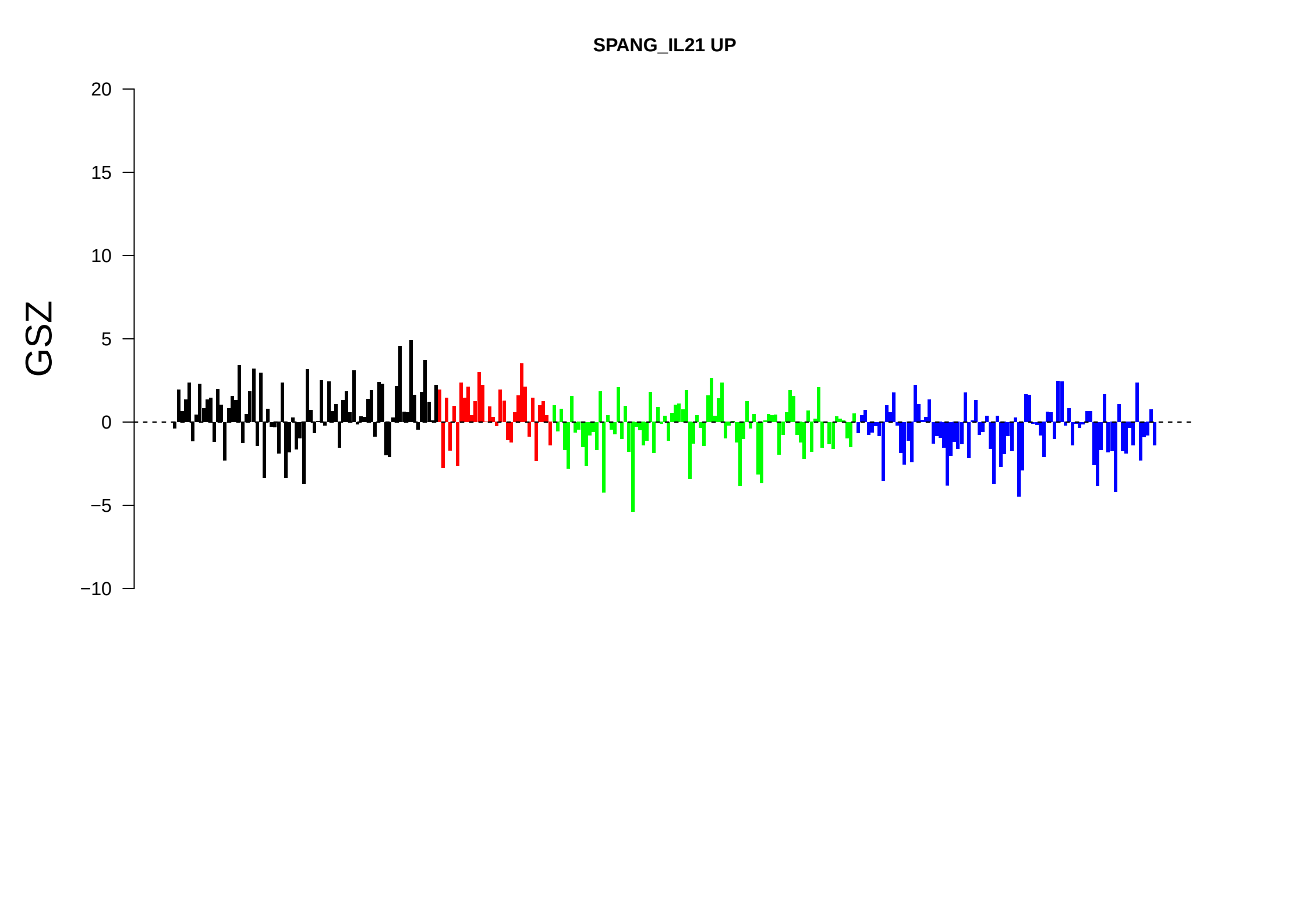
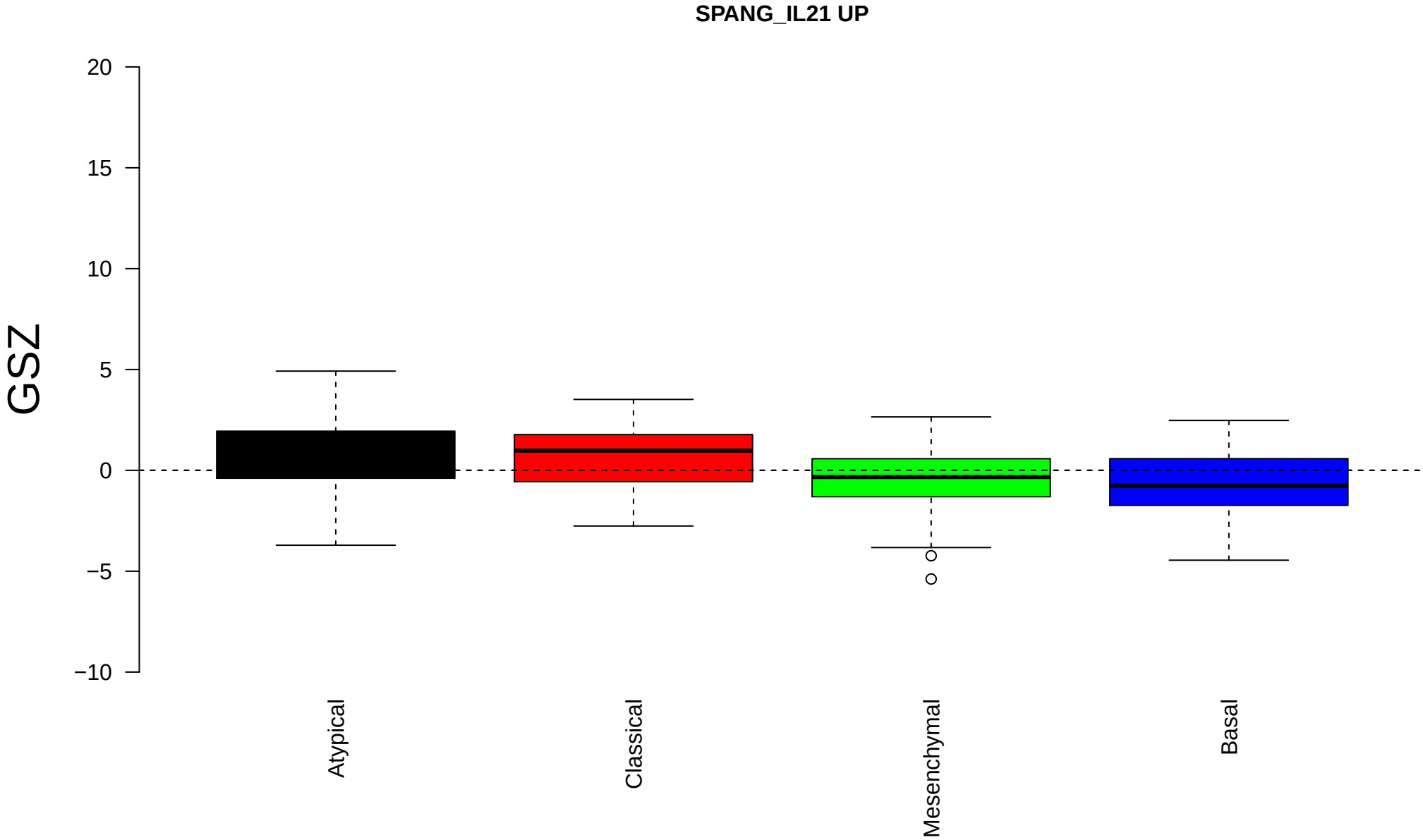
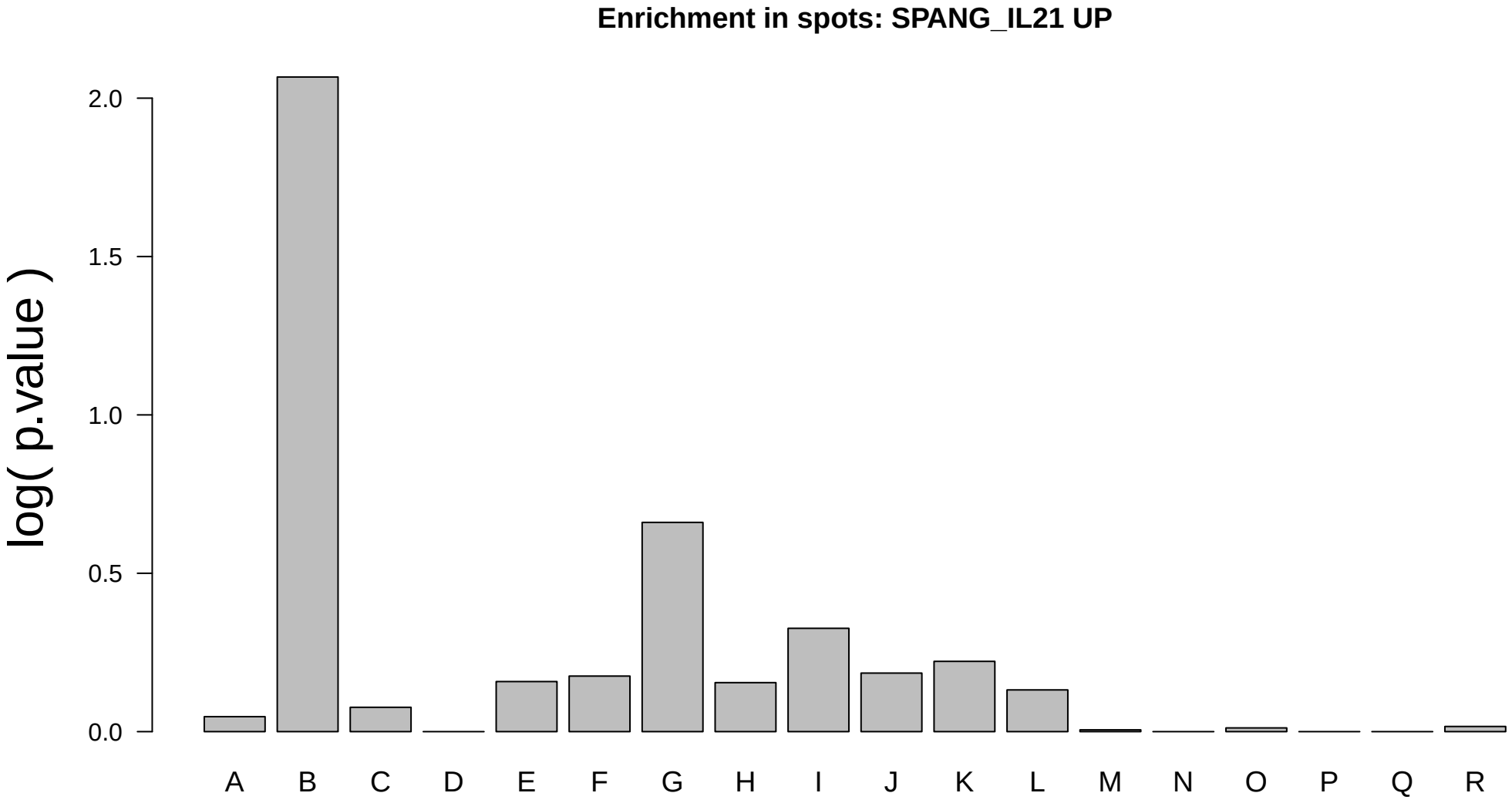
