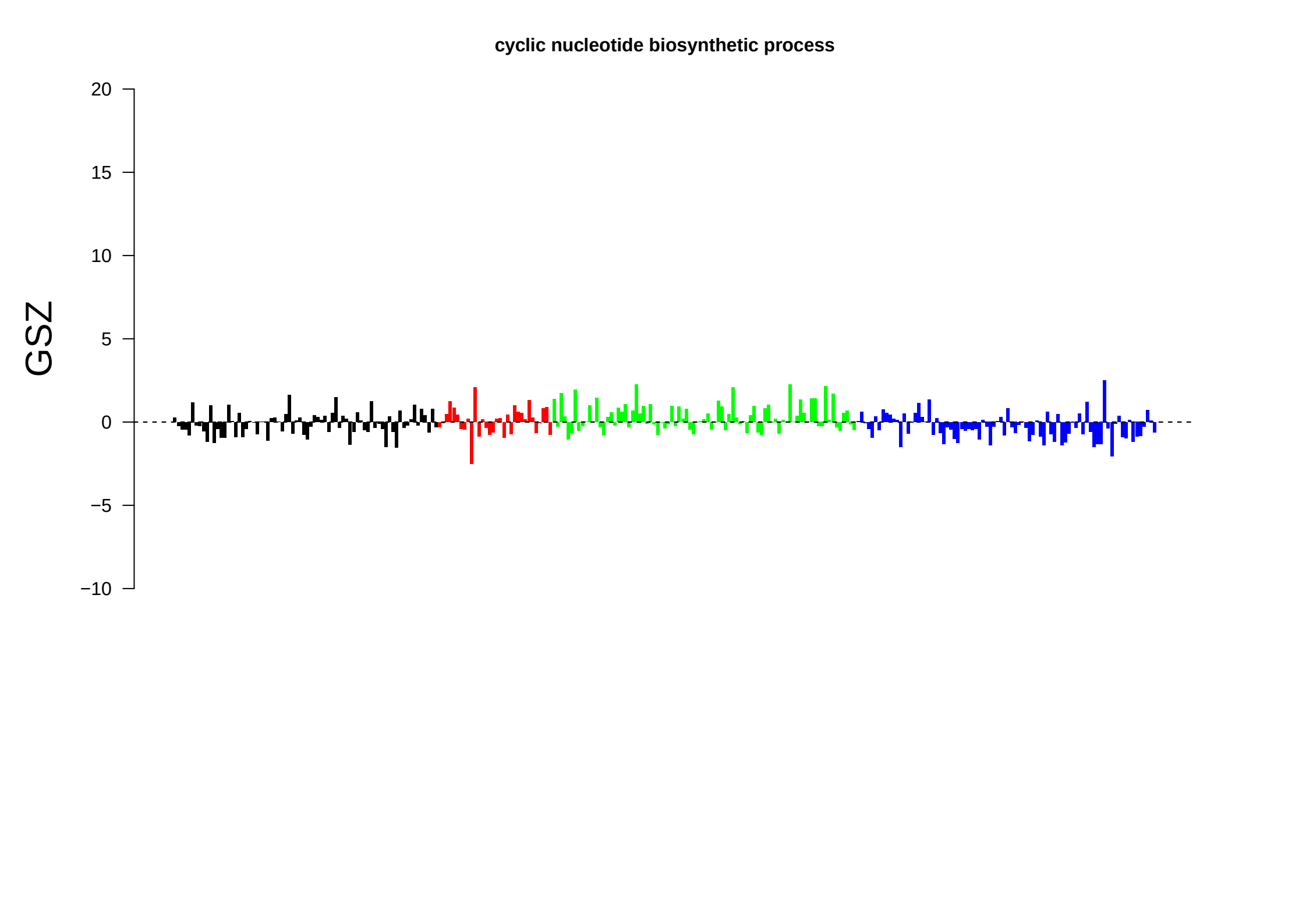
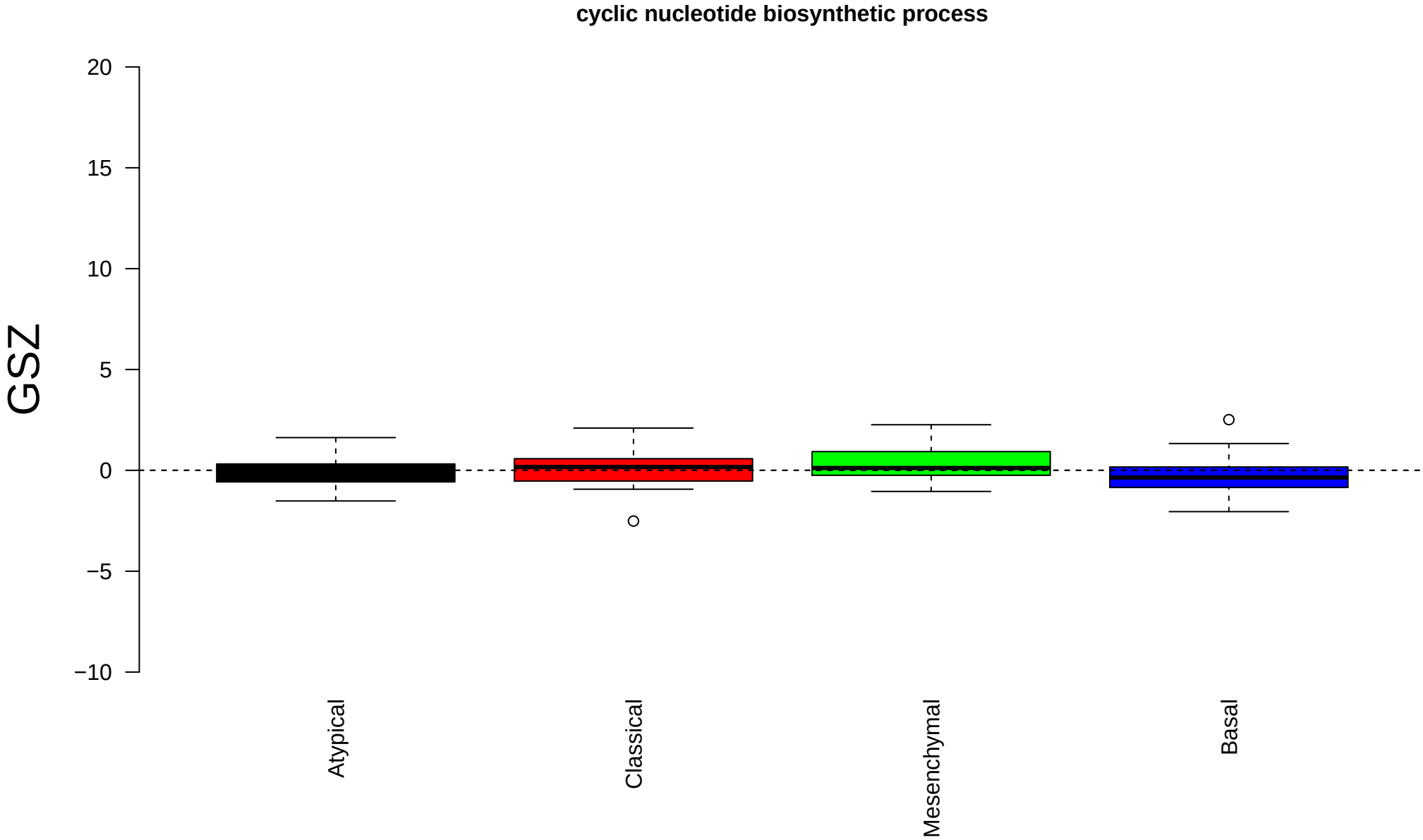
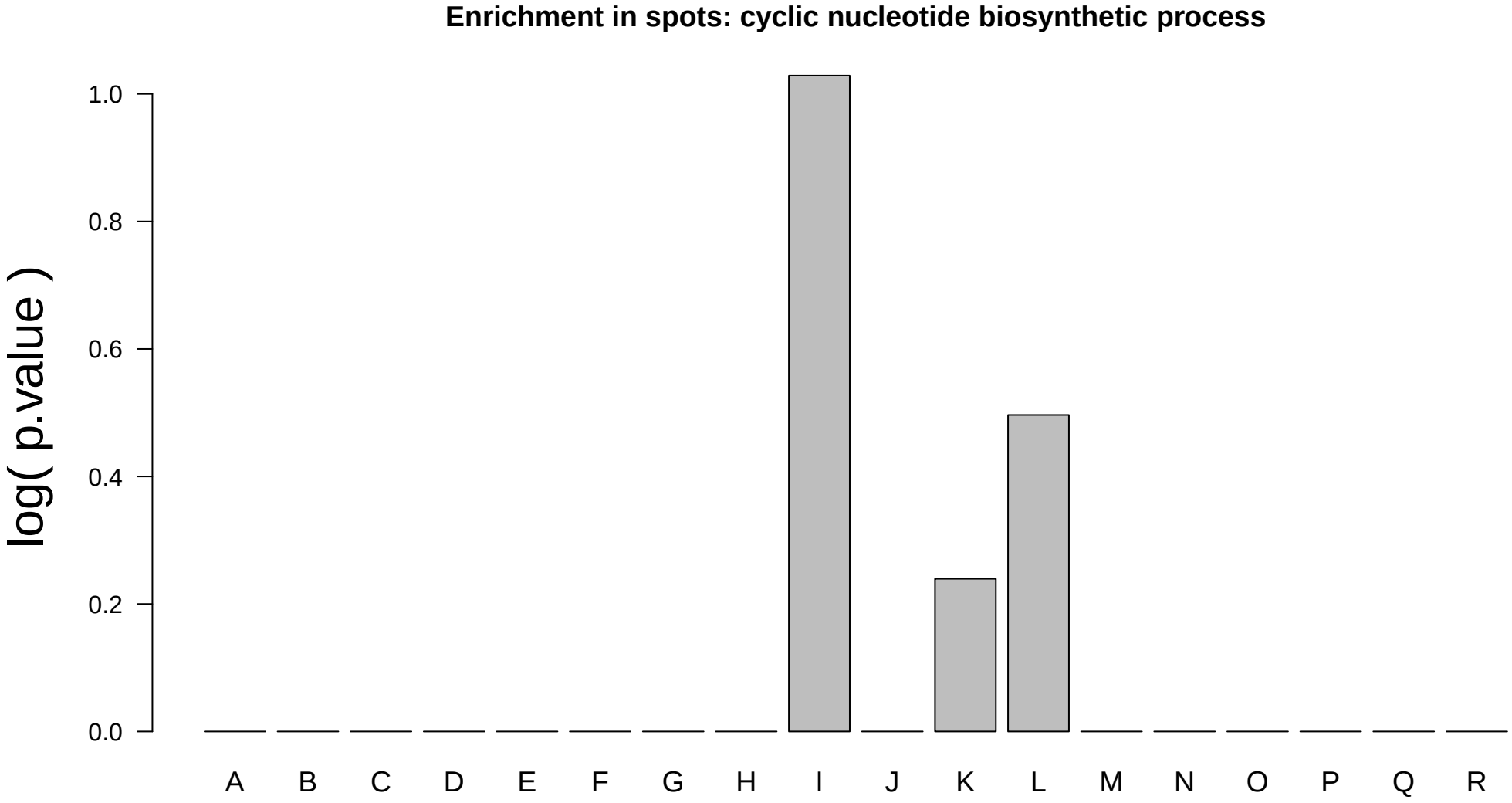
