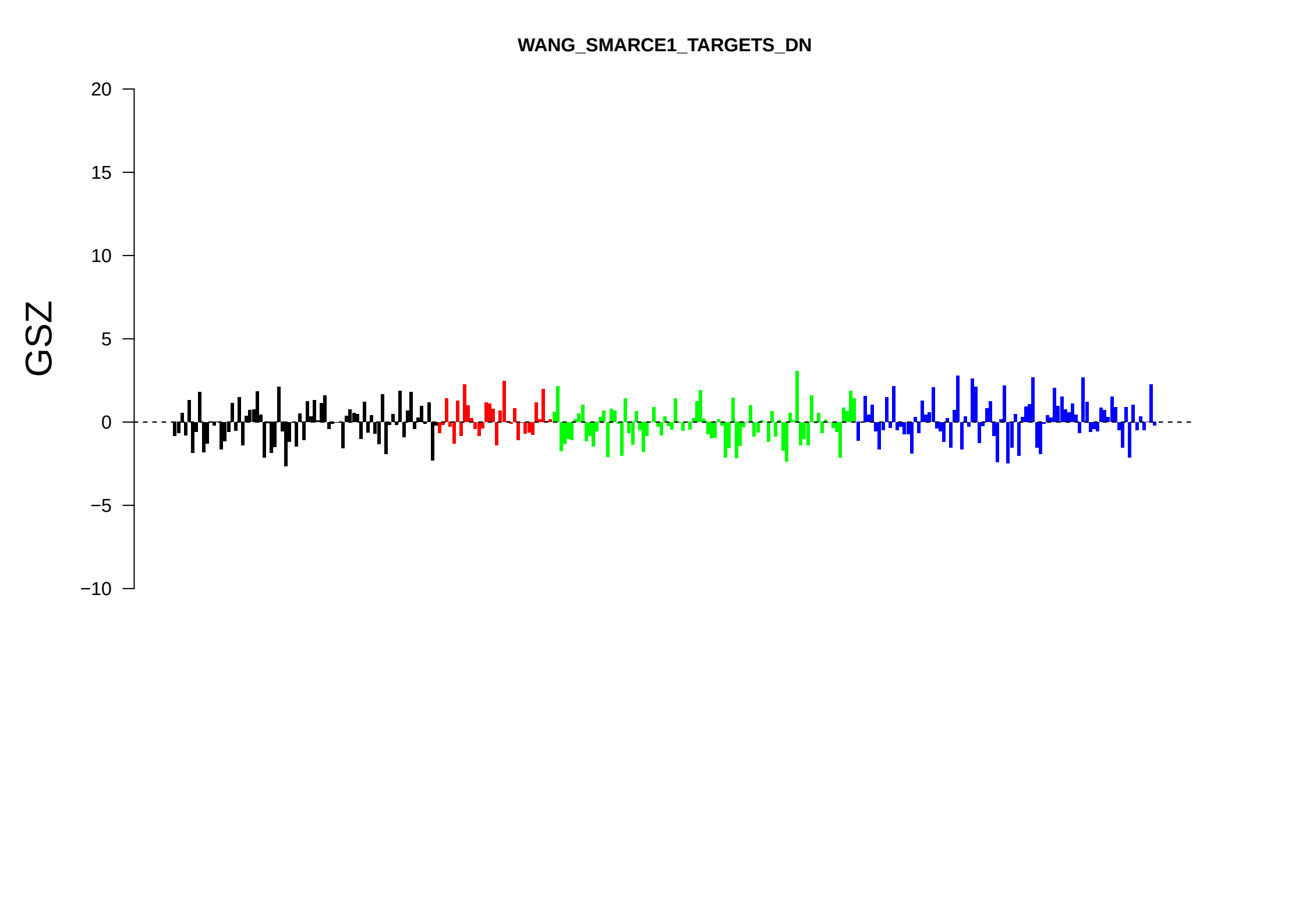
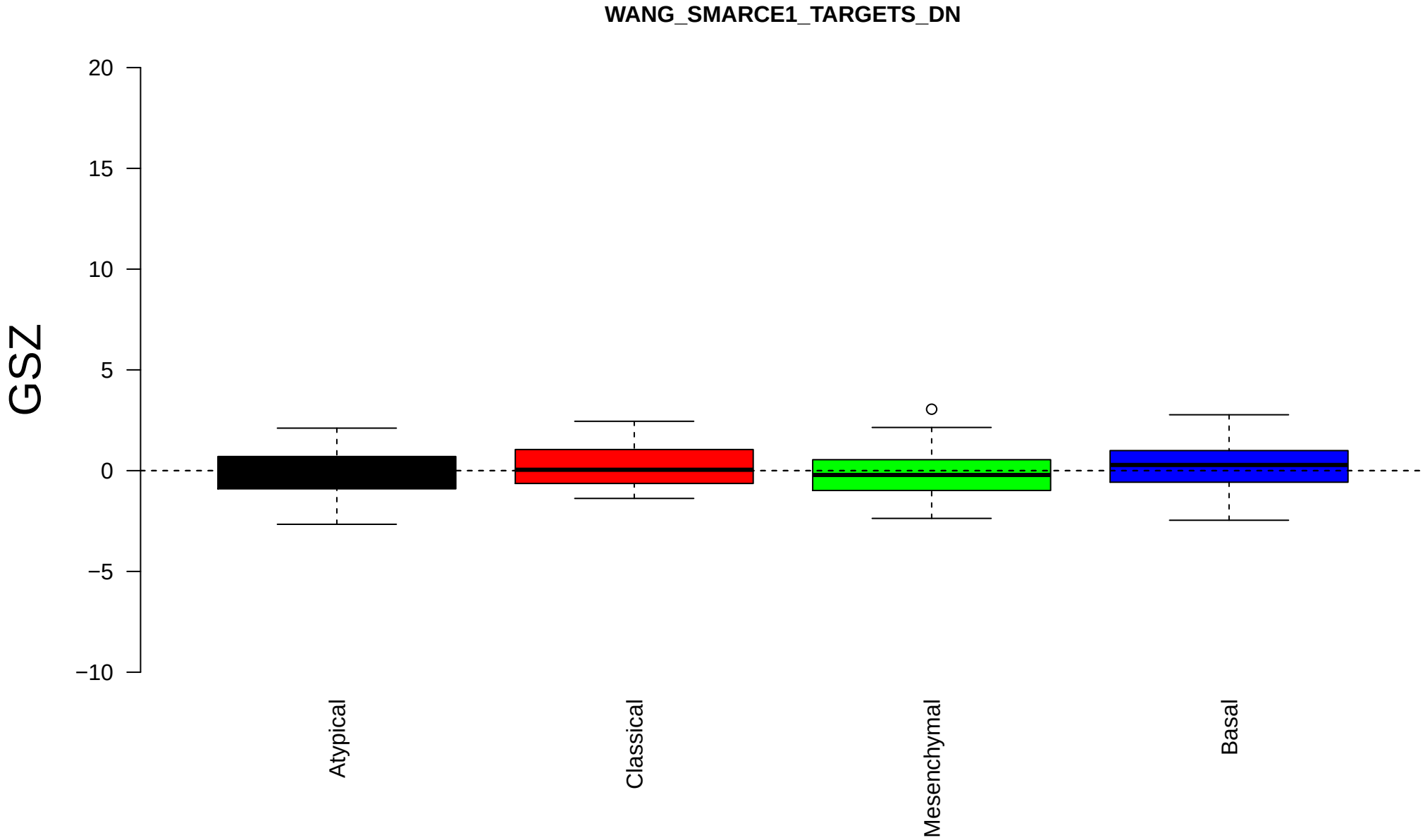
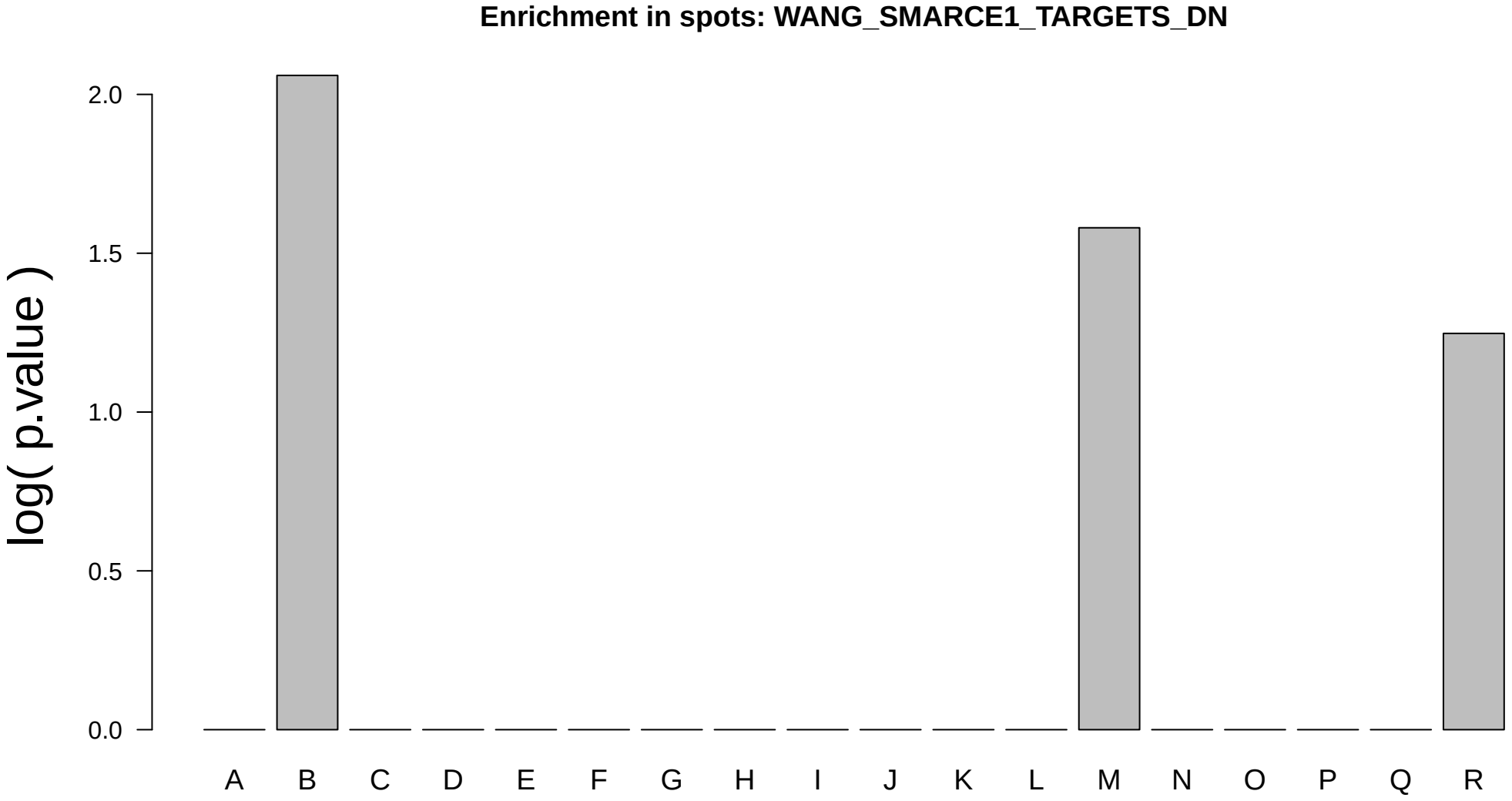