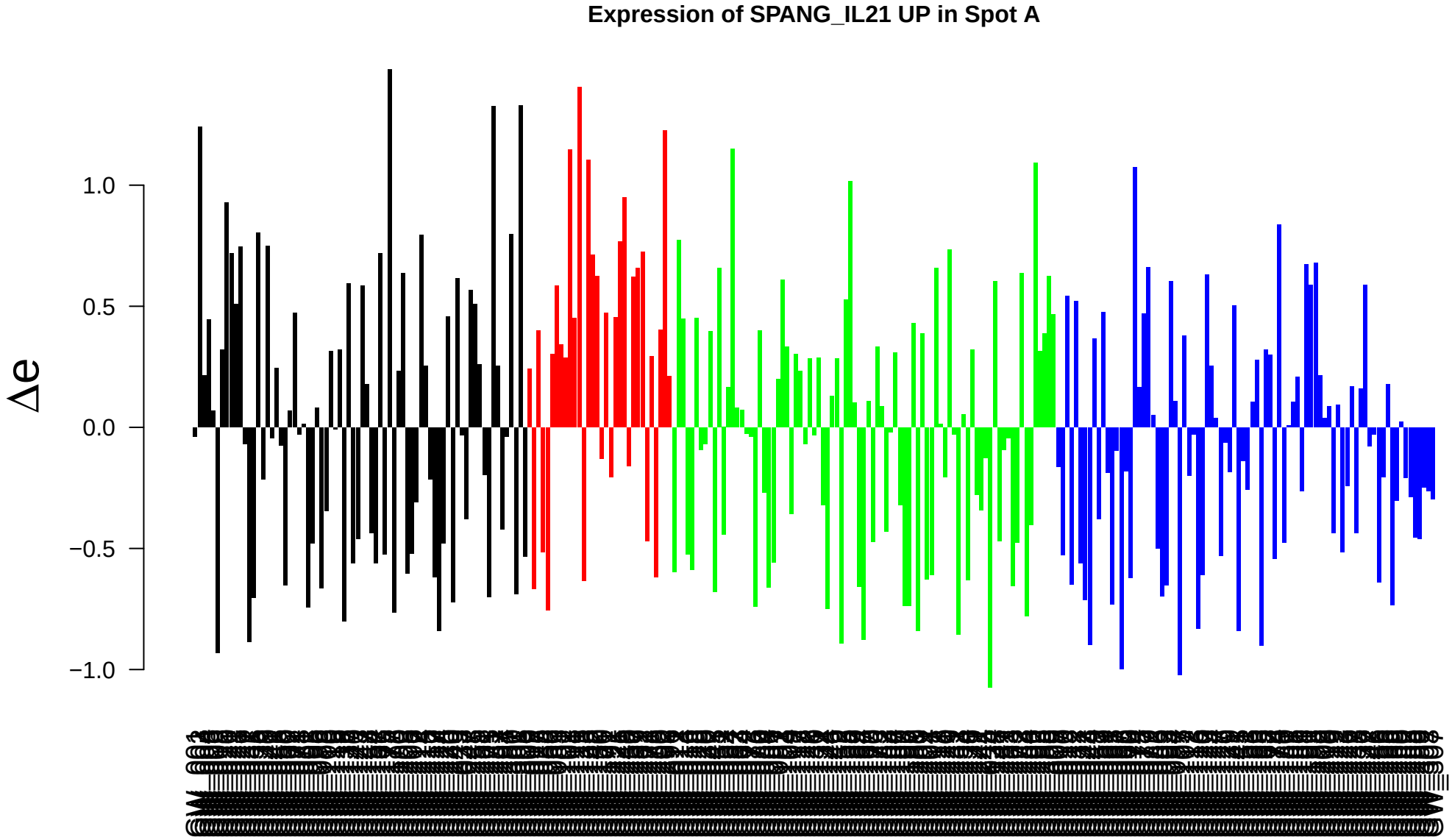
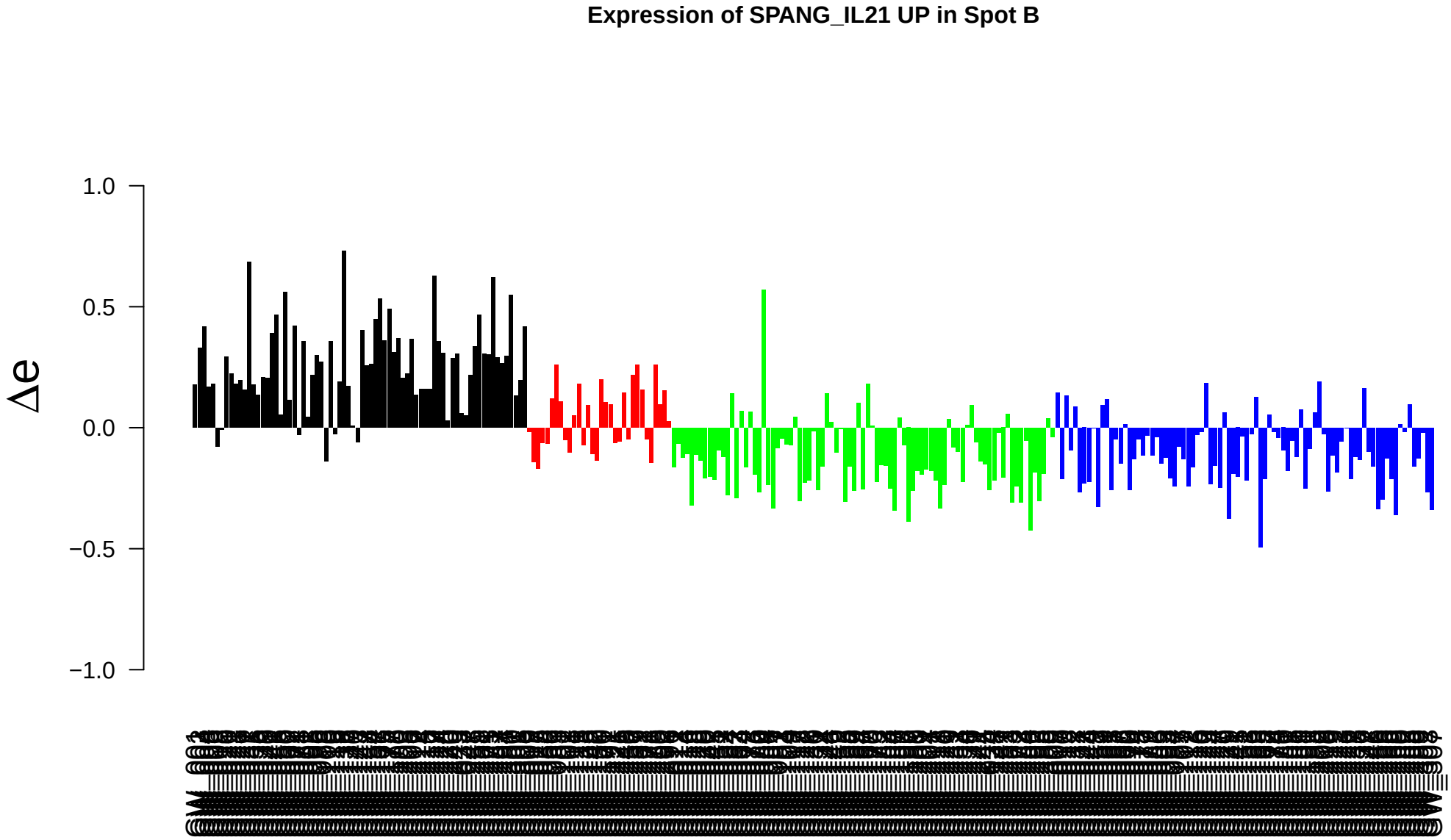
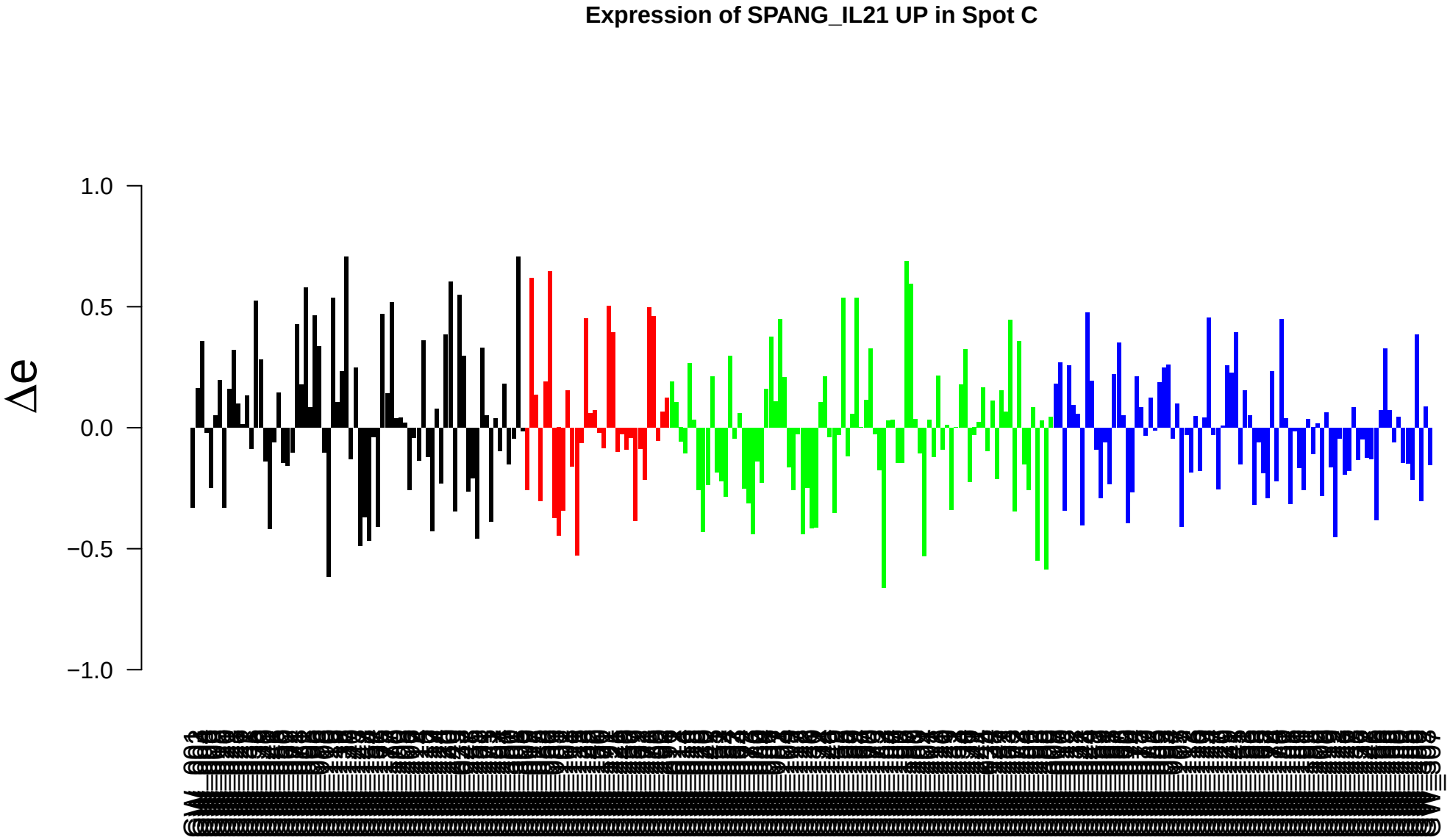

