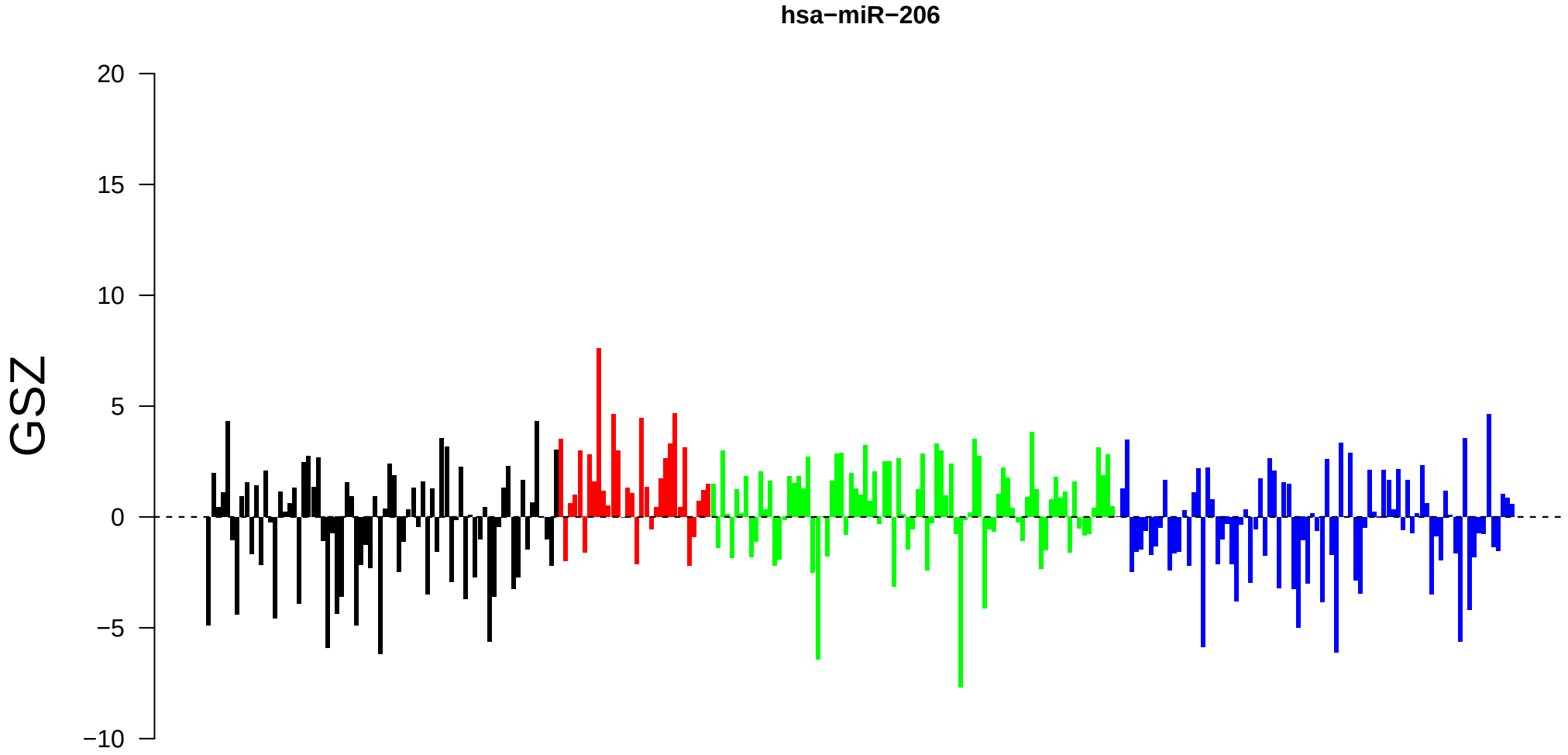
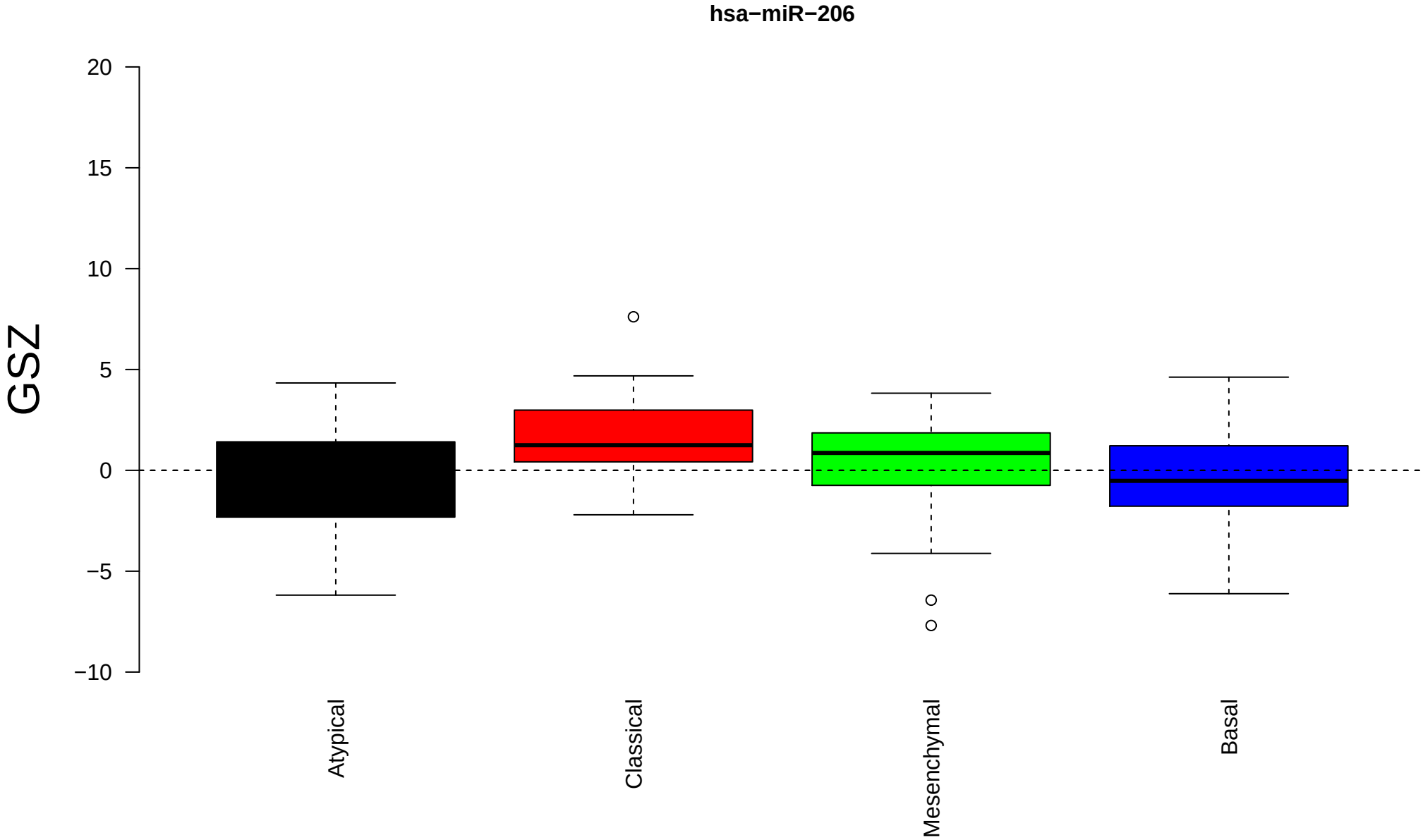
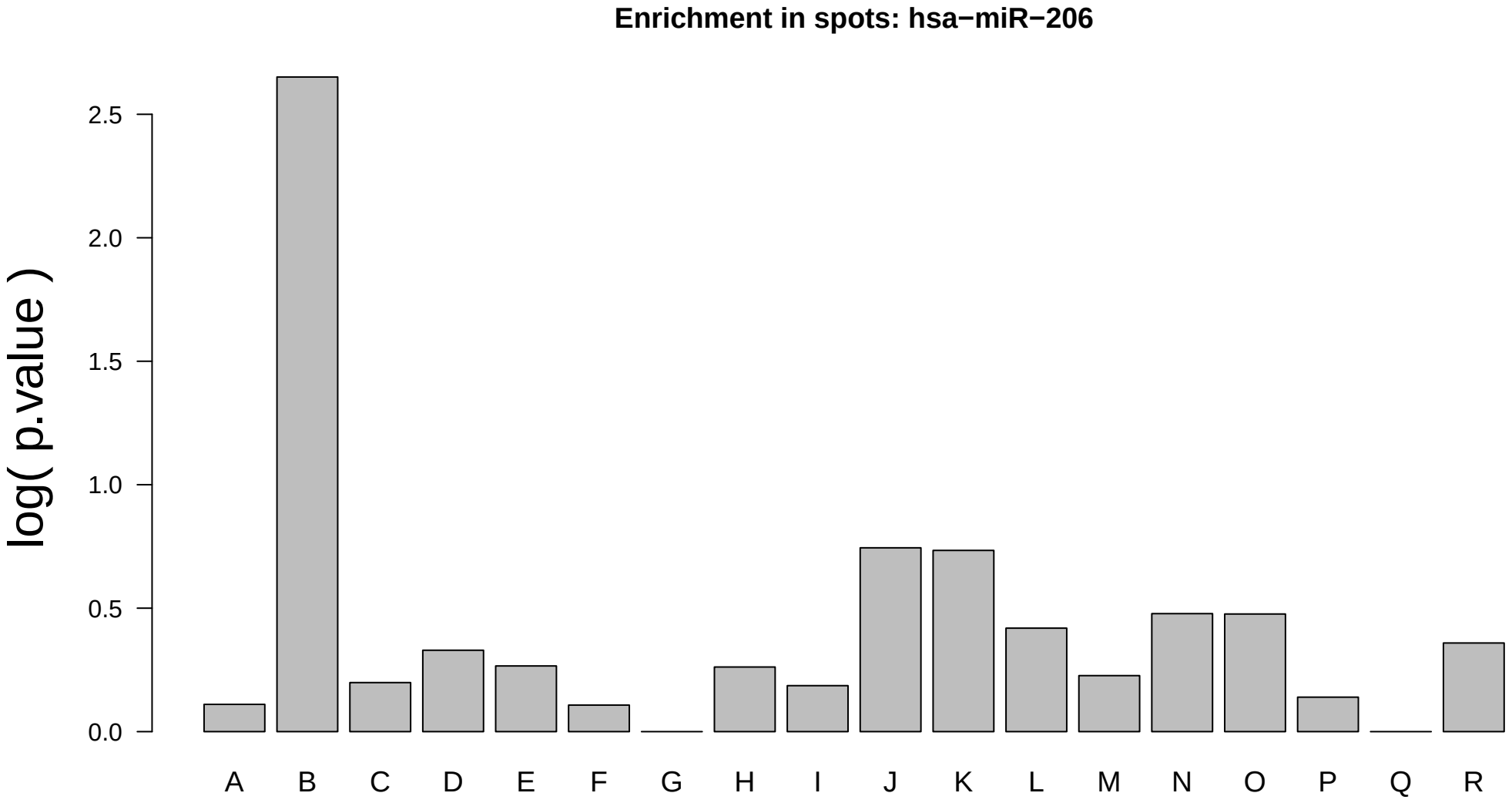
