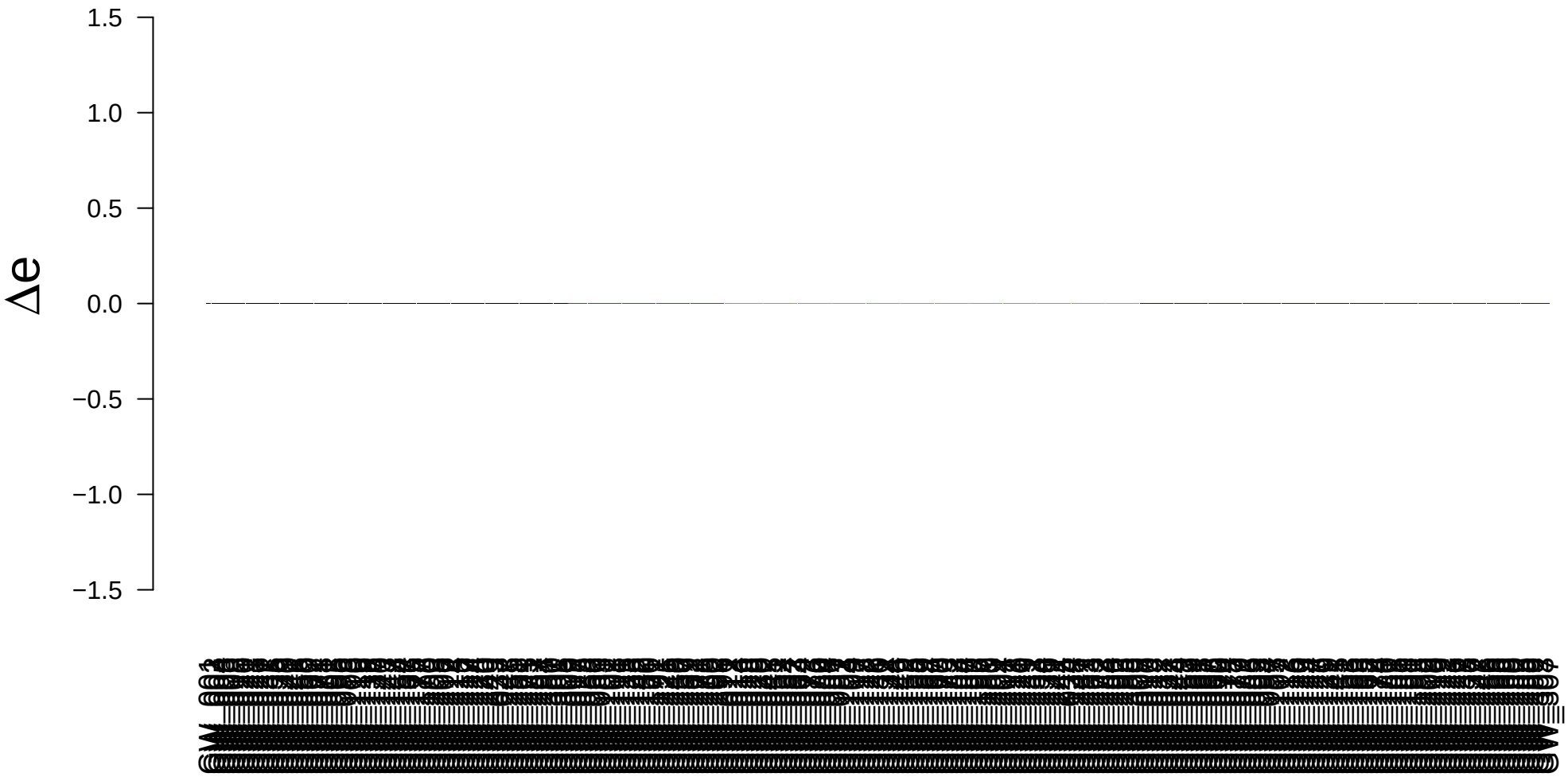

Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot B
Δe
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5

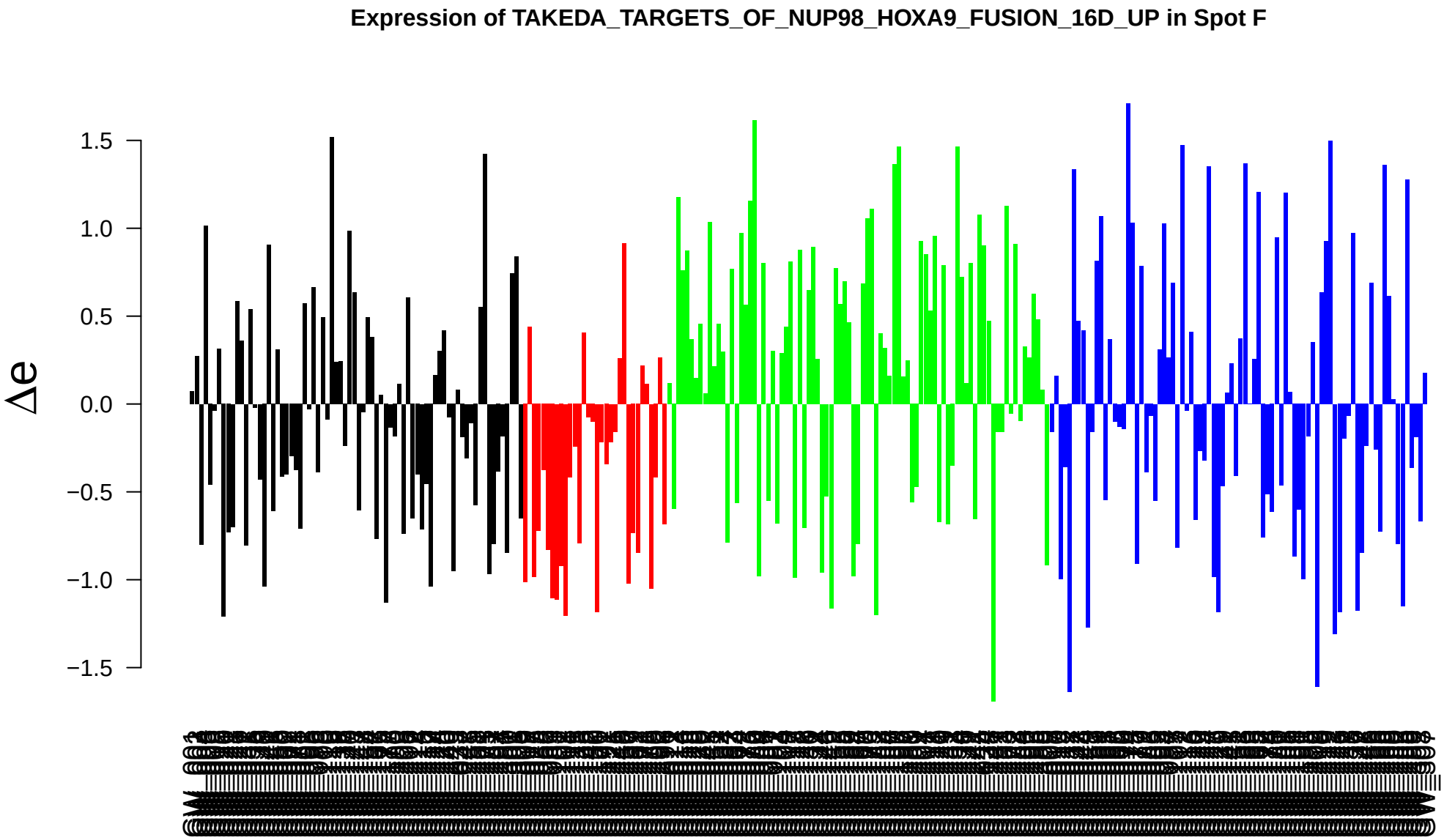

Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot C
Δe
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5