SPANG_IL21 UP
GSZ
20
15
10
5
0
−5
−10

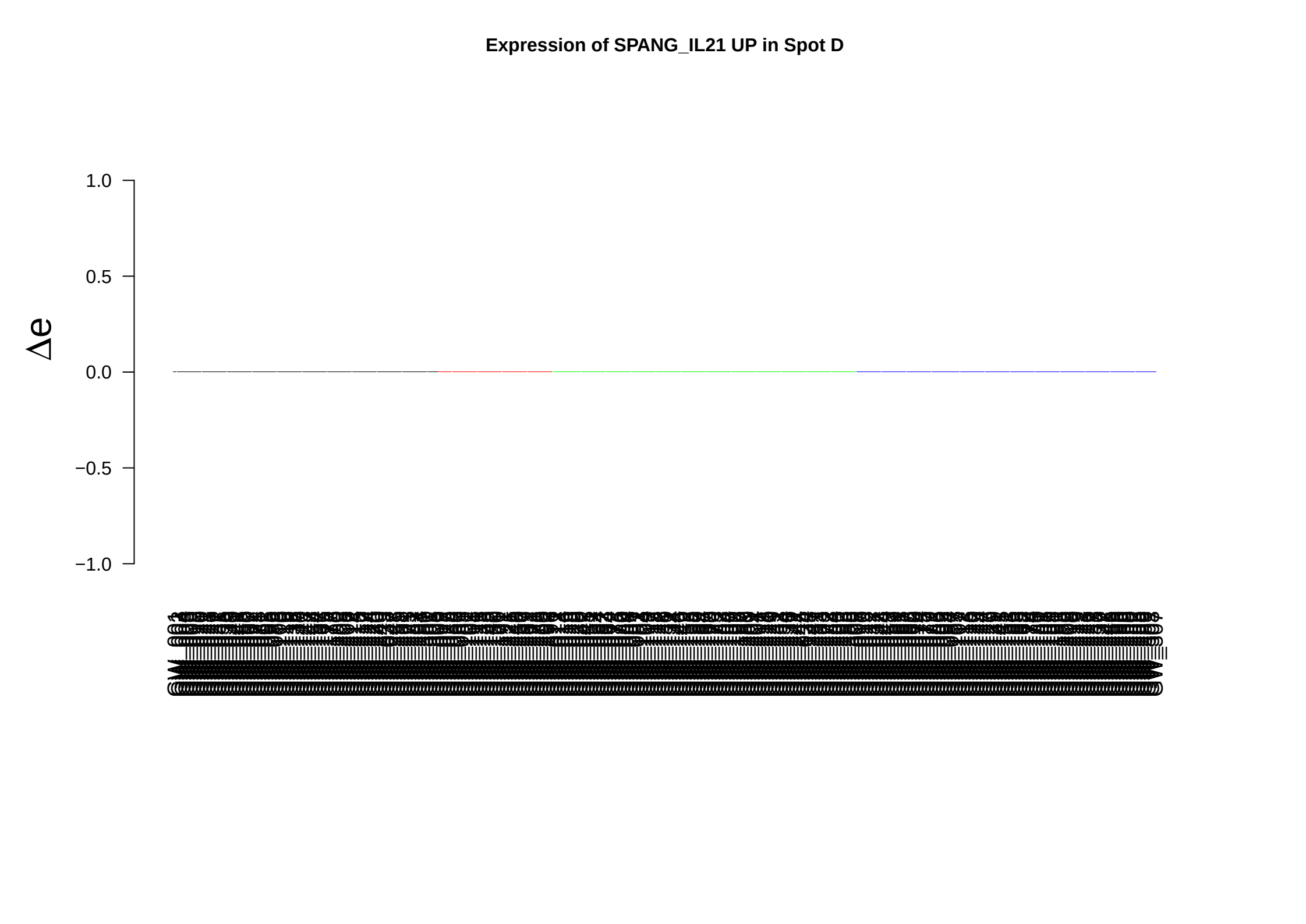

SPANG_IL21 UP
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal

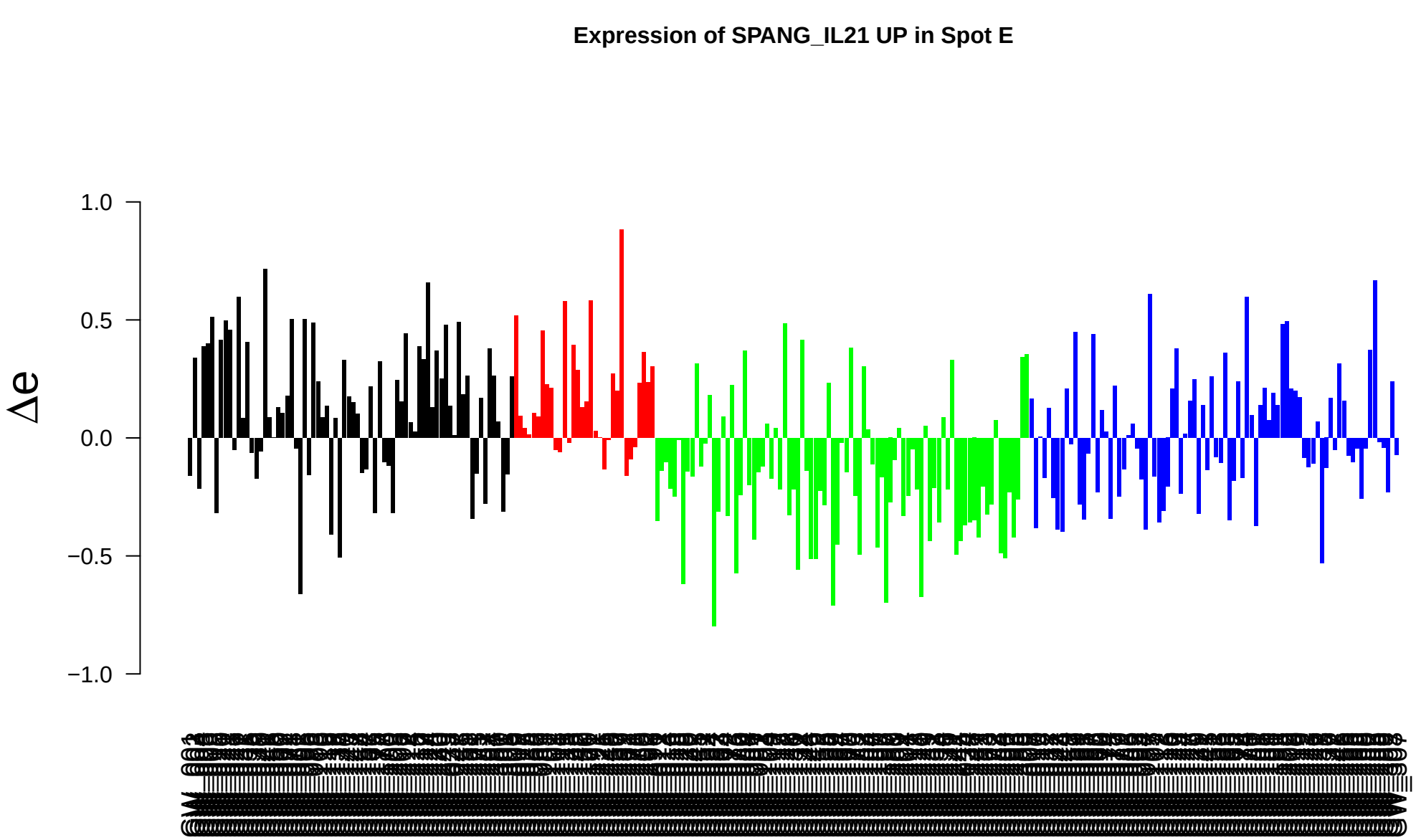

Enrichment in spots: SPANG_IL21 UP
log(p.value)
2.0
1.5
1.0
0.5
0.0
A
B
C
D
E
F
G
H
I
J
K
L
M
N
O
P
Q
R