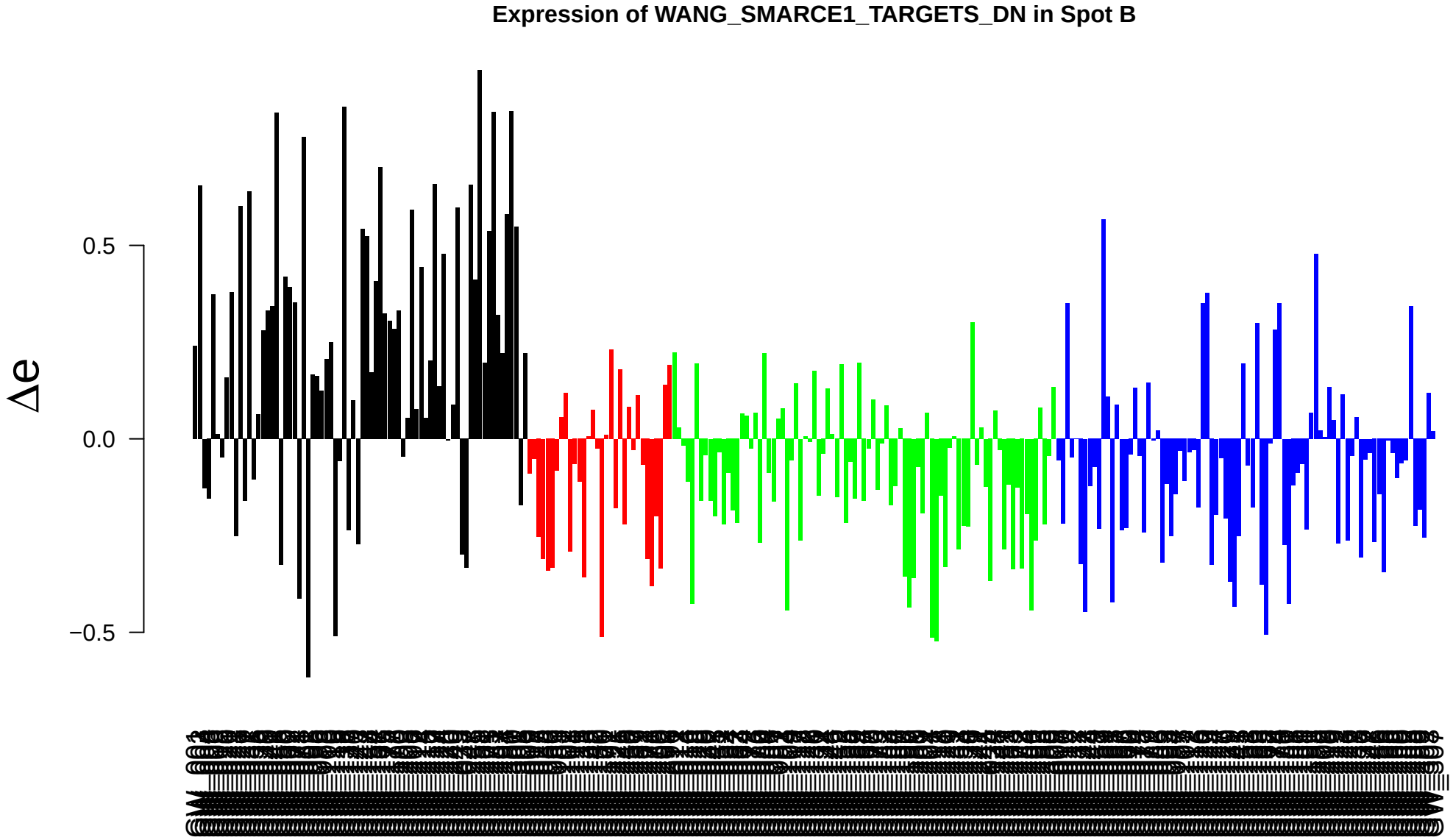
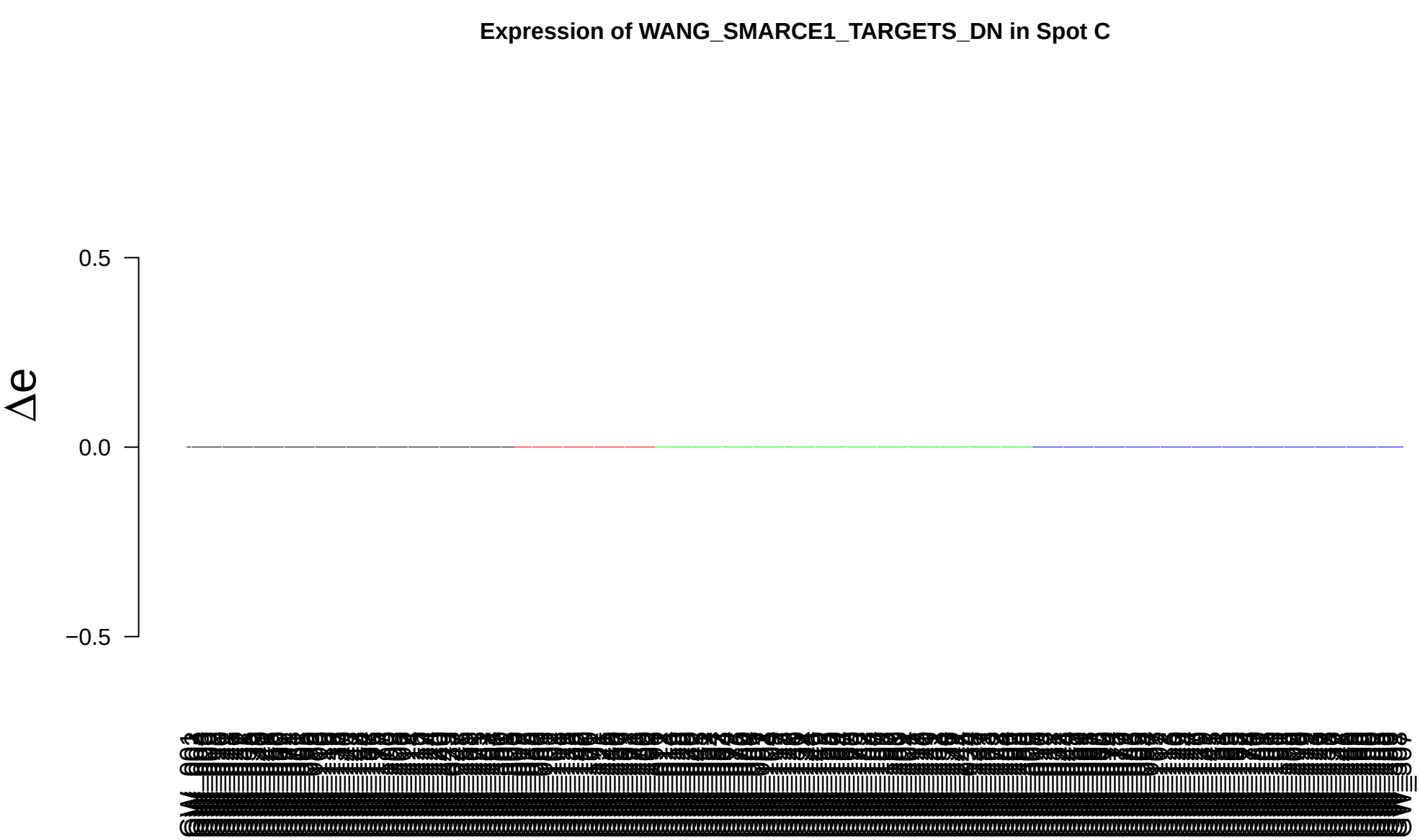