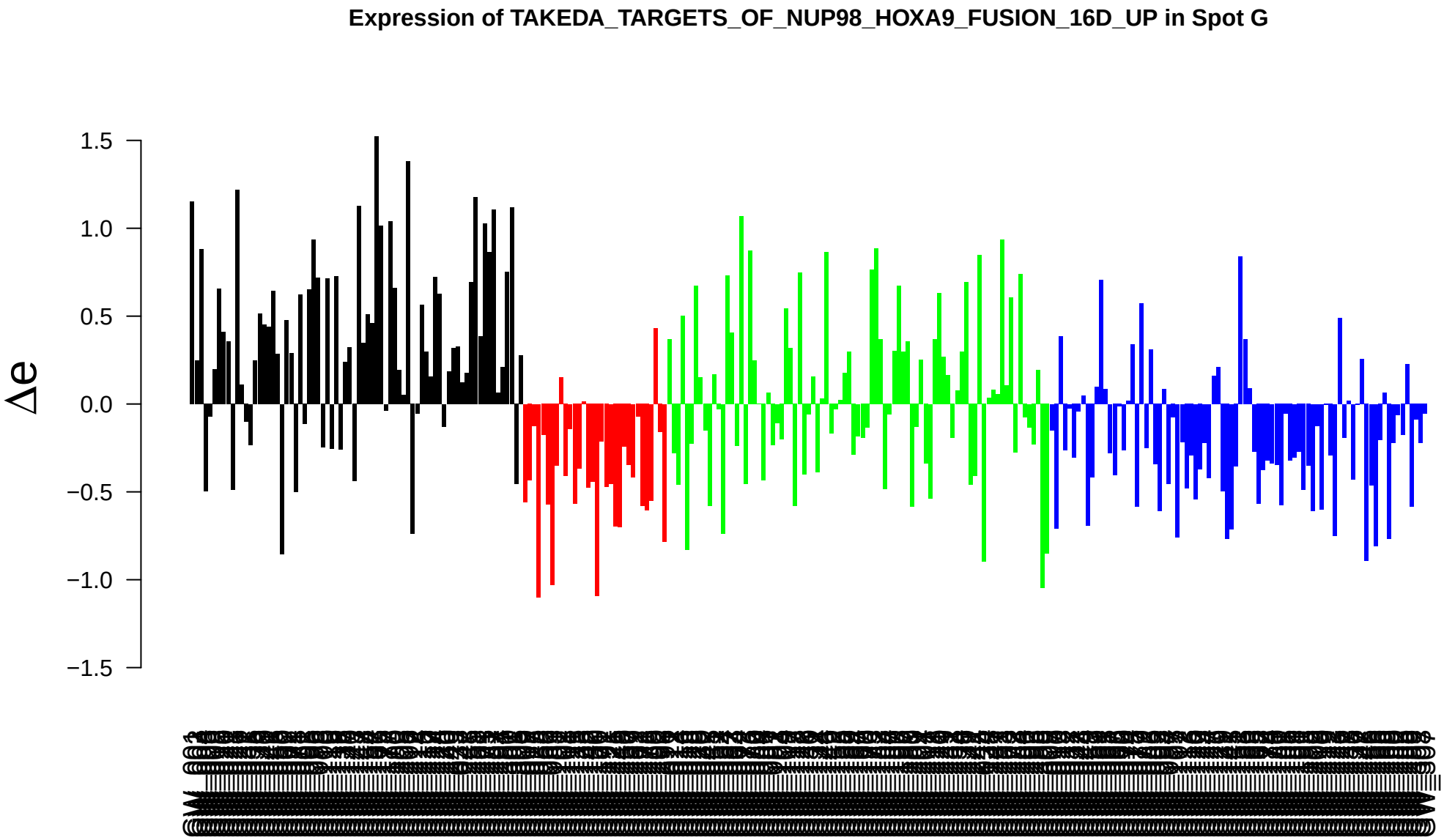
Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot D



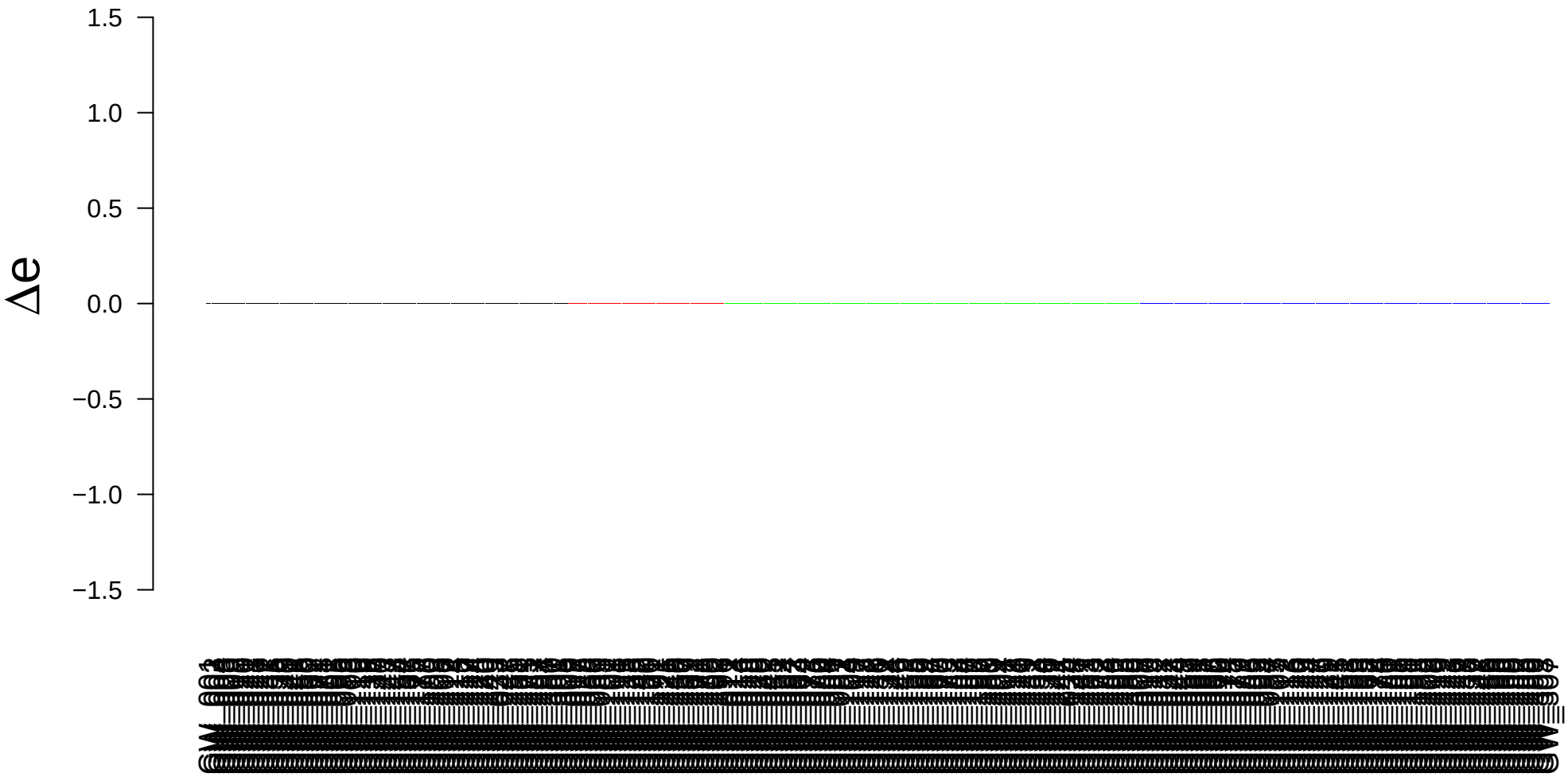
Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot E