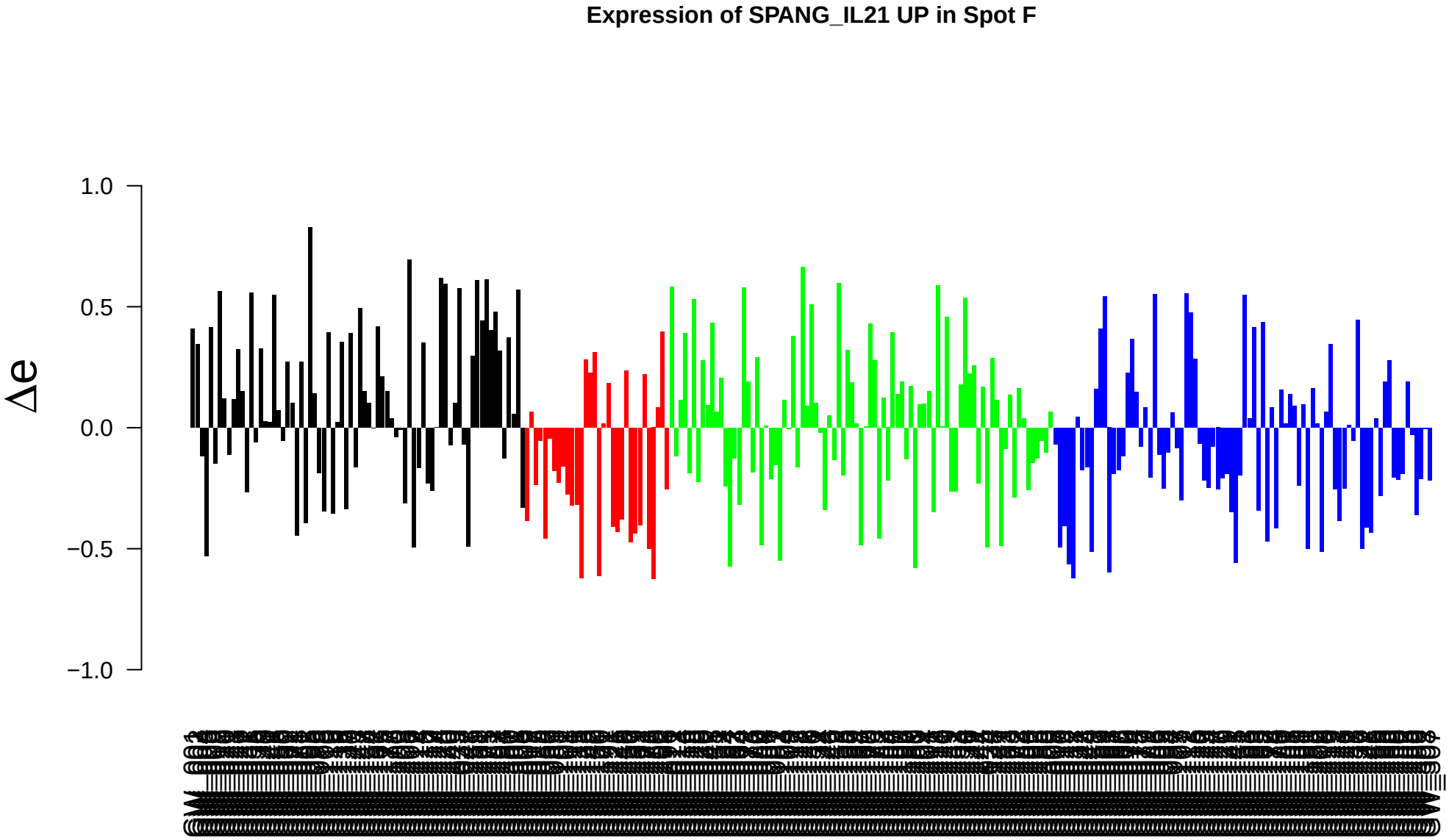

Expression of SPANG_IL21 UP in Spot A
Δe
1.0
0.5
0.0
−0.5
−1.0

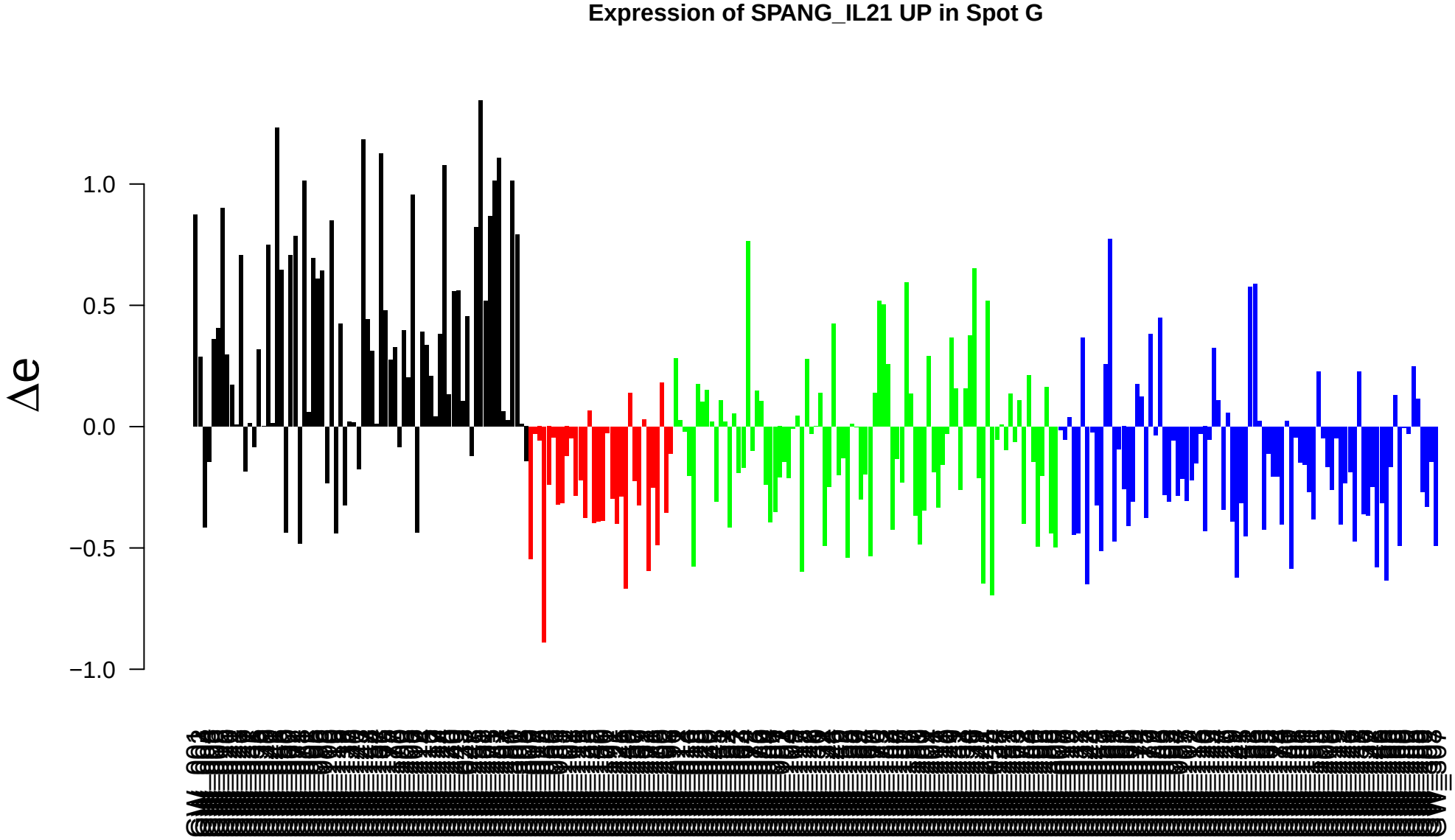

Expression of SPANG_IL21 UP in Spot B
Δe
1.0
0.5
0.0
−0.5
−1.0

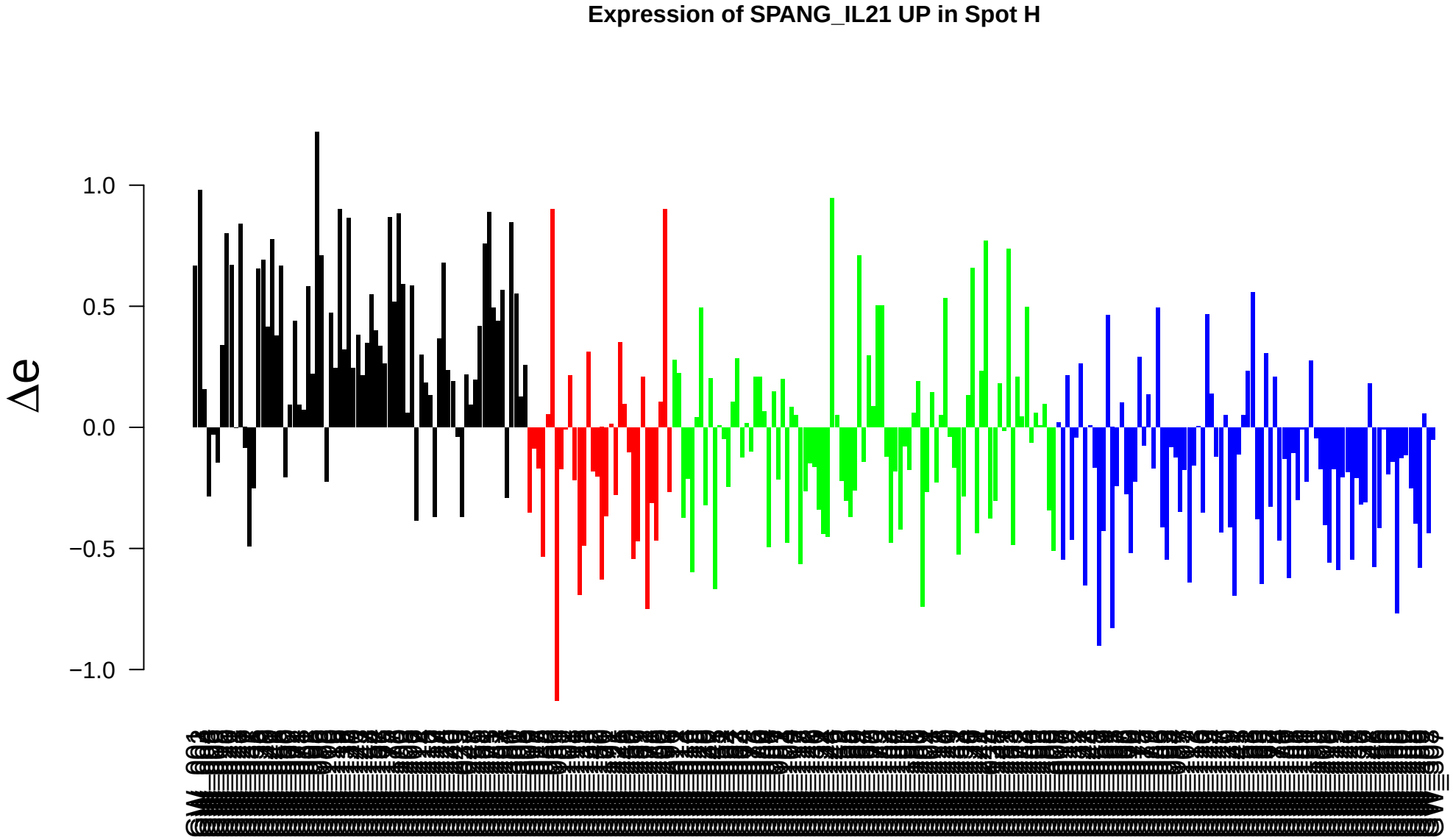

Expression of SPANG_IL21 UP in Spot C
Δe
1.0
0.5
0.0
−0.5
−1.0

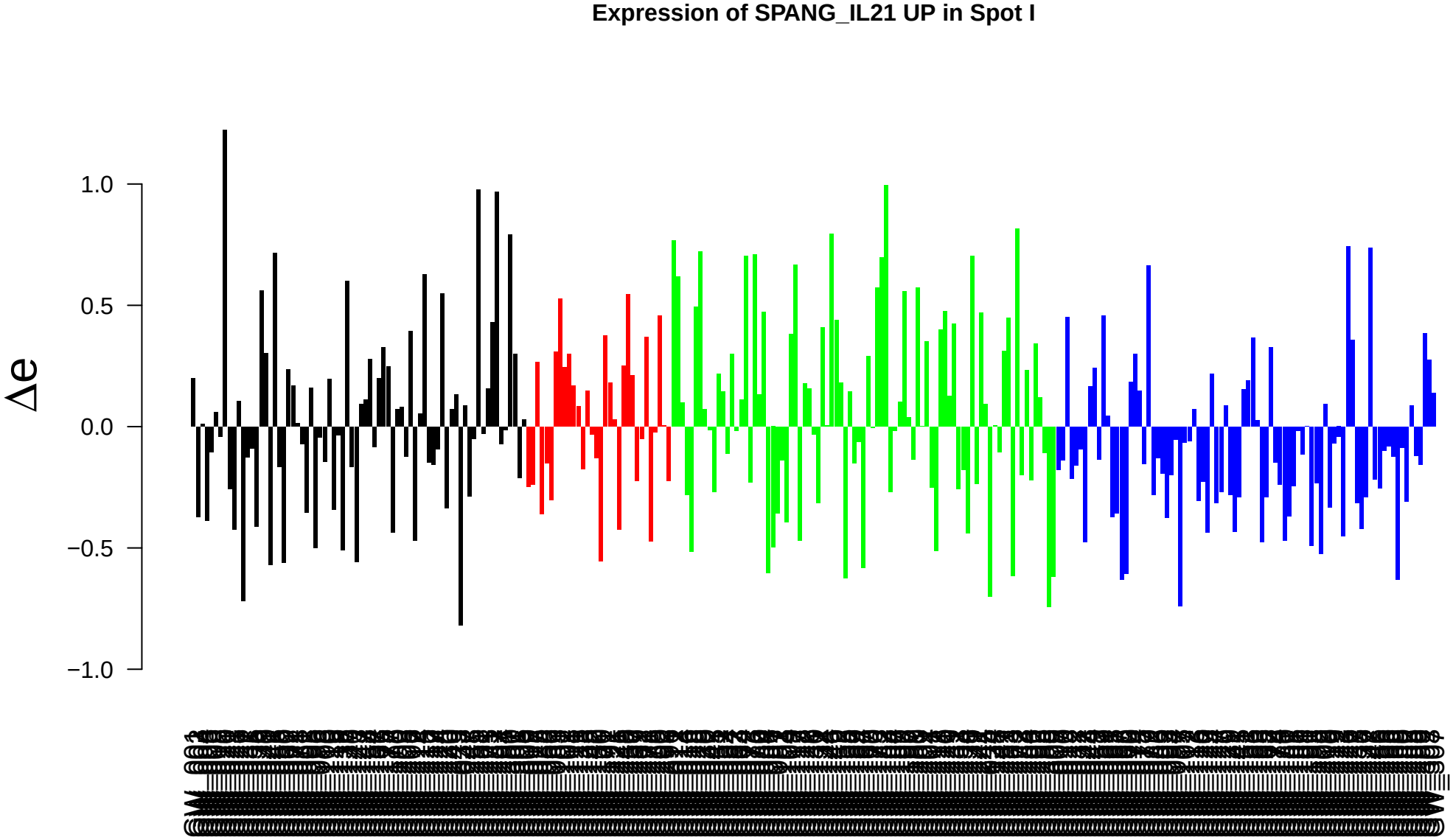

