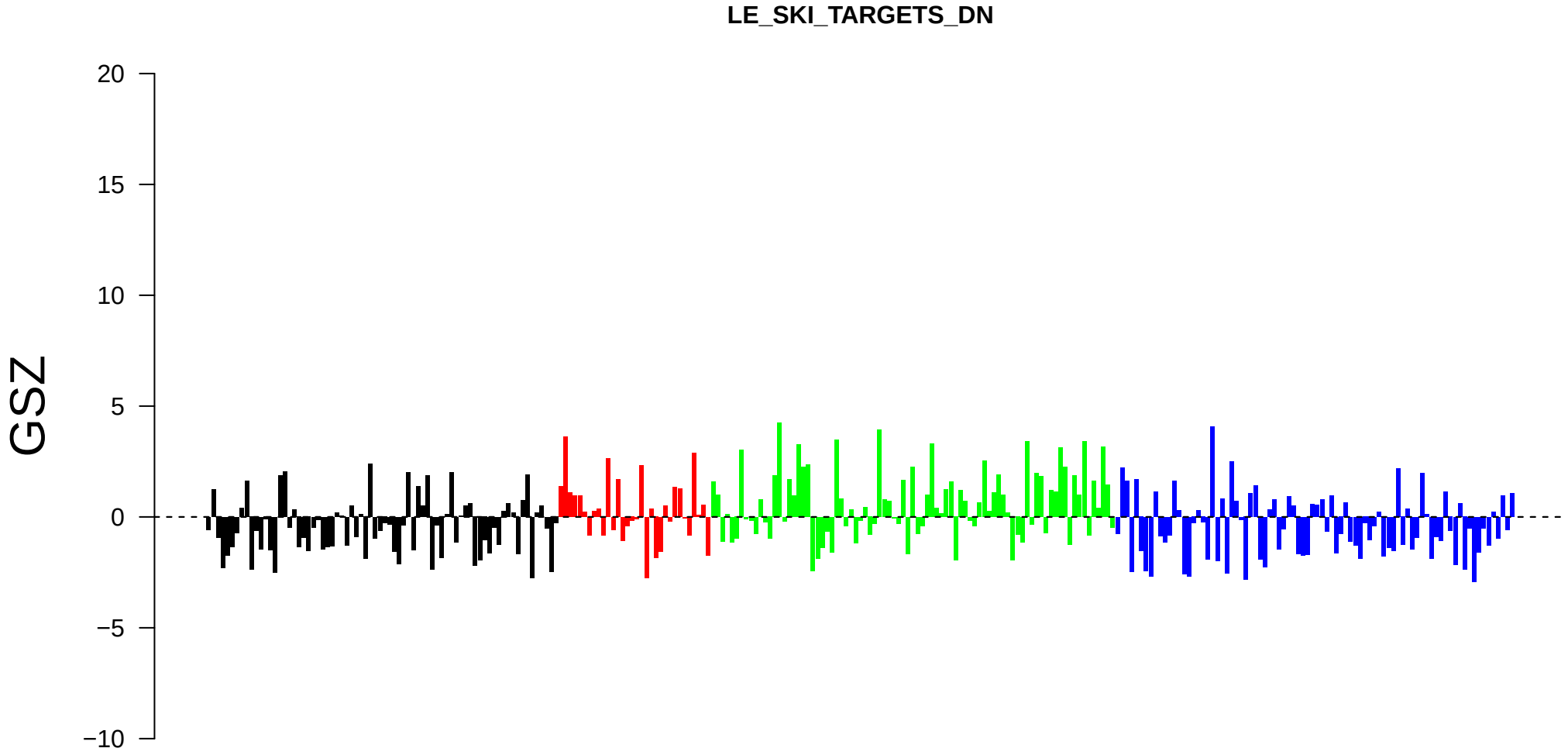
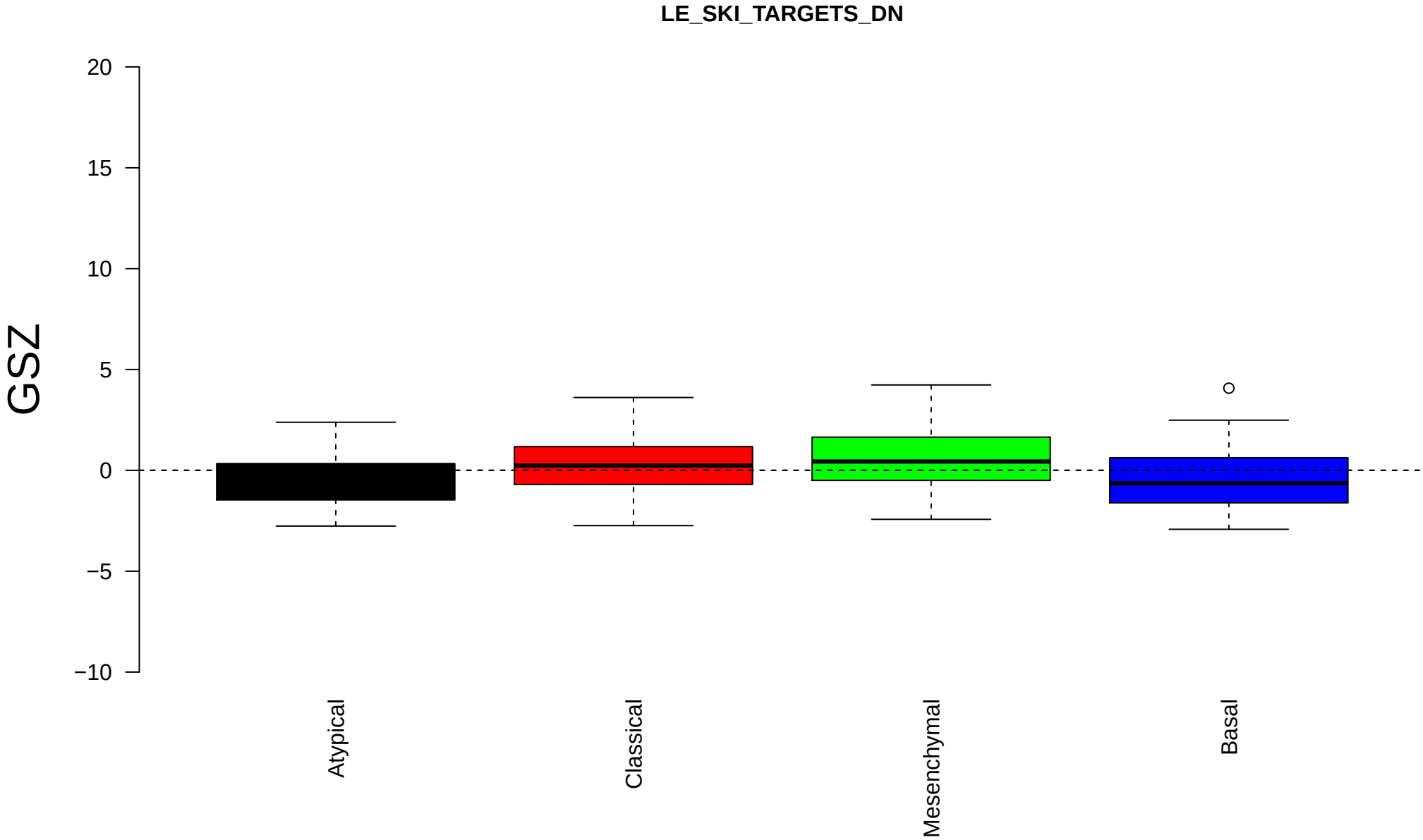
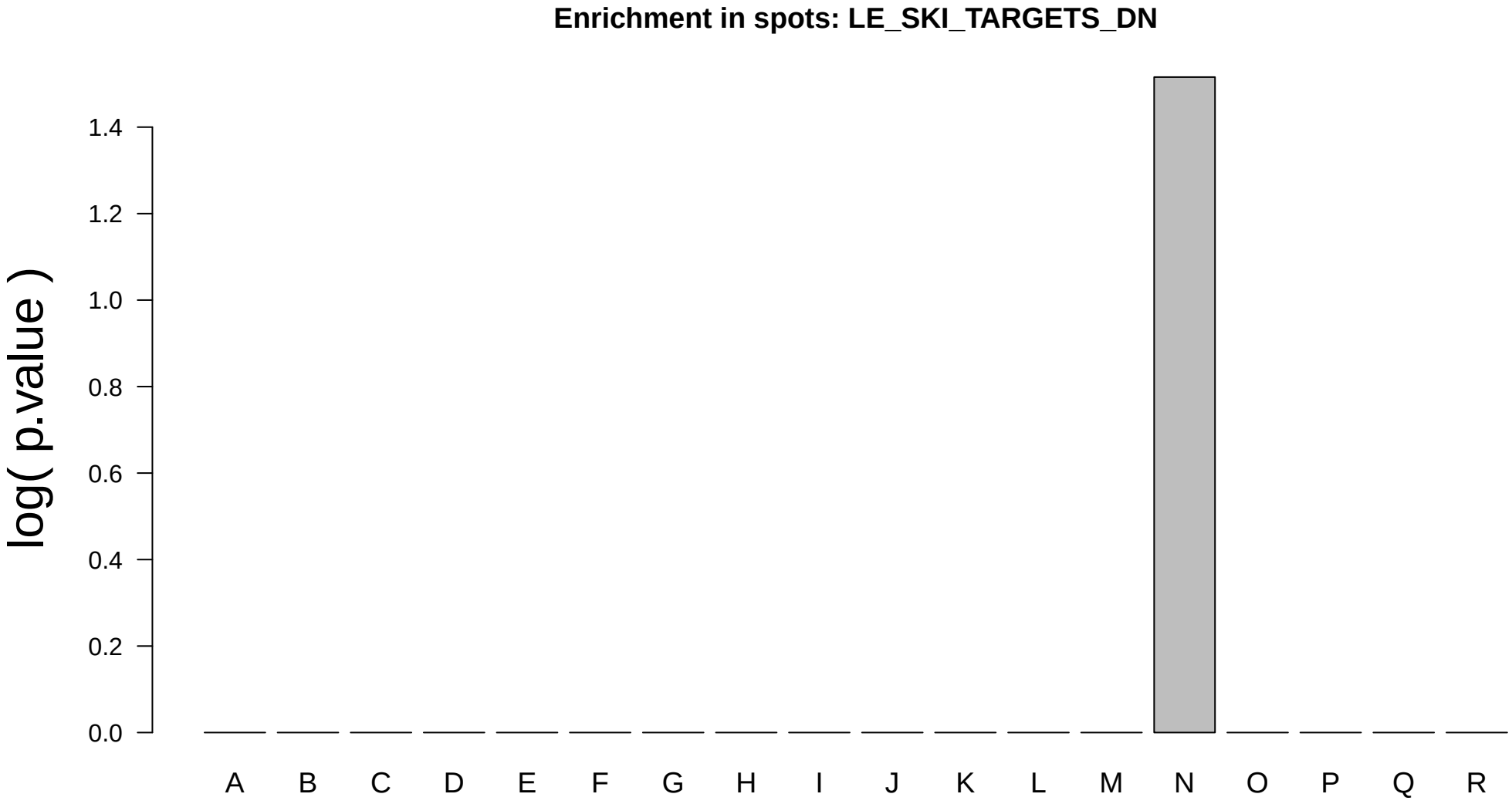