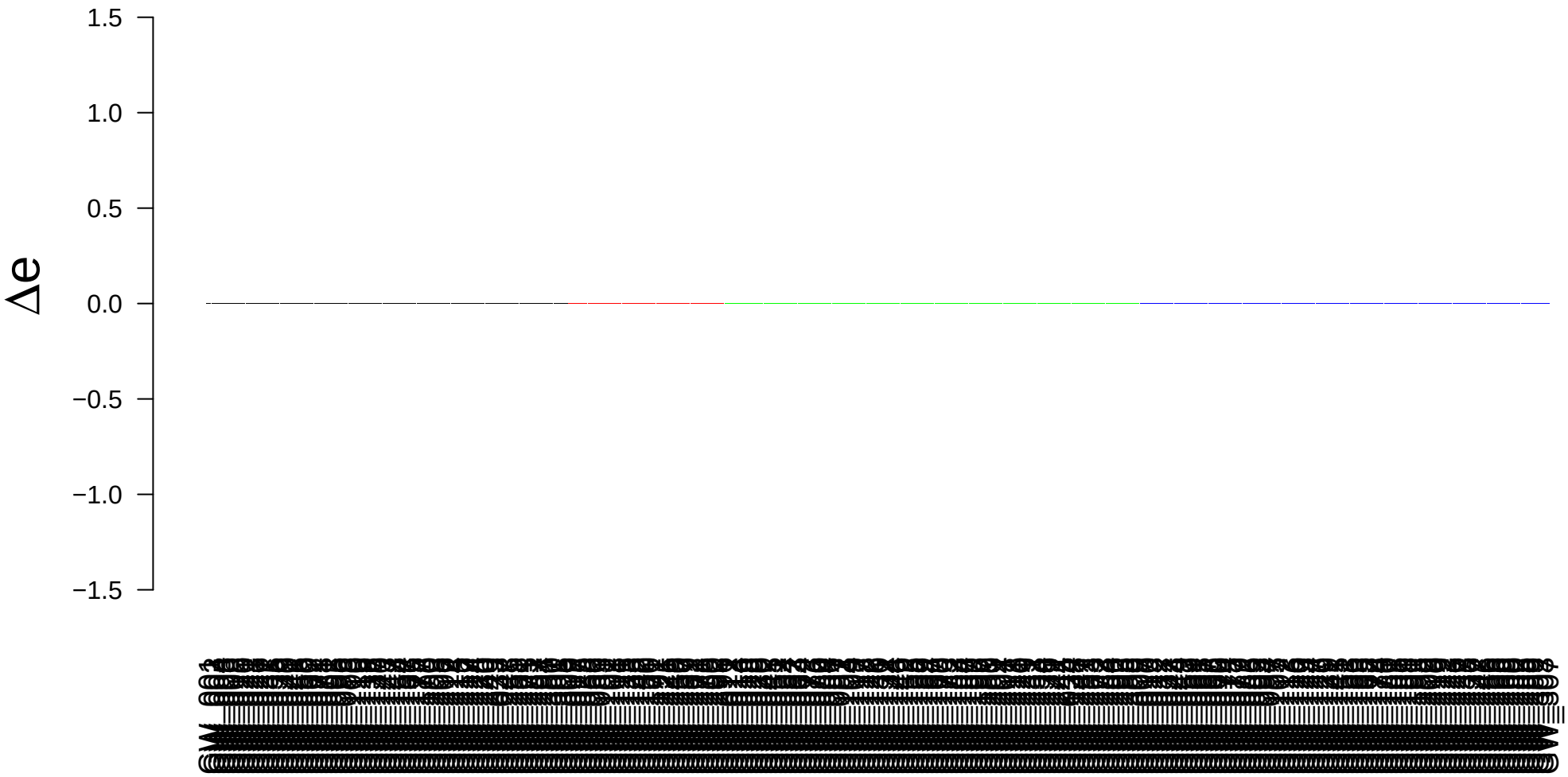

Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot F
Δe
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5


Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot G
Δe
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5