WANG_SMARCE1_TARGETS_DN
GSZ
20
15
10
5
0
−5
−10

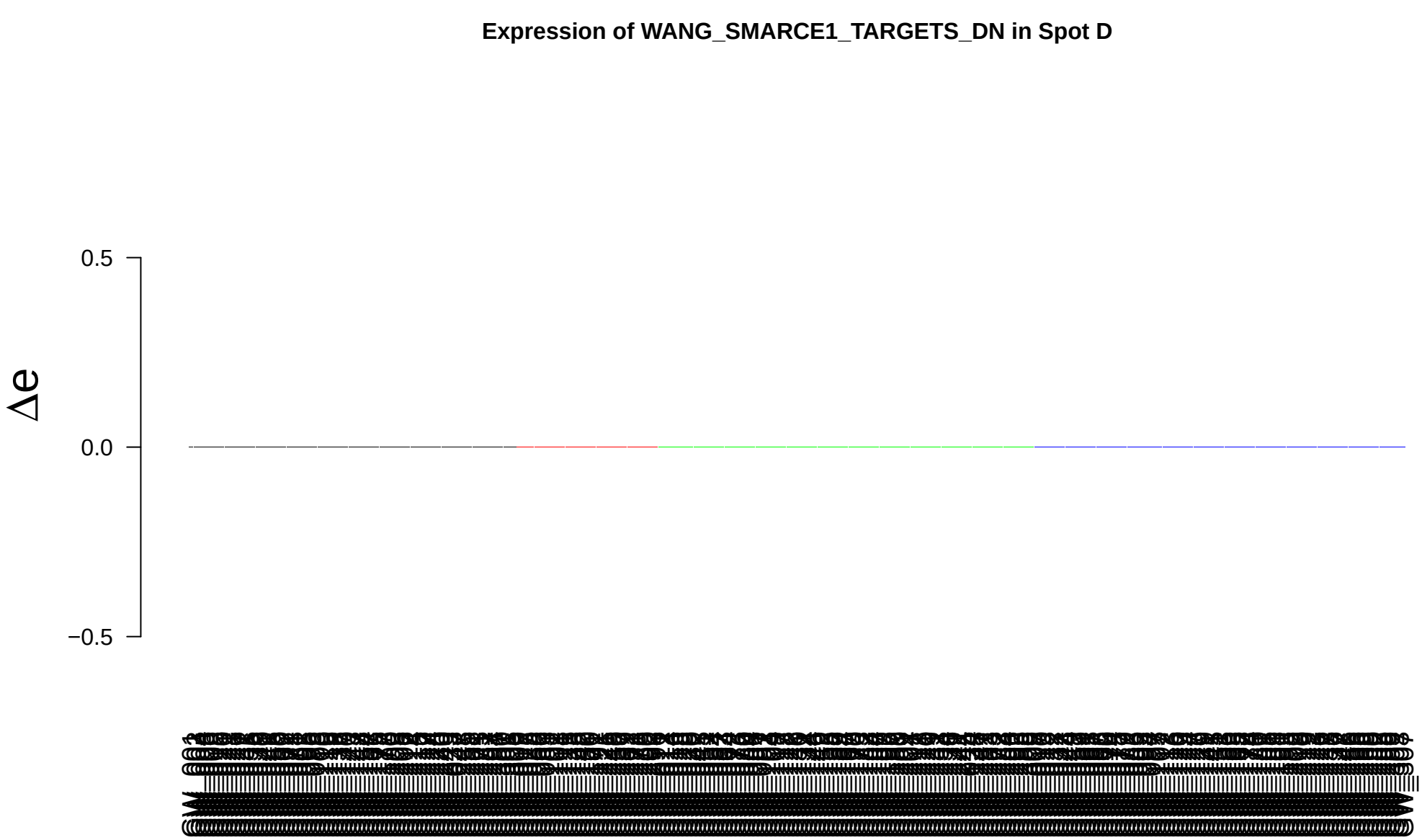

WANG_SMARCE1_TARGETS_DN
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal

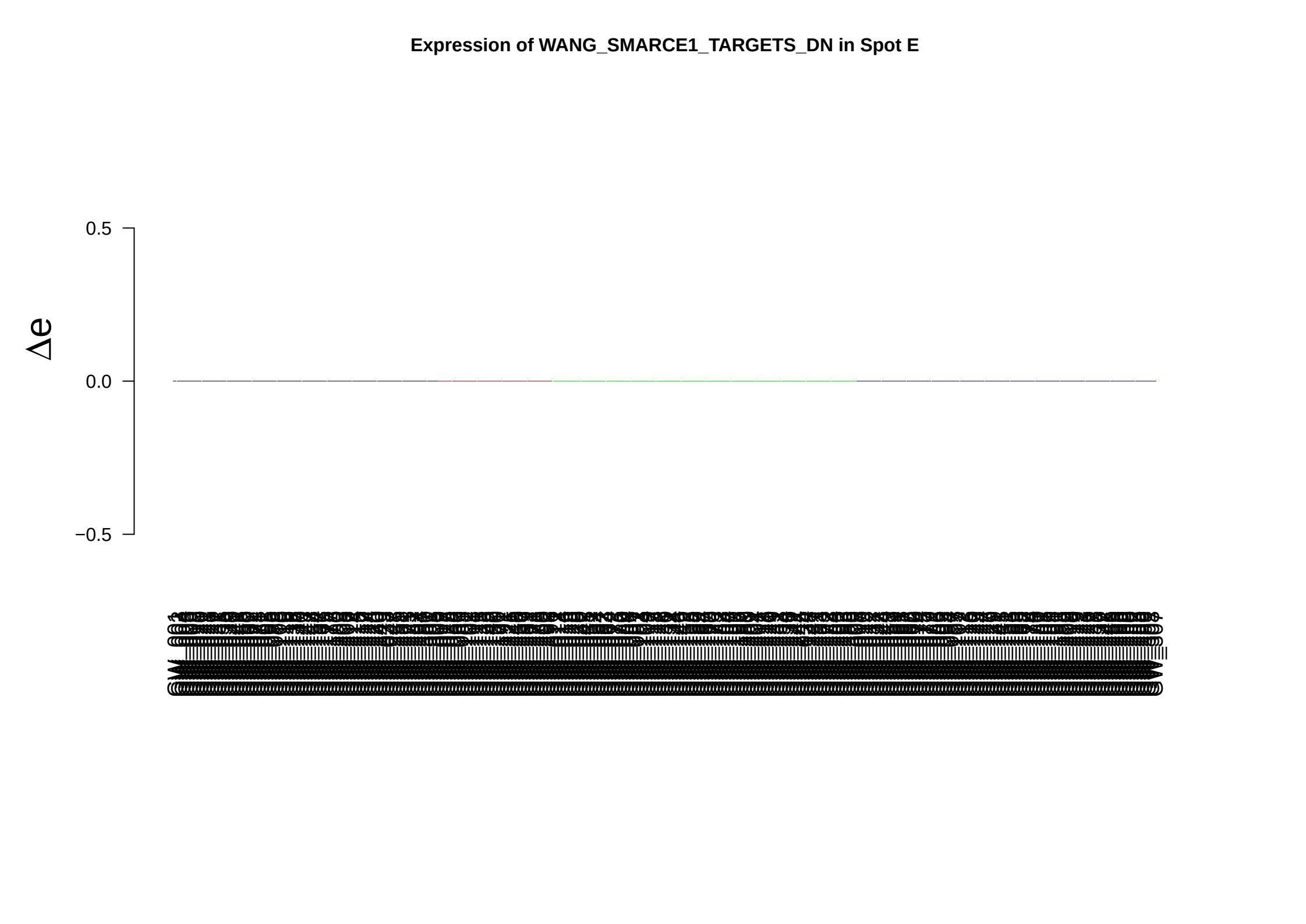

Enrichment in spots: WANG_SMARCE1_TARGETS_DN
log(p.value)
2.0
1.5
1.0
0.5
0.0
A
B
C
D
E
F
G
H
I
J
K
L
M
N
O
P
Q
R