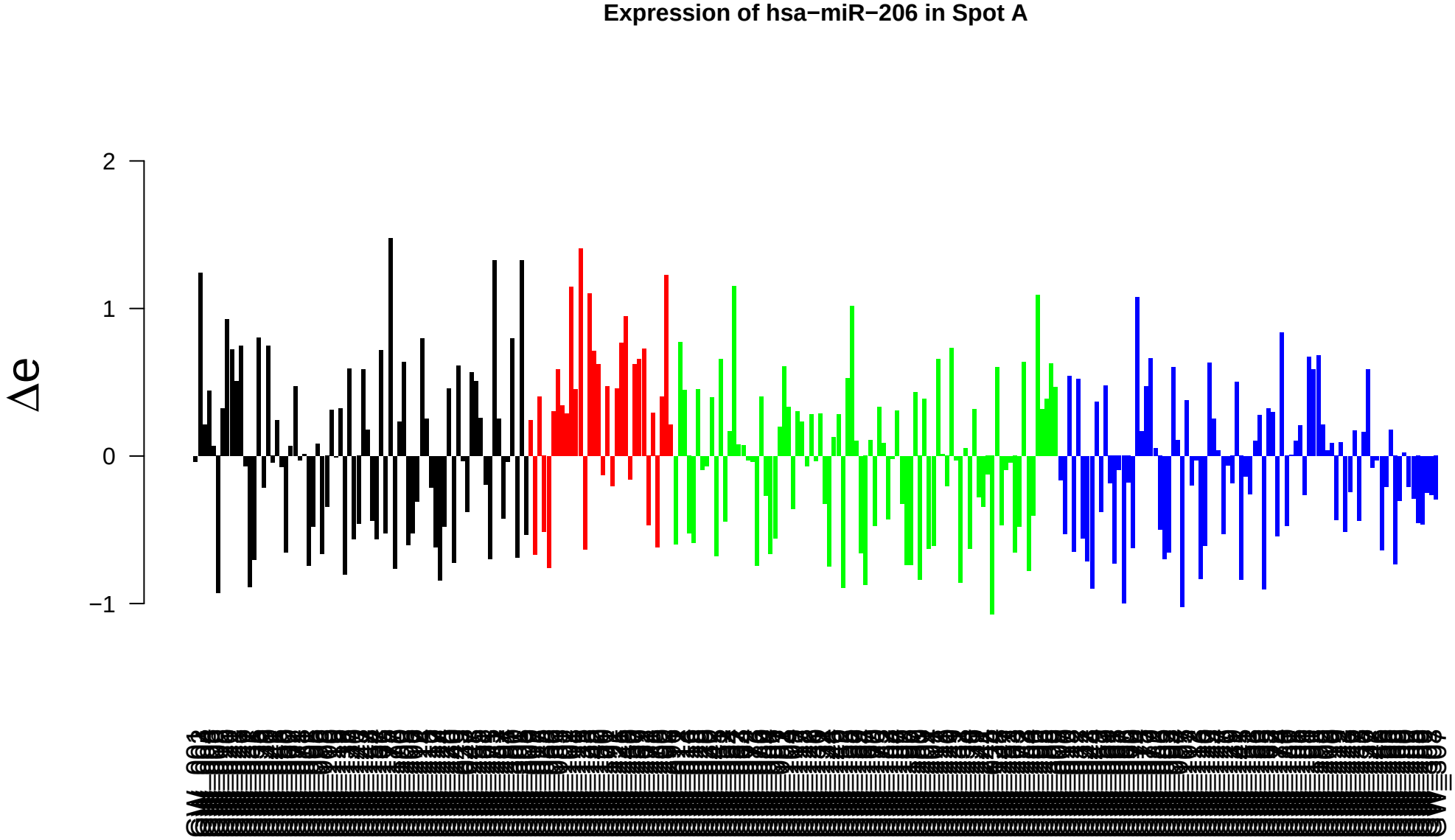
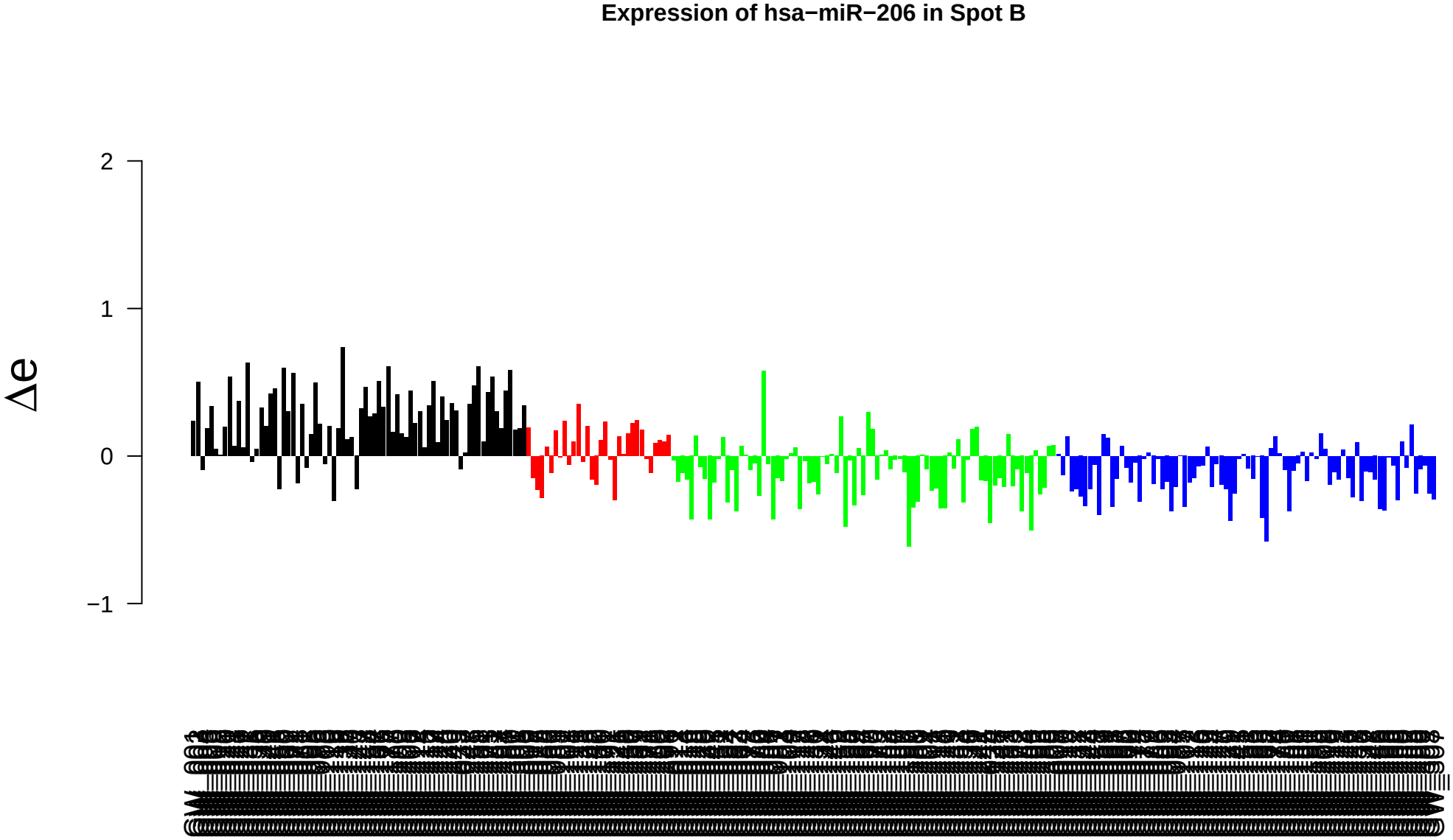
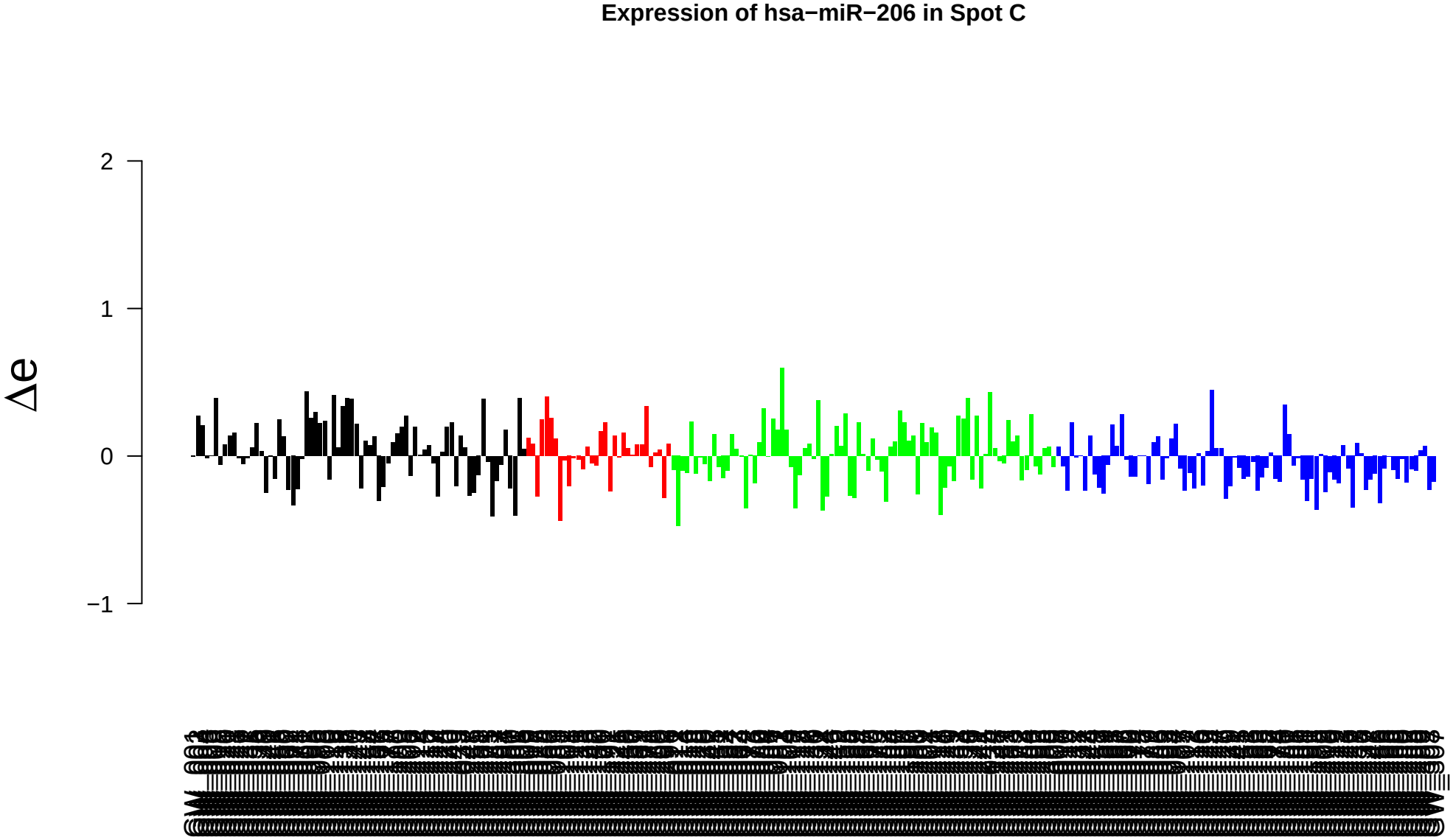