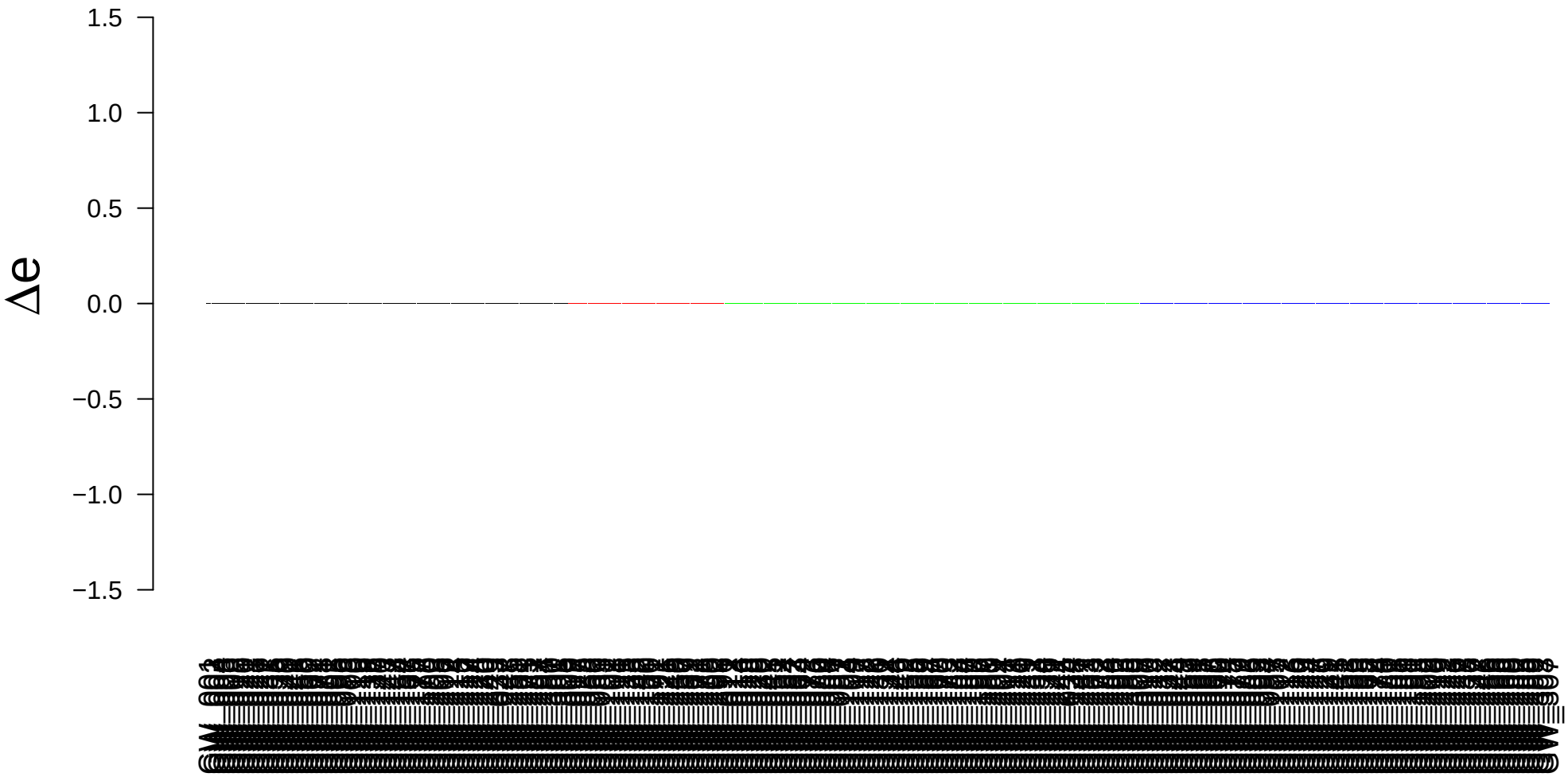
Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot H



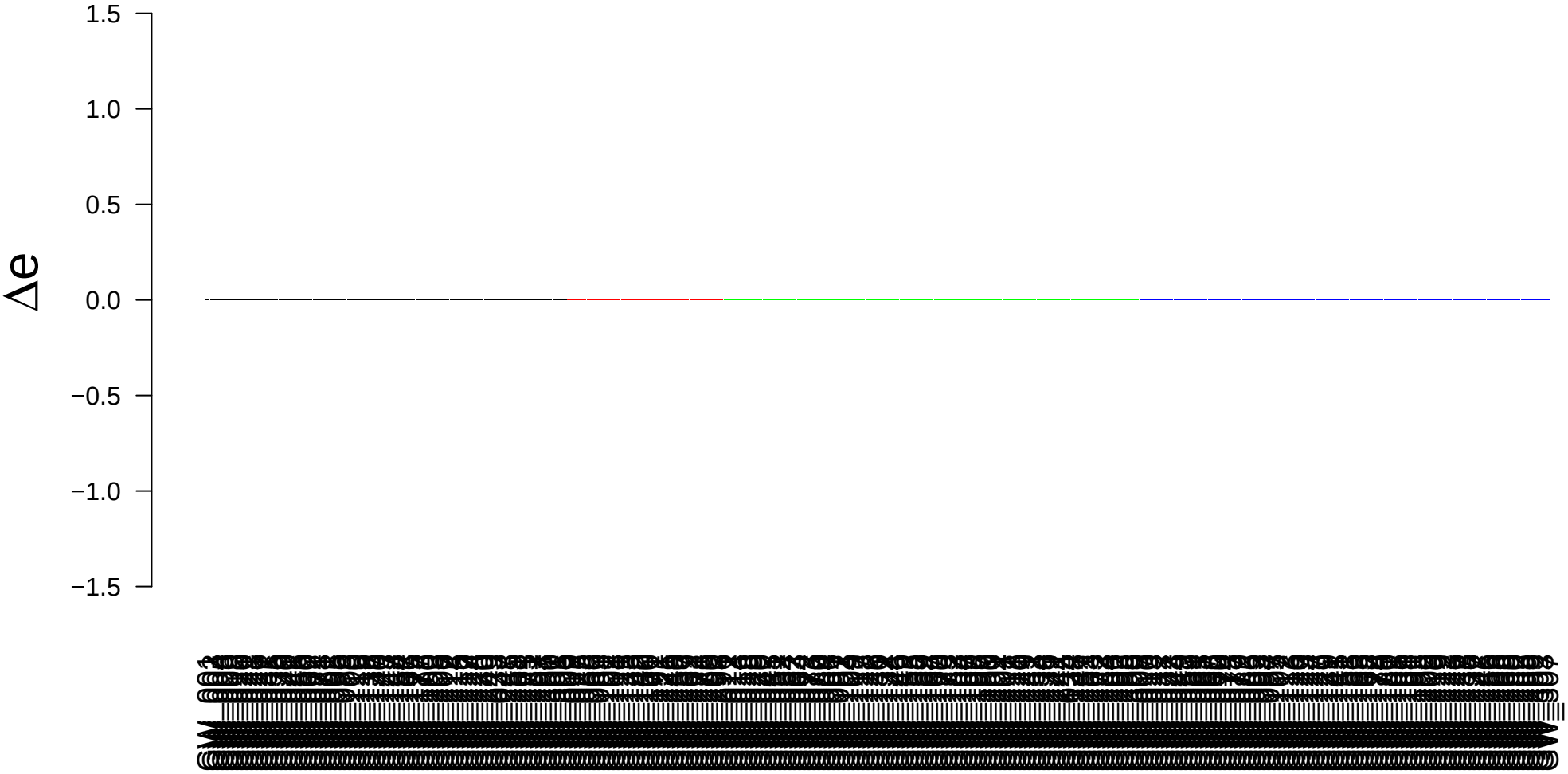
Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot I

