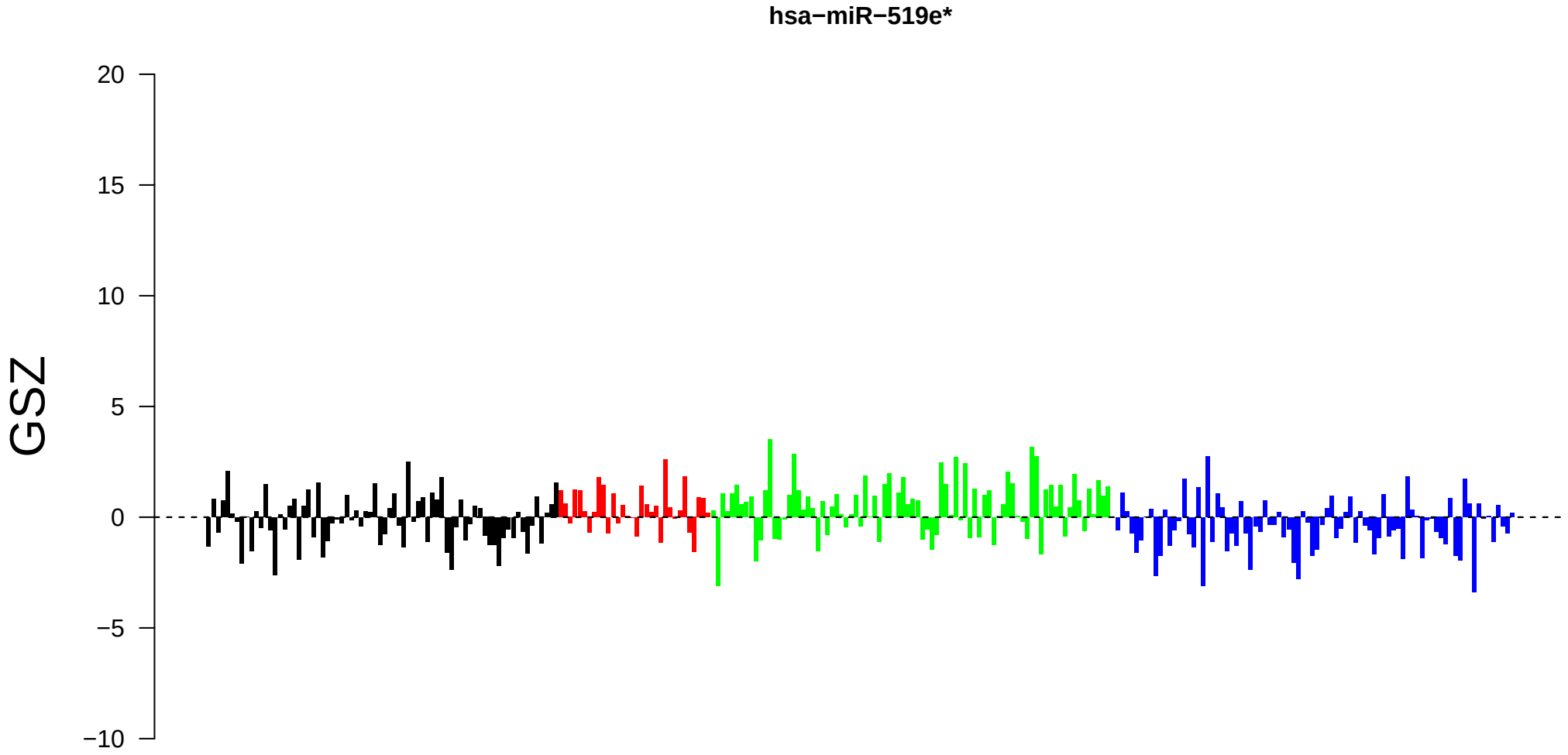
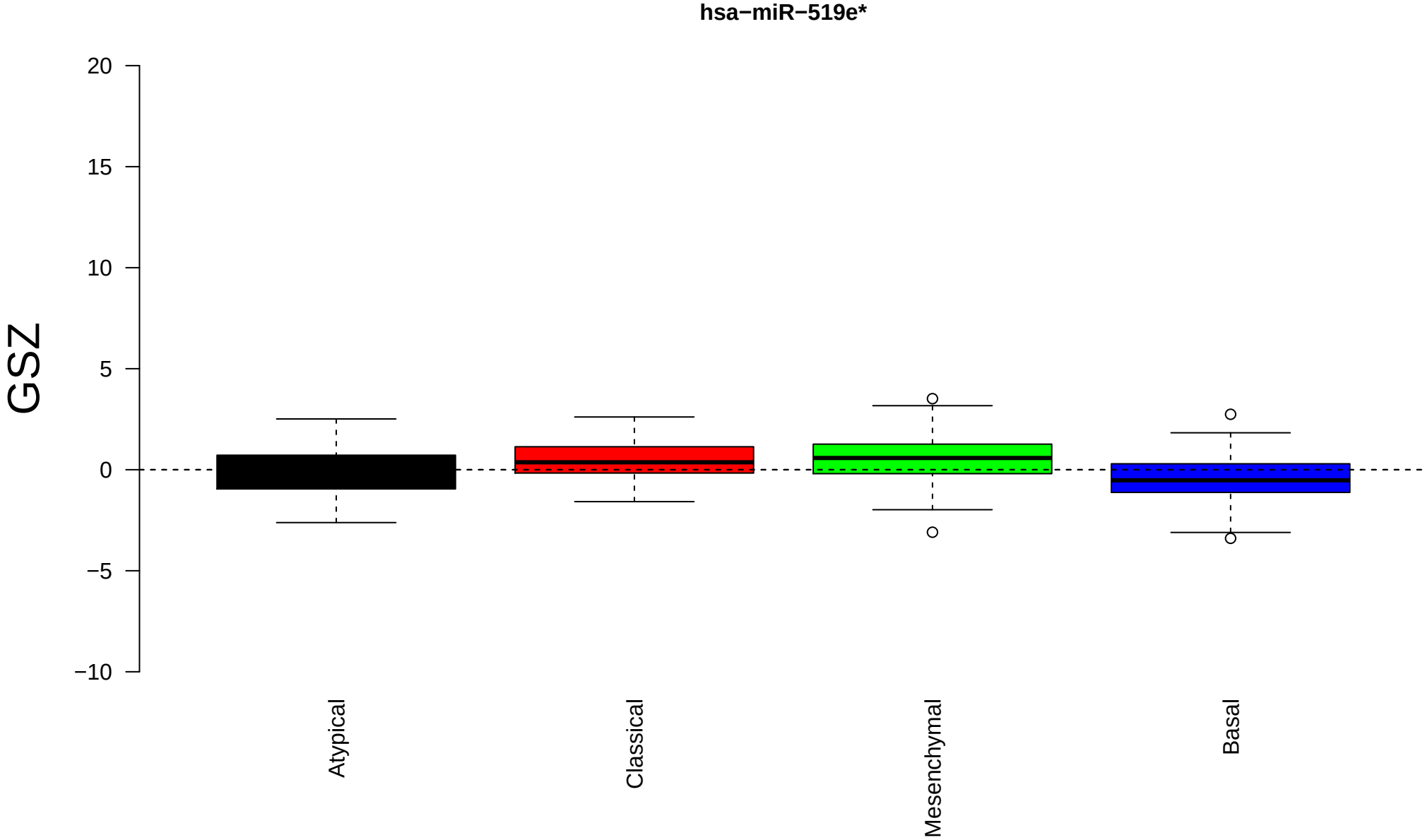
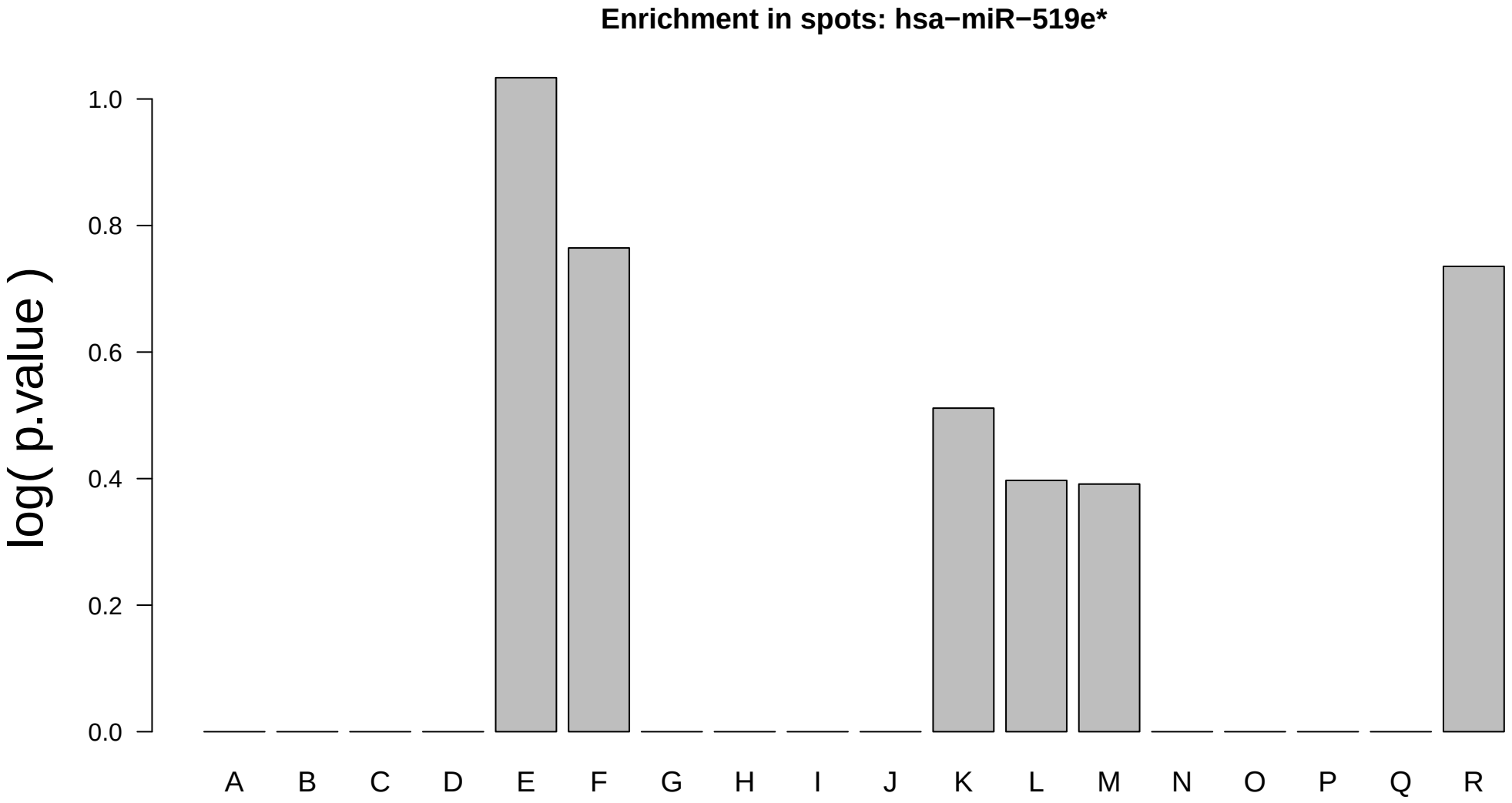