hsa−miR−206
GSZ
20
15
10
5
0
−5
−10

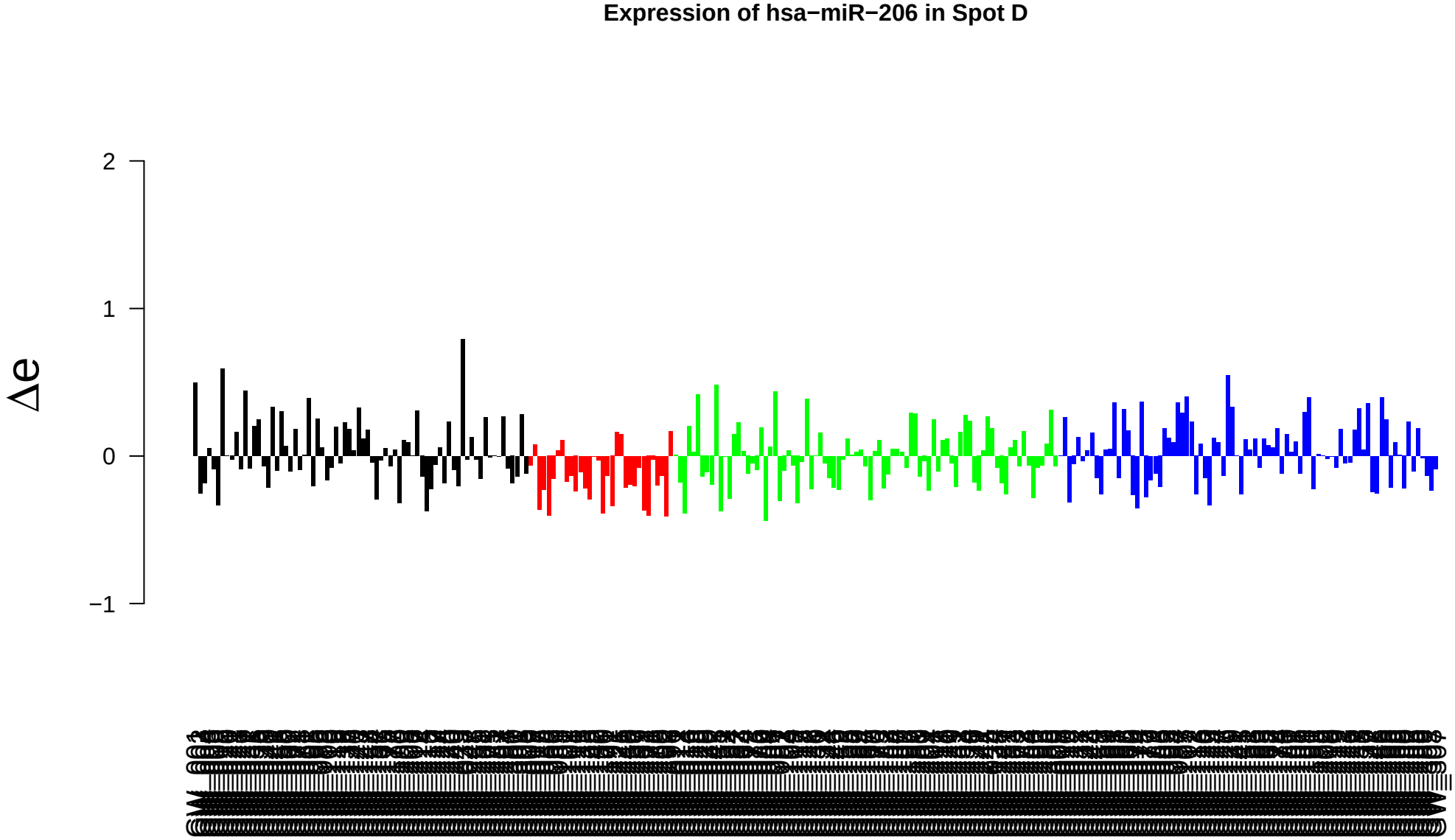

hsa−miR−206
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal

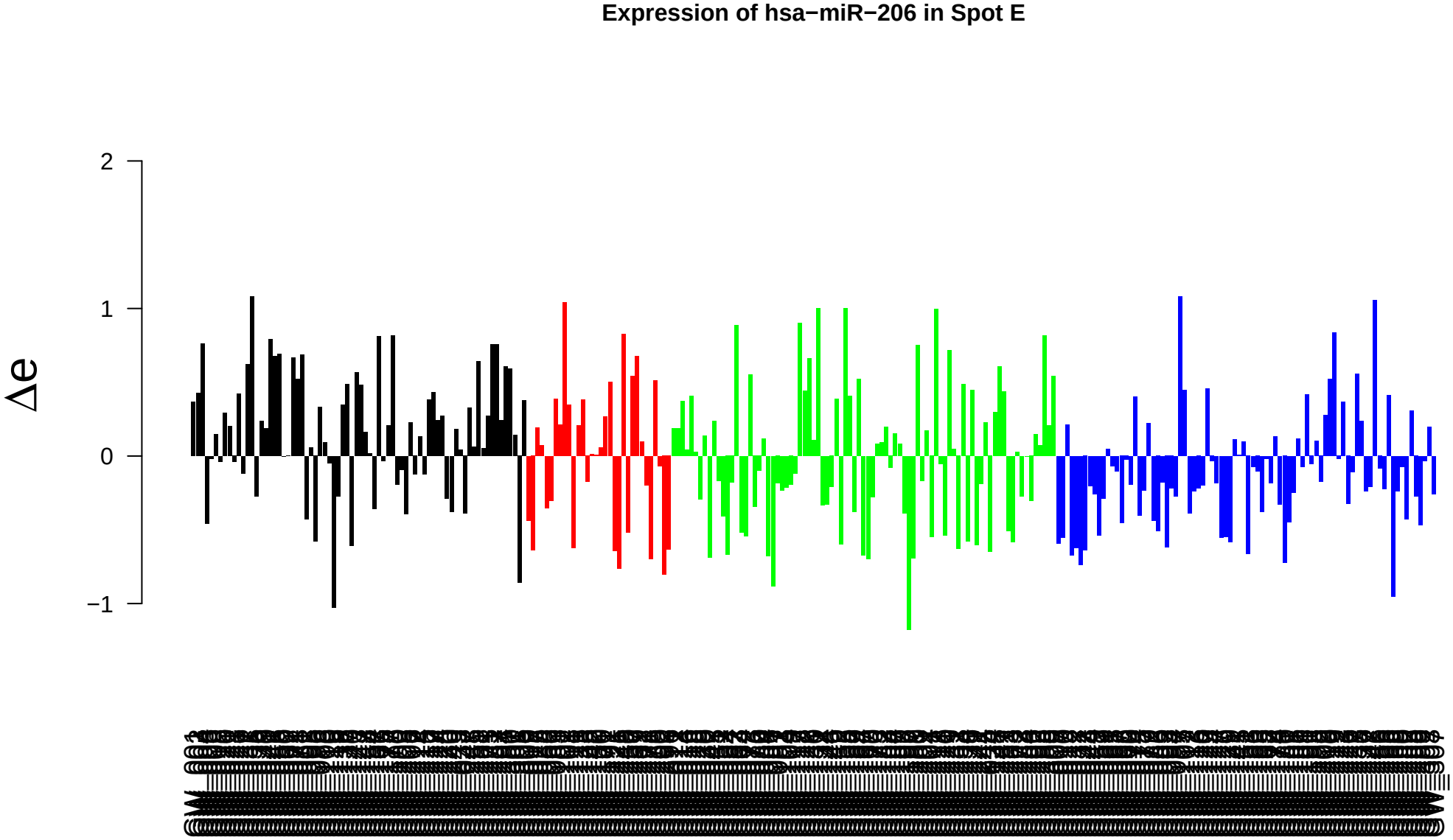

Enrichment in spots: hsa−miR−206
log(p.value)
2.5
2.0
1.5
1.0
0.5
0.0
A
B
C
D
E
F
G
H
I
J
K
L
M
N
O
P
Q
R