Expression of SPANG_IL21 UP in Spot D
Δe
1.0
0.5
0.0
−0.5
−1.0

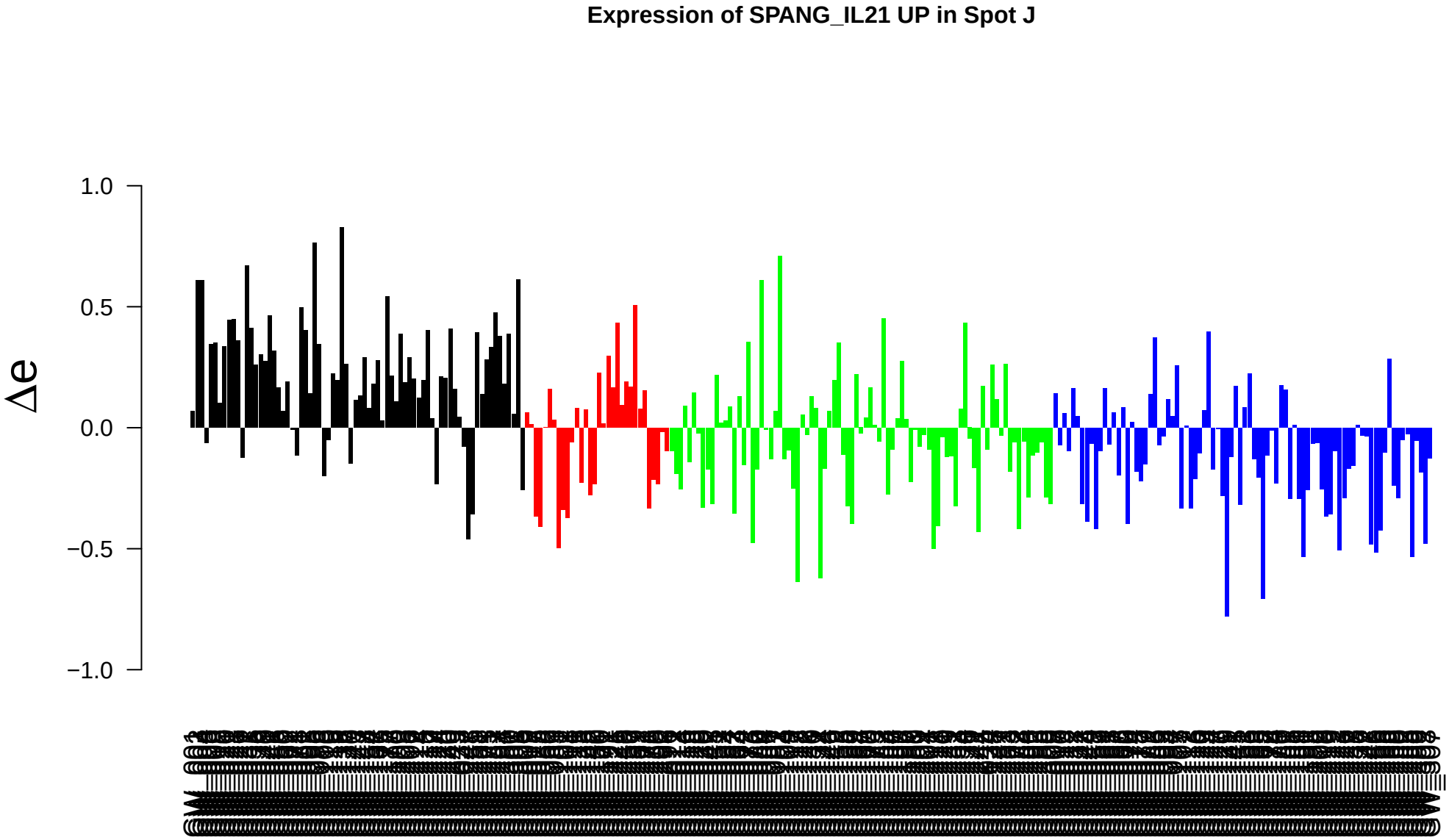

Expression of SPANG_IL21 UP in Spot E
Δe
1.0
0.5
0.0
−0.5
−1.0

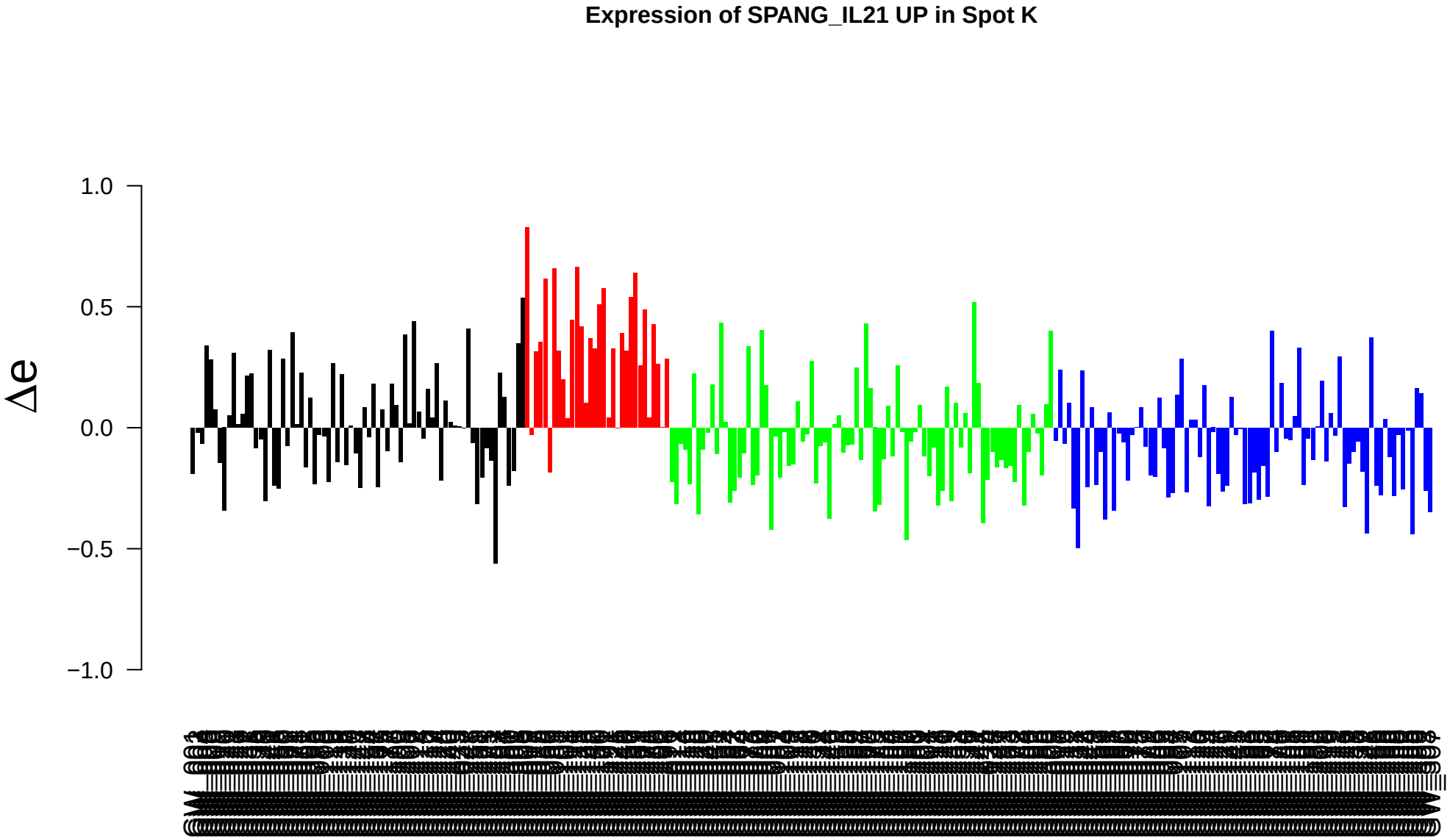

Expression of SPANG_IL21 UP in Spot F
Δe
1.0
0.5
0.0
−0.5
−1.0

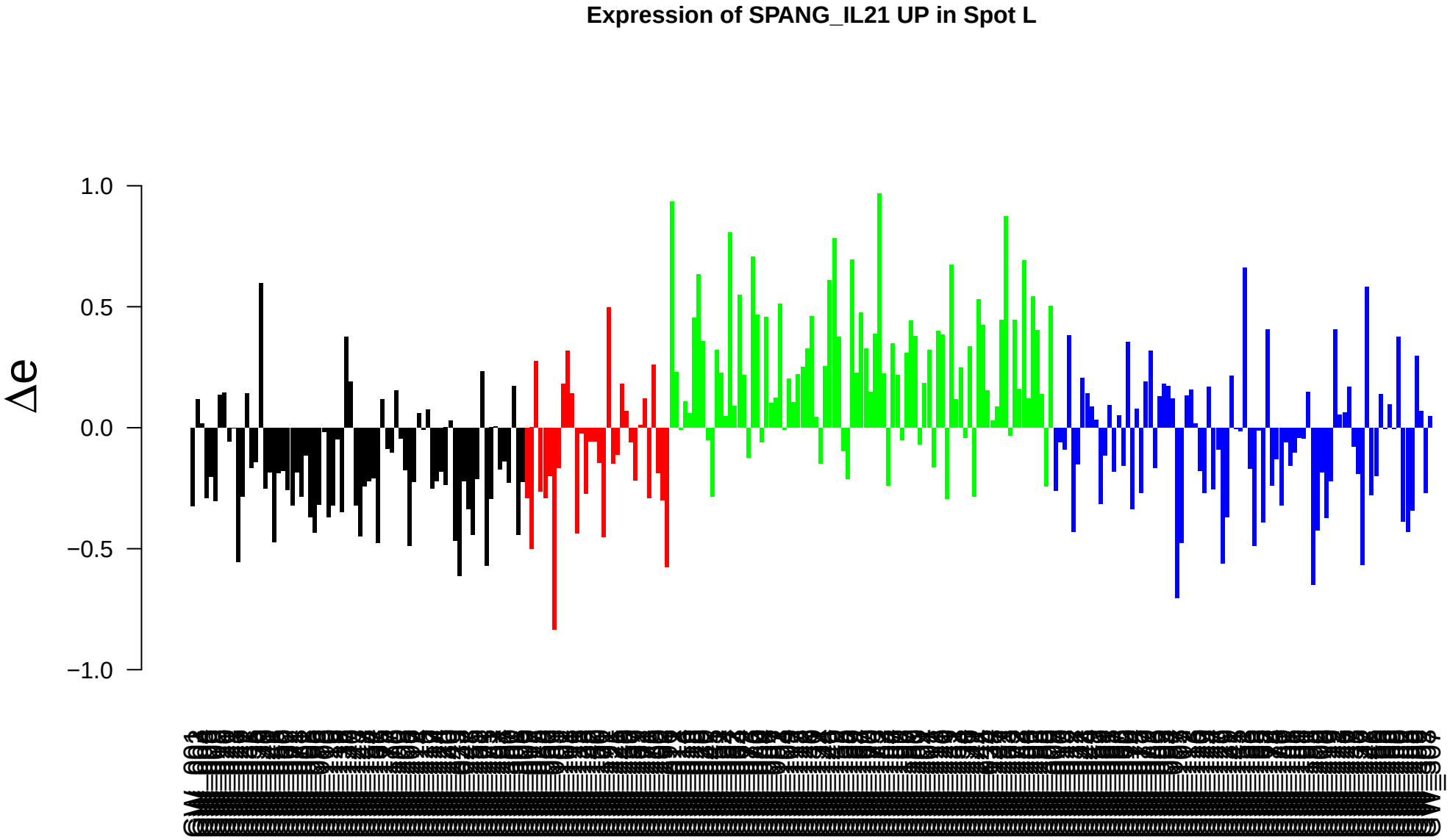