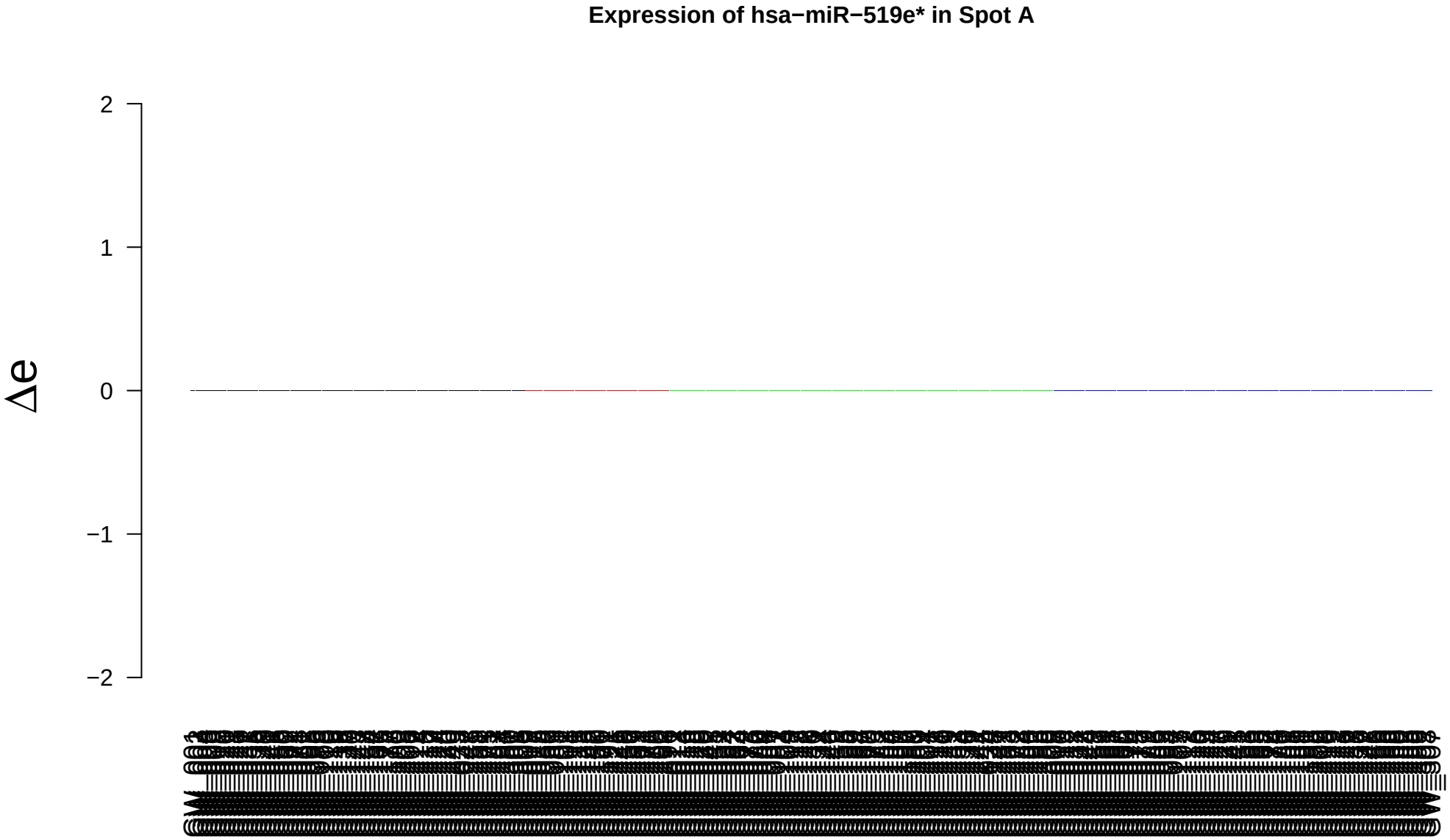
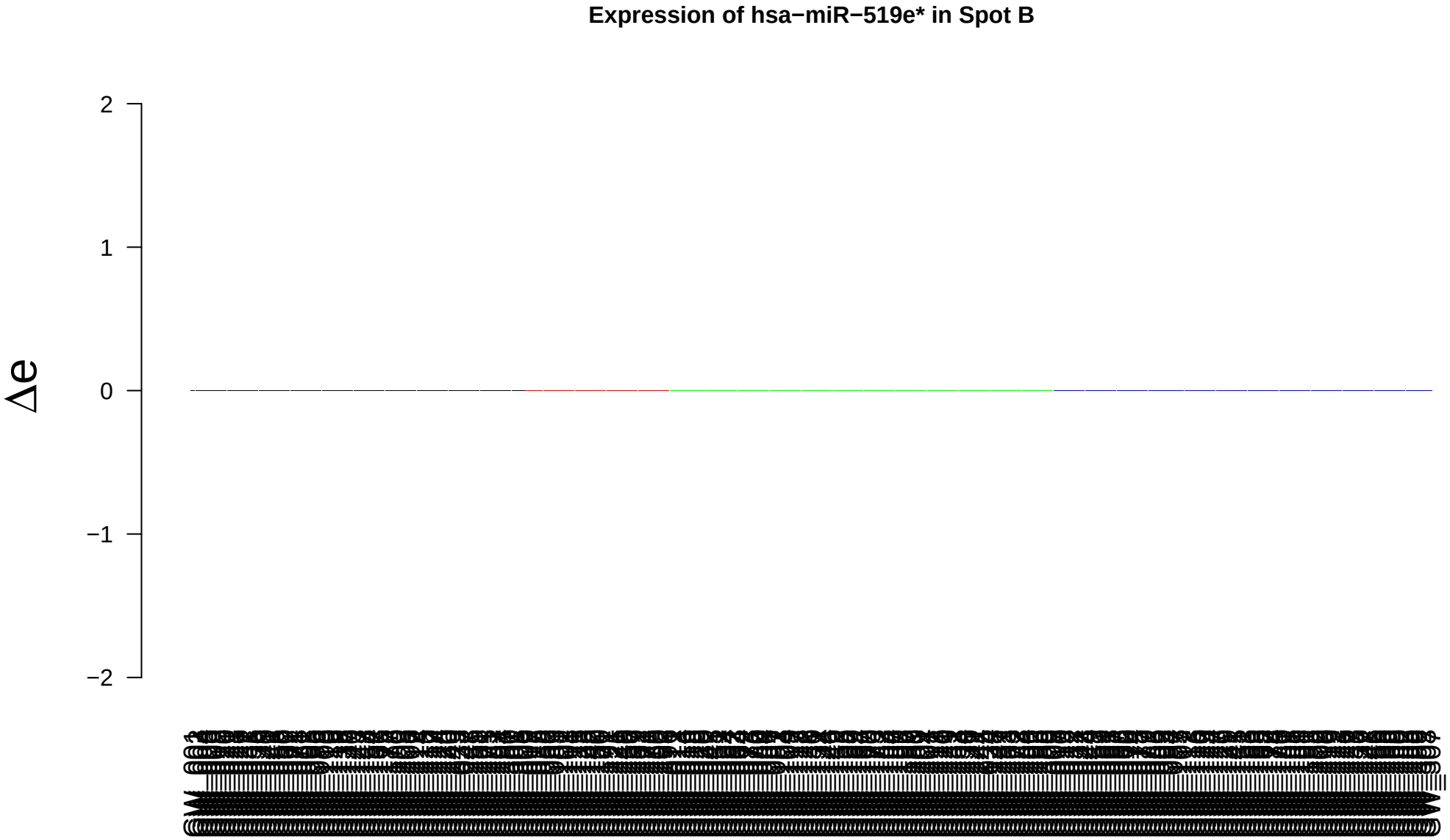
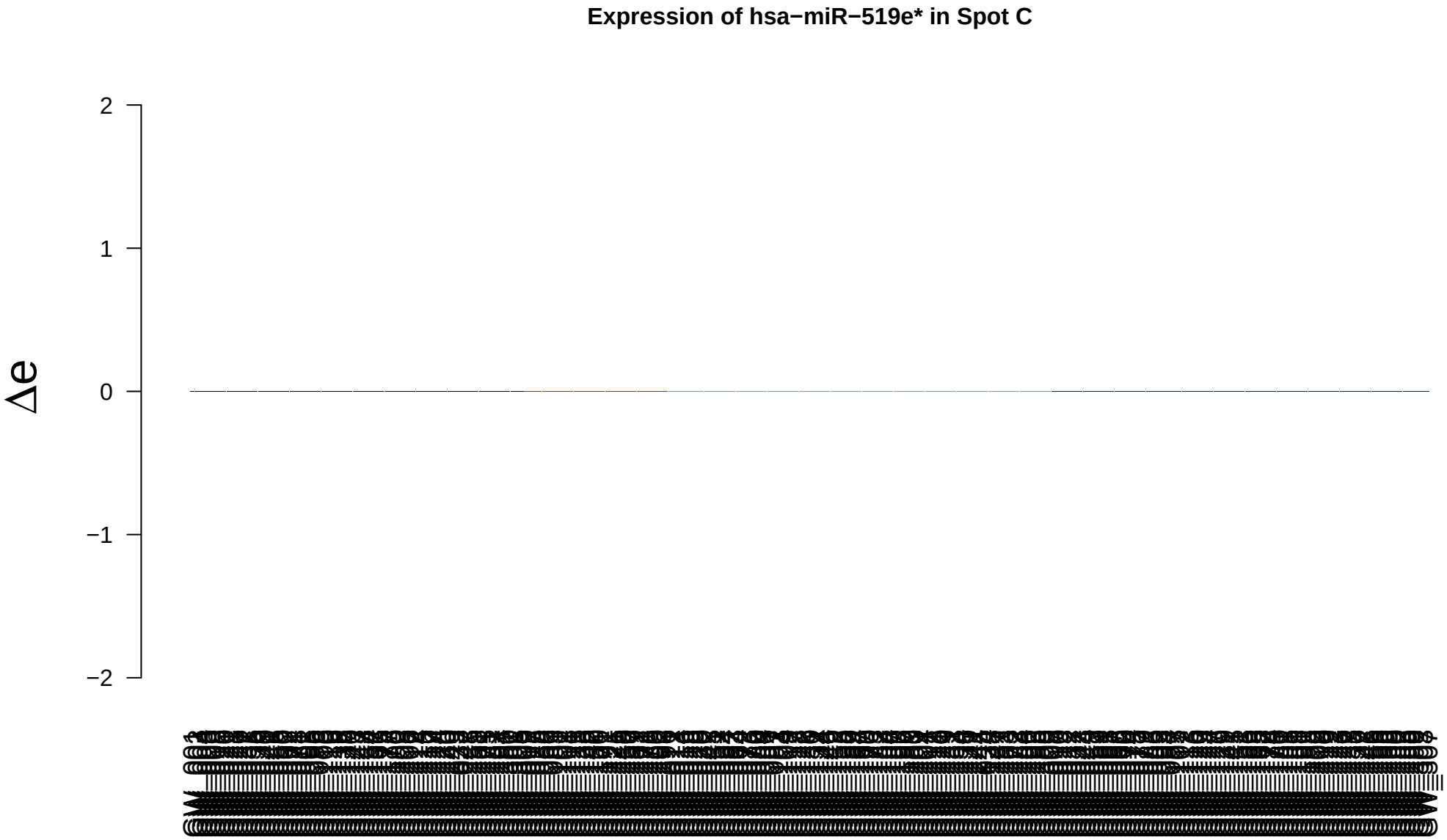