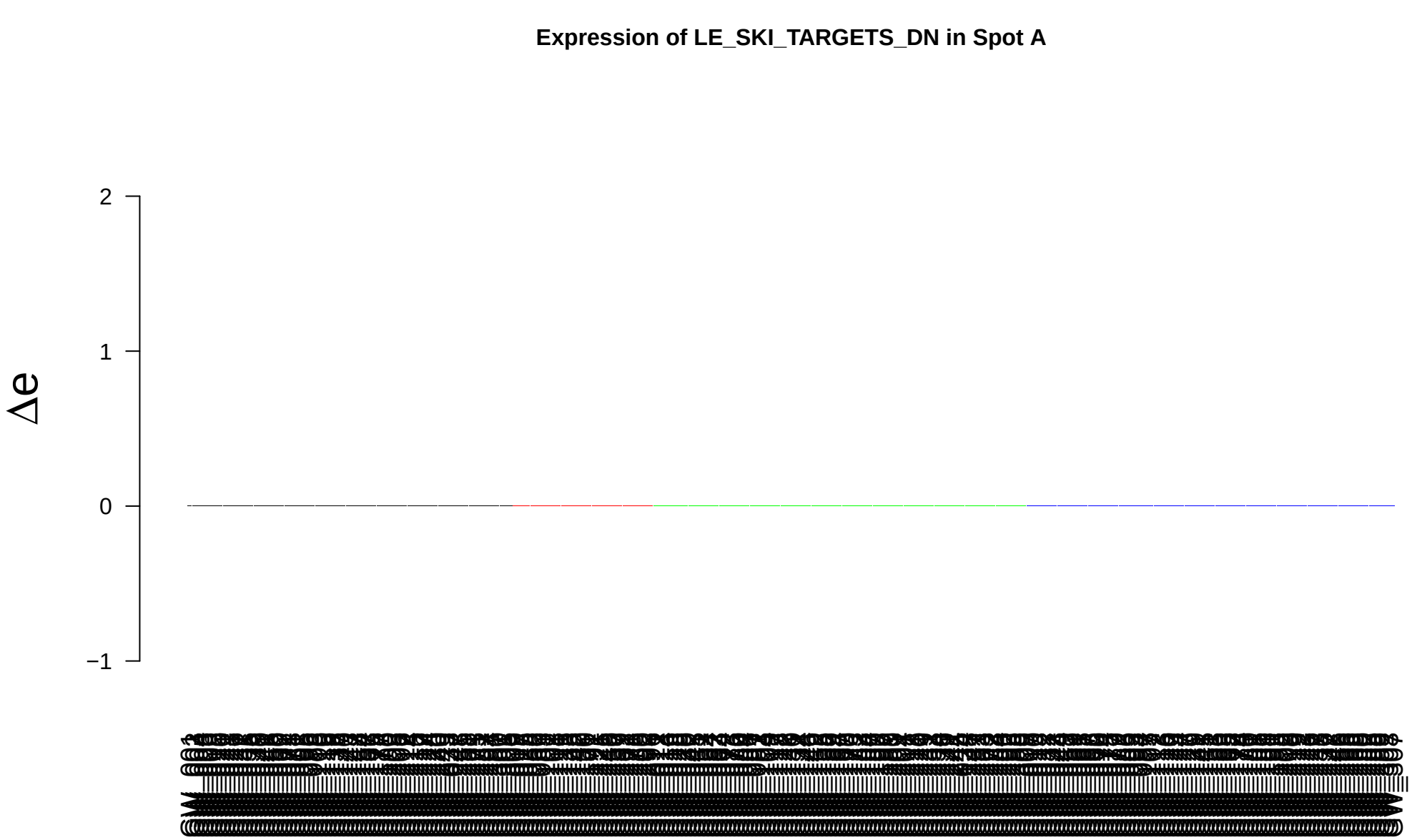
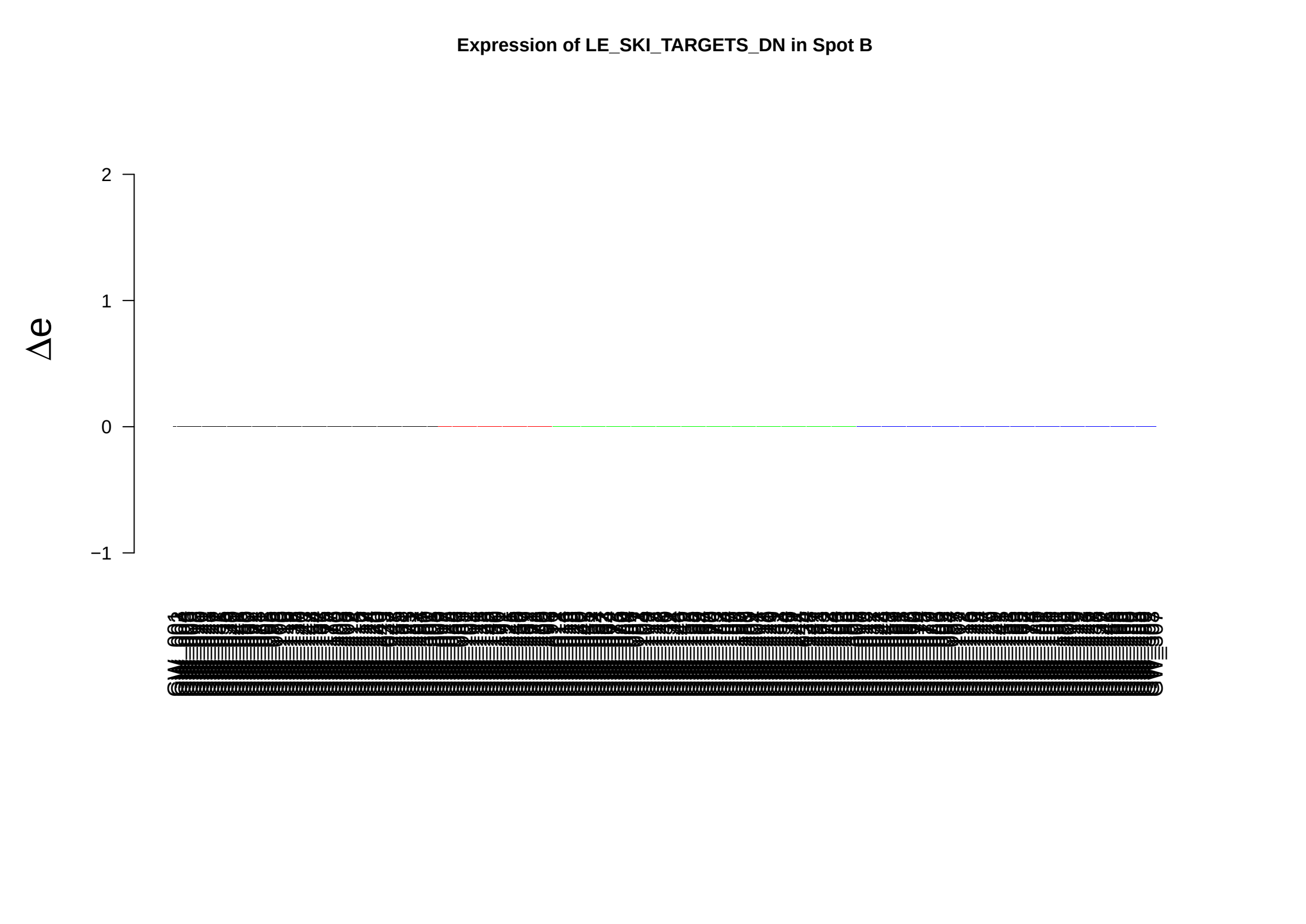
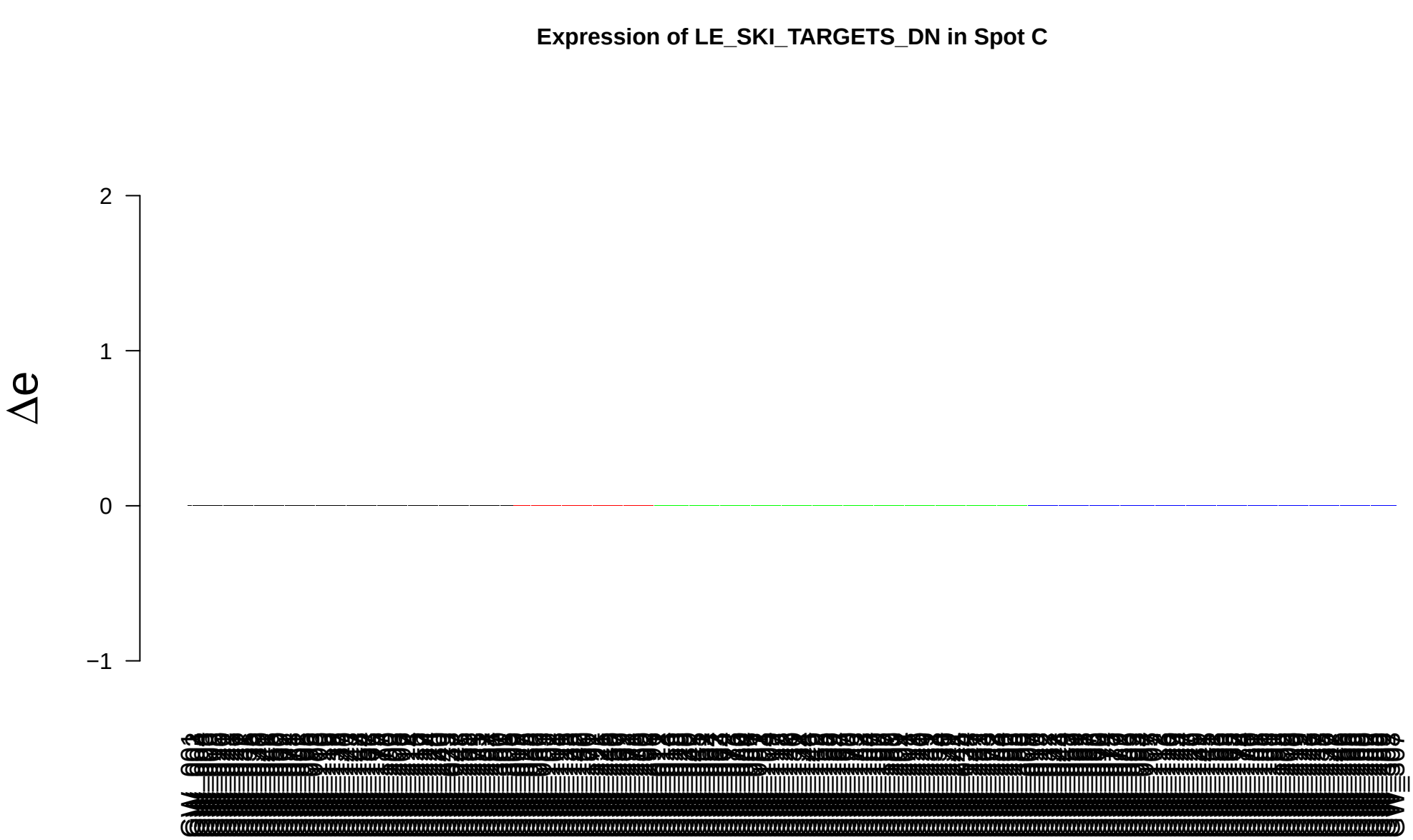