Expression of SPANG_IL21 UP in Spot G
Δe
1.0
0.5
0.0
−0.5
−1.0

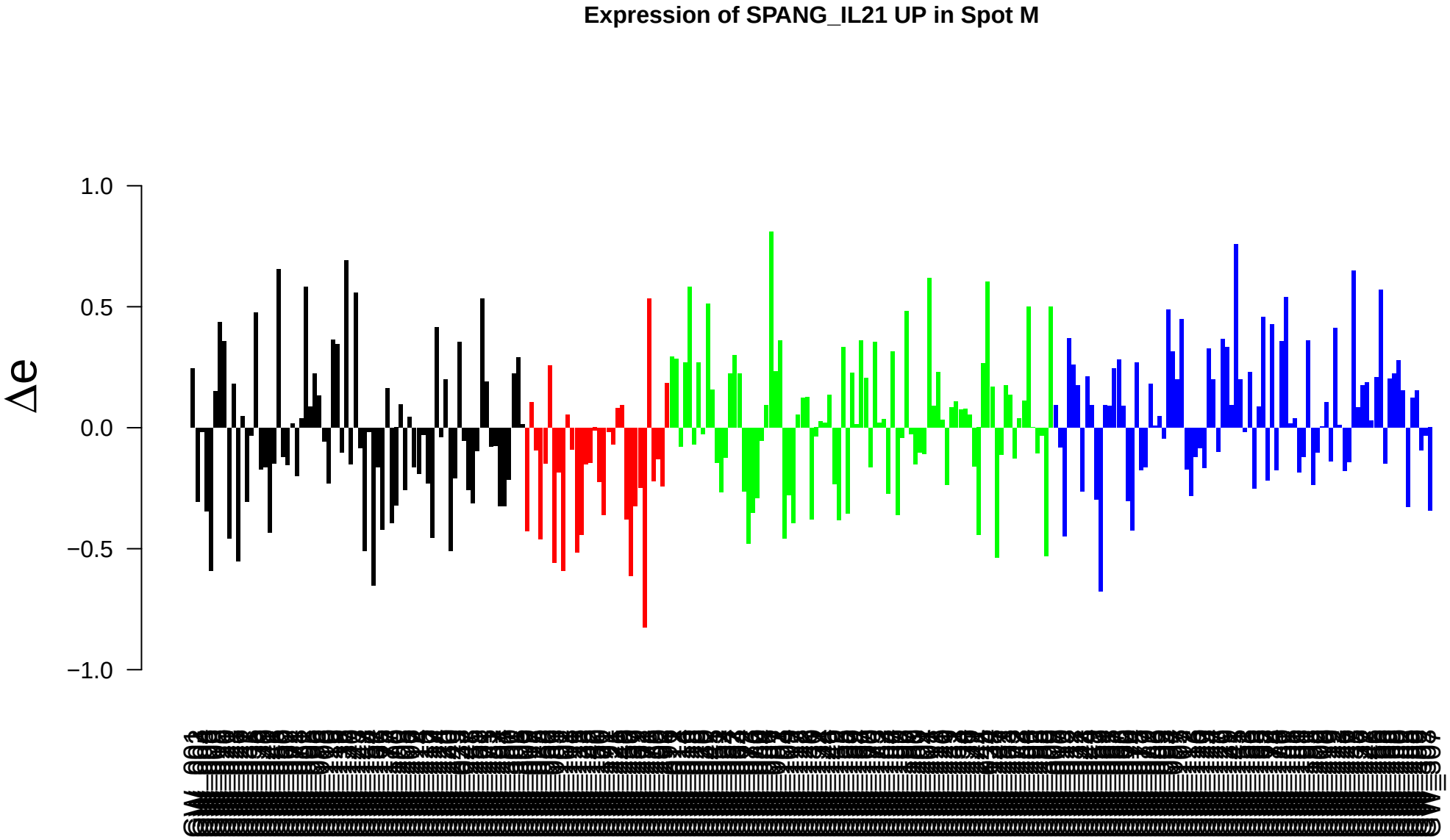

Expression of SPANG_IL21 UP in Spot H
Δe
1.0
0.5
0.0
−0.5
−1.0

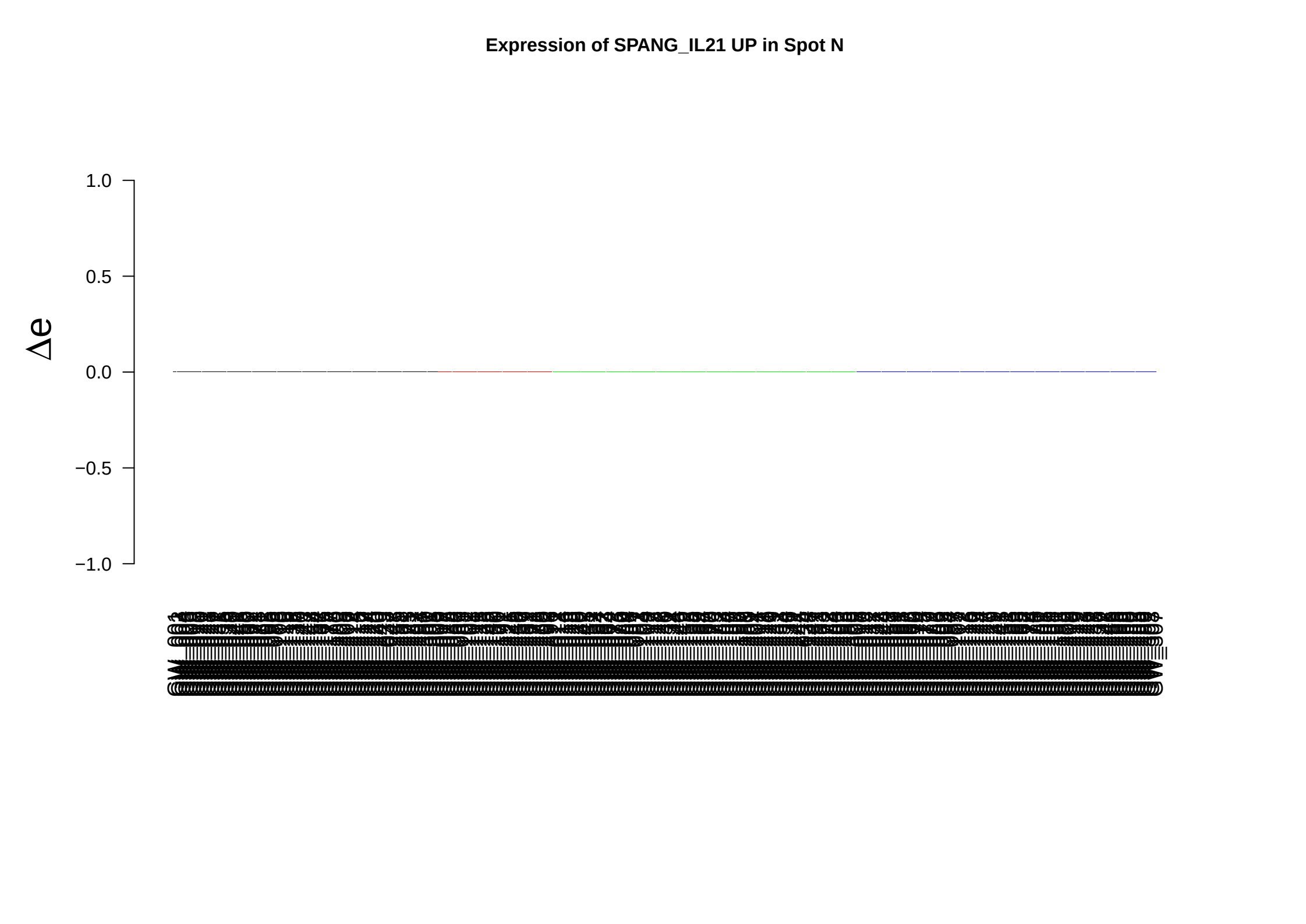

Expression of SPANG_IL21 UP in Spot I
Δe
1.0
0.5
0.0
−0.5
−1.0

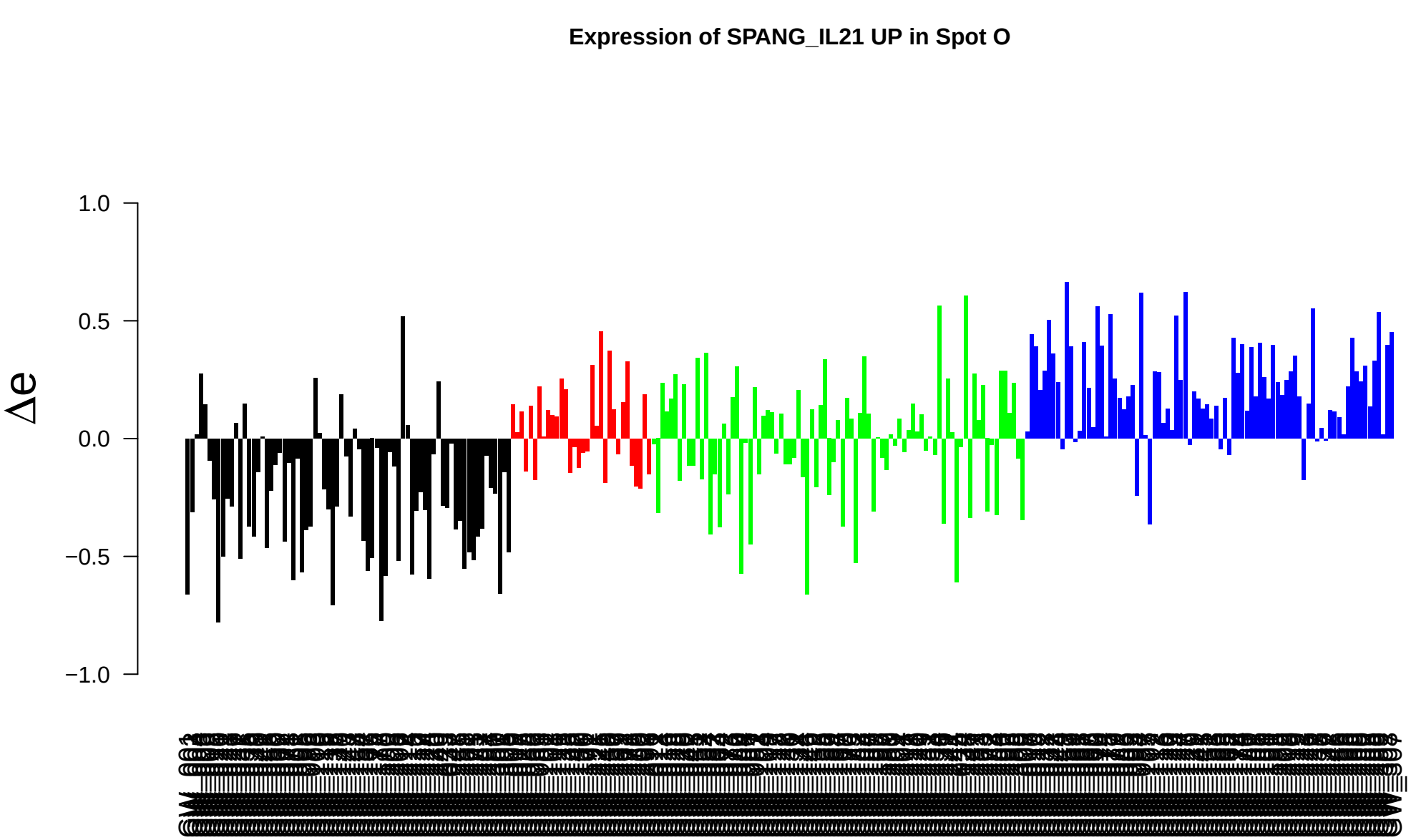

Expression of SPANG_IL21 UP in Spot J