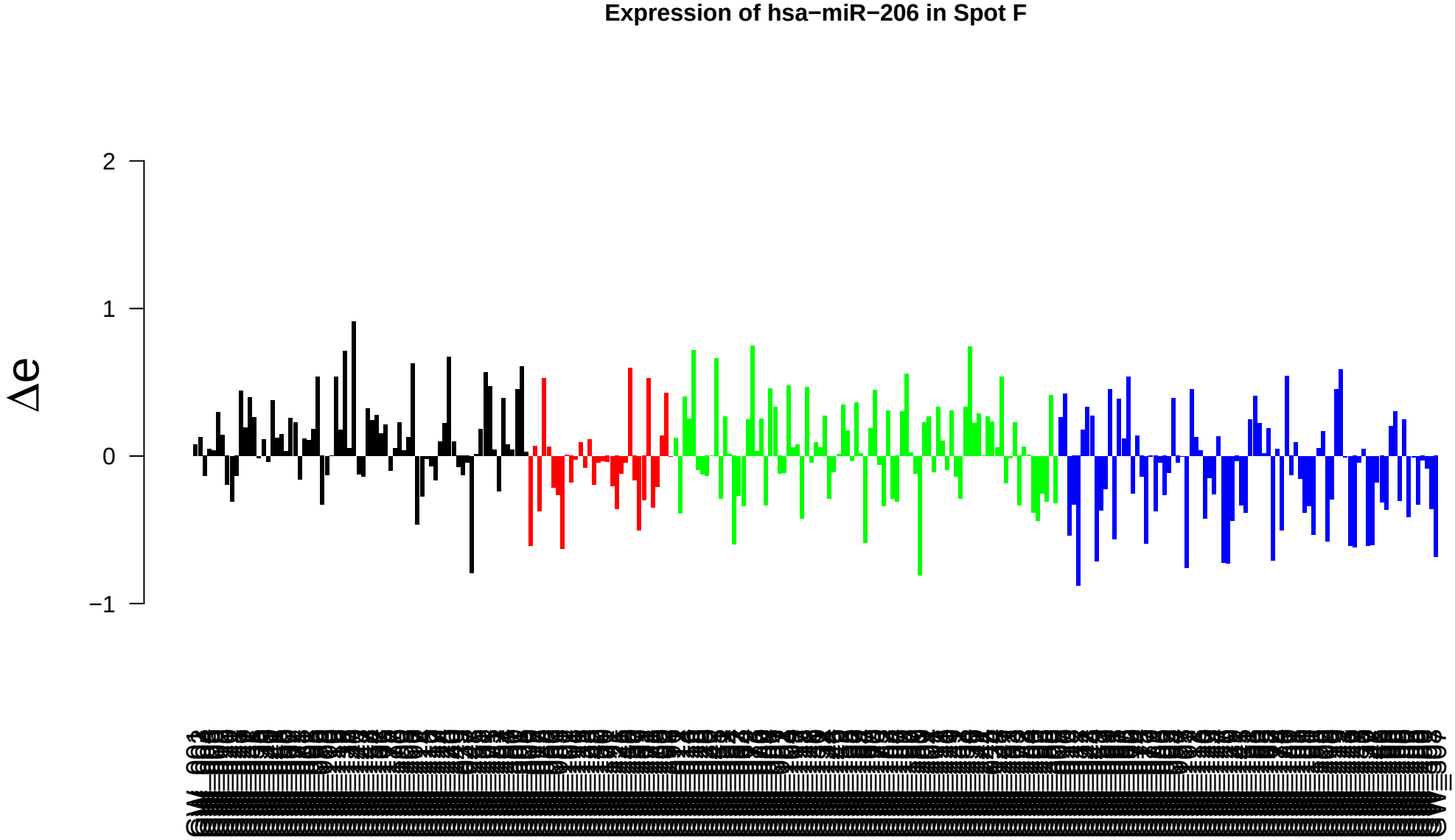

Expression of hsa−miR−206 in Spot A
Δe
2
1
0
−1

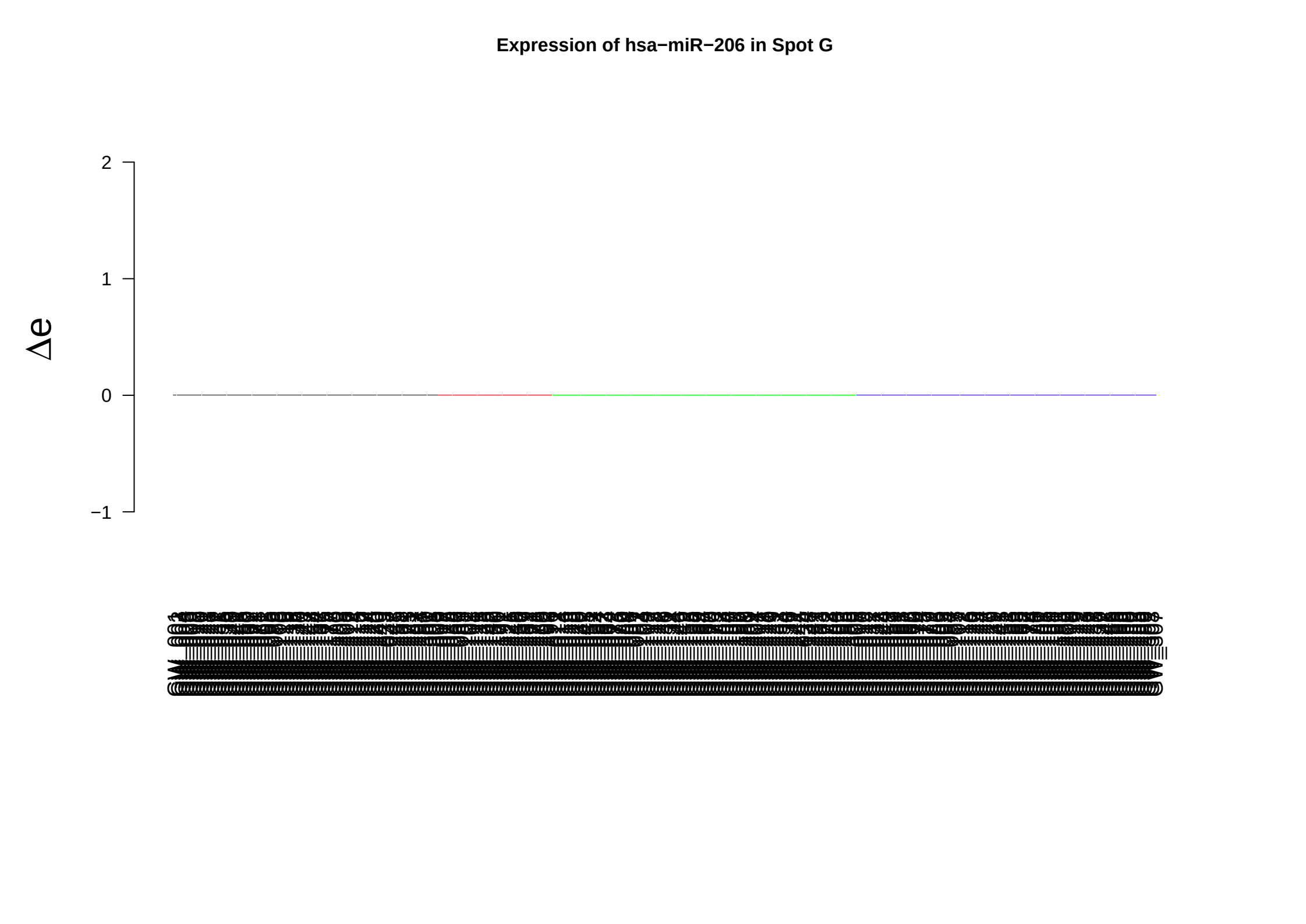

Expression of hsa−miR−206 in Spot B
Δe
2
1
0
−1

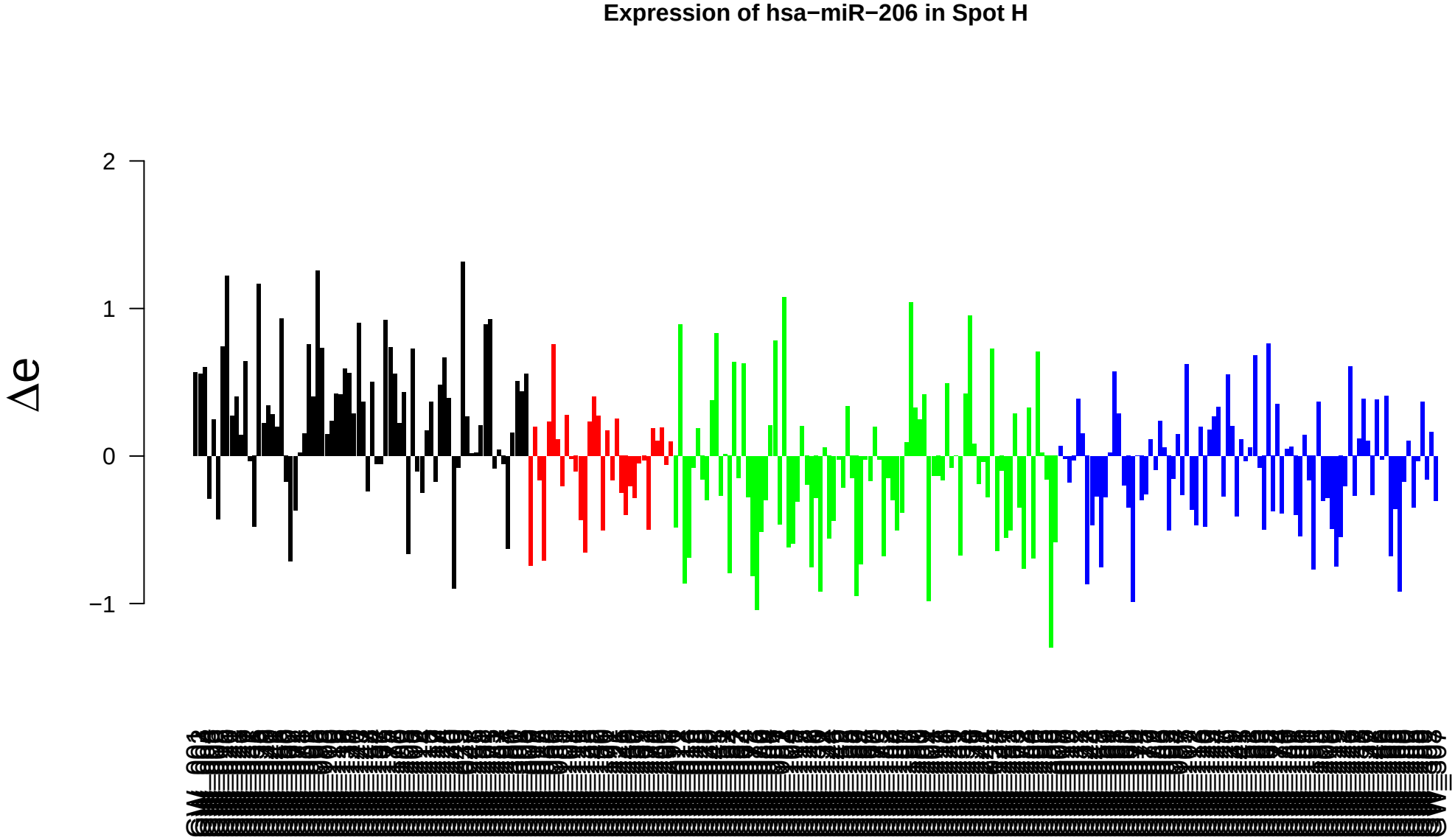

Expression of hsa−miR−206 in Spot C
Δe
2
1
0
−1