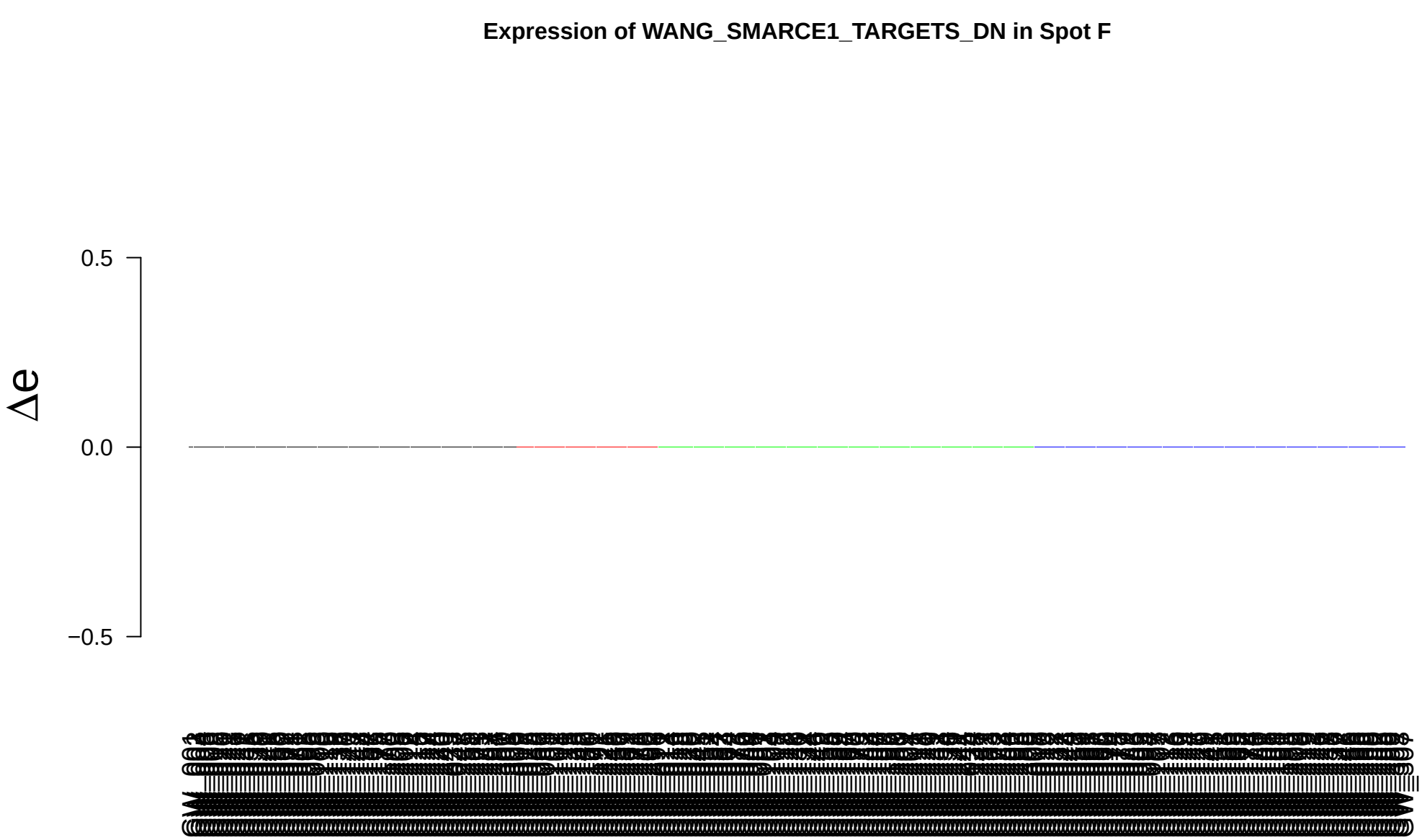
Expression of WANG_SMARCE1_TARGETS_DN in Spot A
Δe
0.5
0.0
−0.5