Δe
1.0
0.5
0.0
−0.5
−1.0

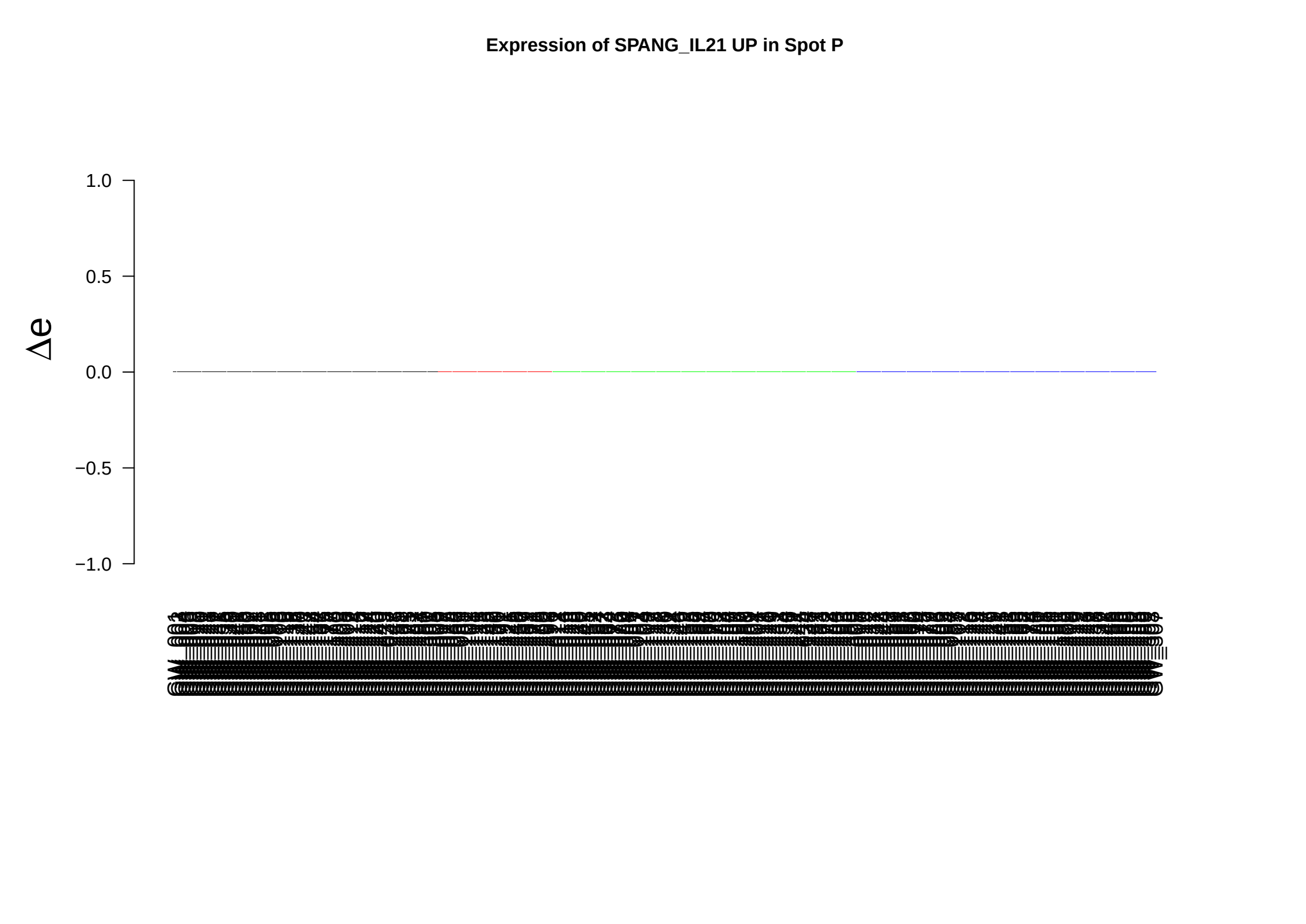

Expression of SPANG_IL21 UP in Spot K
Δe
1.0
0.5
0.0
−0.5
−1.0

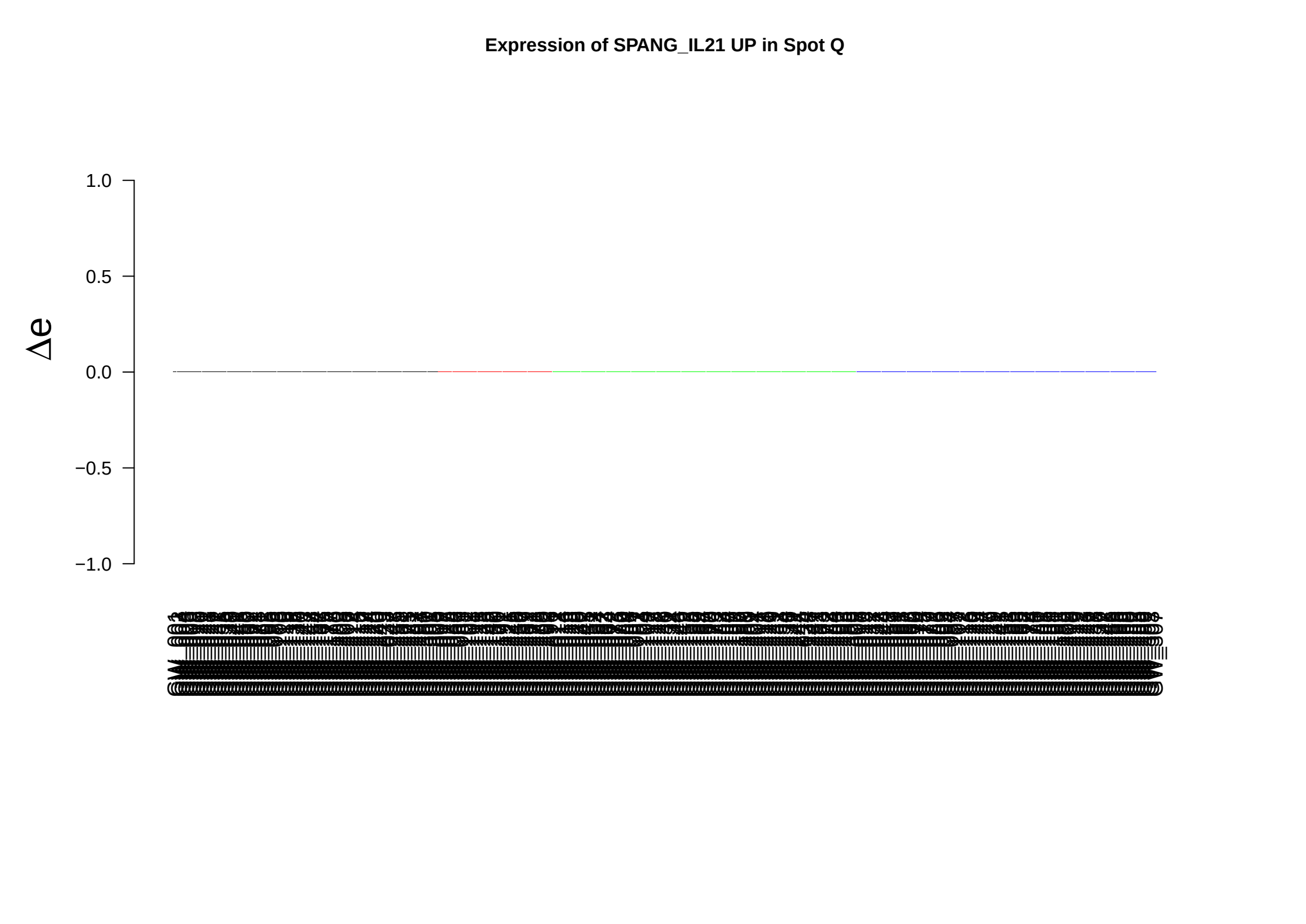

Expression of SPANG_IL21 UP in Spot L
Δe
1.0
0.5
0.0
−0.5
−1.0

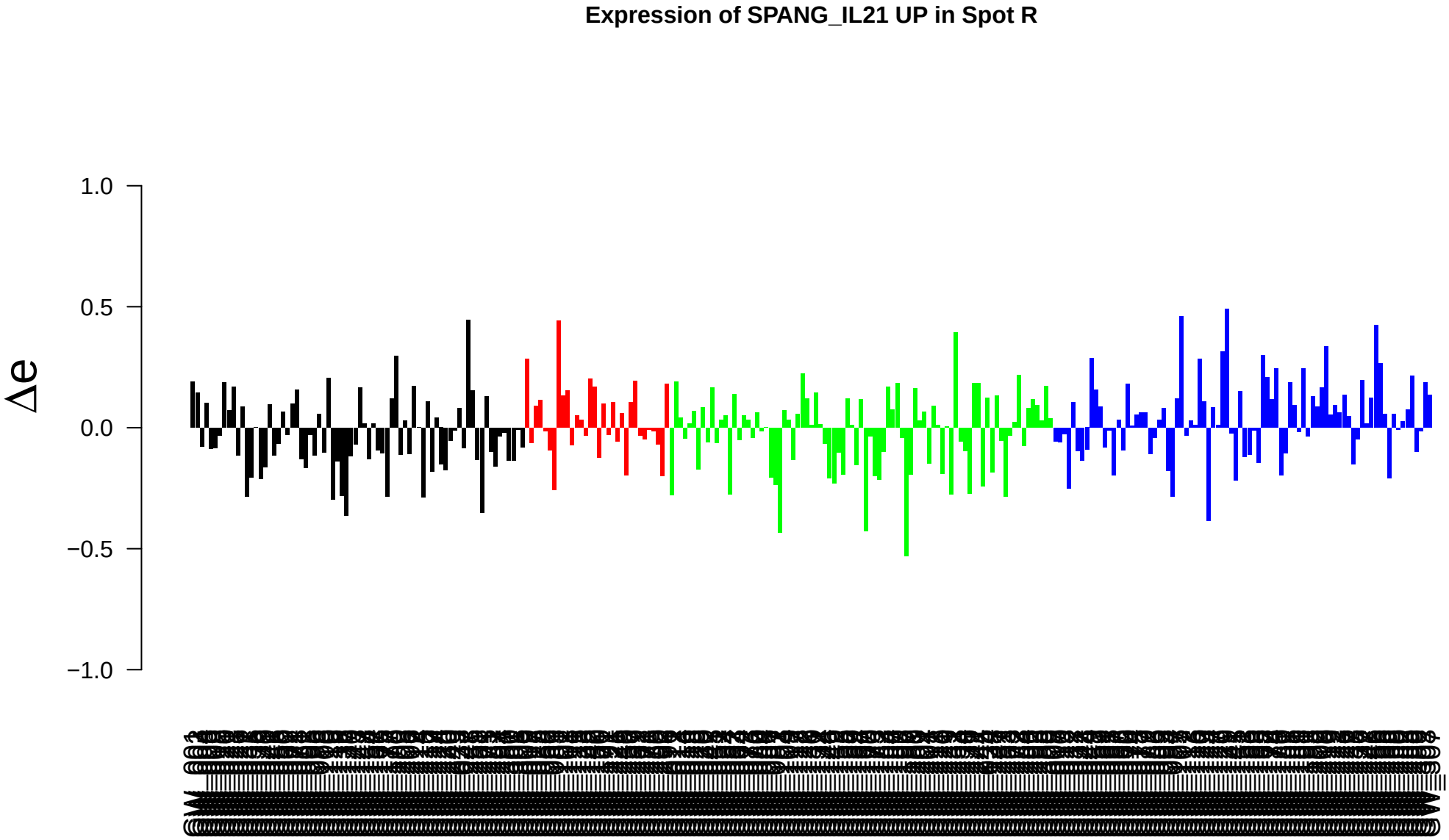

Expression of SPANG_IL21 UP in Spot M
Δe
1.0
0.5
0.0
−0.5
−1.0


Expression of SPANG_IL21 UP in Spot N
Δe
1.0
0.5
0.0
−0.5
−1.0


Expression of SPANG_IL21 UP in Spot O
Δe
1.0
0.5
0.0
−0.5
−1.0


Expression of SPANG_IL21 UP in Spot P
Δe
1.0
0.5
0.0
−0.5
−1.0


Expression of SPANG_IL21 UP in Spot Q
Δe
1.0
0.5
0.0
−0.5
−1.0


Expression of SPANG_IL21 UP in Spot R
Δe
1.0
0.5
0.0
−0.5
−1.0