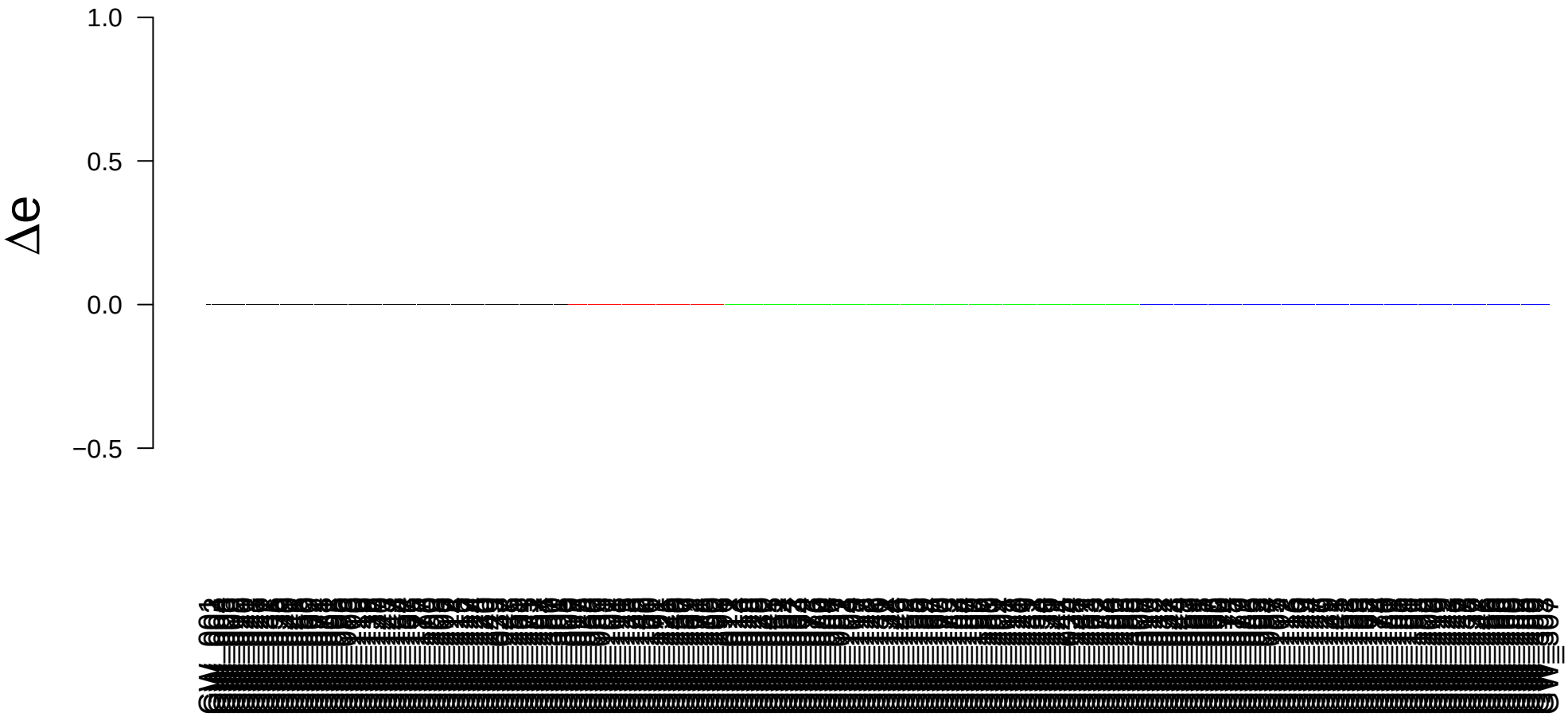

cyclic nucleotide biosynthetic process
GSZ
20
15
10
5
0
−5
−10