hsa−miR−519e*
GSZ
20
15
10
5
0
−5
−10

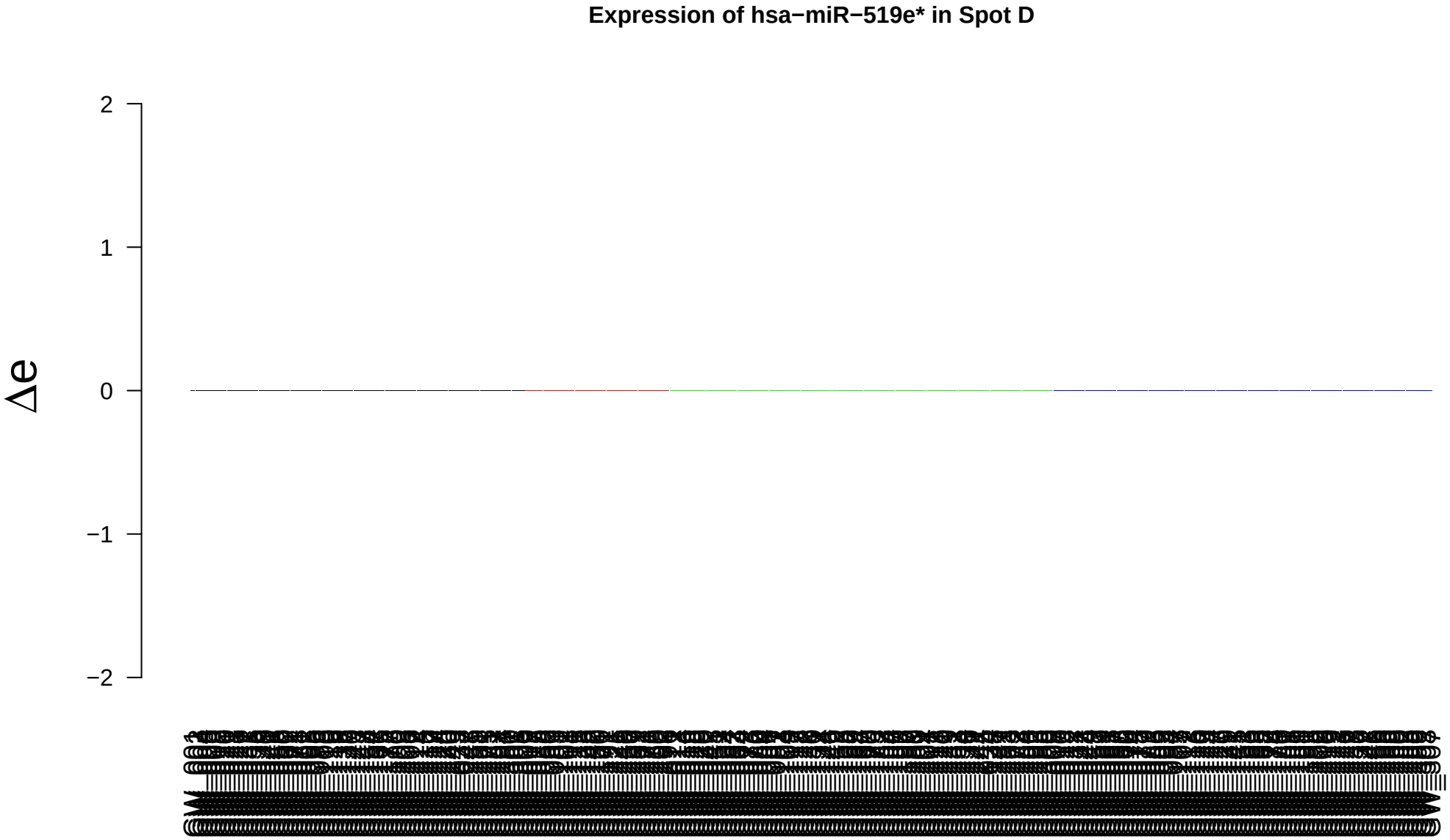

hsa−miR−519e*
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal

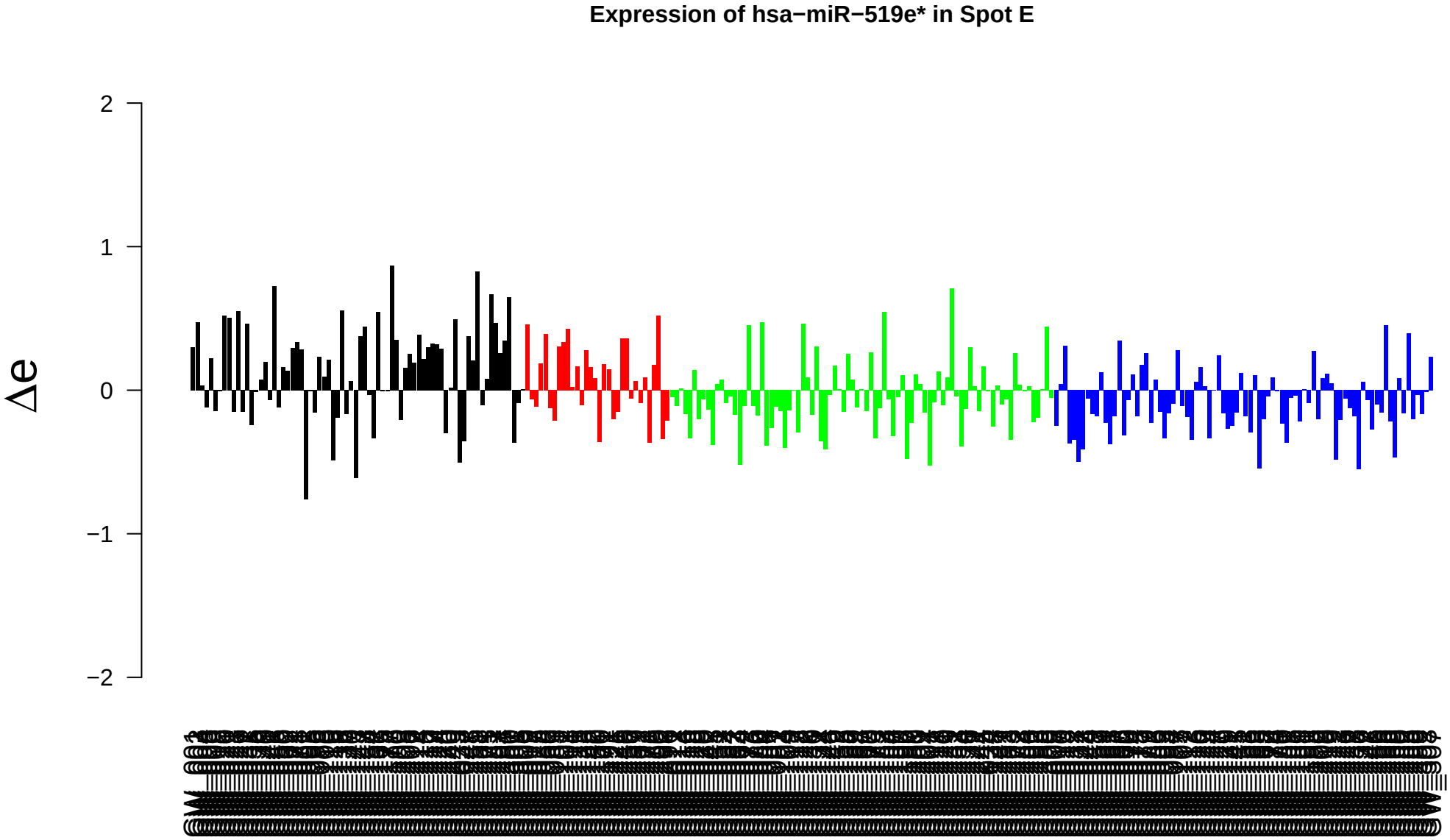

Enrichment in spots: hsa−miR−519e*
log(p.value)
1.0
0.8
0.6
0.4
0.2
0.0
A
B
C
D
E
F
G
H
I
J
K
L
M
N
O
P
Q
R