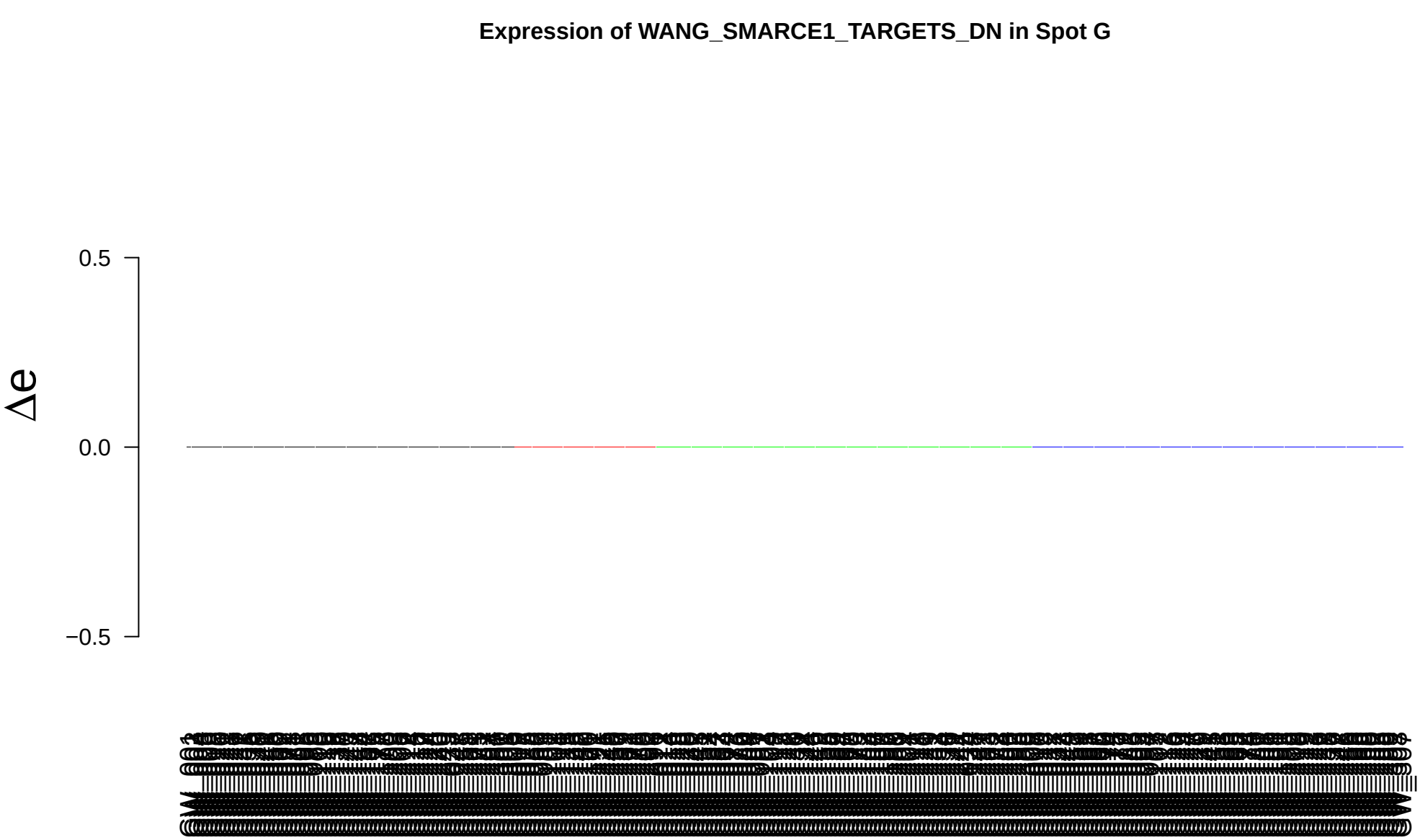

Expression of WANG_SMARCE1_TARGETS_DN in Spot B
Δe
0.5
0.0
−0.5


Expression of WANG_SMARCE1_TARGETS_DN in Spot C
Δe
0.5
0.0
−0.5

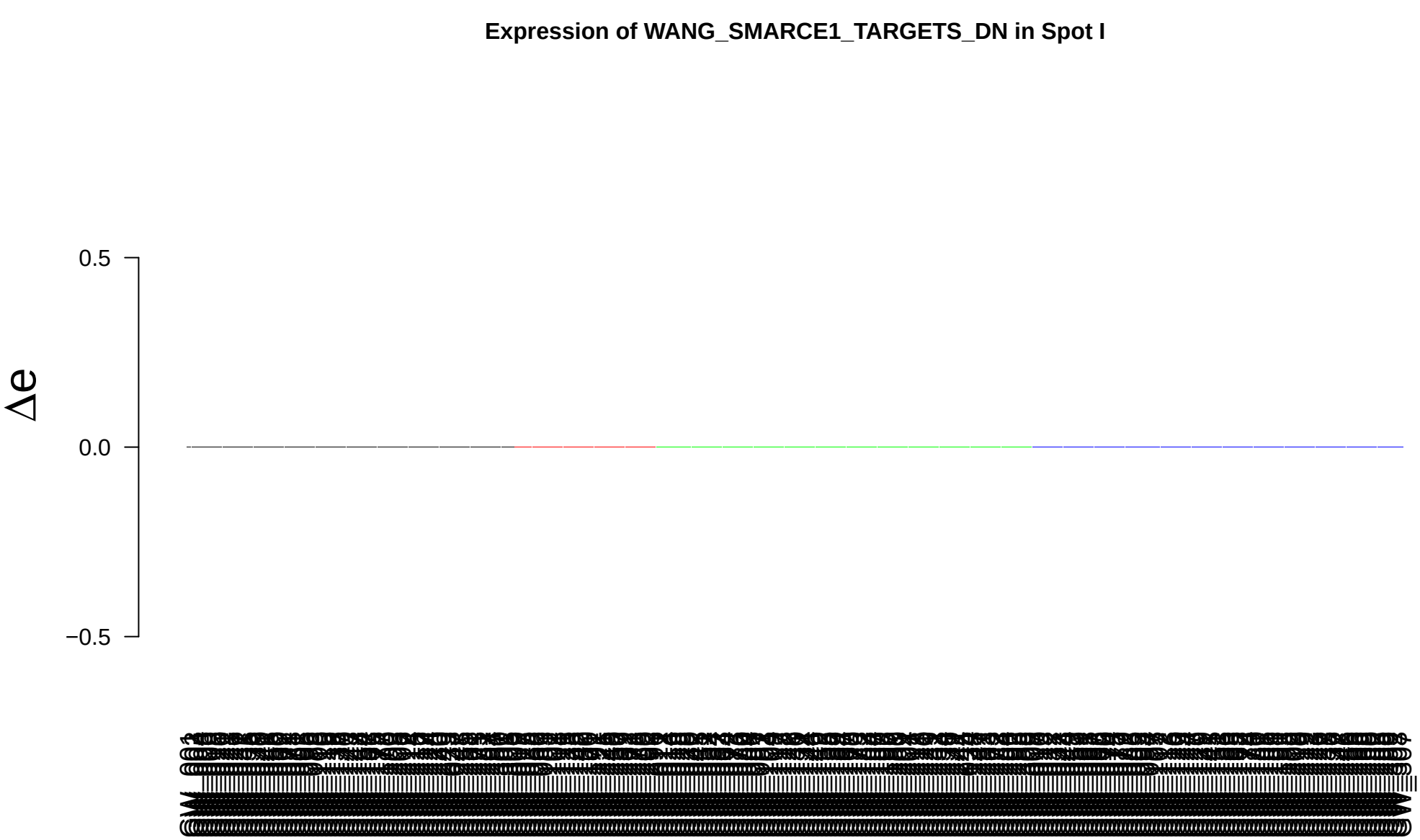

Expression of WANG_SMARCE1_TARGETS_DN in Spot D
Δe
0.5
0.0
−0.5

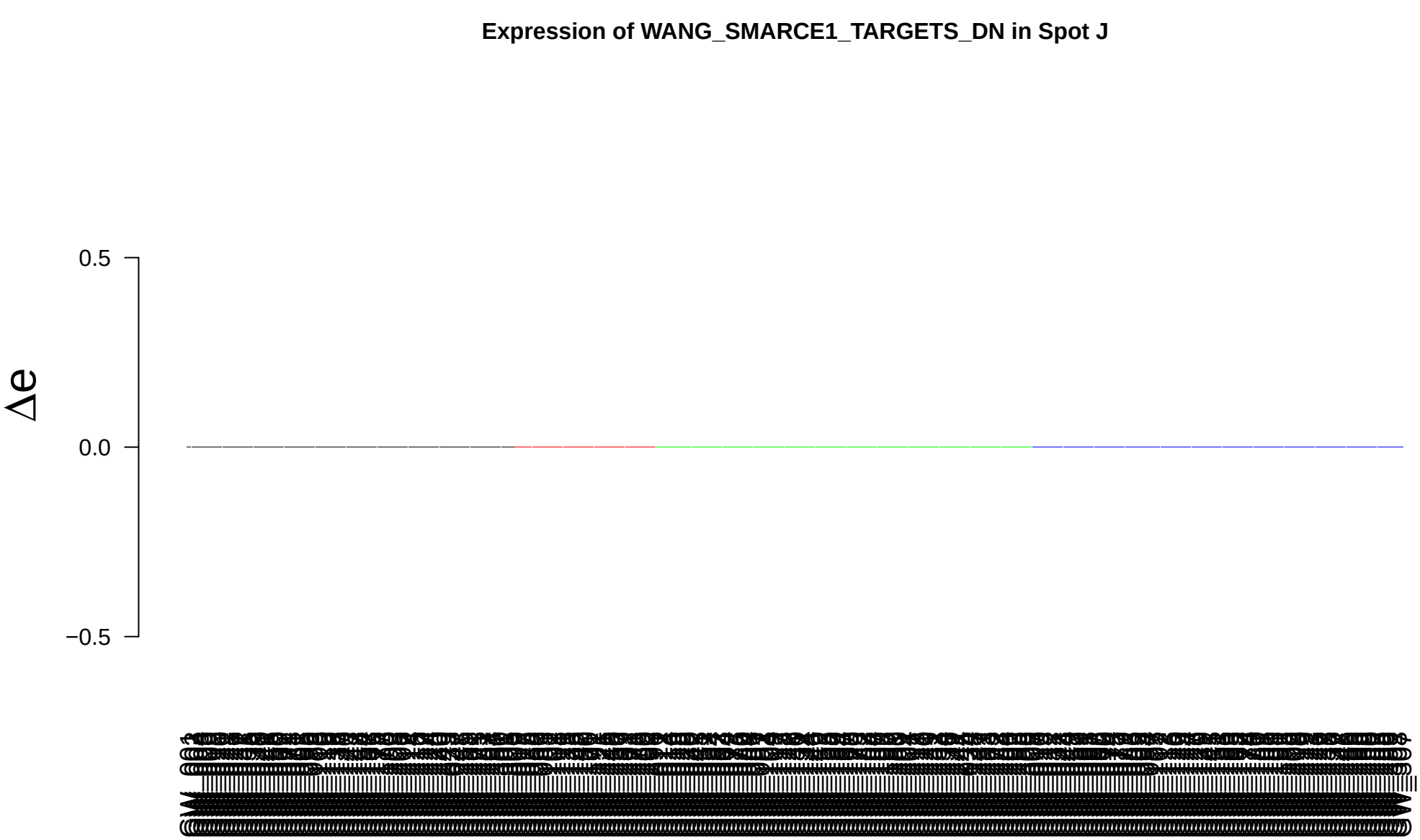