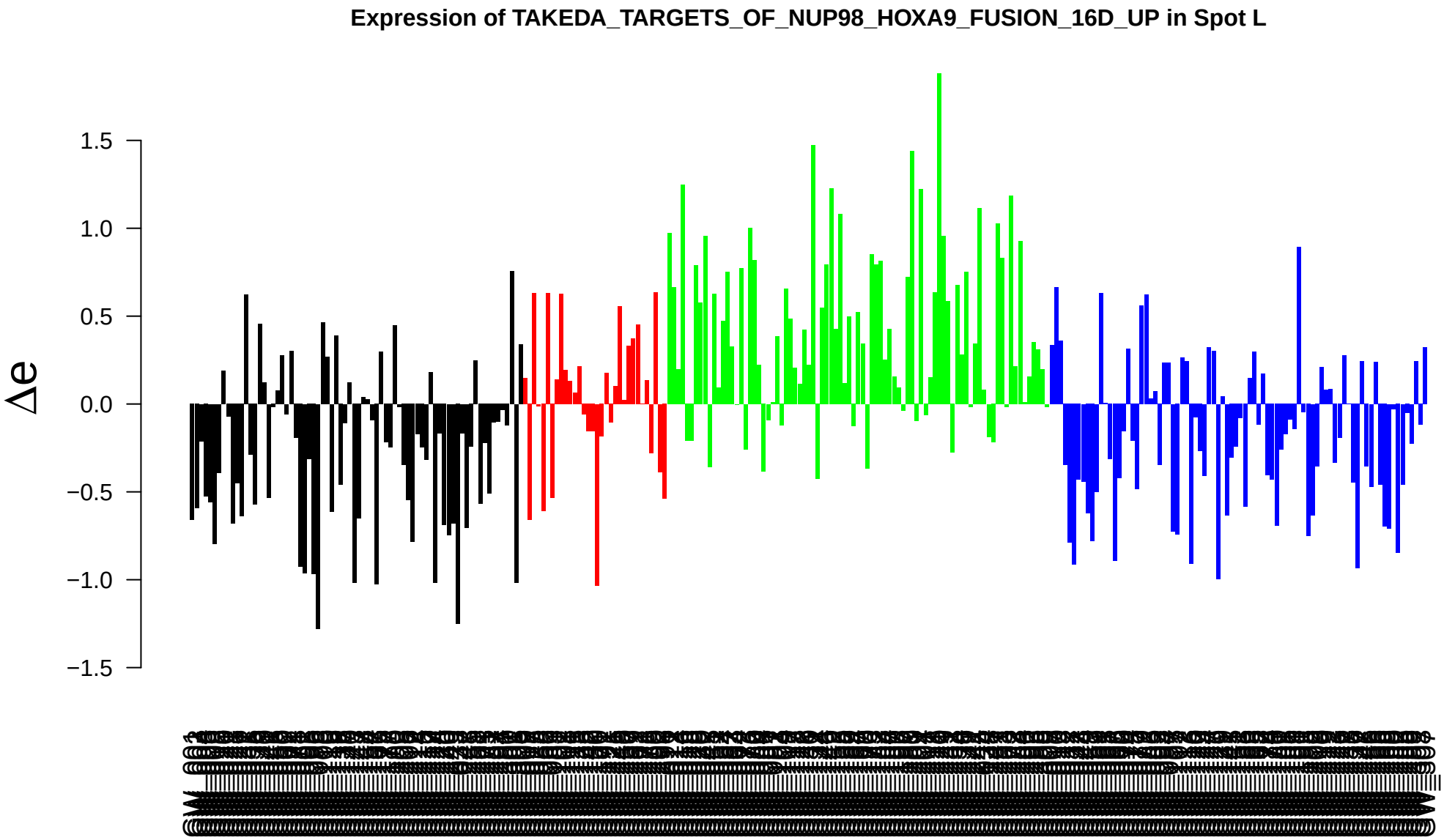


Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot J

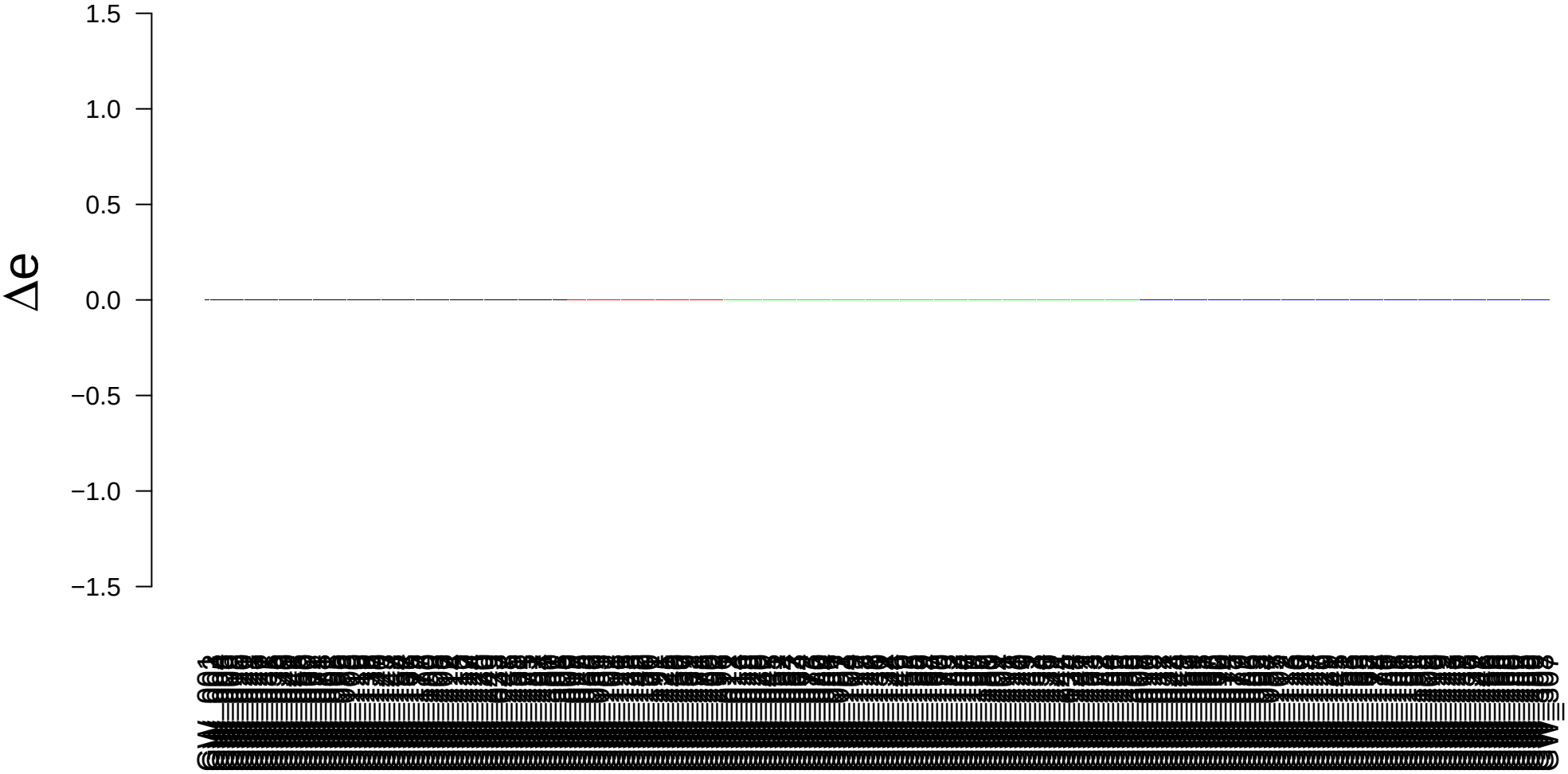


Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot K

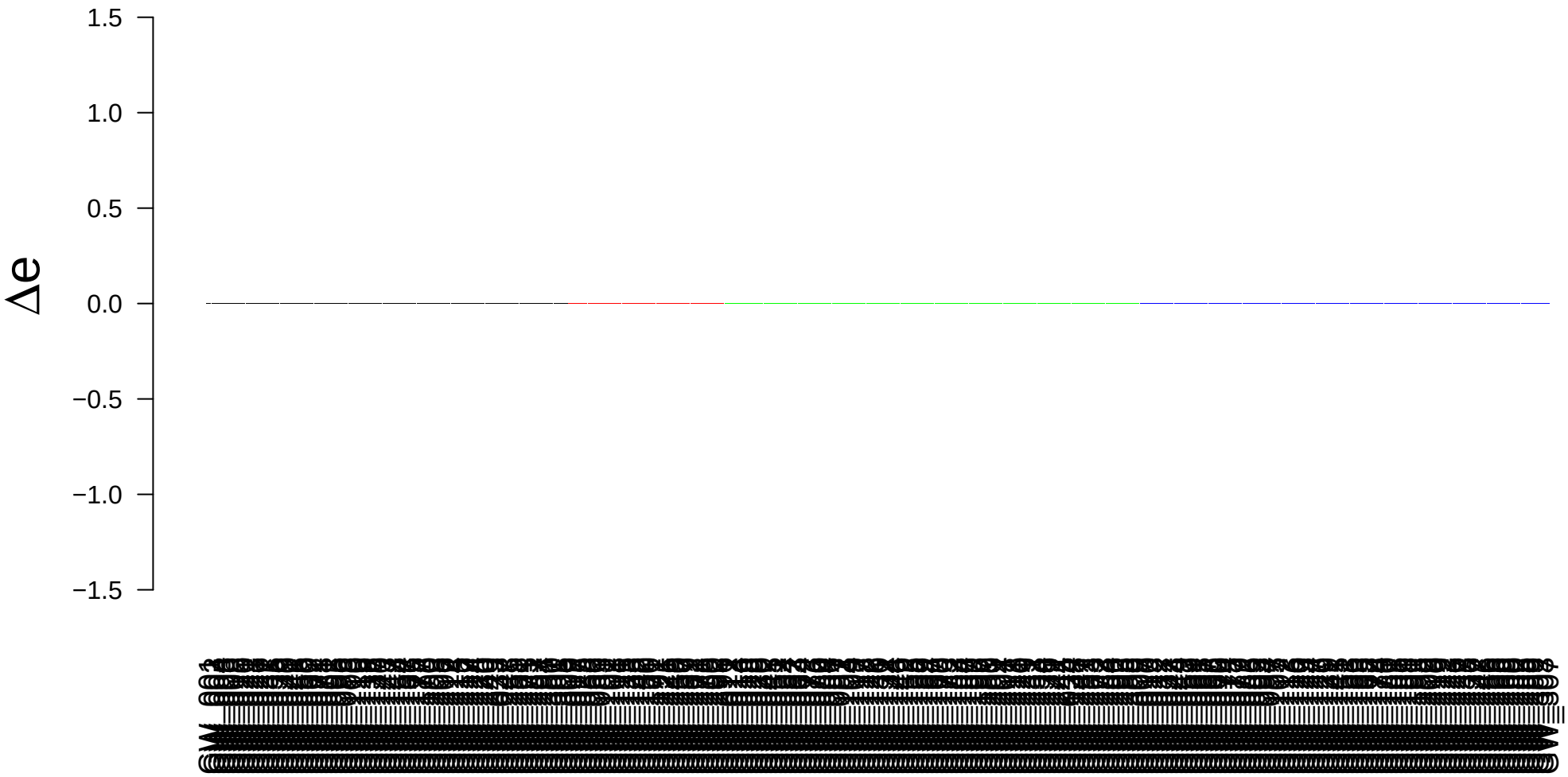



Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot L
Δe
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5

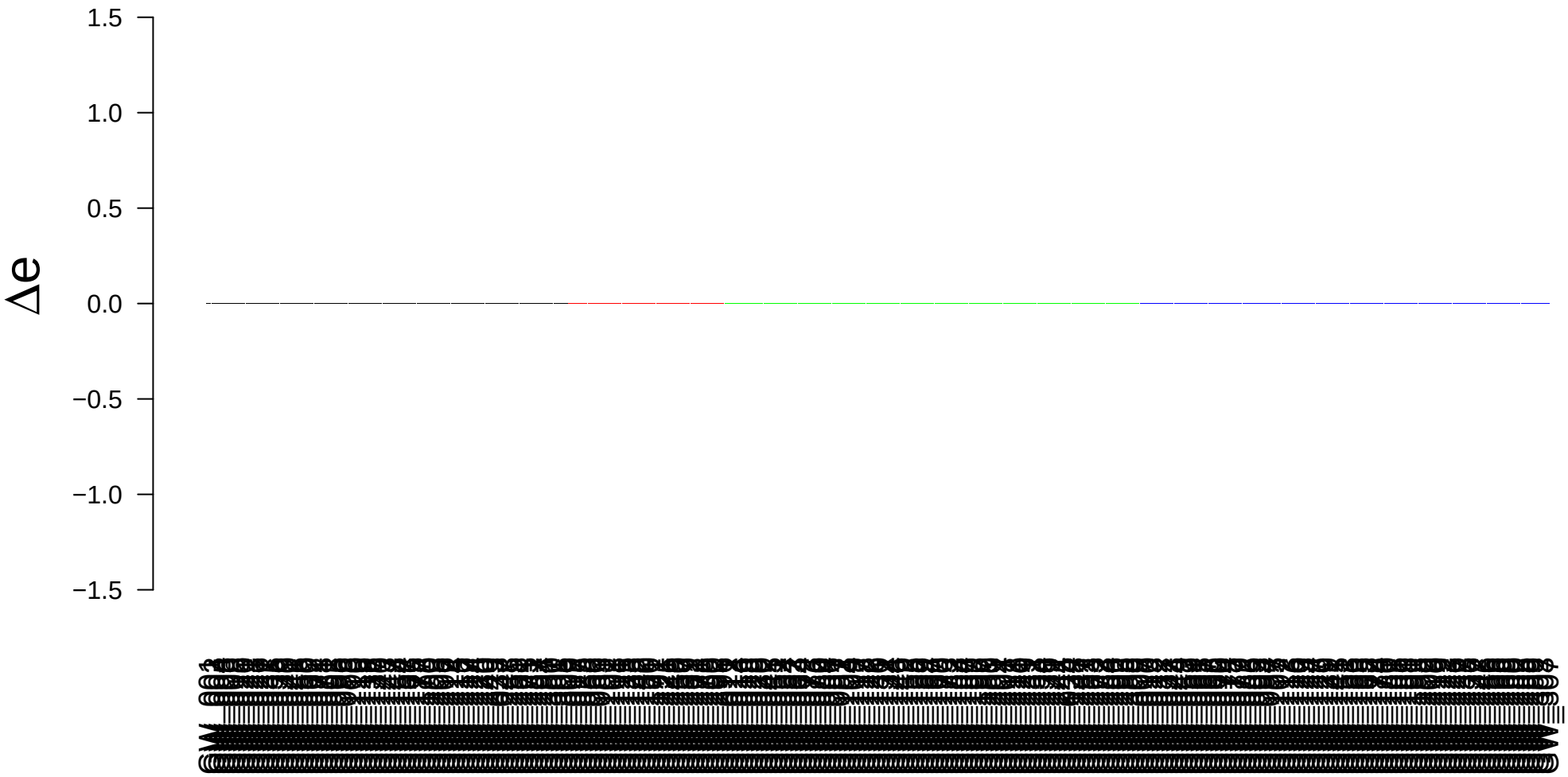
Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot M