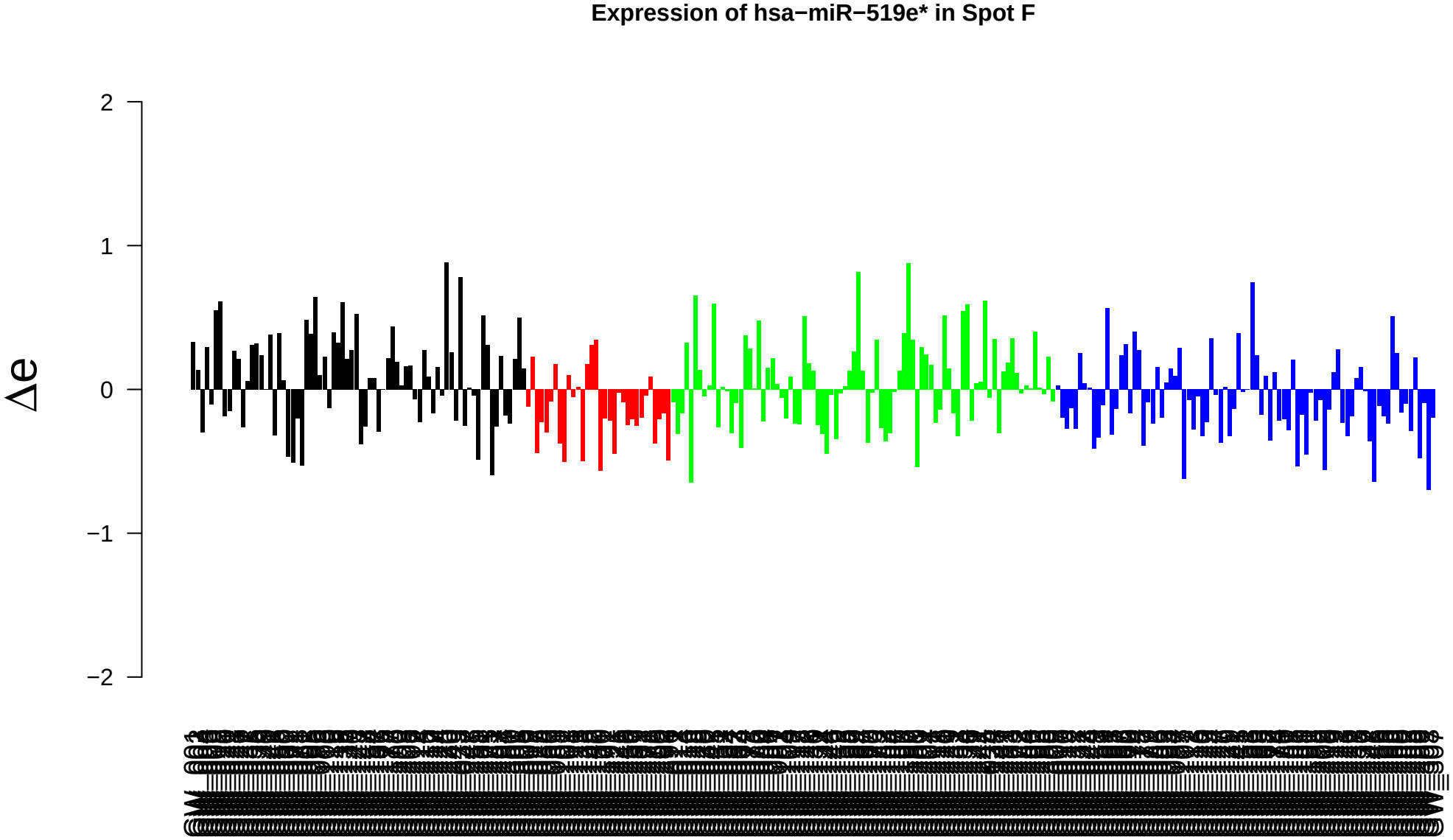

Expression of hsa−miR−519e* in Spot A
Δe
2
1
0
−1
−2

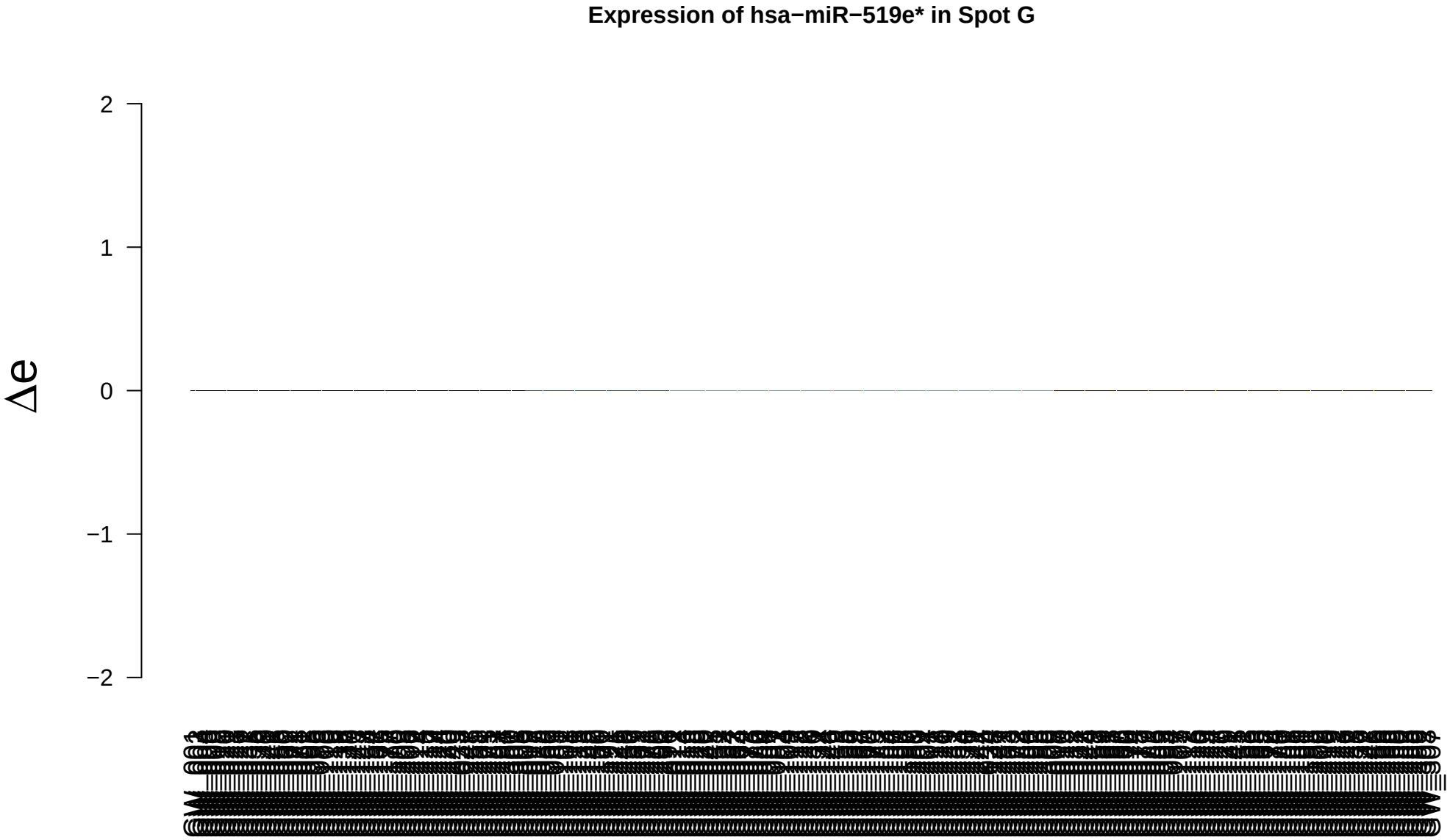

Expression of hsa−miR−519e* in Spot B
Δe
2
1
0
−1
−2

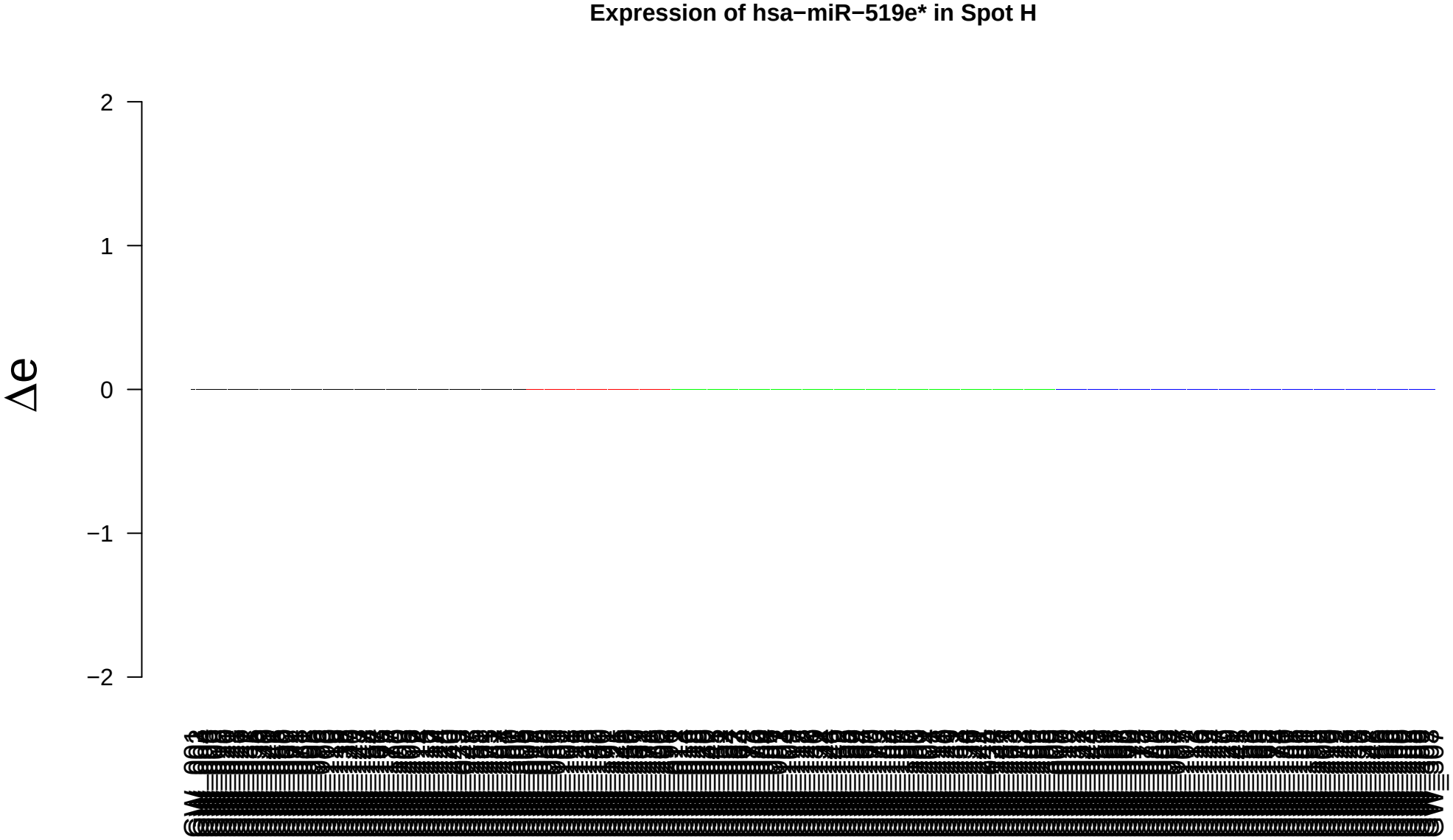

Expression of hsa−miR−519e* in Spot C
Δe
2
1
0
−1
−2