LE_SKI_TARGETS_DN
GSZ
20
15
10
5
0
−5
−10

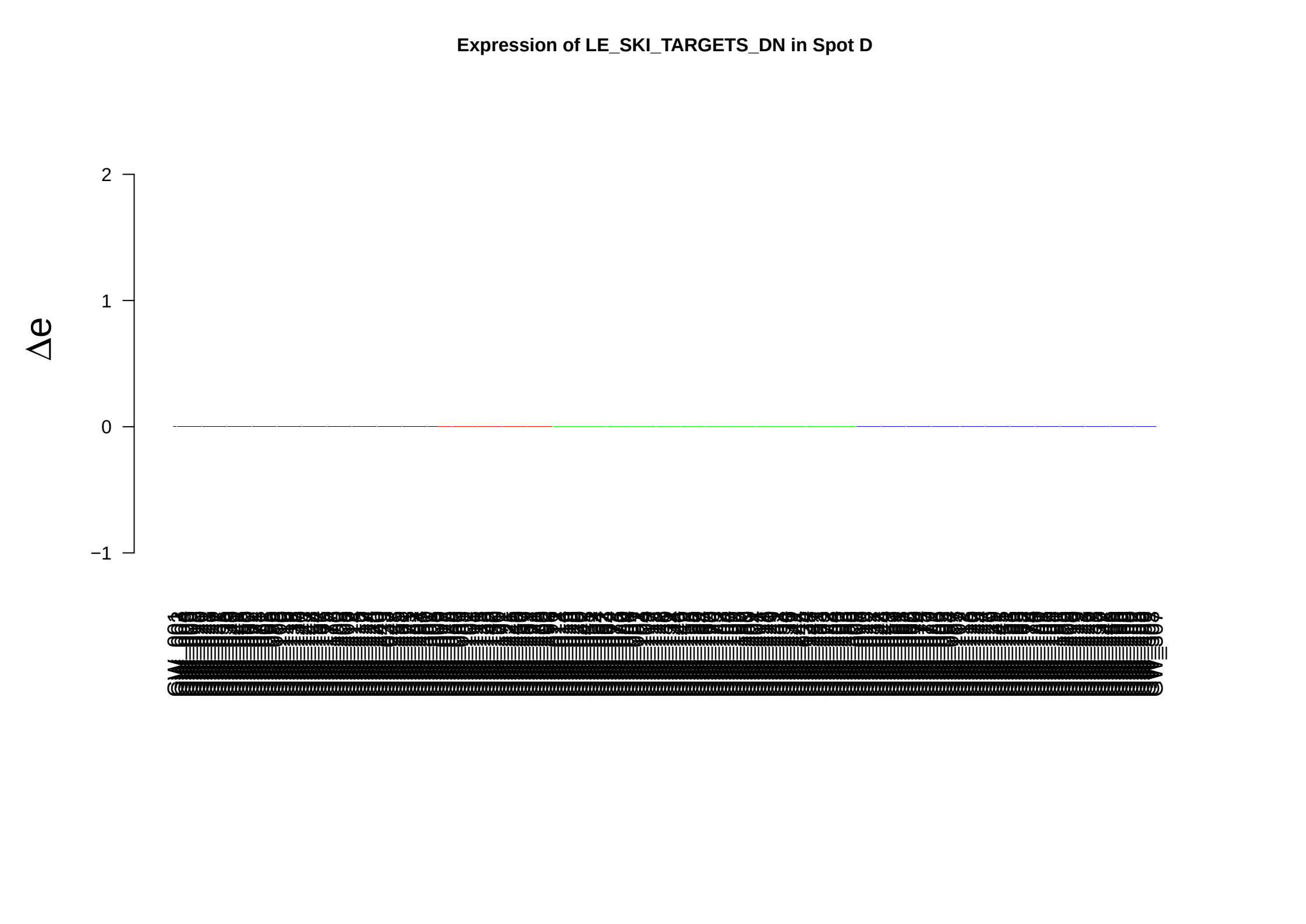

LE_SKI_TARGETS_DN
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal

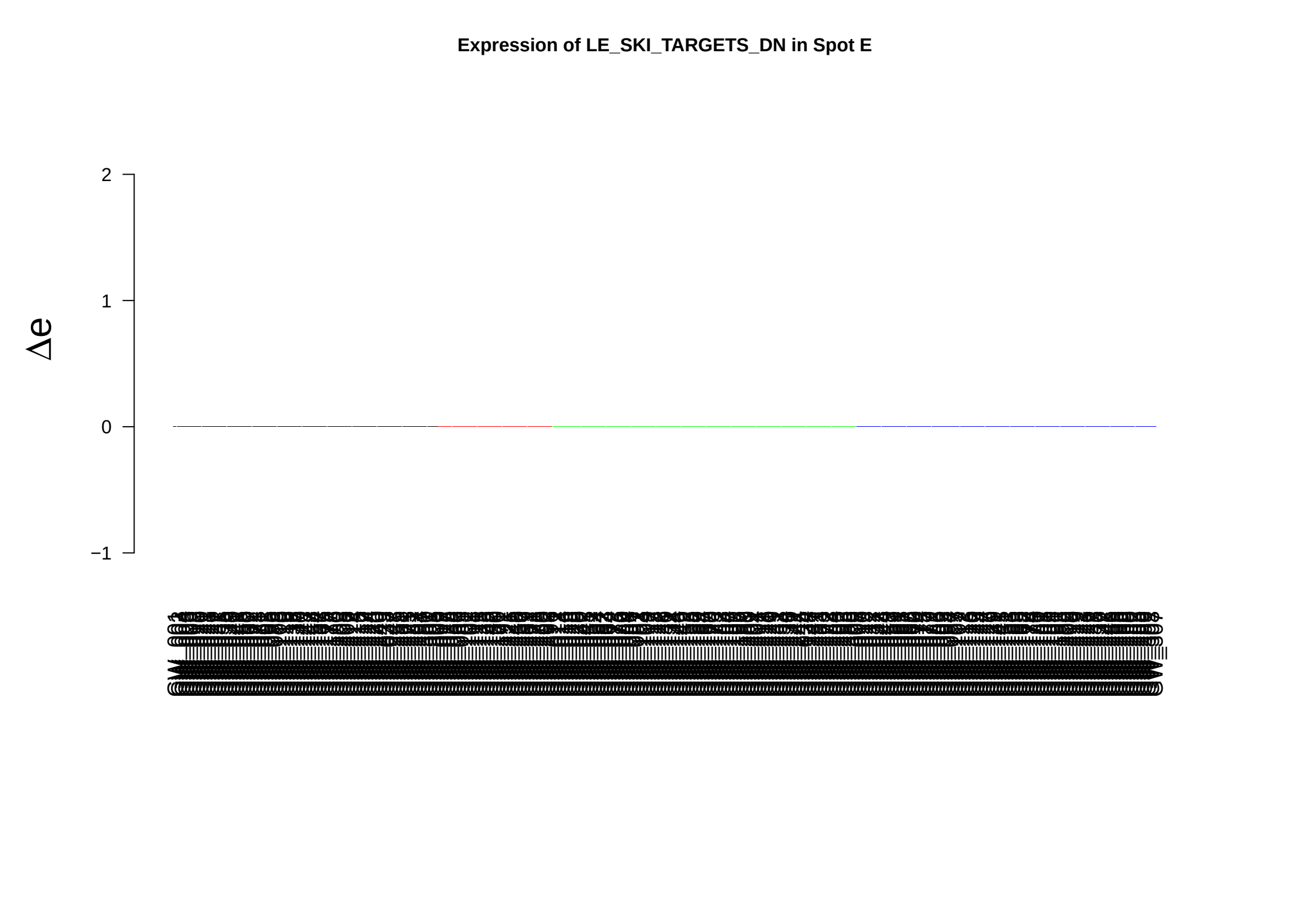

Enrichment in spots: LE_SKI_TARGETS_DN
log(p.value)
1.4
1.2
1.0
0.8
0.6
0.4
0.2
0.0
A
B
C
D
E
F
G
H
I
J
K
L
M
N
O
P
Q
R