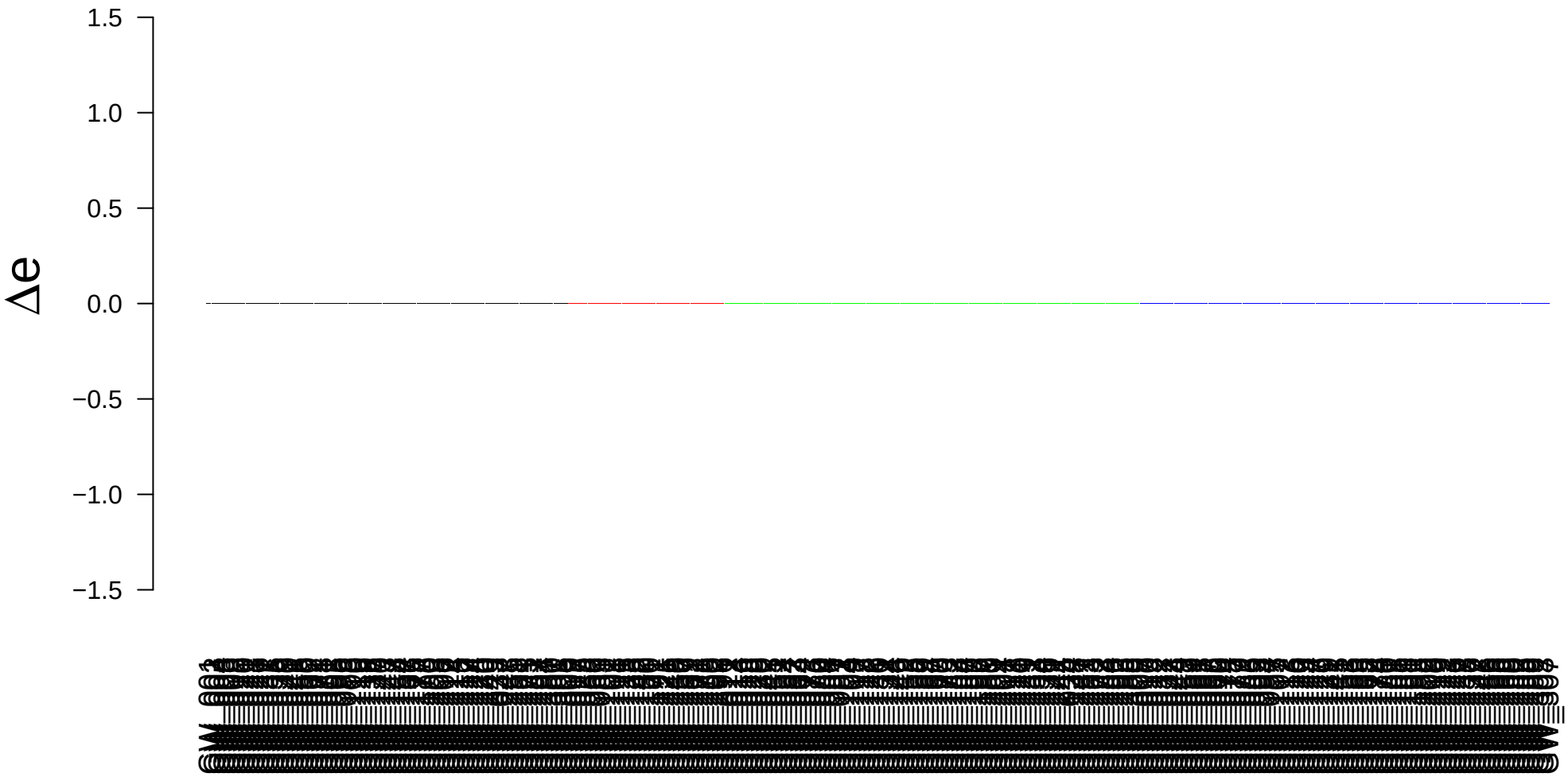
Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot N