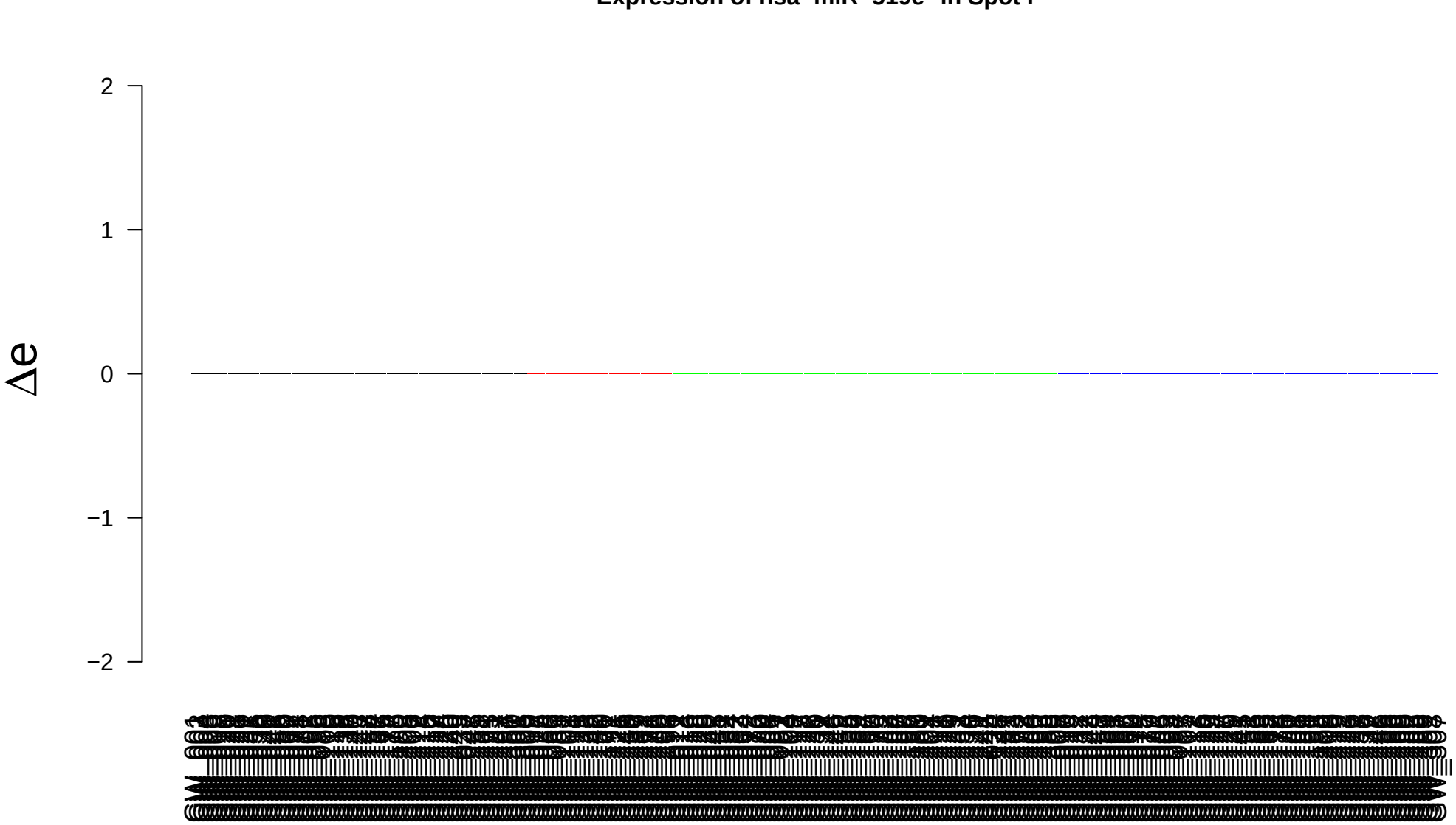

Expression of hsa−miR−519e* in Spot D
Δe
2
1
0
−1
−2

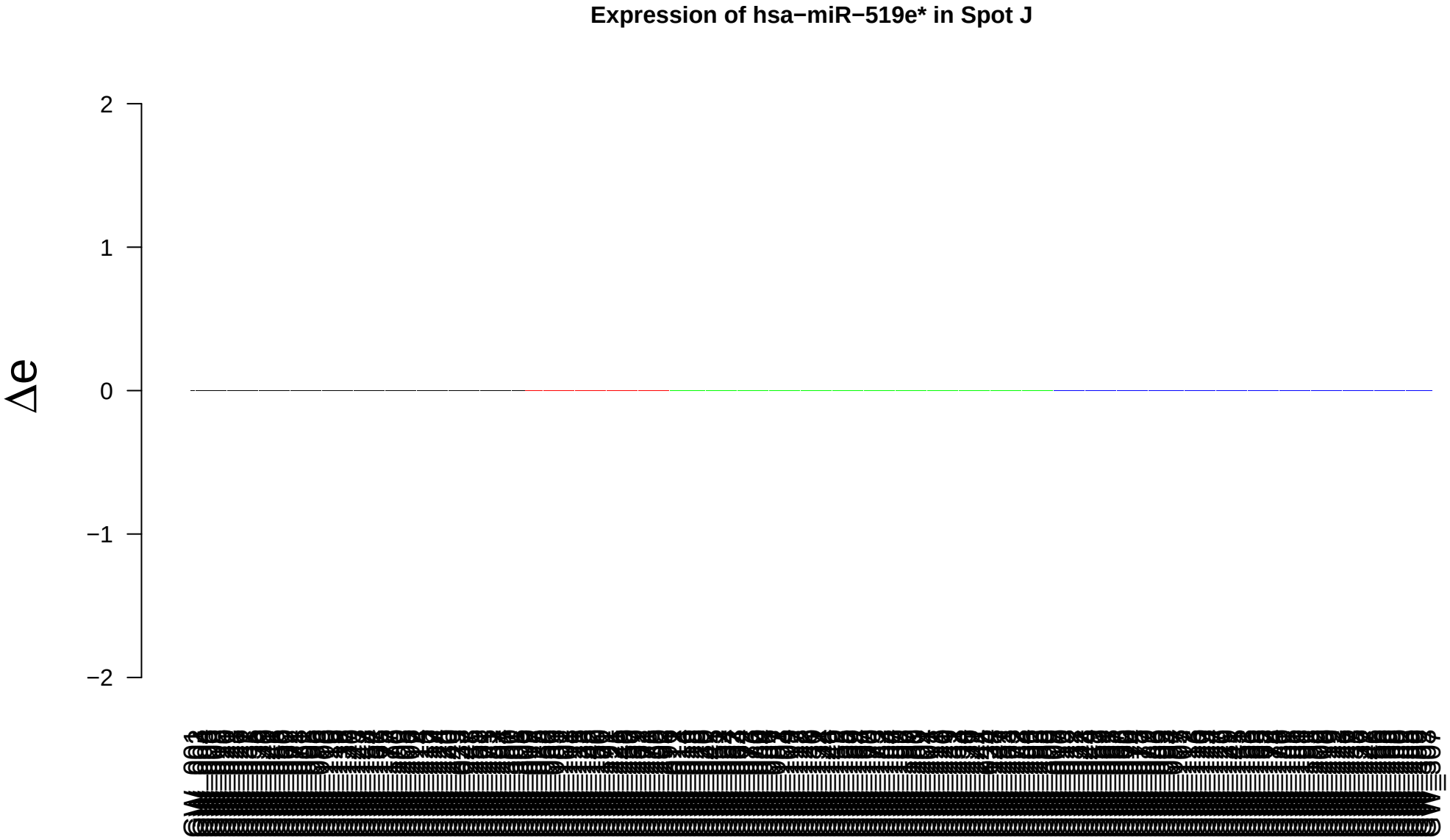

Expression of hsa−miR−519e* in Spot E
Δe
2
1
0
−1
−2

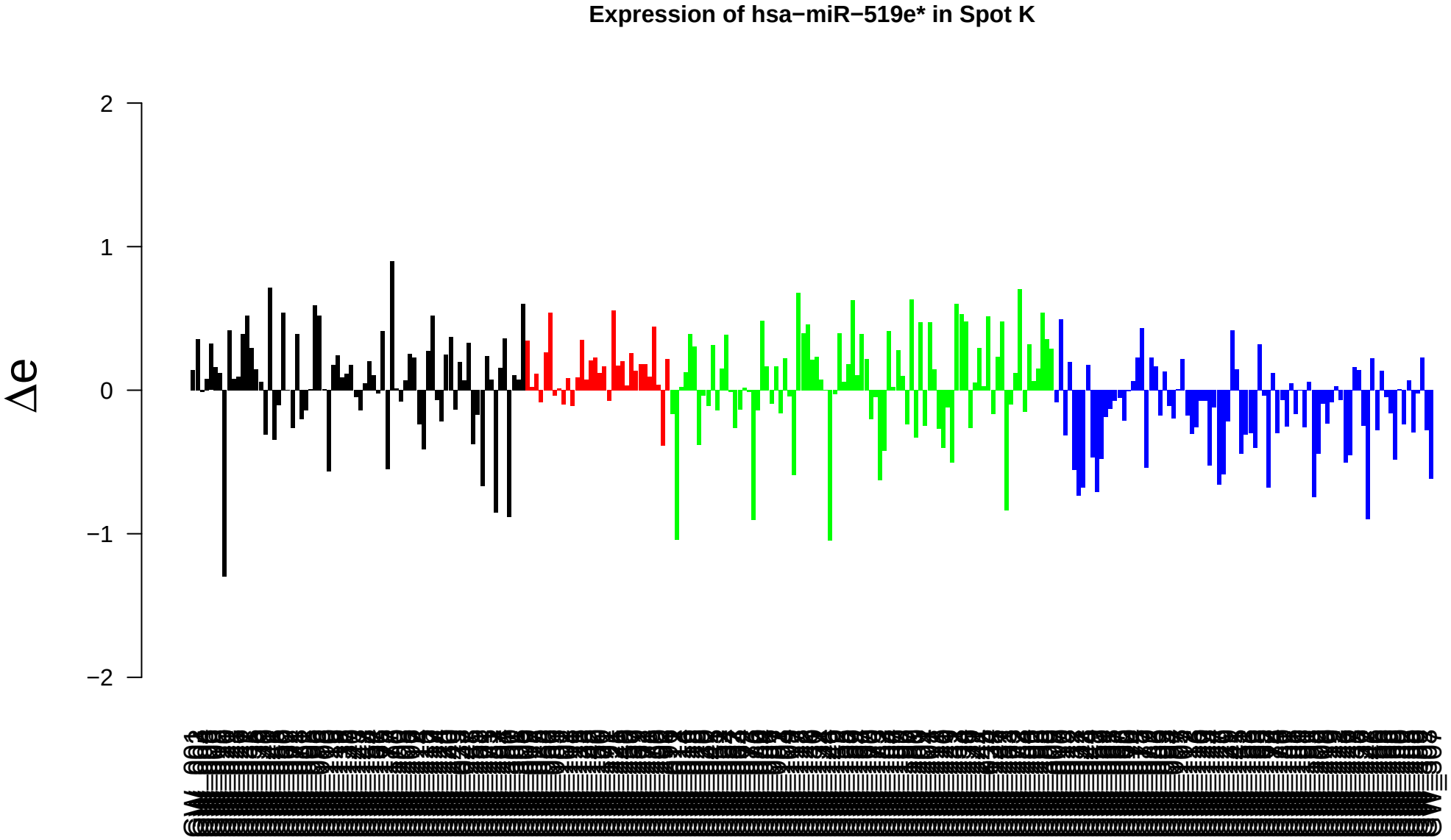

Expression of hsa−miR−519e* in Spot F
Δe
2
1
0
−1
−2

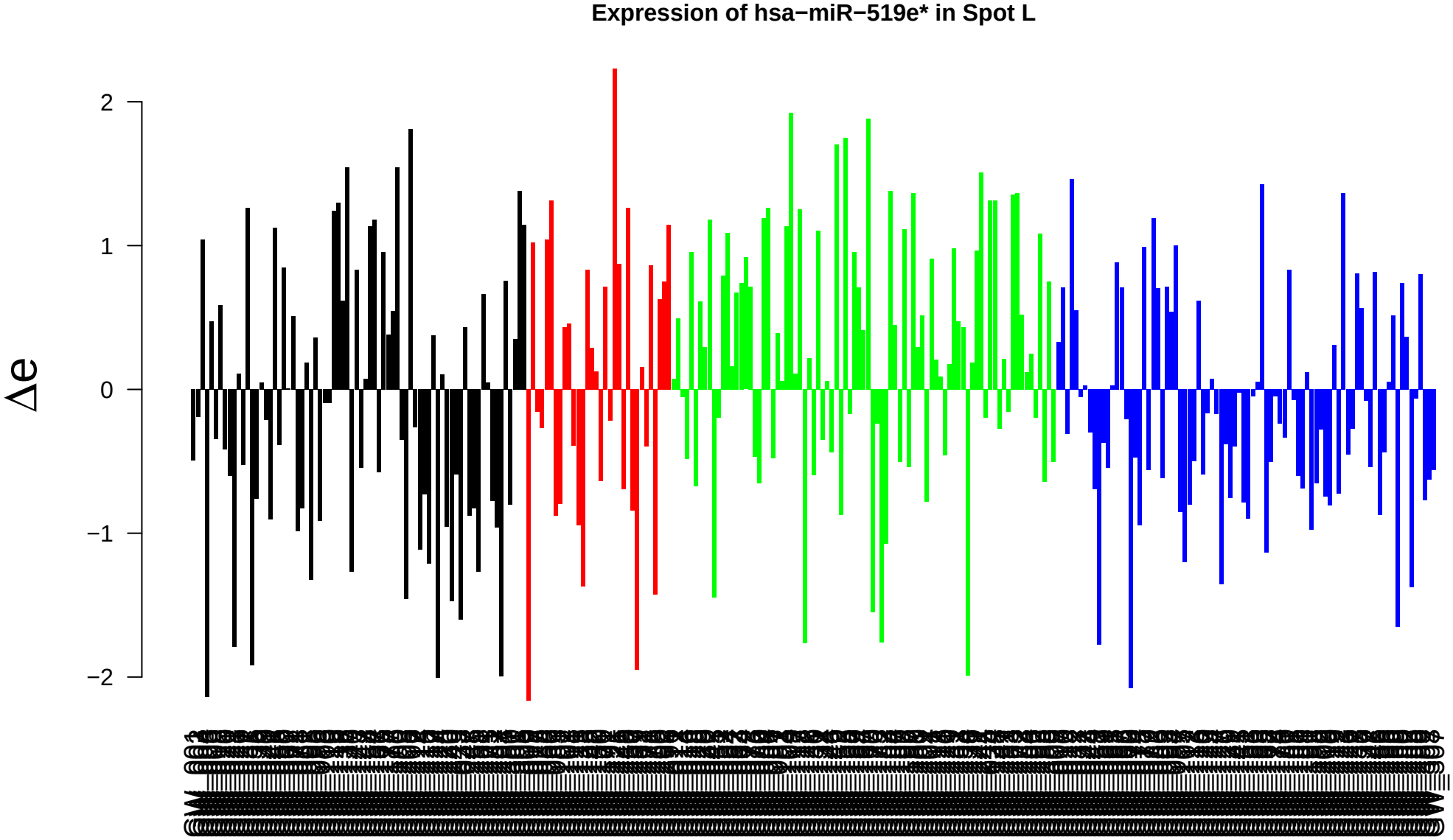