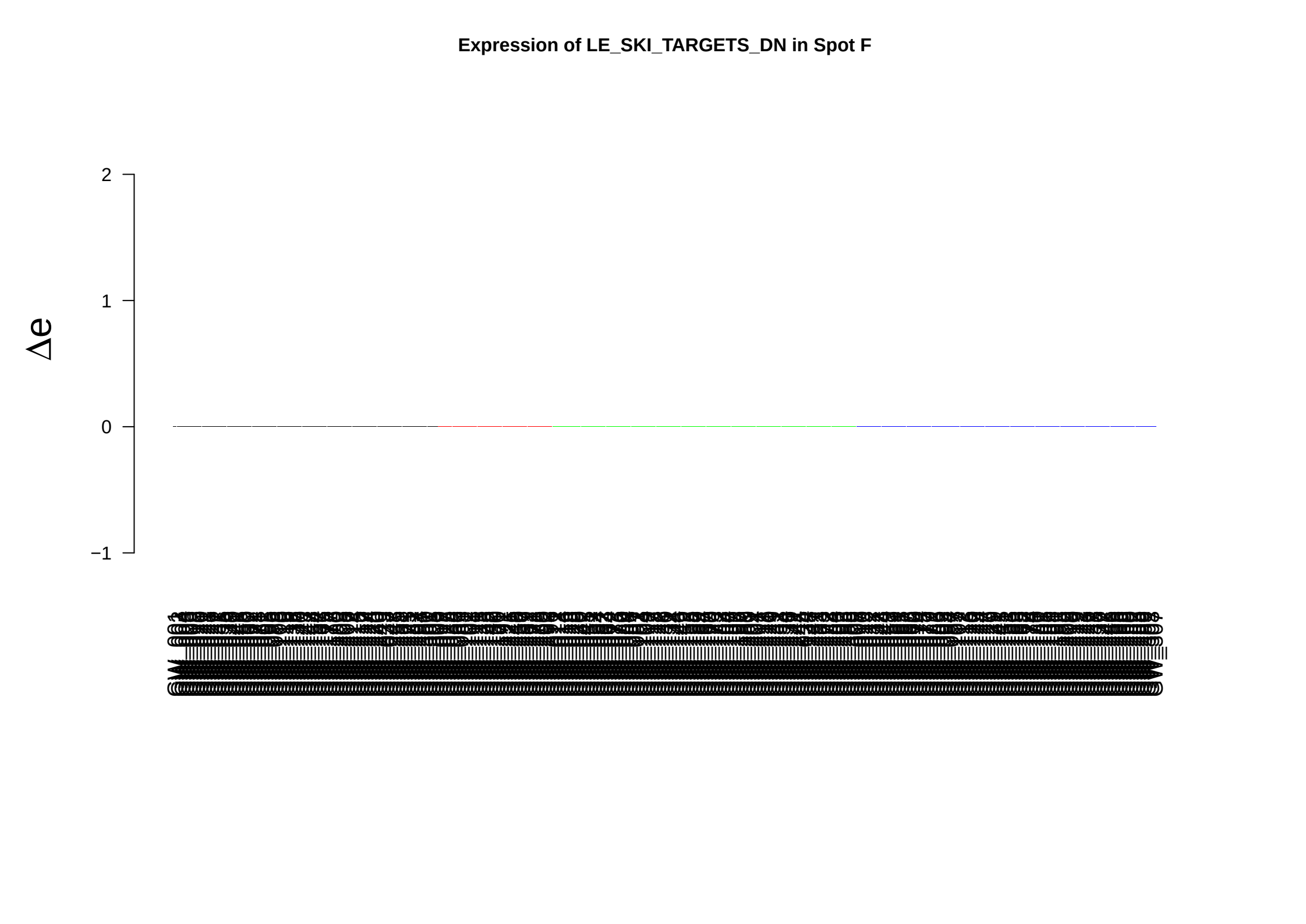

Expression of LE_SKI_TARGETS_DN in Spot A
Δe
2
1
0
−1

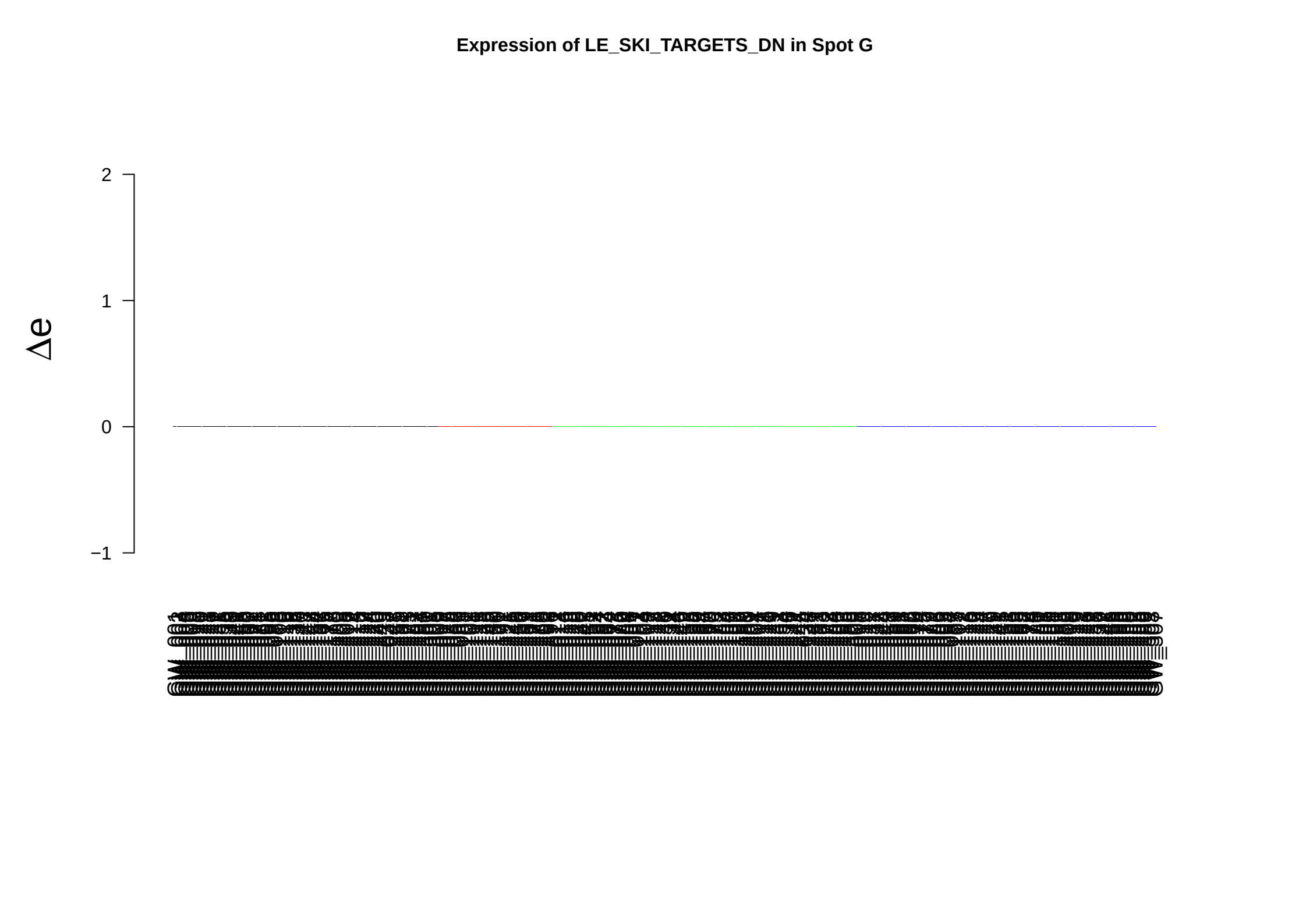

Expression of LE_SKI_TARGETS_DN in Spot B
Δe
2
1
0
−1

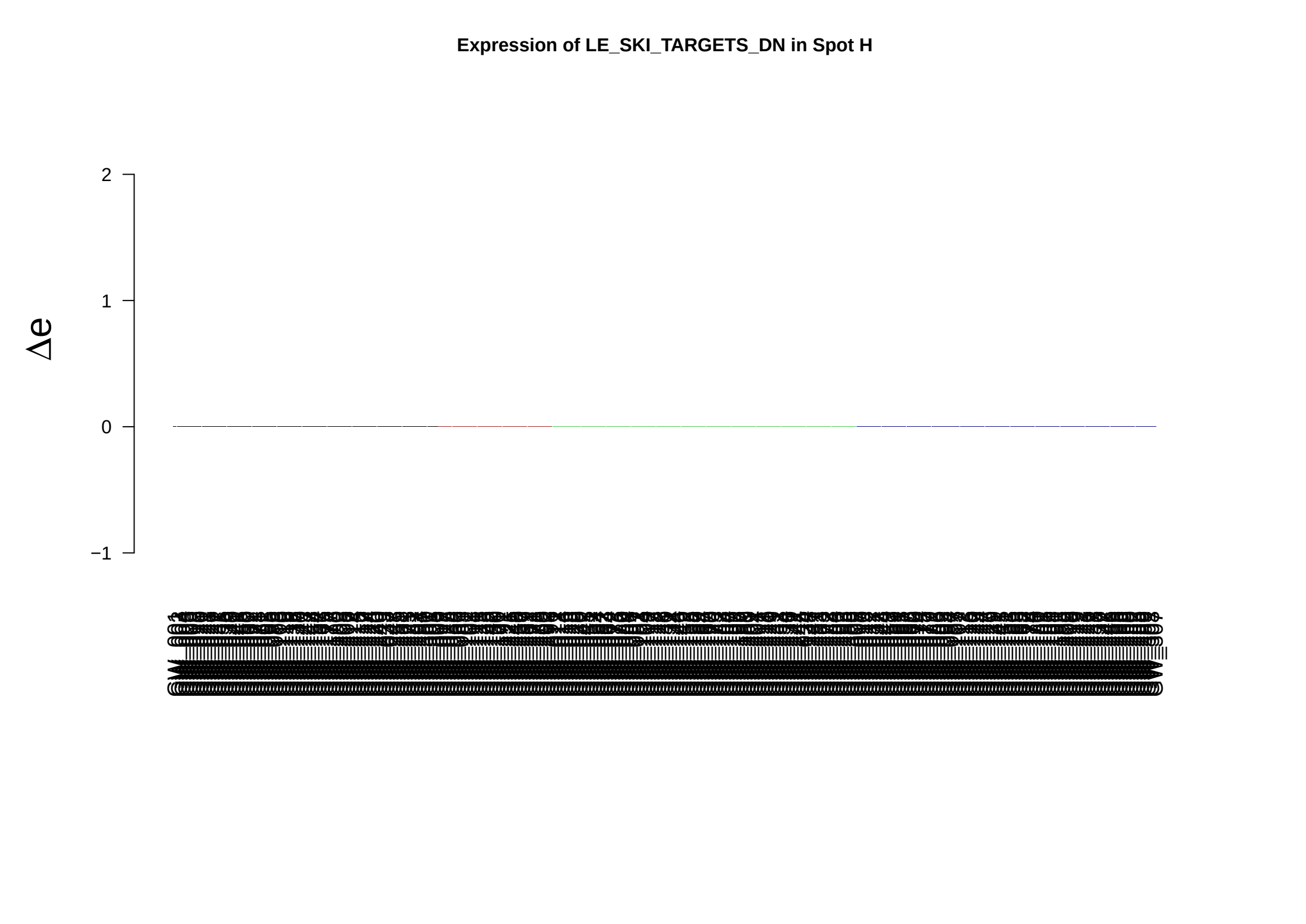

Expression of LE_SKI_TARGETS_DN in Spot C
Δe
2
1
0
−1