Expression of WANG_SMARCE1_TARGETS_DN in Spot E
Δe
0.5
0.0
−0.5


Expression of WANG_SMARCE1_TARGETS_DN in Spot F
Δe
0.5
0.0
−0.5

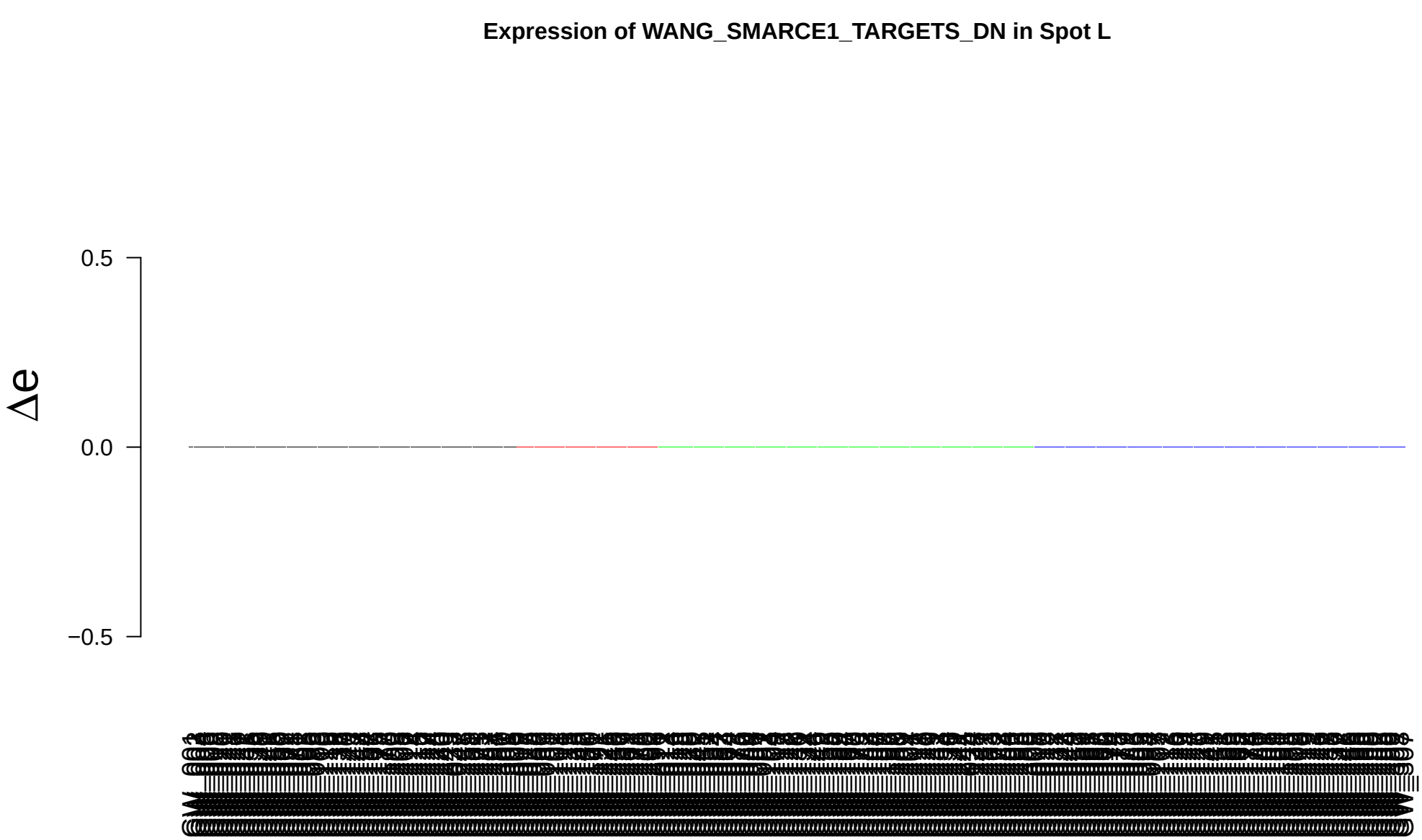

Expression of WANG_SMARCE1_TARGETS_DN in Spot G
Δe
0.5
0.0
−0.5

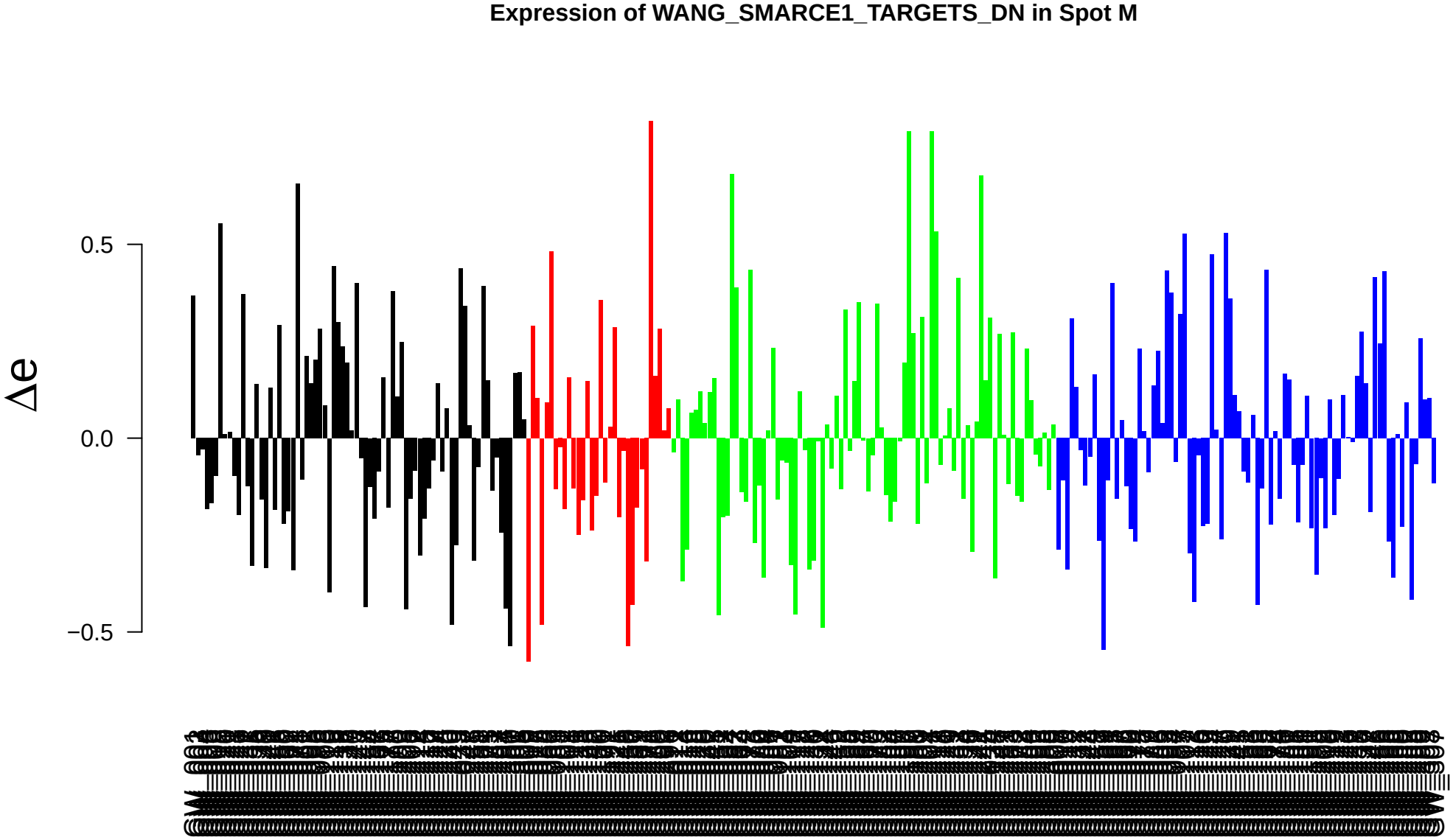
Expression of WANG_SMARCE1_TARGETS_DN in Spot H
Δe
0.5
0.0
−0.5