Expression of hsa−miR−519e* in Spot G
Δe
2
1
0
−1
−2

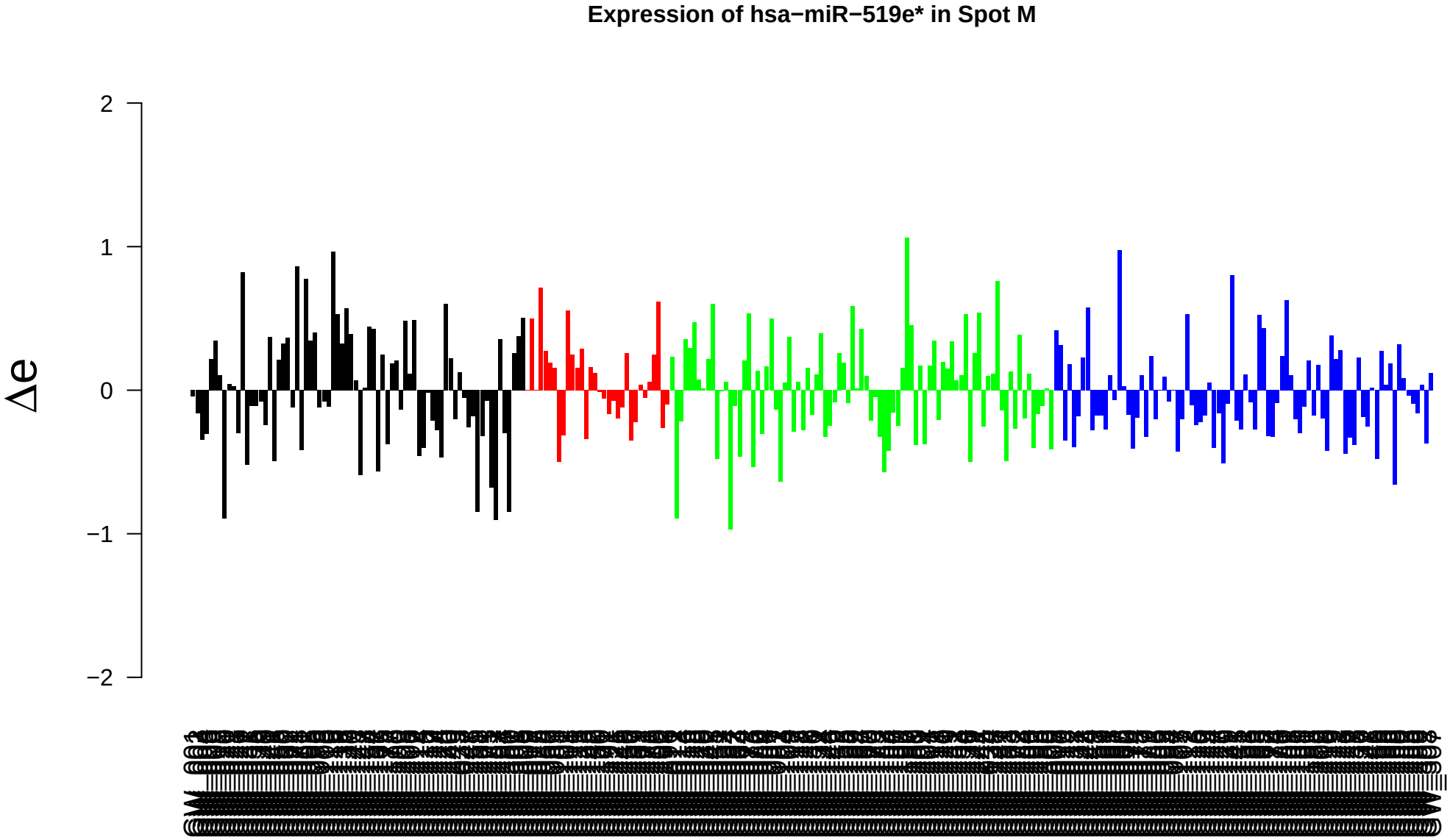

Expression of hsa−miR−519e* in Spot H
Δe
2
1
0
−1
−2

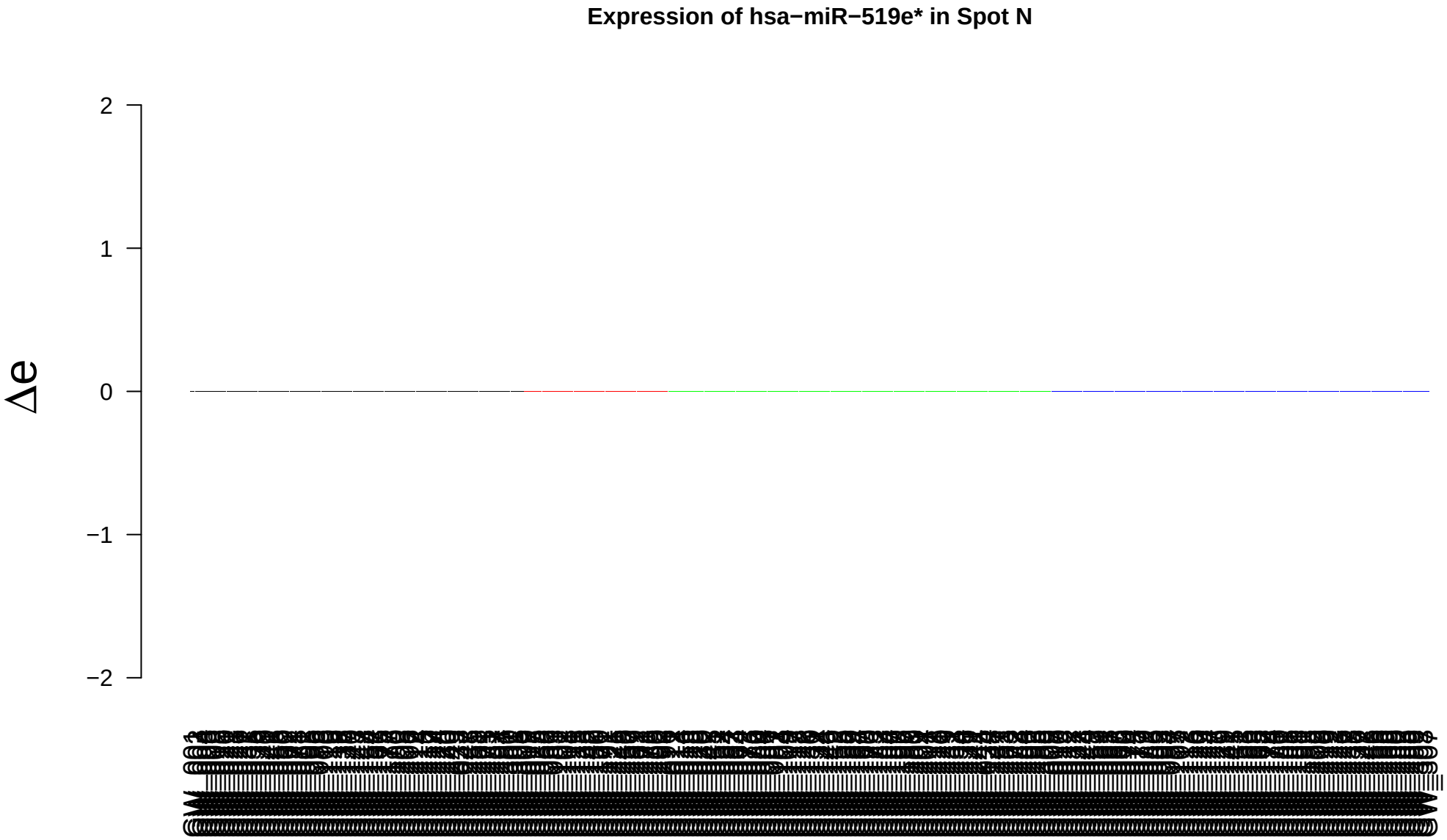

Expression of hsa−miR−519e* in Spot I
Δe
2
1
0
−1
−2

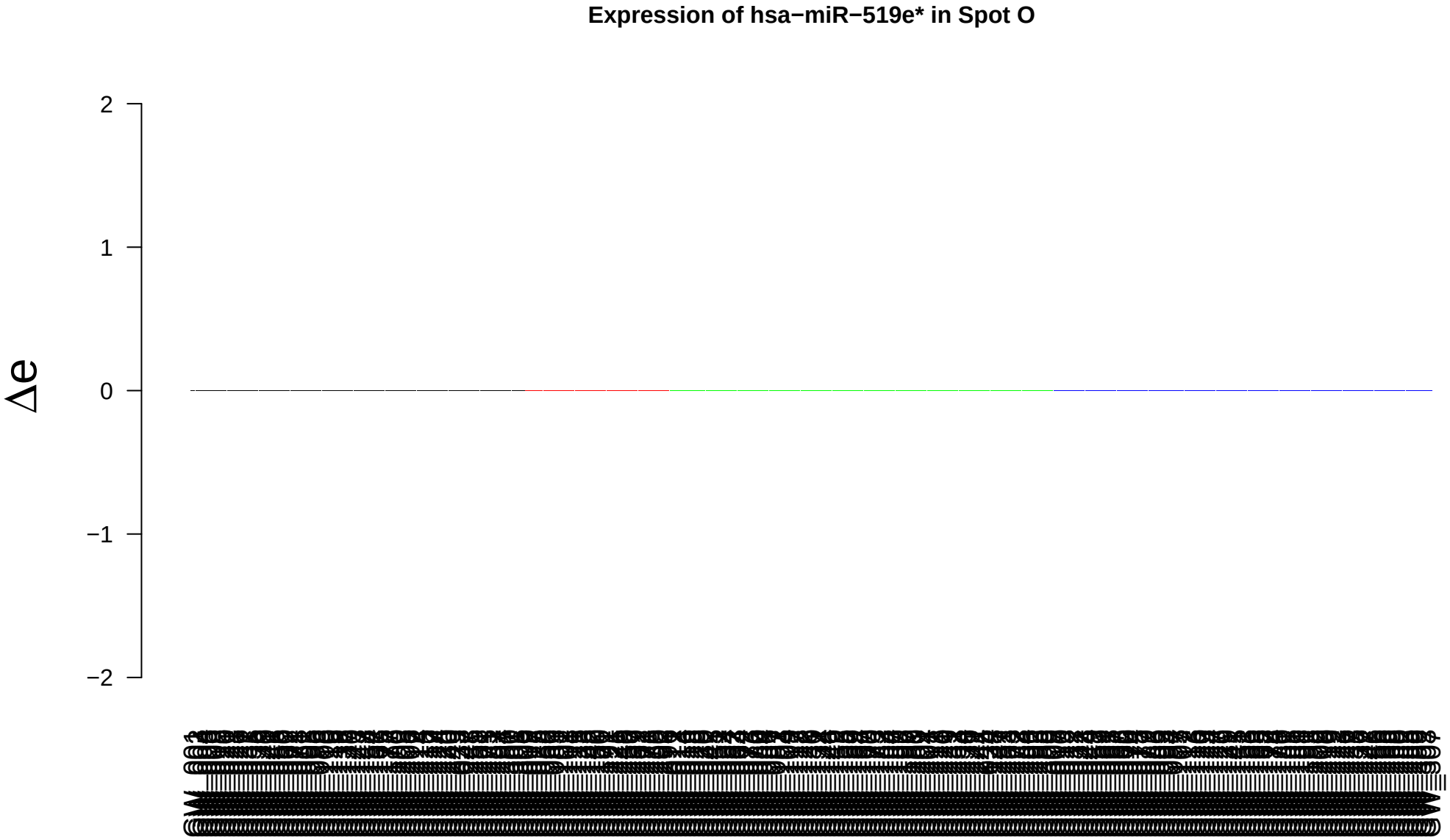