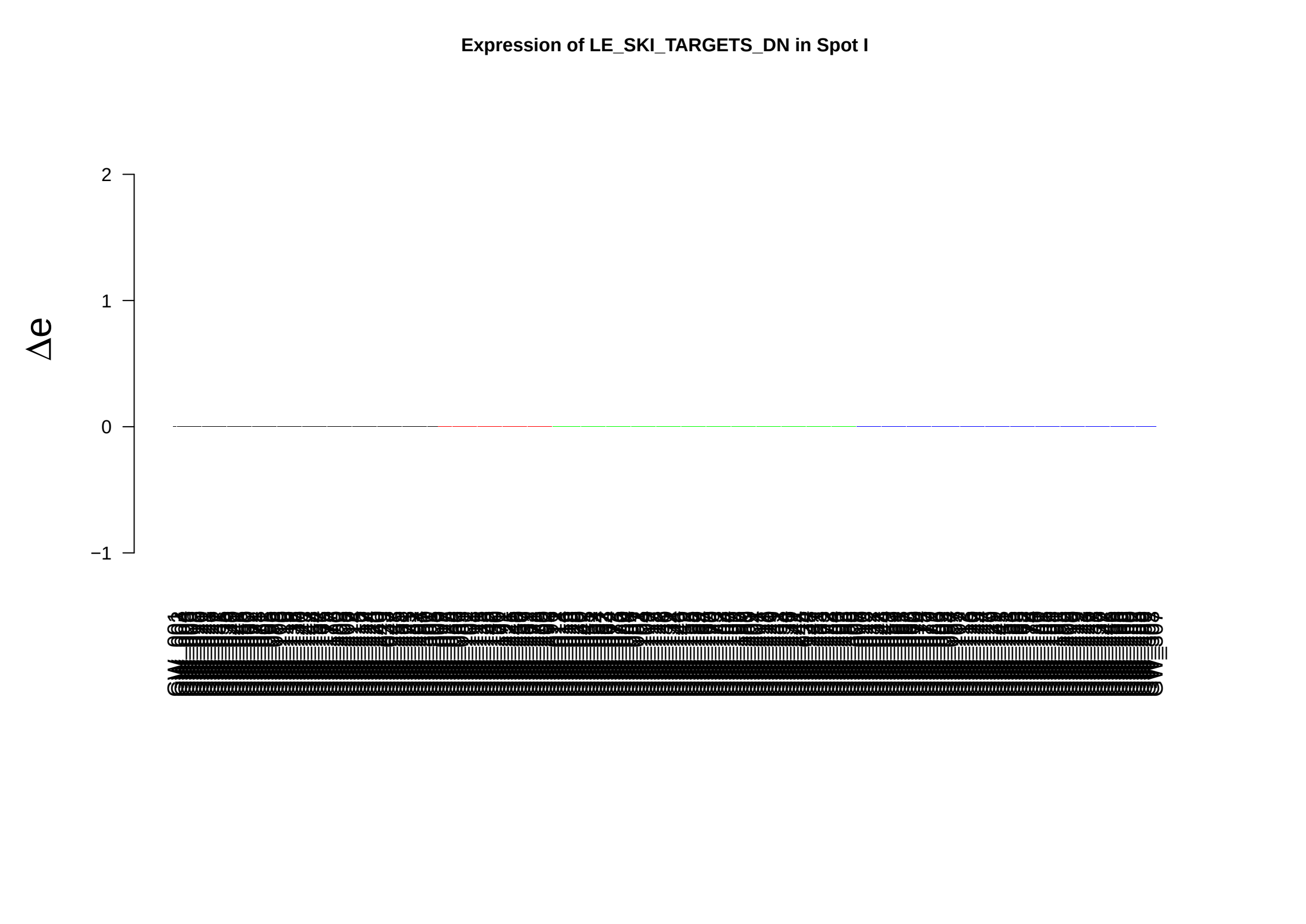

Expression of LE_SKI_TARGETS_DN in Spot D
Δe
2
1
0
−1

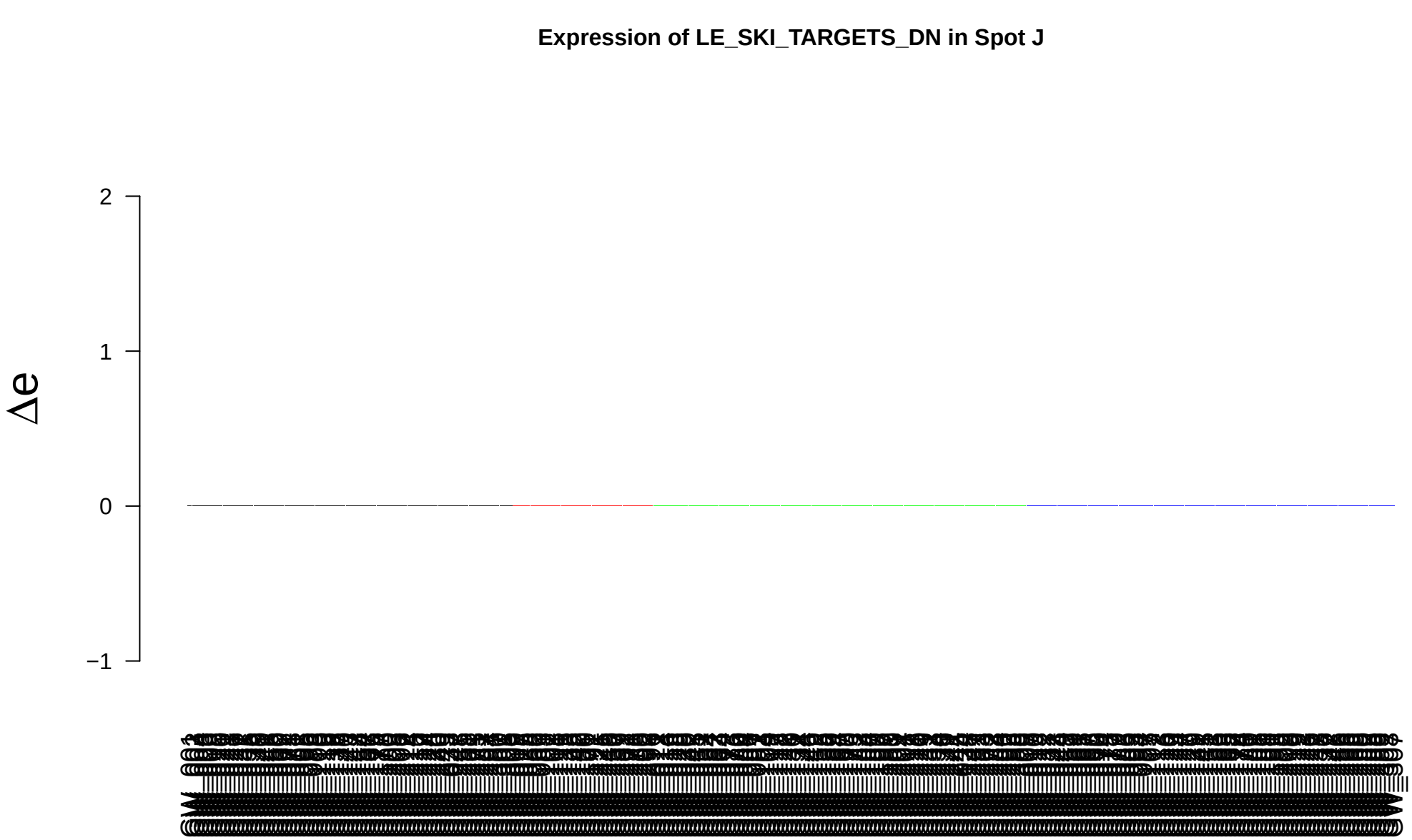

Expression of LE_SKI_TARGETS_DN in Spot E
Δe
2
1
0
−1

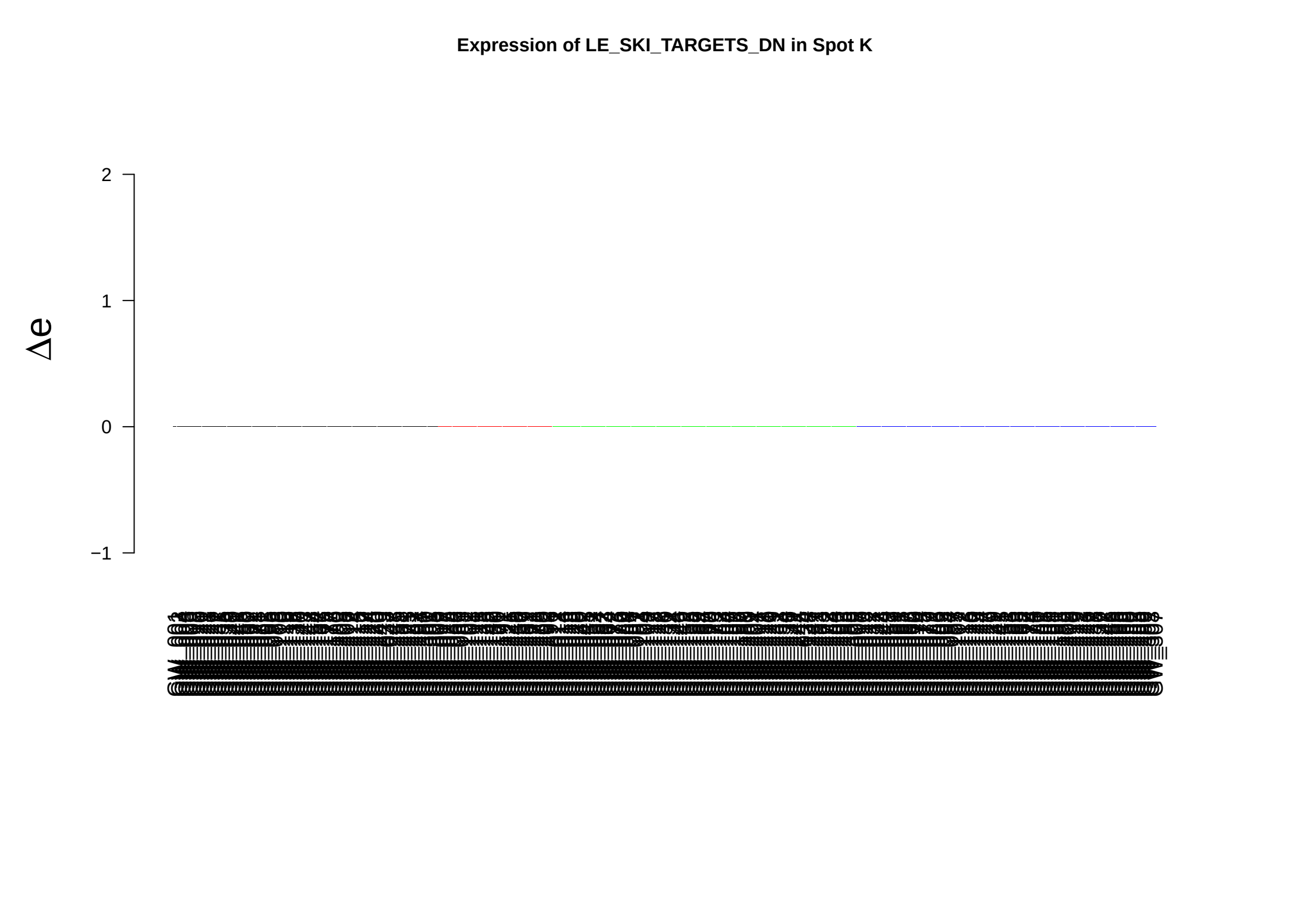

Expression of LE_SKI_TARGETS_DN in Spot F
Δe
2
1
0
−1

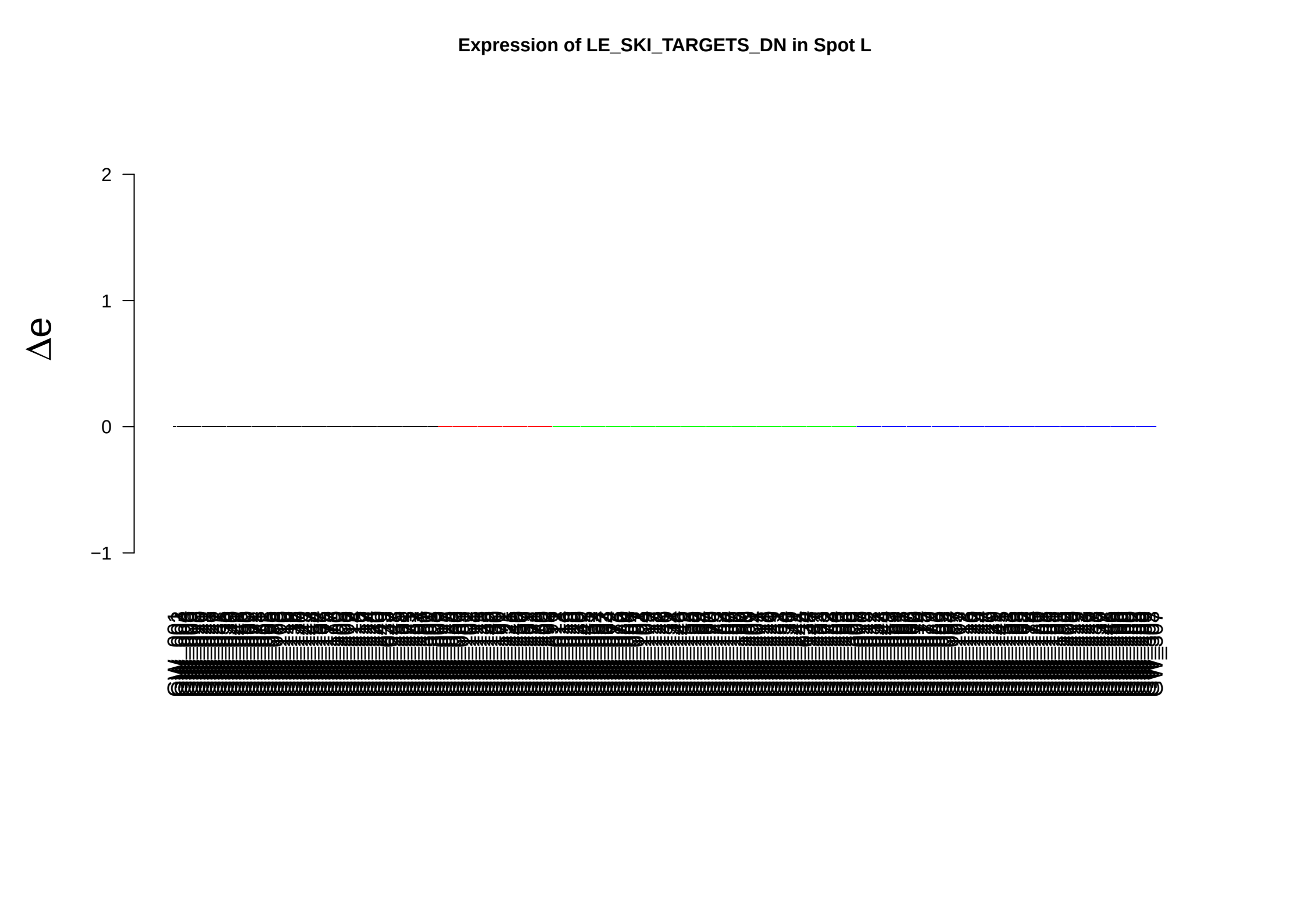