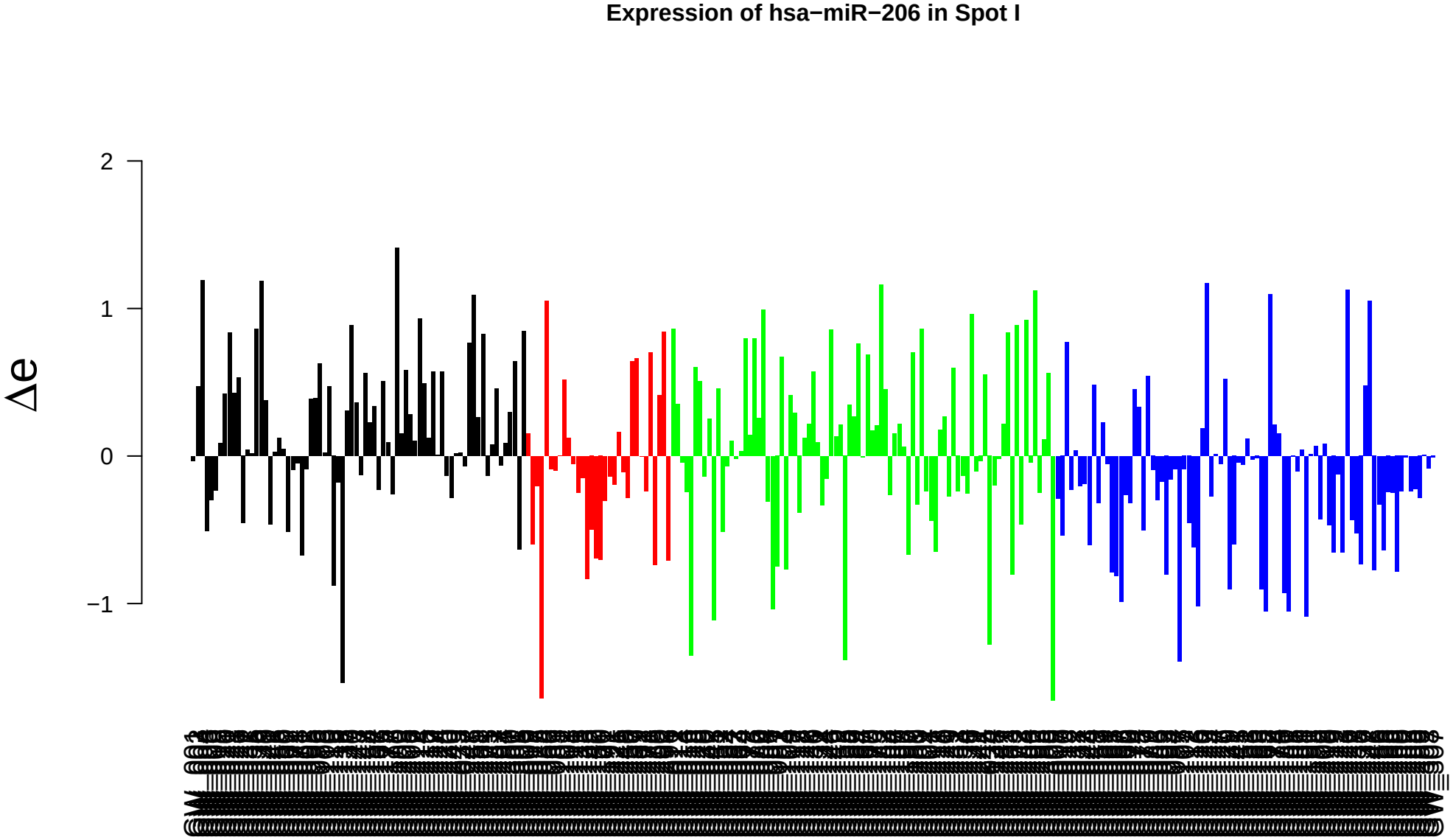

Expression of hsa−miR−206 in Spot D
Δe
2
1
0
−1

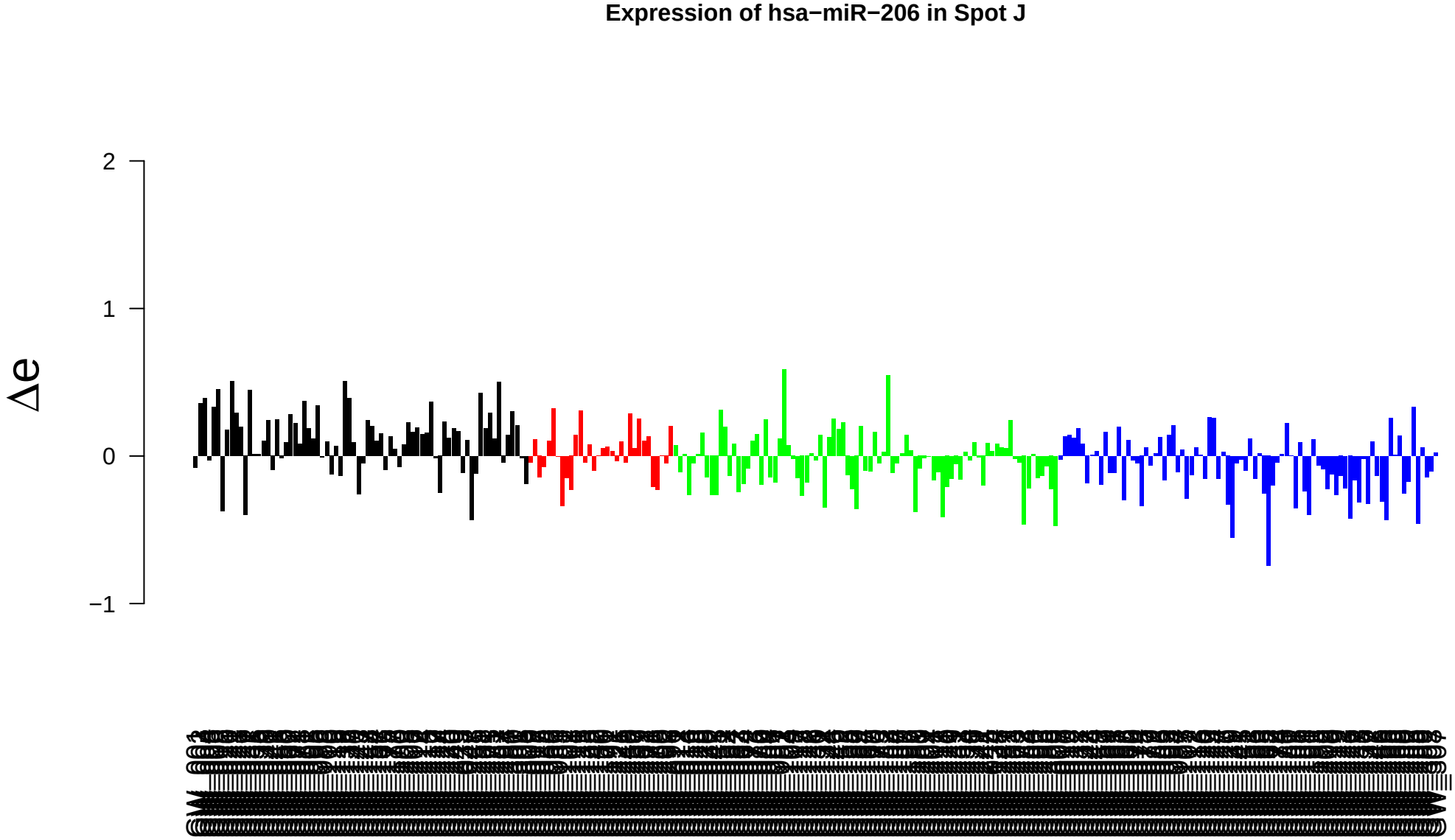

Expression of hsa−miR−206 in Spot E
Δe
2
1
0
−1

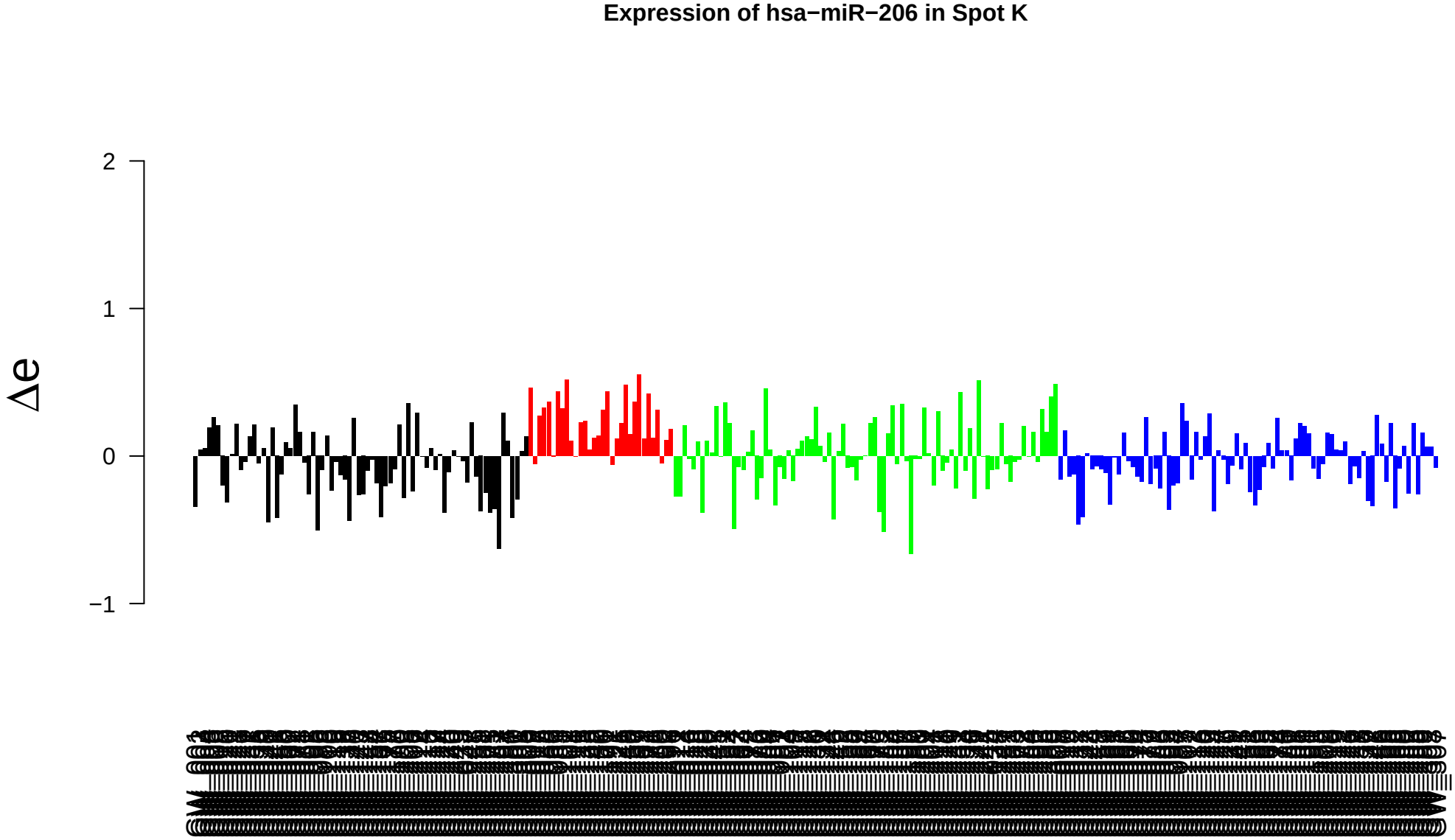

Expression of hsa−miR−206 in Spot F
Δe
2
1
0
−1

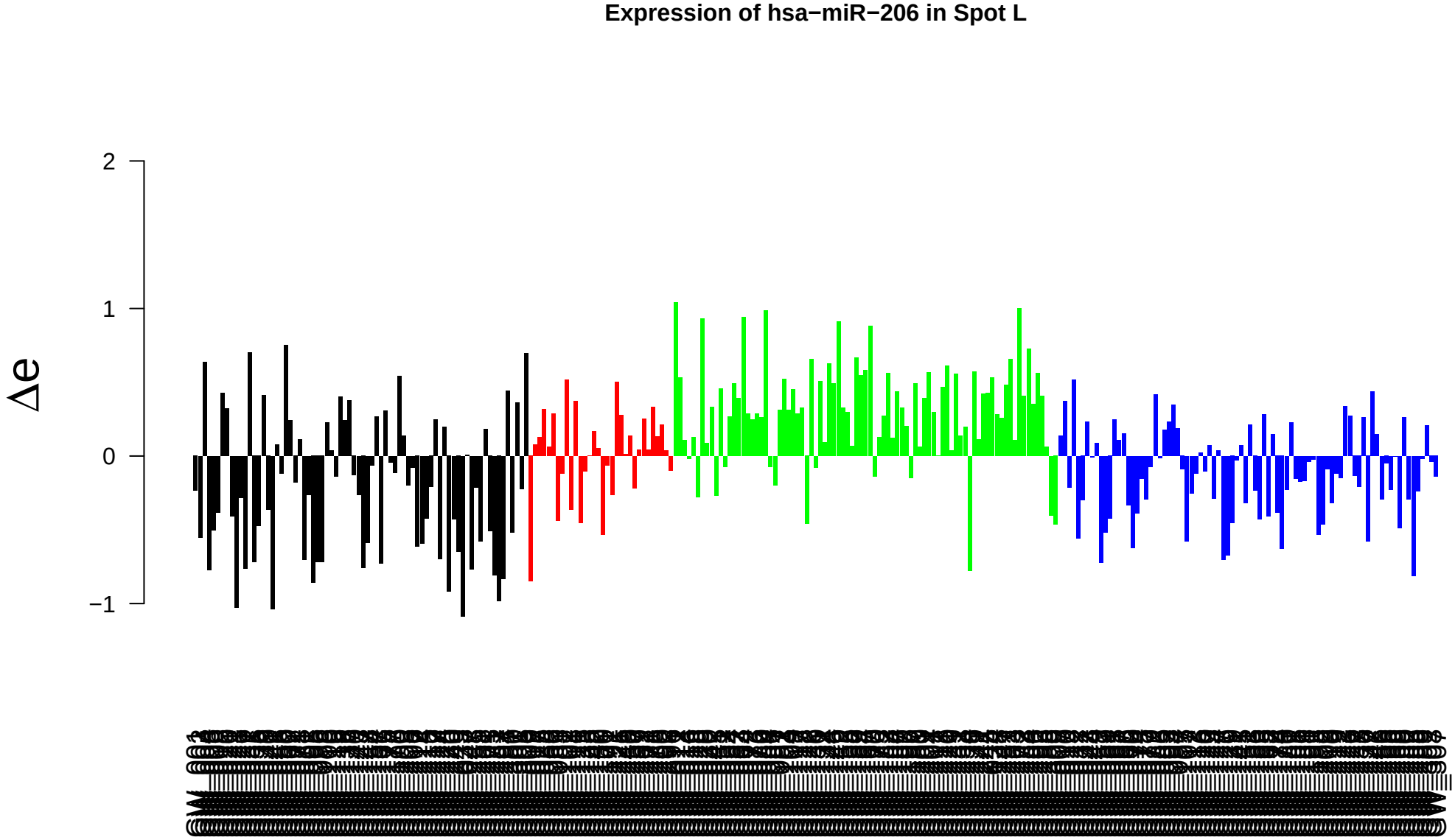