Expression of hsa−miR−519e* in Spot J
Δe
2
1
0
−1
−2

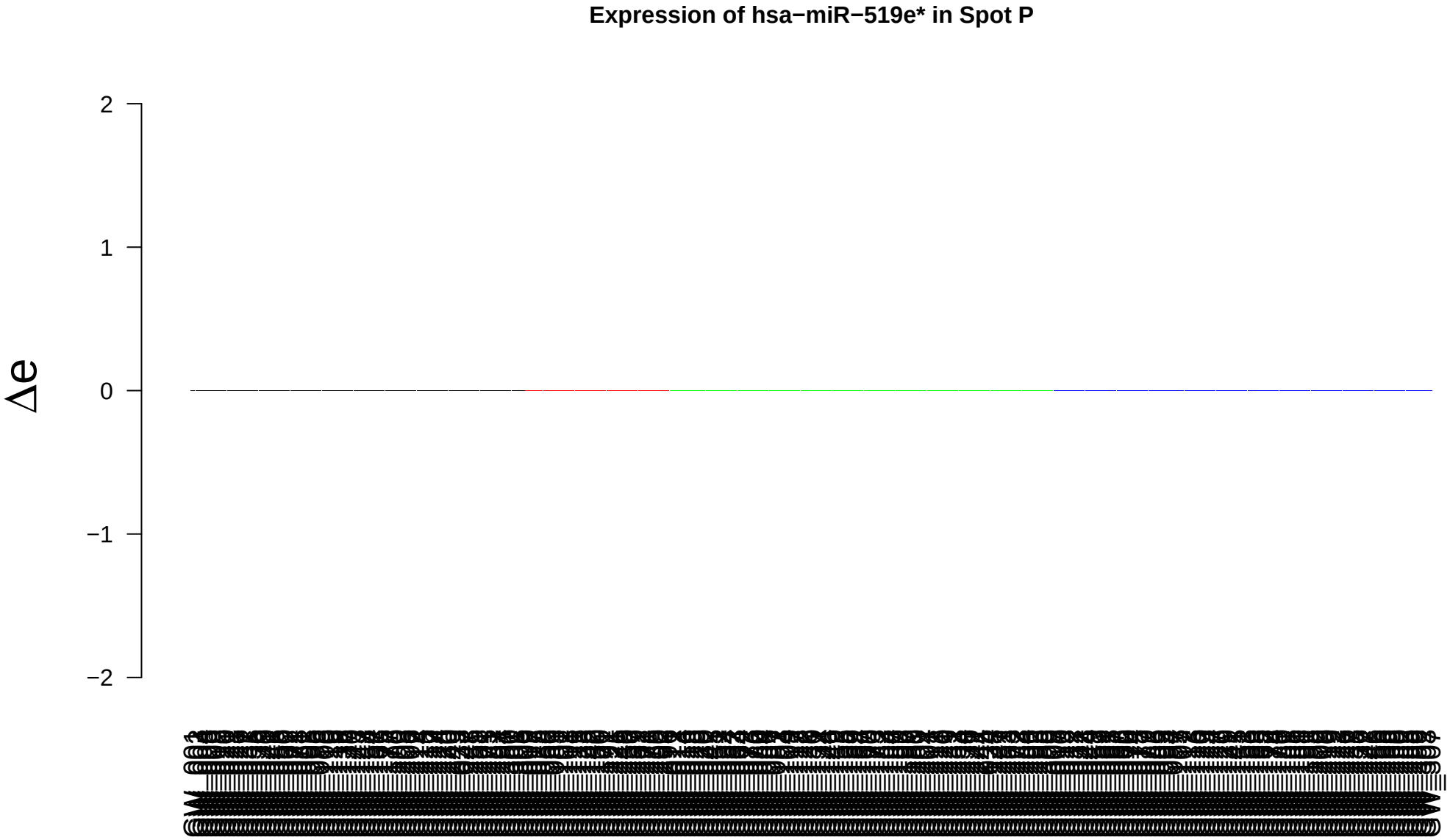

Expression of hsa−miR−519e* in Spot K
Δe
2
1
0
−1
−2

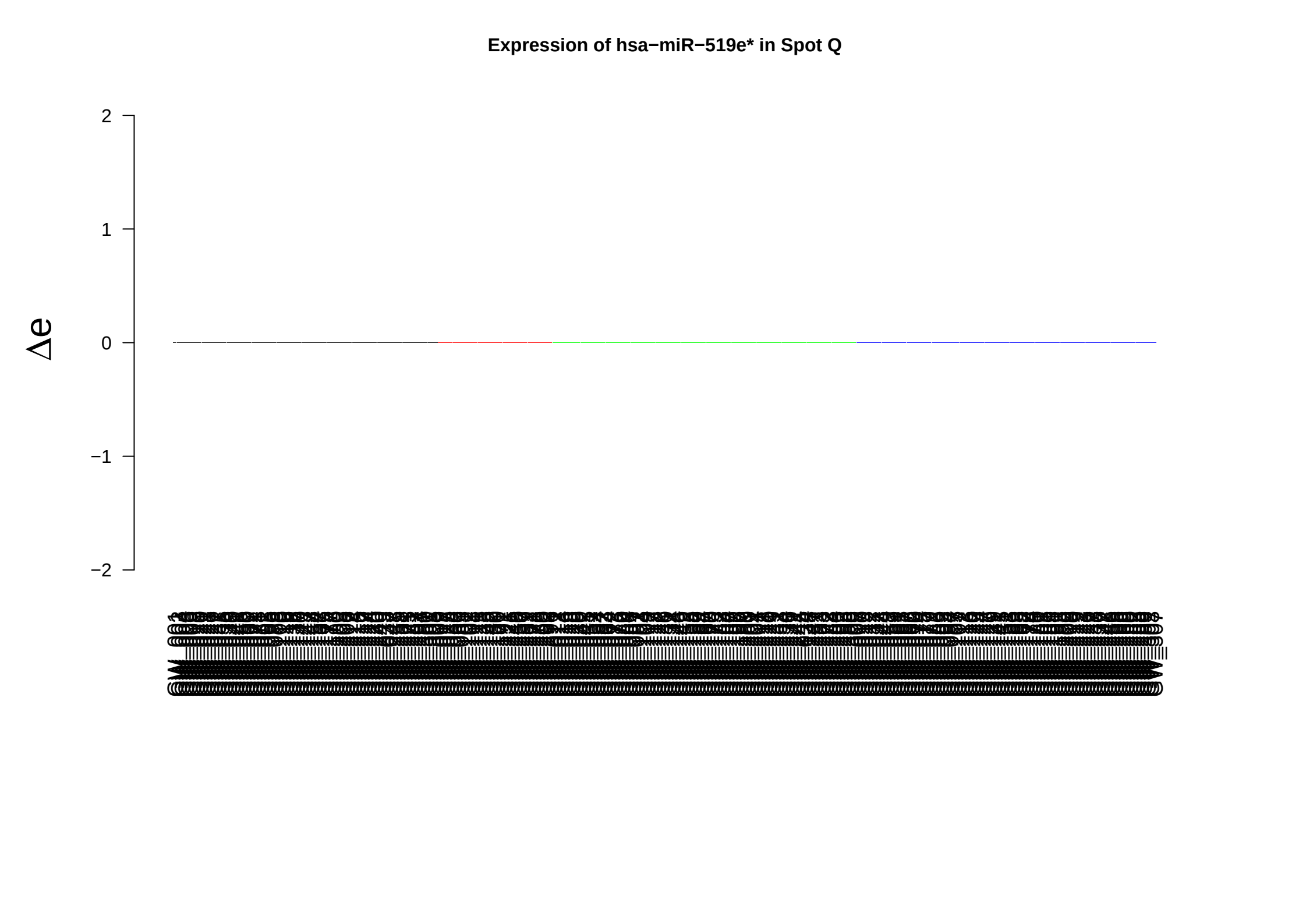

Expression of hsa−miR−519e* in Spot L
Δe
2
1
0
−1
−2

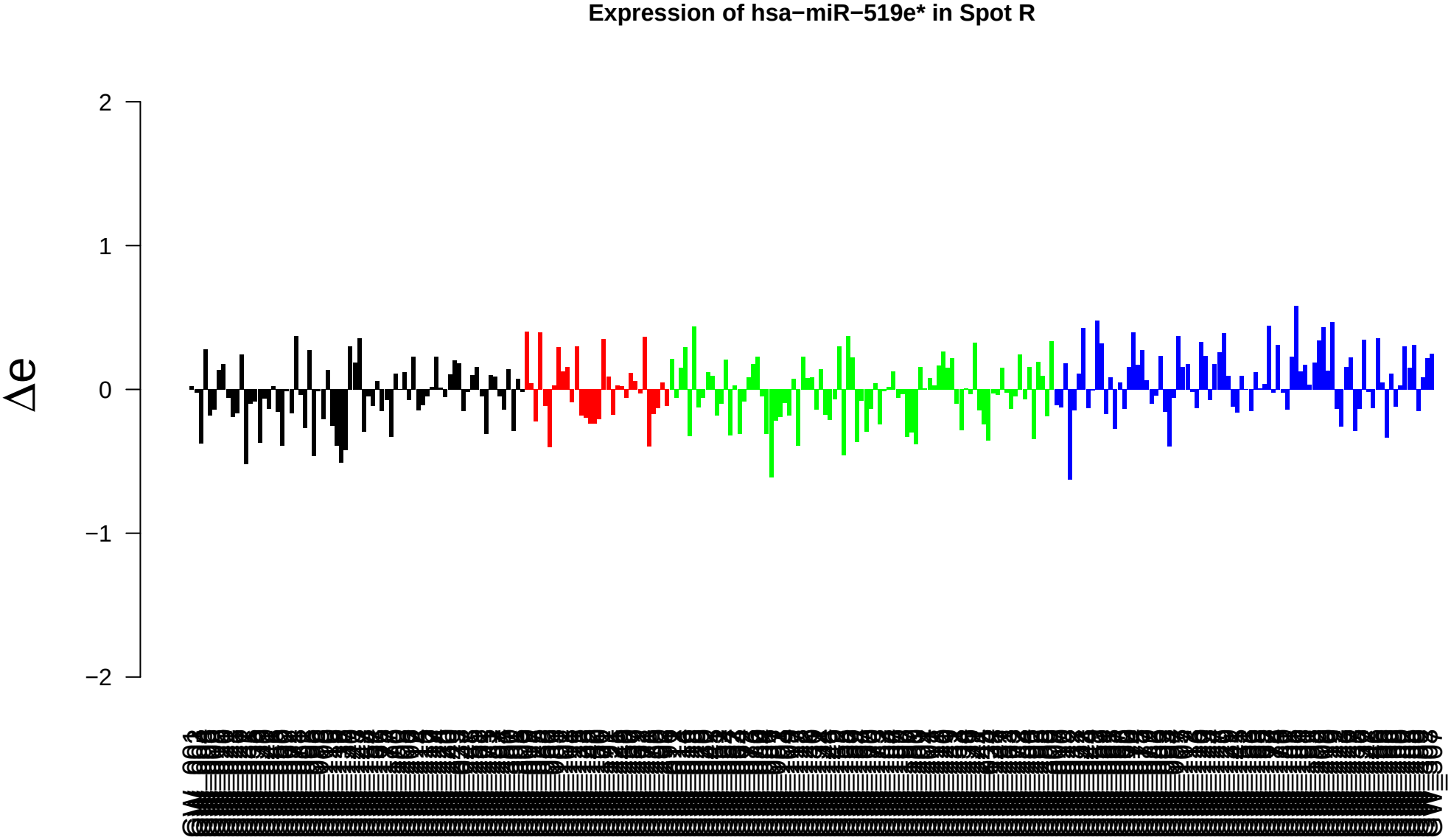

Expression of hsa−miR−519e* in Spot M
Δe
2
1
0
−1
−2


Expression of hsa−miR−519e* in Spot N
Δe
2
1
0
−1
−2


Expression of hsa−miR−519e* in Spot O
Δe
2
1
0
−1
−2


Expression of hsa−miR−519e* in Spot P
Δe
2
1
0
−1
−2


Expression of hsa−miR−519e* in Spot Q
Δe
2
1
0
−1
−2


Expression of hsa−miR−519e* in Spot R
Δe
2
1
0
−1
−2