Expression of LE_SKI_TARGETS_DN in Spot G
Δe
2
1
0
−1

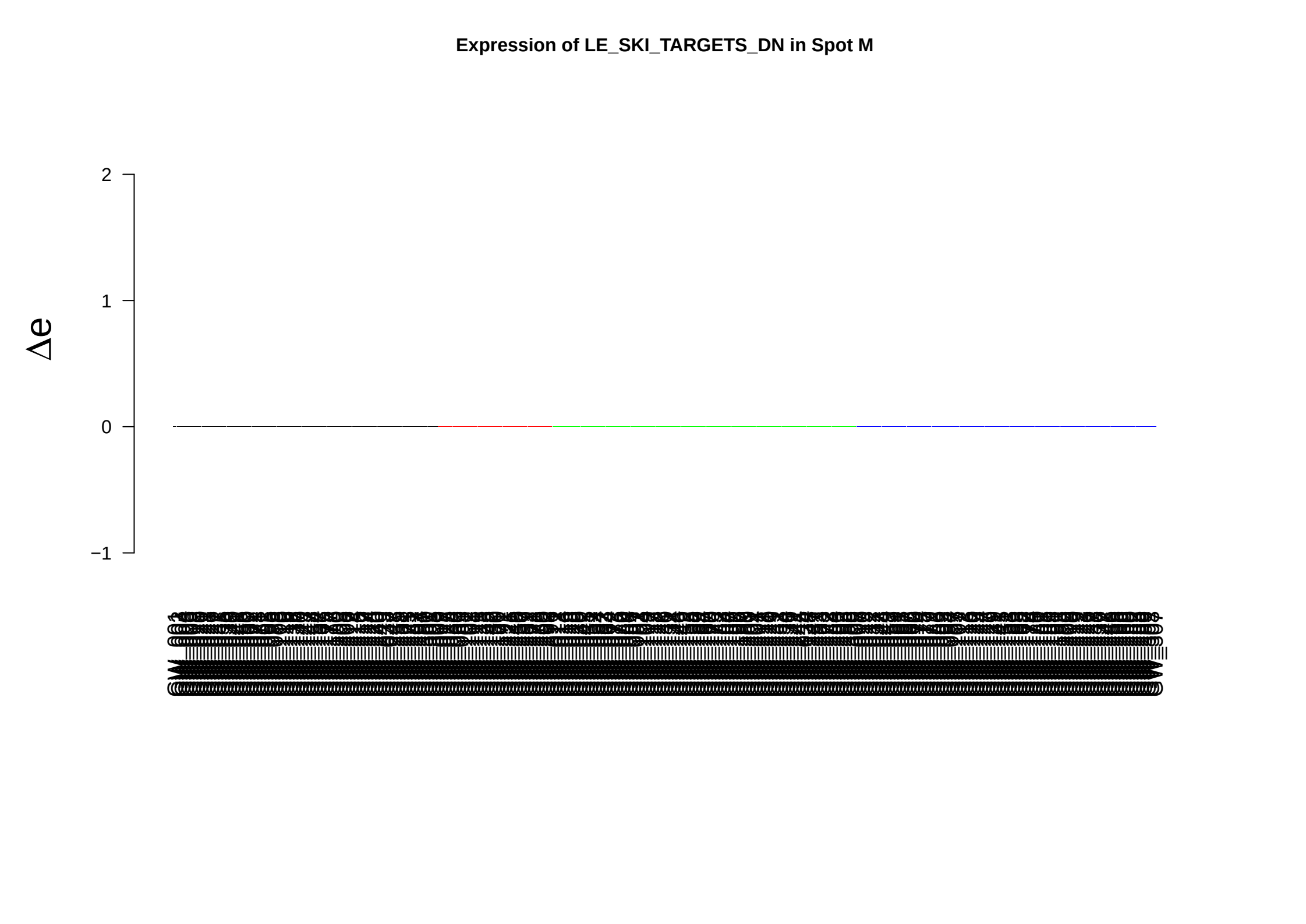

Expression of LE_SKI_TARGETS_DN in Spot H
Δe
2
1
0
−1

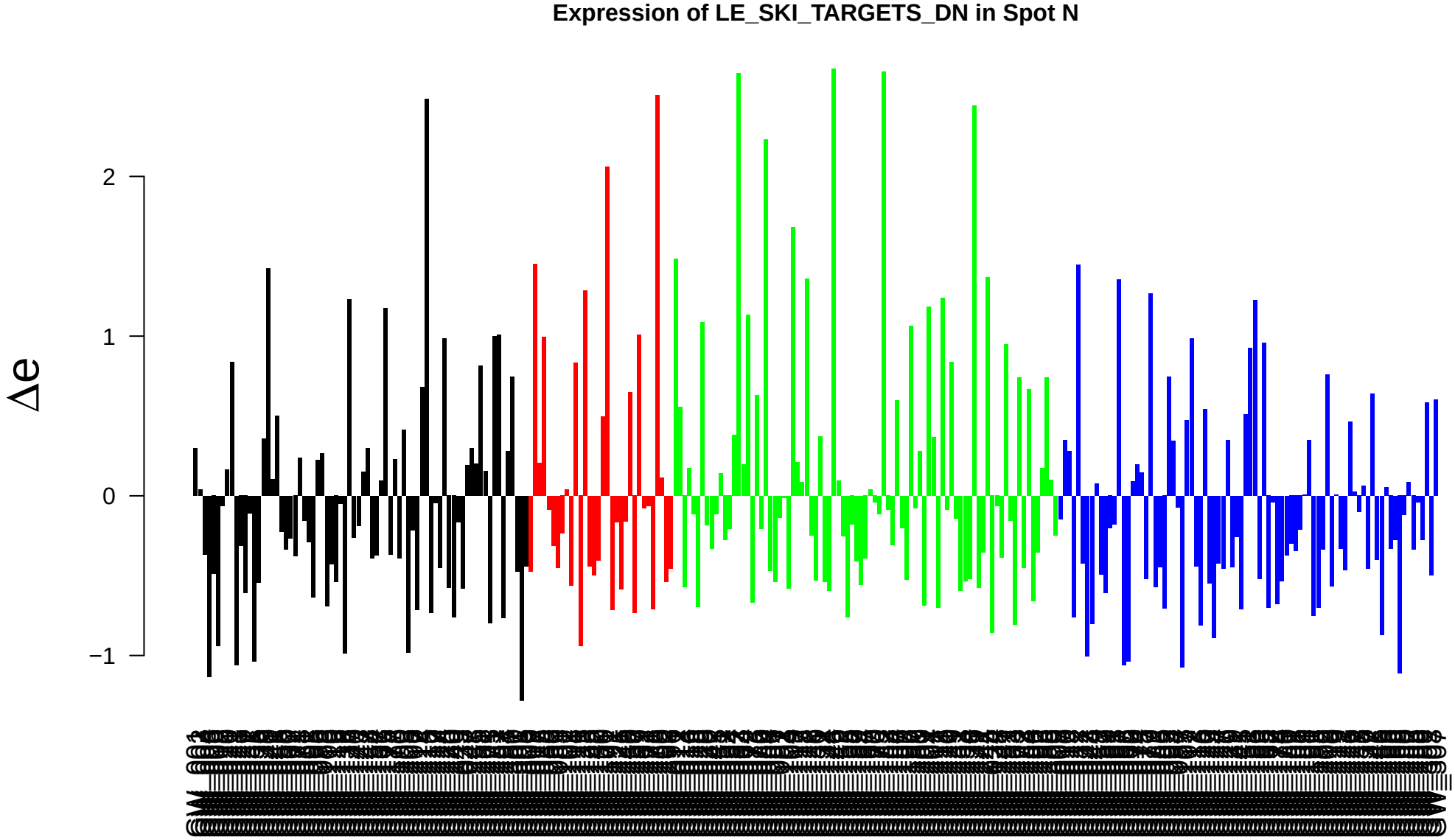

Expression of LE_SKI_TARGETS_DN in Spot I
Δe
2
1
0
−1

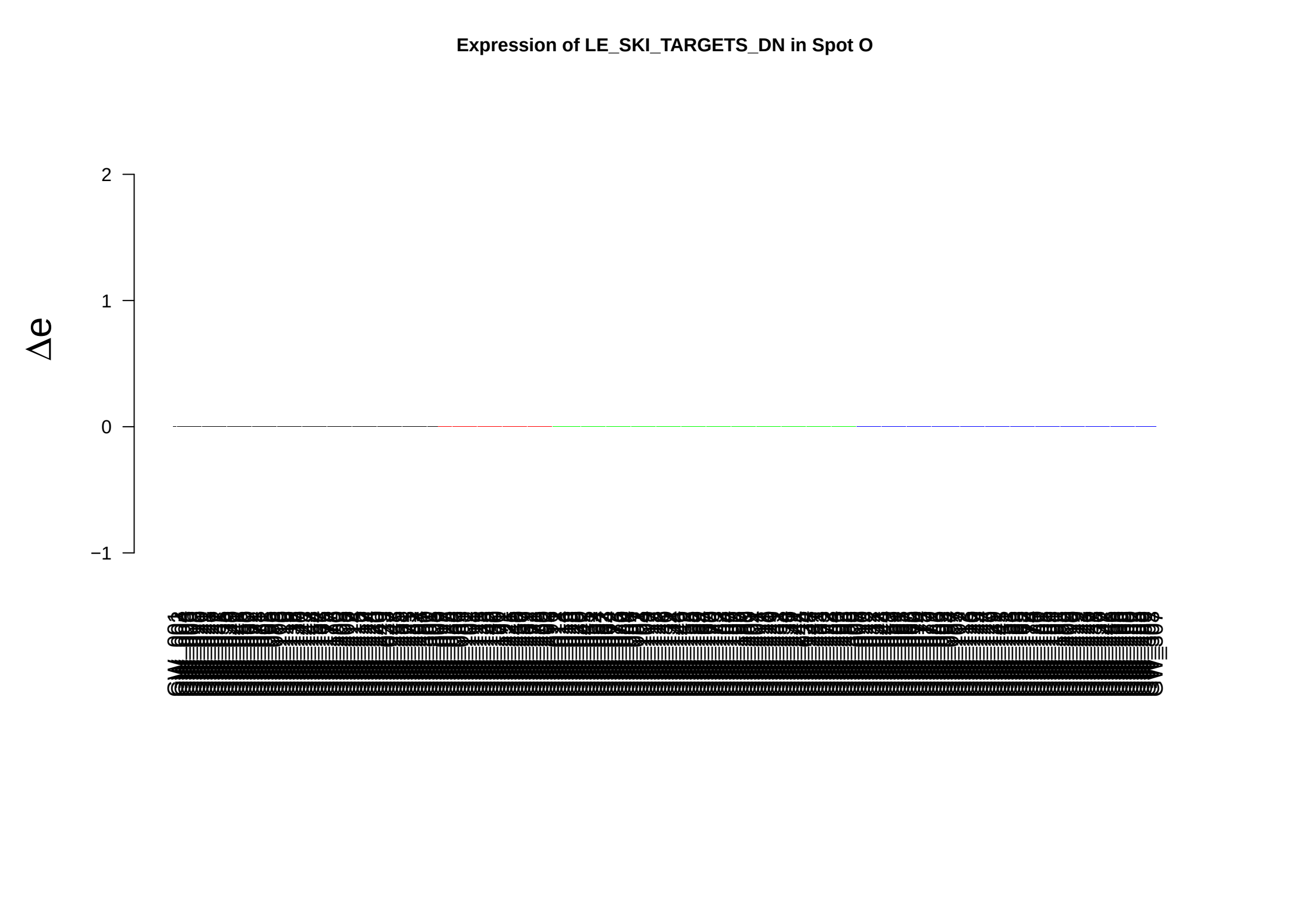