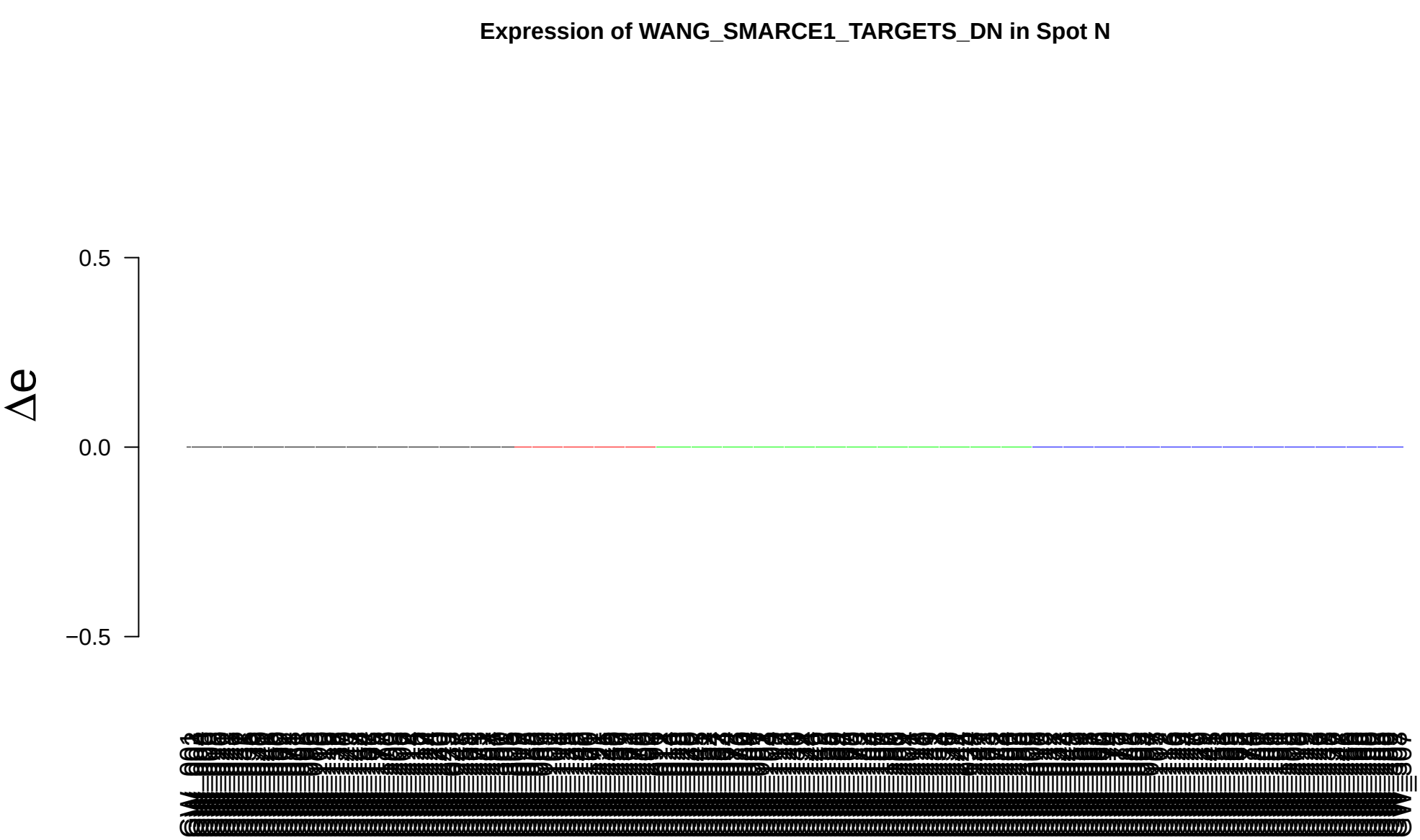

Expression of WANG_SMARCE1_TARGETS_DN in Spot I
Δe
0.5
0.0
−0.5

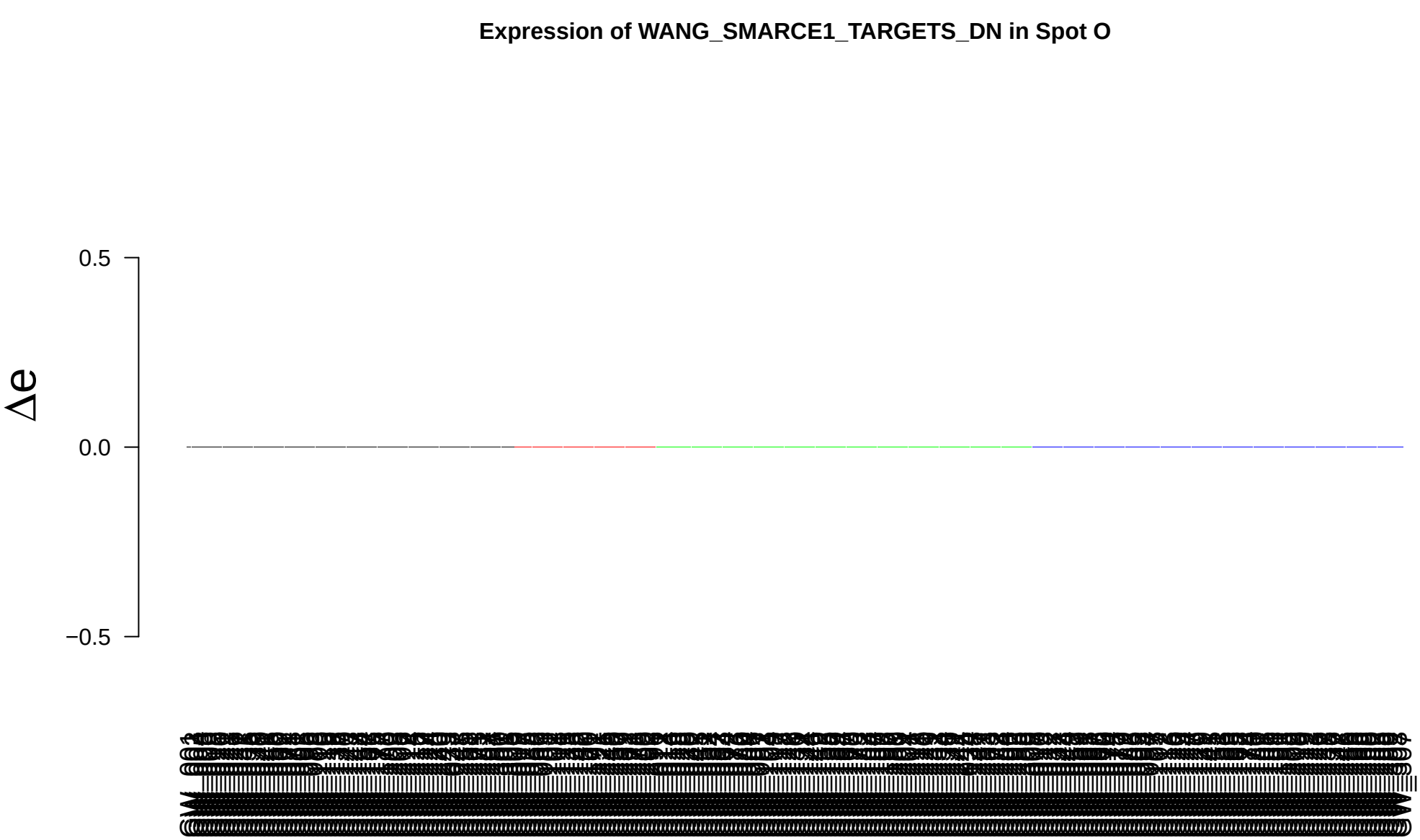

Expression of WANG_SMARCE1_TARGETS_DN in Spot J
Δe
0.5
0.0
−0.5

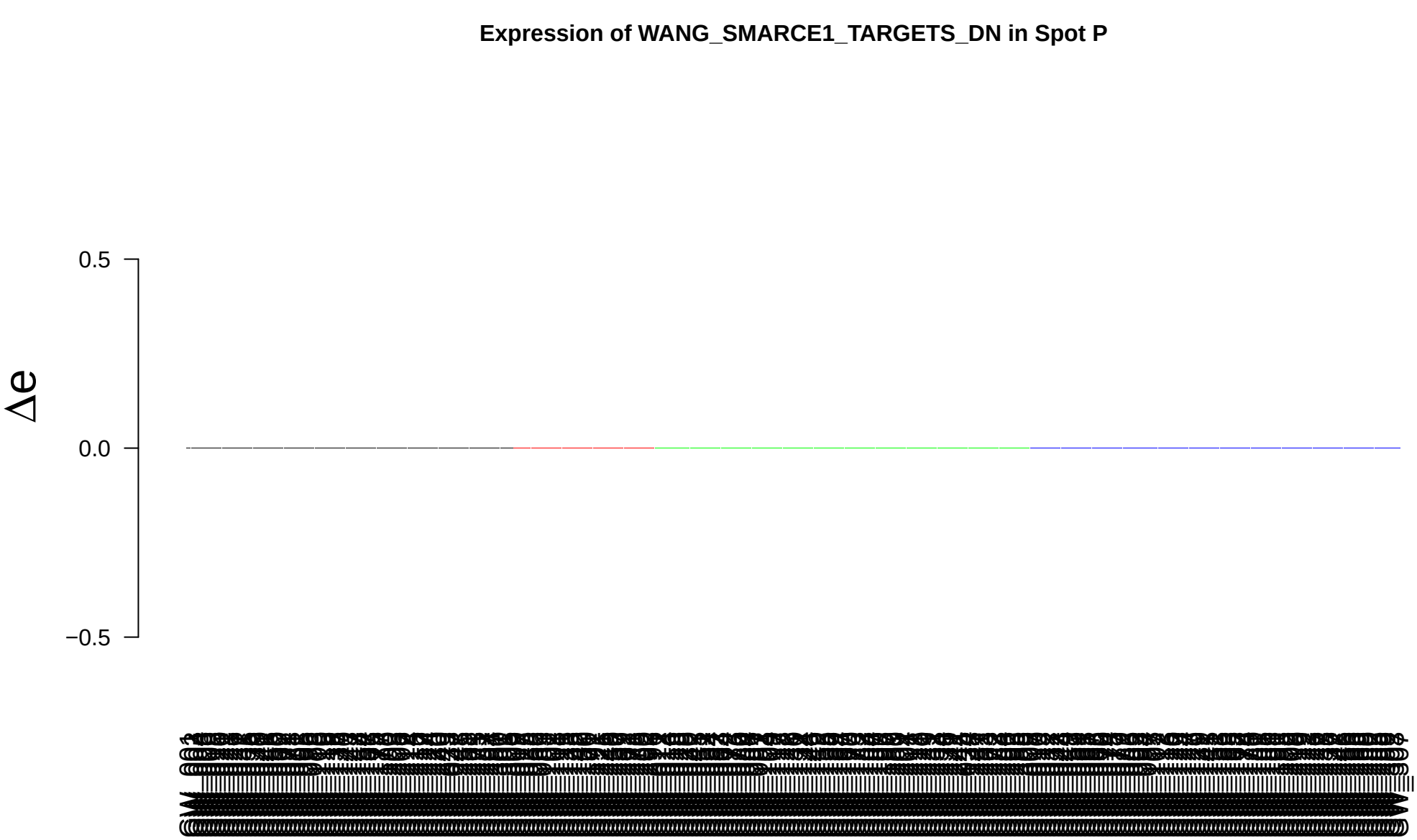
Expression of WANG_SMARCE1_TARGETS_DN in Spot K
Δe
0.5
0.0
−0.5