Expression of hsa−miR−206 in Spot G
Δe
2
1
0
−1

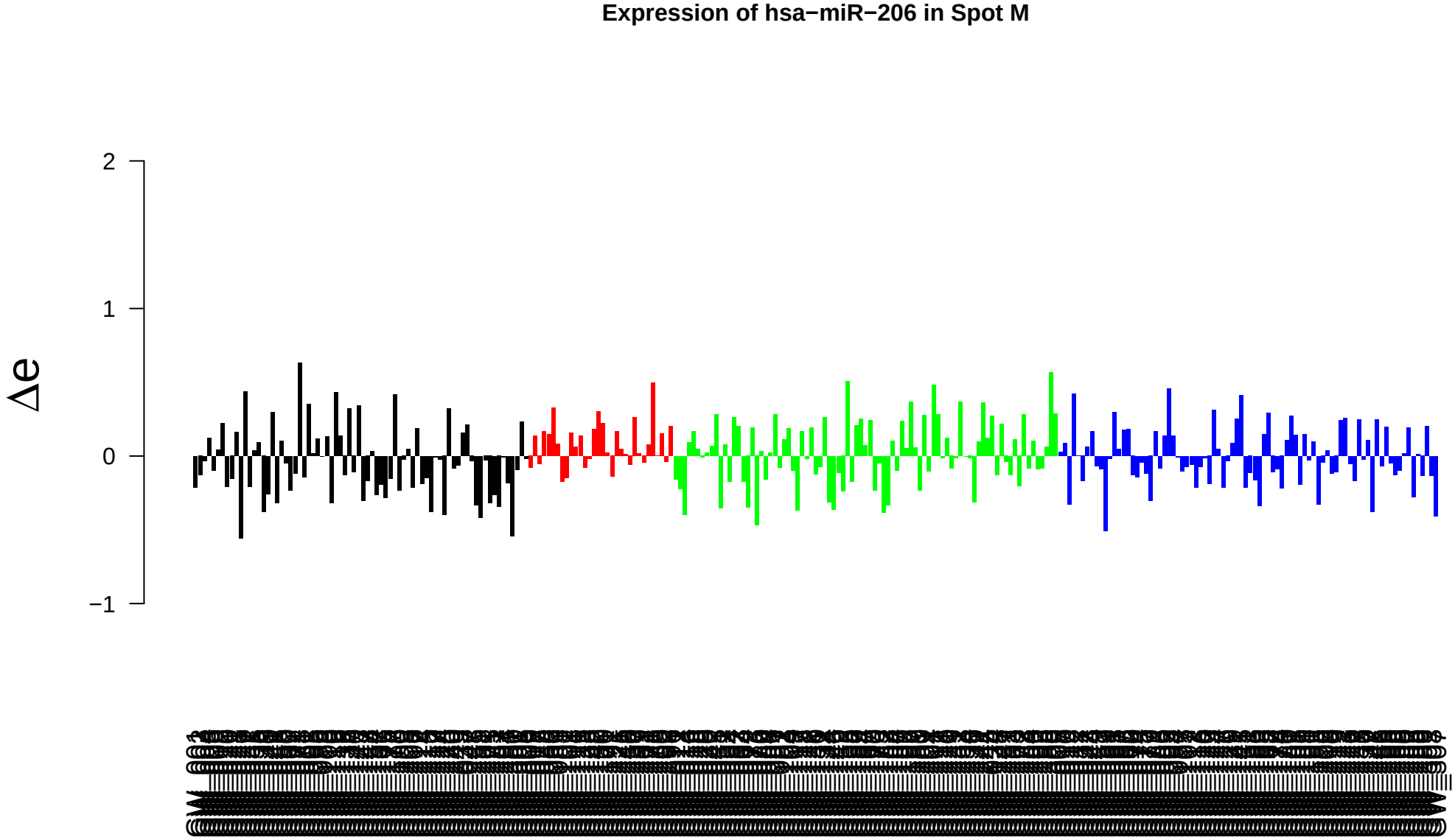

Expression of hsa−miR−206 in Spot H
Δe
2
1
0
−1

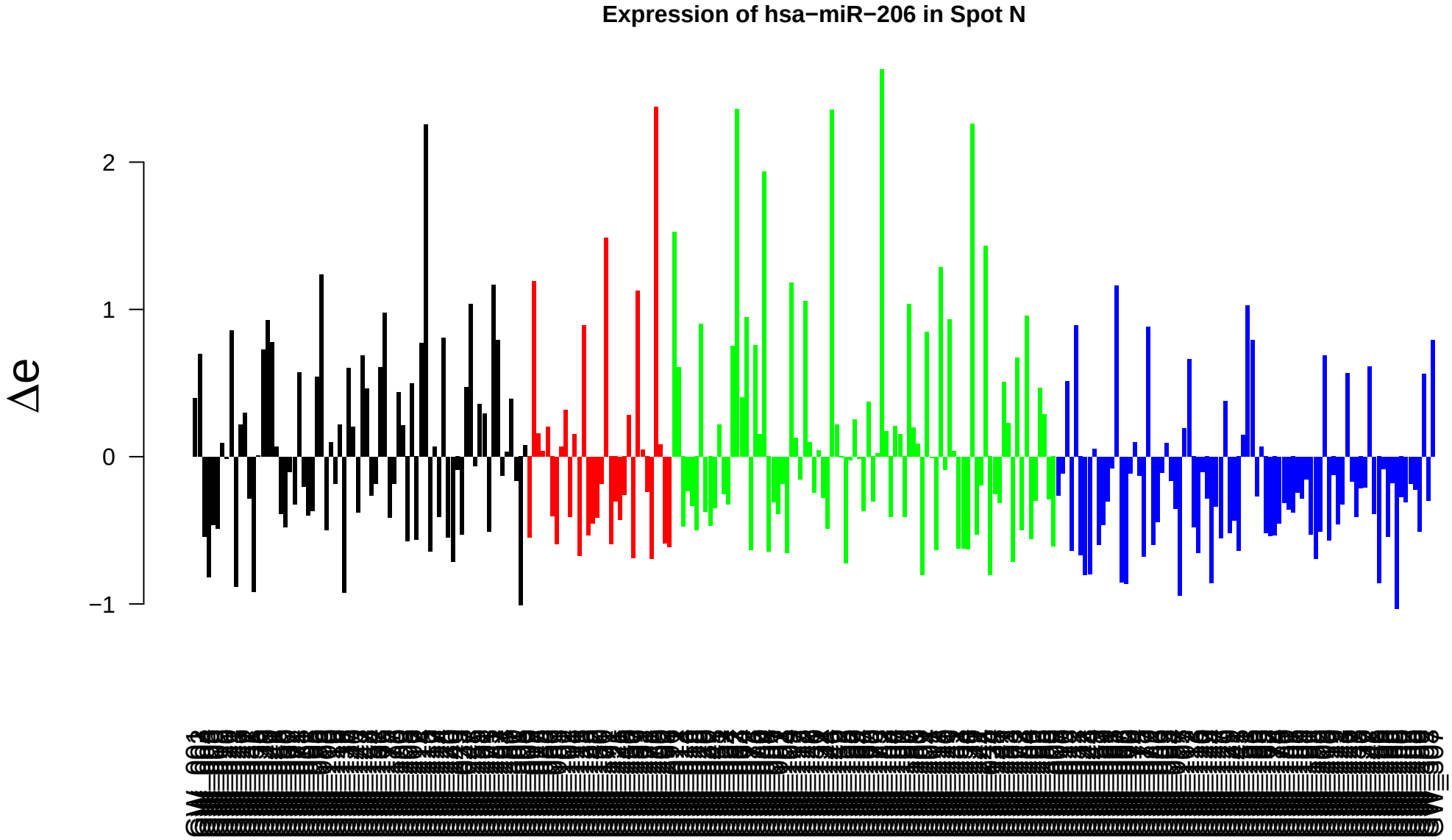

Expression of hsa−miR−206 in Spot I
Δe
2
1
0
−1

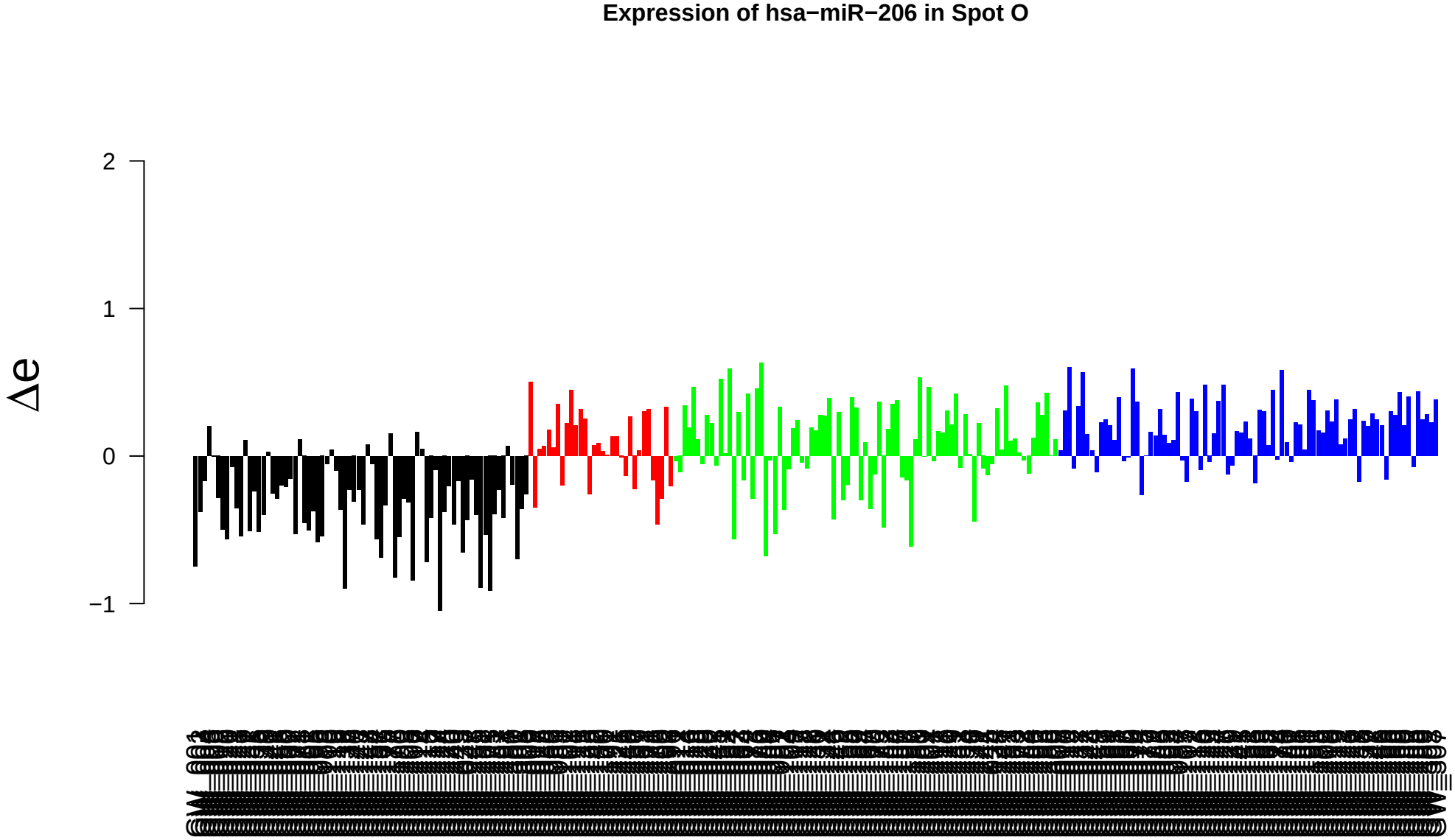