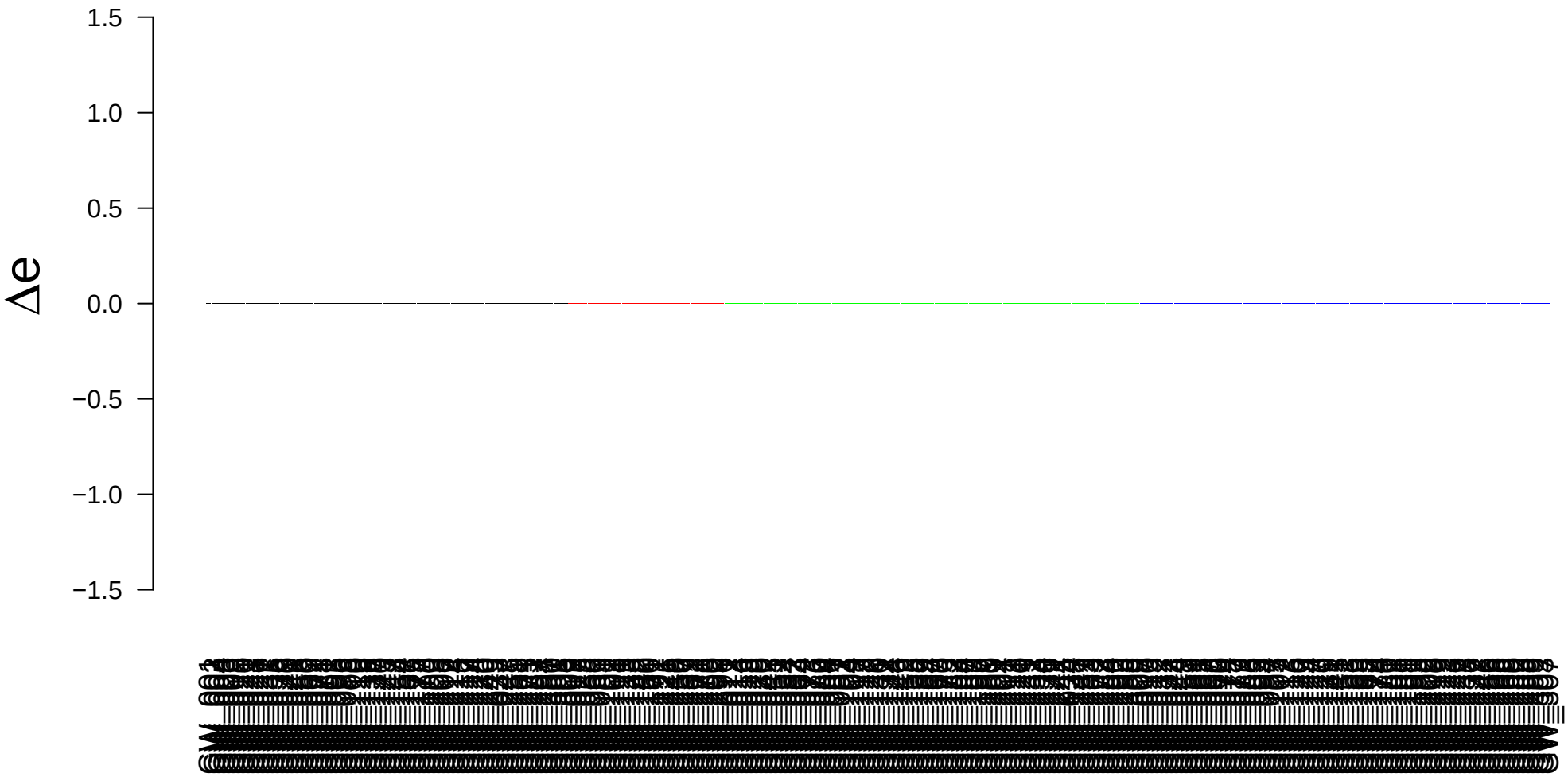
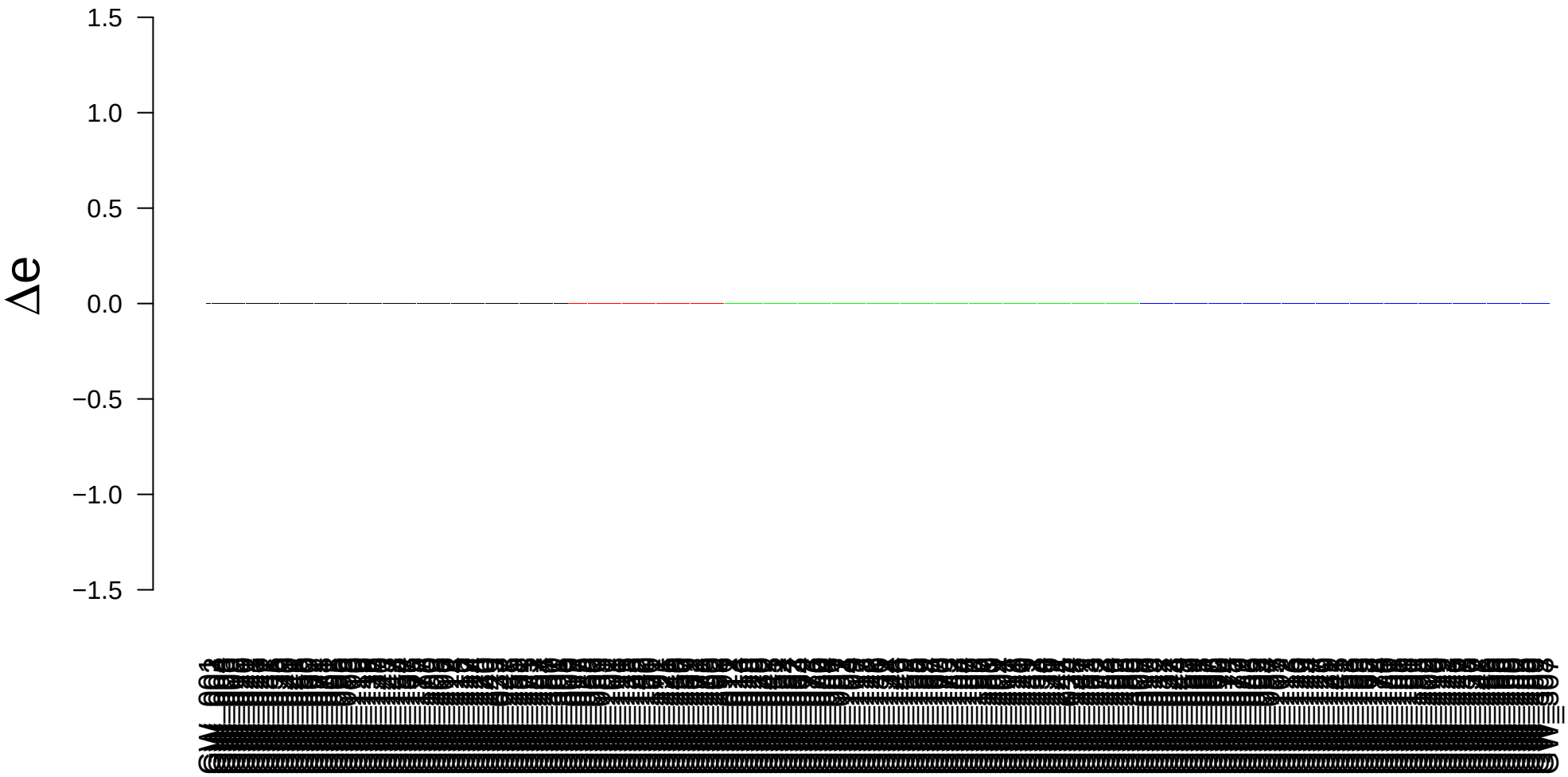
Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot O



Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot P



Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot Q



Expression of TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_16D_UP in Spot R