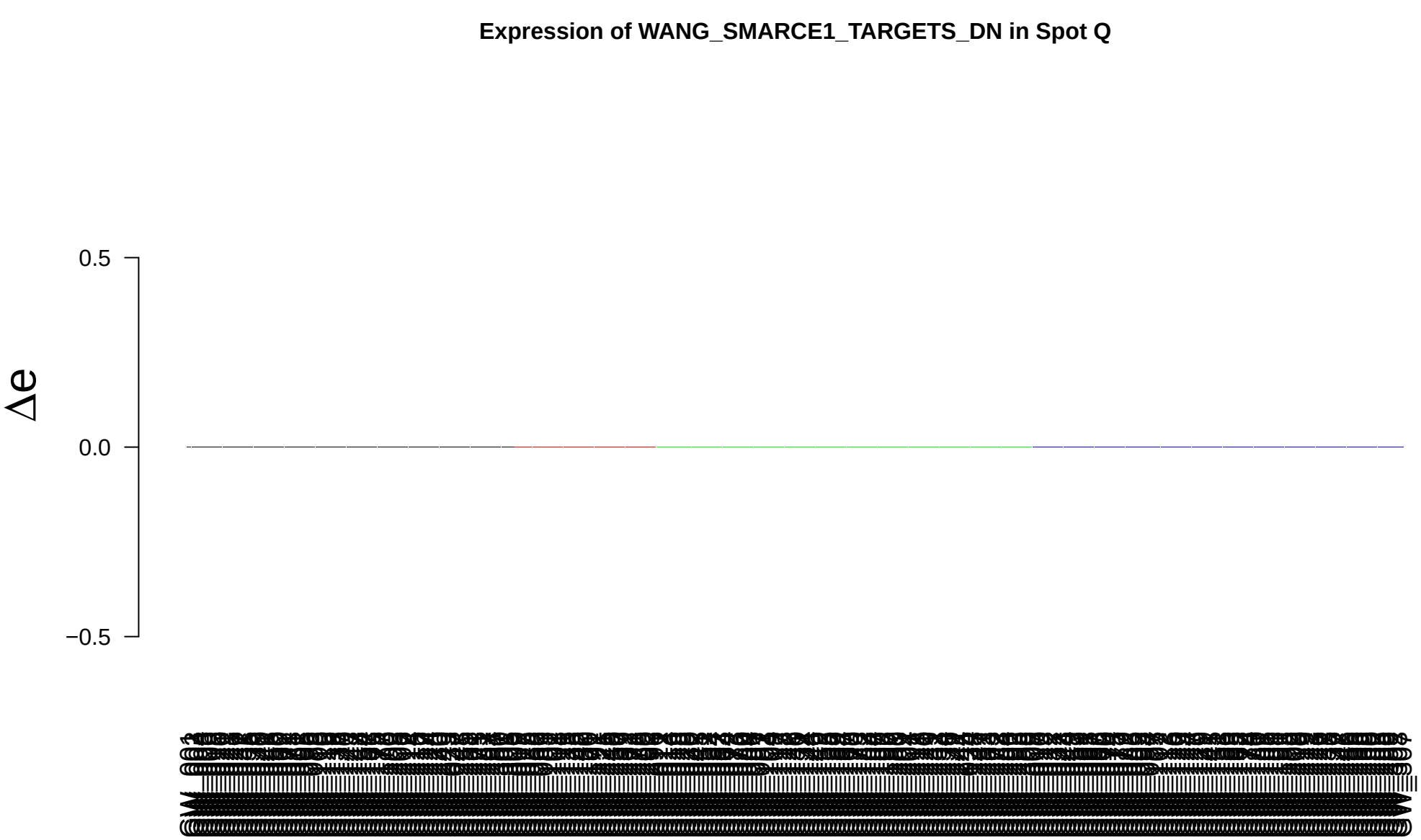

Expression of WANG_SMARCE1_TARGETS_DN in Spot L
Δe
0.5
0.0
−0.5

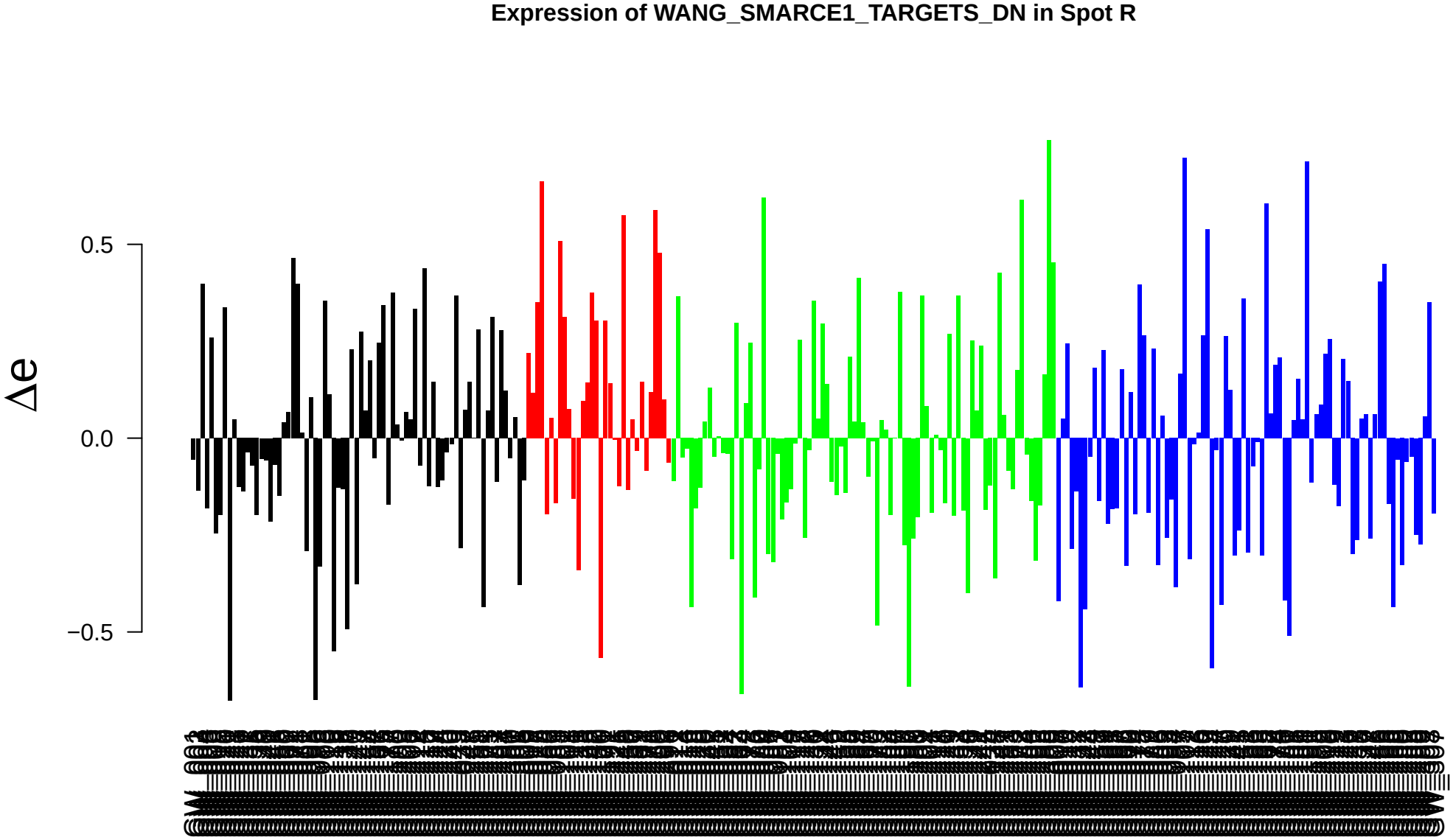

Expression of WANG_SMARCE1_TARGETS_DN in Spot M
Δe
0.5
0.0
−0.5


Expression of WANG_SMARCE1_TARGETS_DN in Spot N
Δe
0.5
0.0
−0.5


Expression of WANG_SMARCE1_TARGETS_DN in Spot O
Δe
0.5
0.0
−0.5


Expression of WANG_SMARCE1_TARGETS_DN in Spot P
Δe
0.5
0.0
−0.5


Expression of WANG_SMARCE1_TARGETS_DN in Spot Q
Δe
0.5
0.0
−0.5


Expression of WANG_SMARCE1_TARGETS_DN in Spot R
Δe
0.5
0.0
−0.5