Expression of hsa−miR−206 in Spot J
Δe
2
1
0
−1

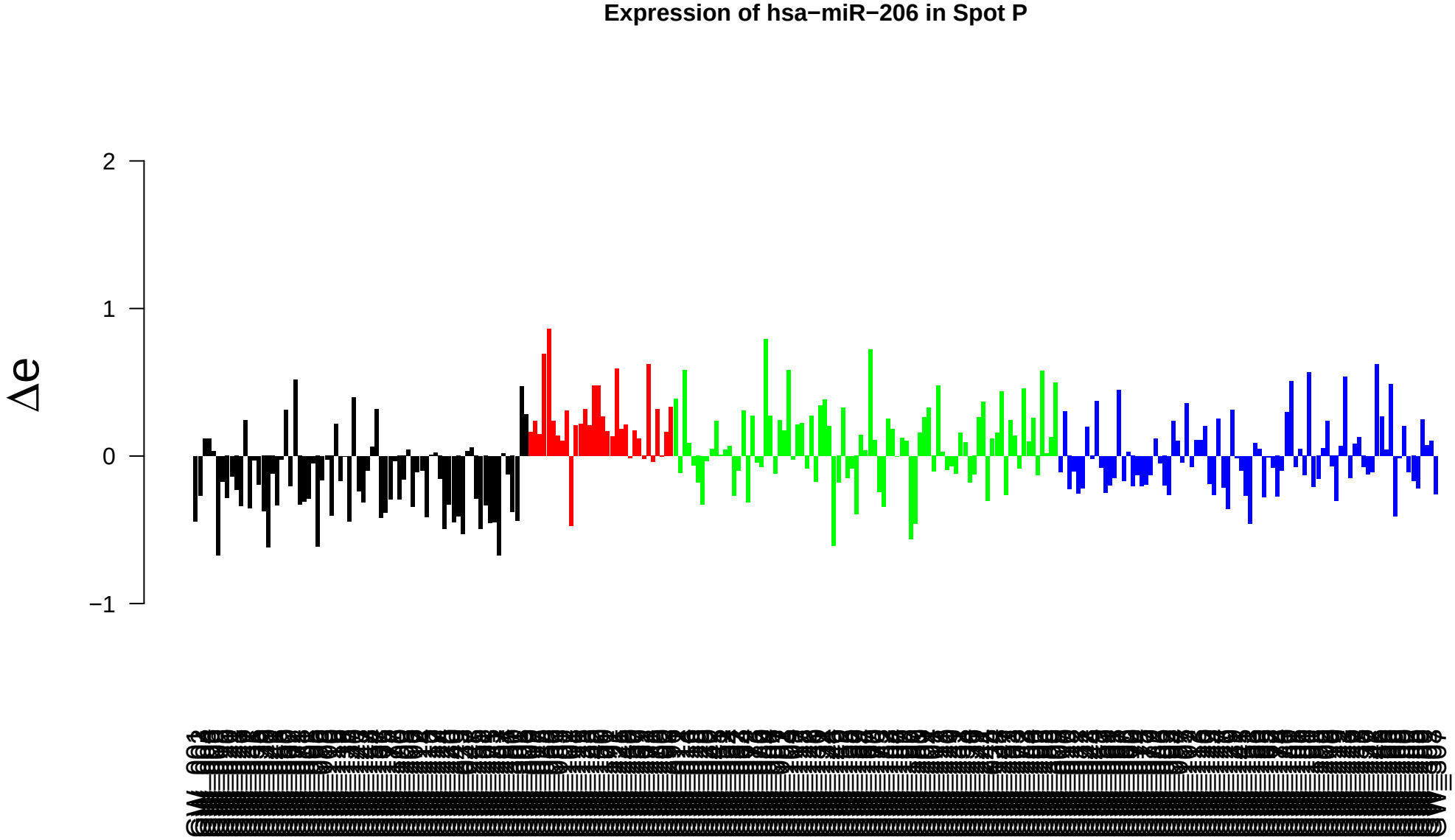

Expression of hsa−miR−206 in Spot K
Δe
2
1
0
−1

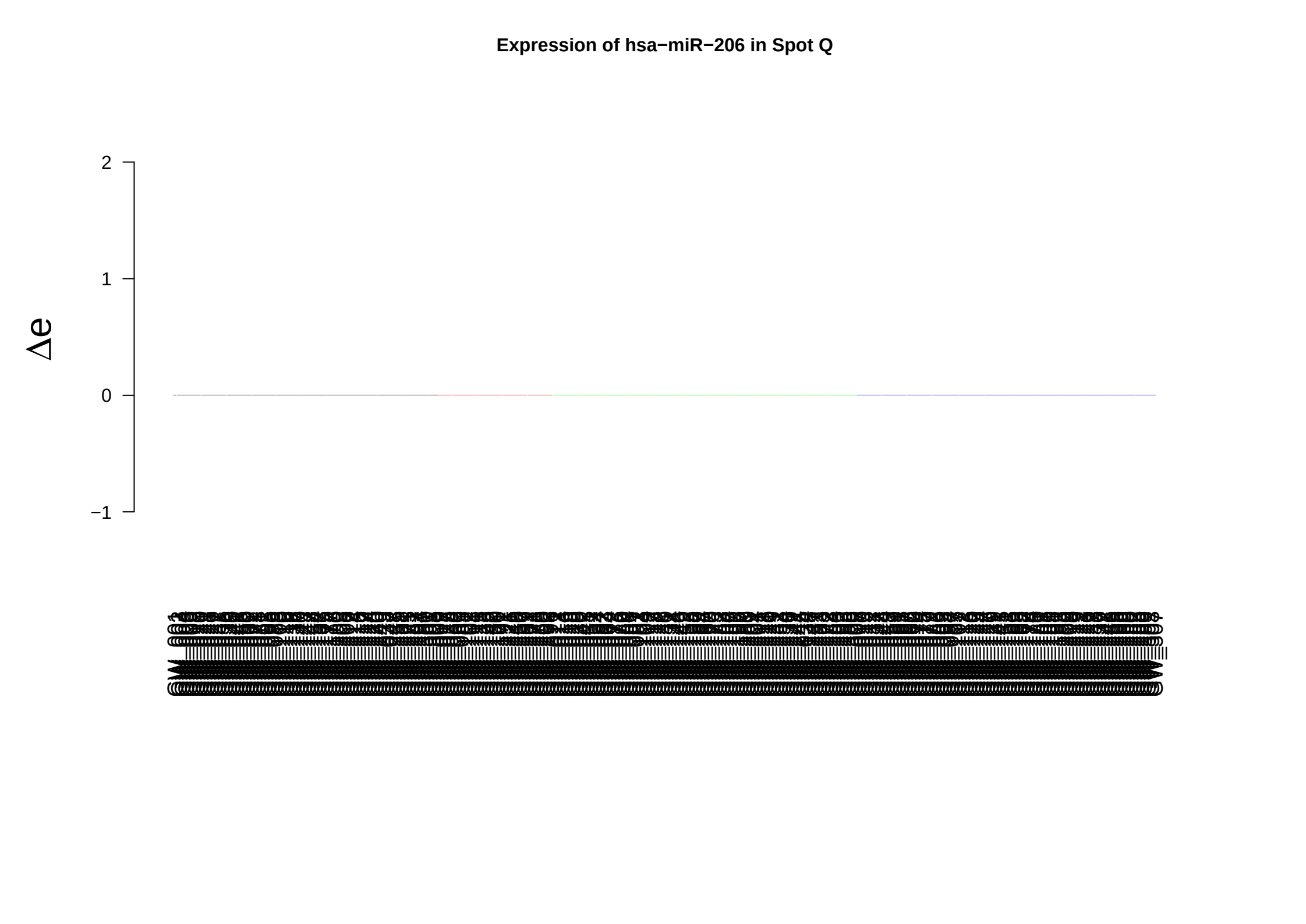

Expression of hsa−miR−206 in Spot L
Δe
2
1
0
−1

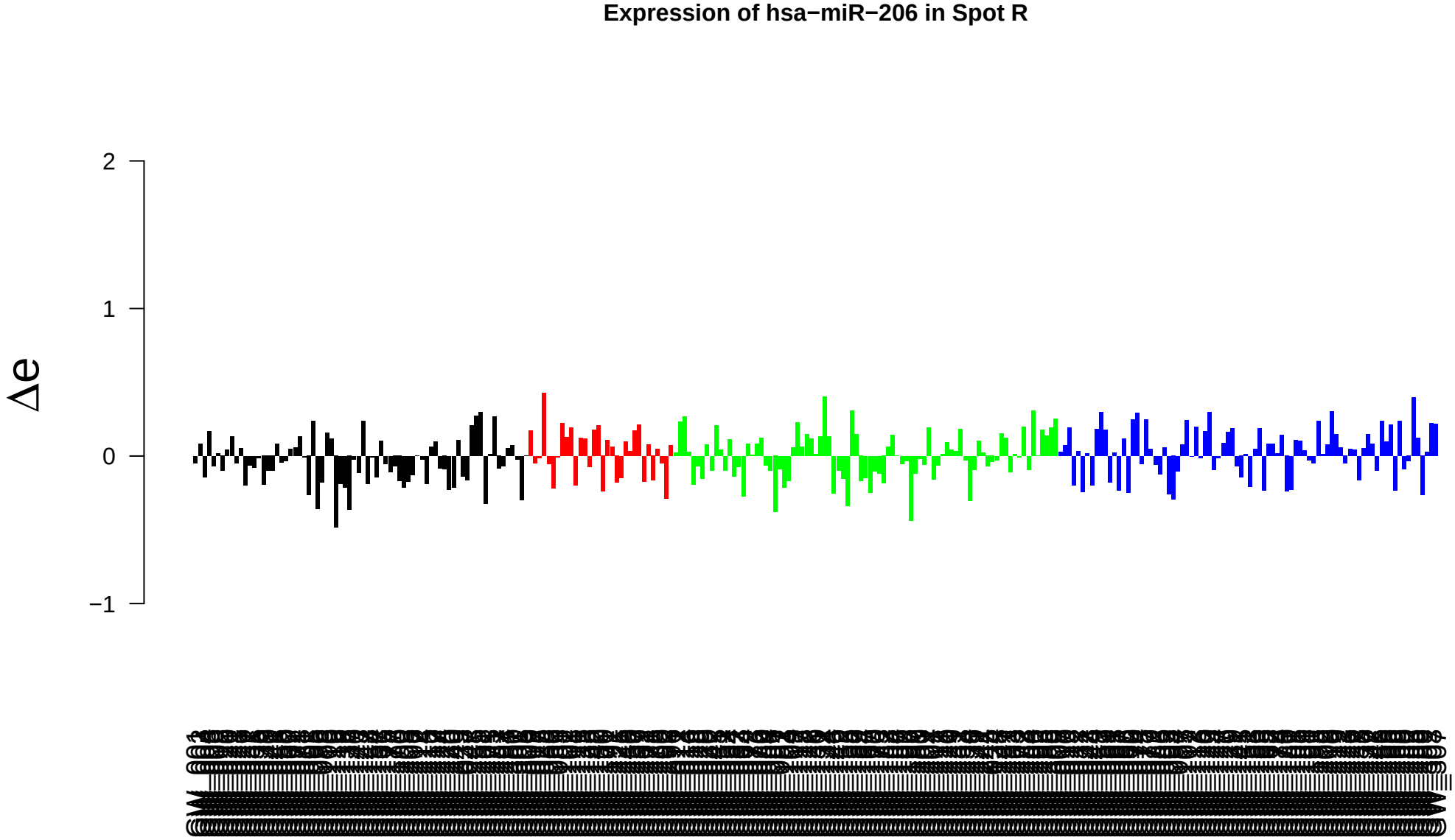

Expression of hsa−miR−206 in Spot M
Δe
2
1
0
−1


Expression of hsa−miR−206 in Spot N
Δe
2
1
0
−1


Expression of hsa−miR−206 in Spot O
Δe
2
1
0
−1


Expression of hsa−miR−206 in Spot P
Δe
2
1
0
−1


Expression of hsa−miR−206 in Spot Q
Δe
2
1
0
−1


Expression of hsa−miR−206 in Spot R
Δe
2
1
0
−1