Expression of LE_SKI_TARGETS_DN in Spot J
Δe
2
1
0
−1

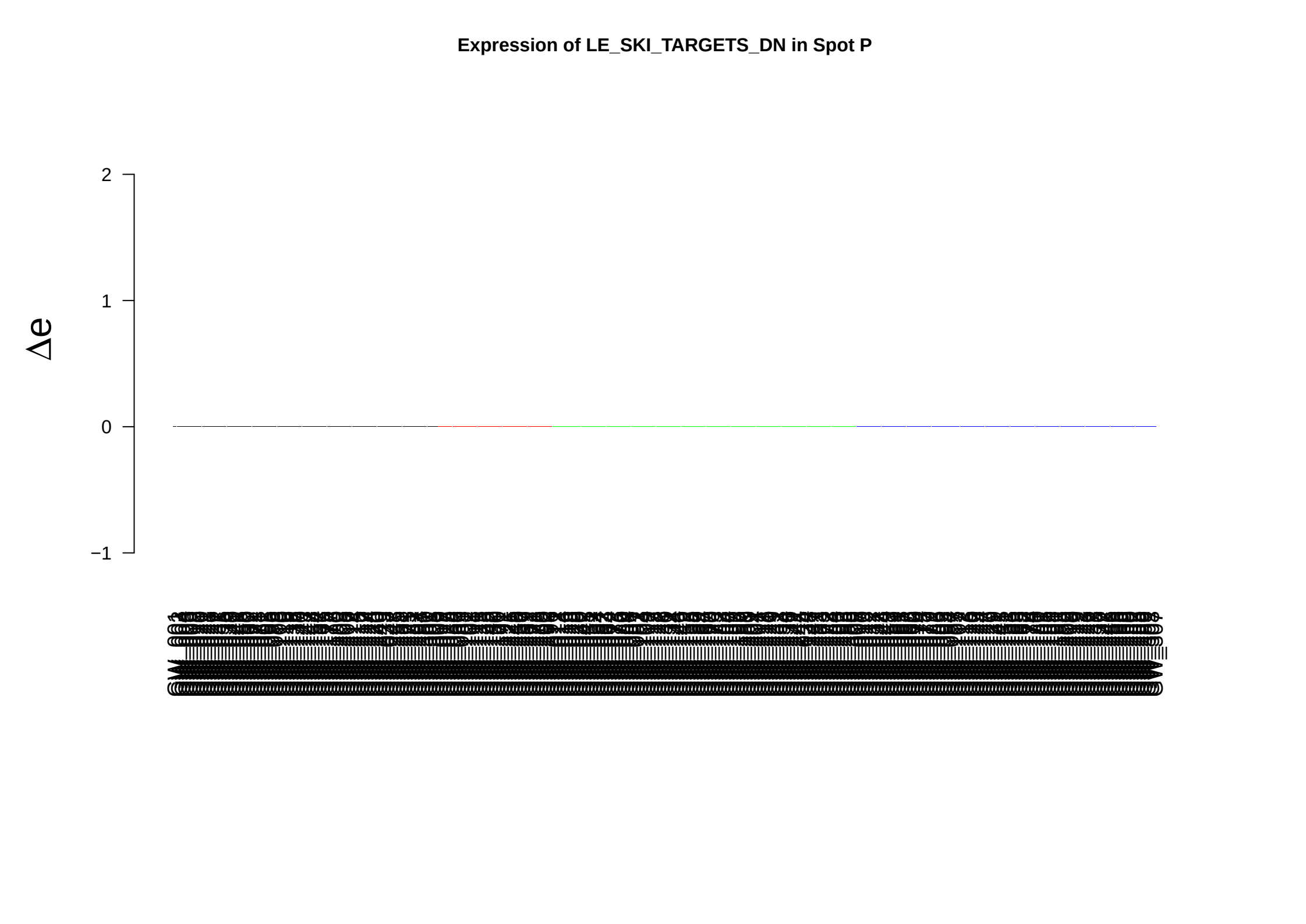

Expression of LE_SKI_TARGETS_DN in Spot K
Δe
2
1
0
−1

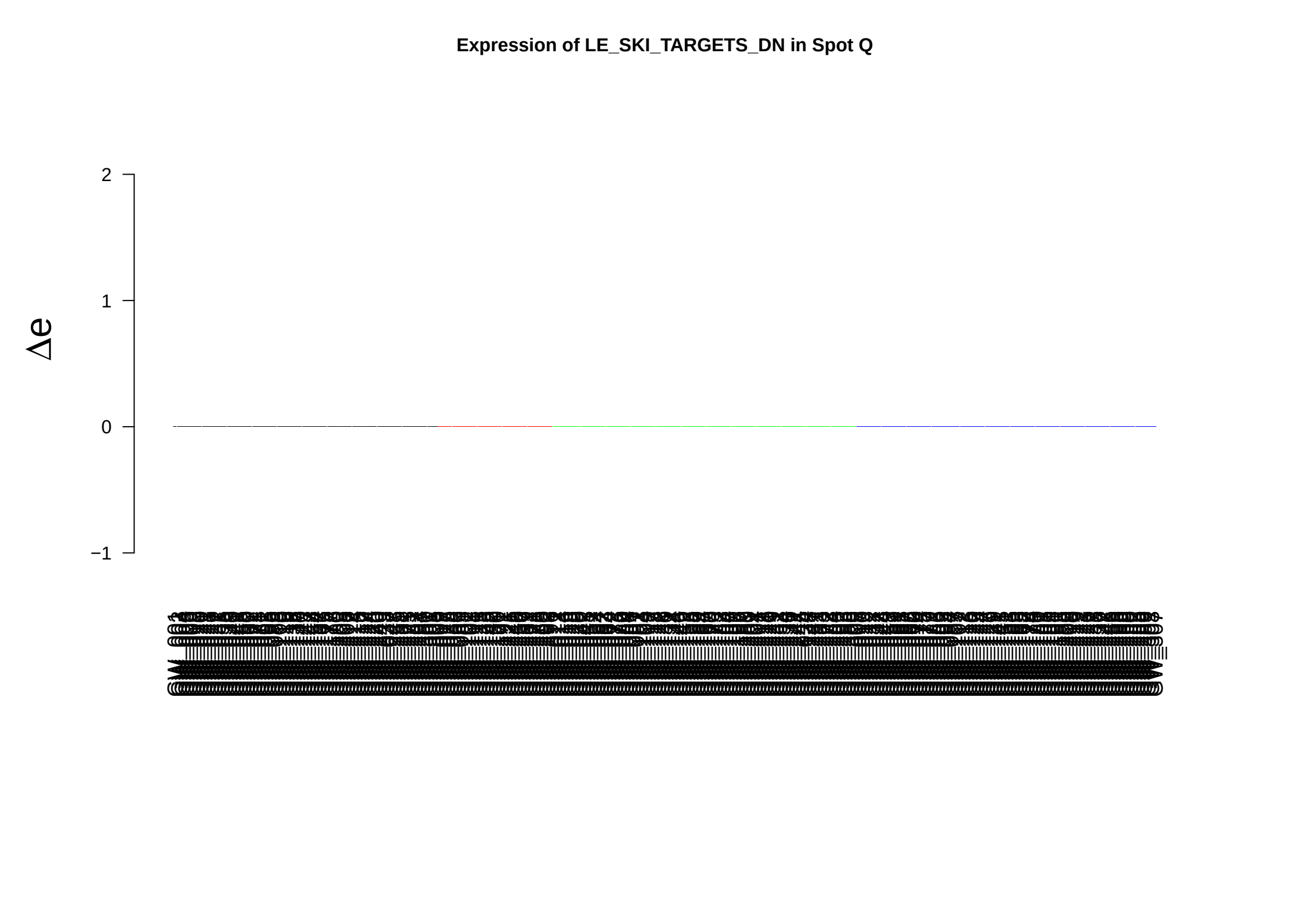

Expression of LE_SKI_TARGETS_DN in Spot L
Δe
2
1
0
−1

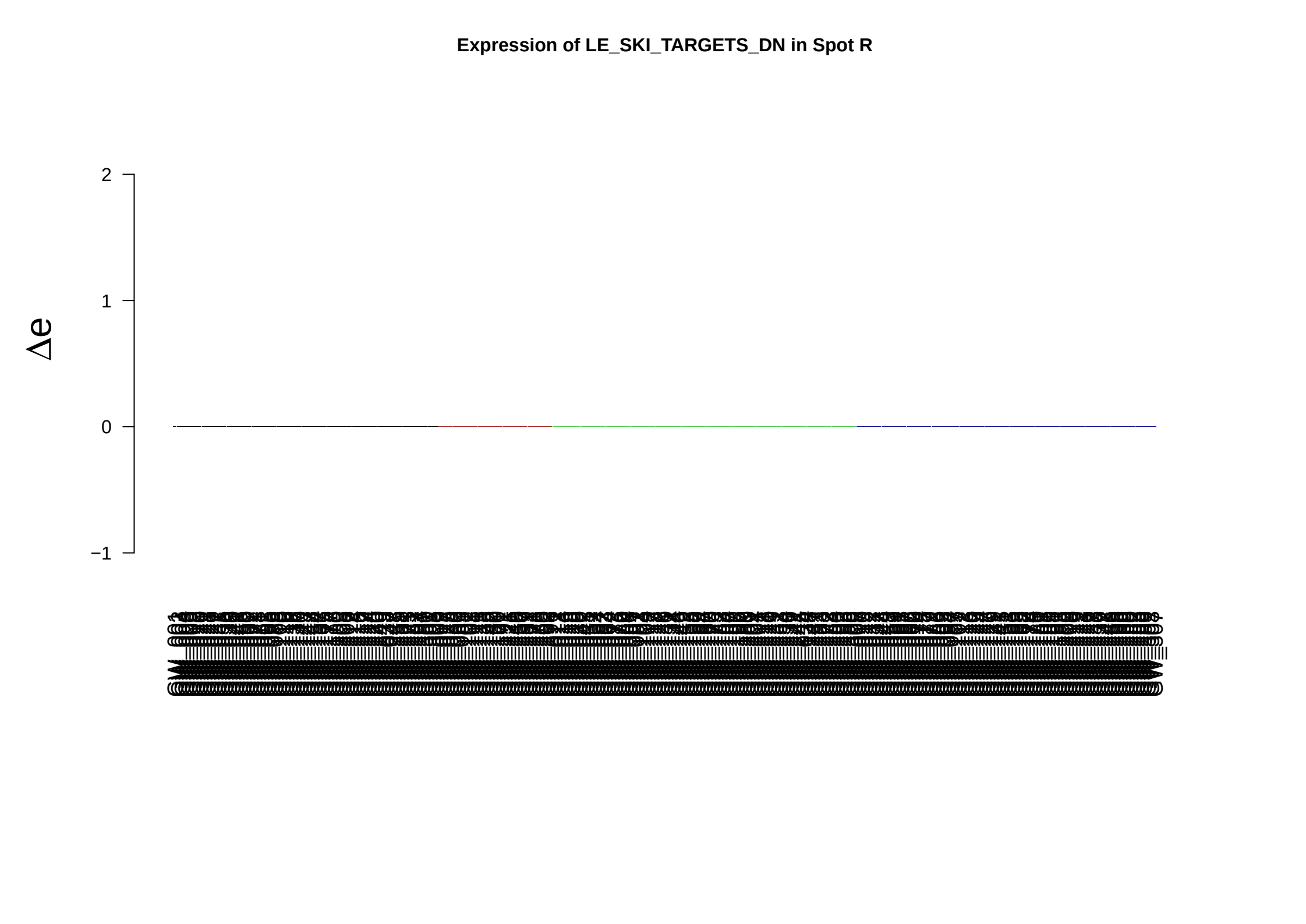

Expression of LE_SKI_TARGETS_DN in Spot M
Δe
2
1
0
−1


Expression of LE_SKI_TARGETS_DN in Spot N
Δe
2
1
0
−1


Expression of LE_SKI_TARGETS_DN in Spot O
Δe
2
1
0
−1


Expression of LE_SKI_TARGETS_DN in Spot P
Δe
2
1
0
−1


Expression of LE_SKI_TARGETS_DN in Spot Q
Δe
2
1
0
−1


Expression of LE_SKI_TARGETS_DN in Spot R
Δe
2
1
0
−1