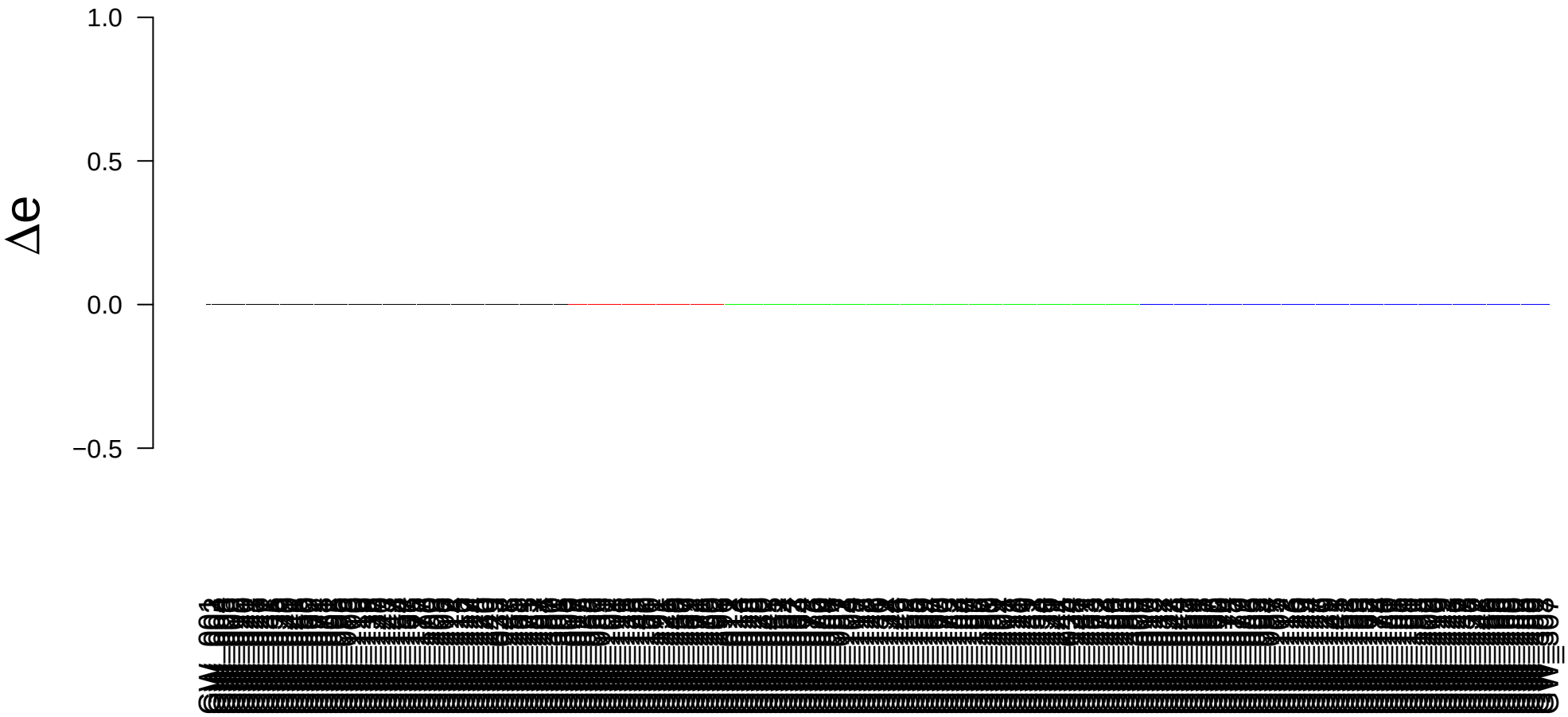

cyclic nucleotide biosynthetic process
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal


Enrichment in spots: cyclic nucleotide biosynthetic process
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

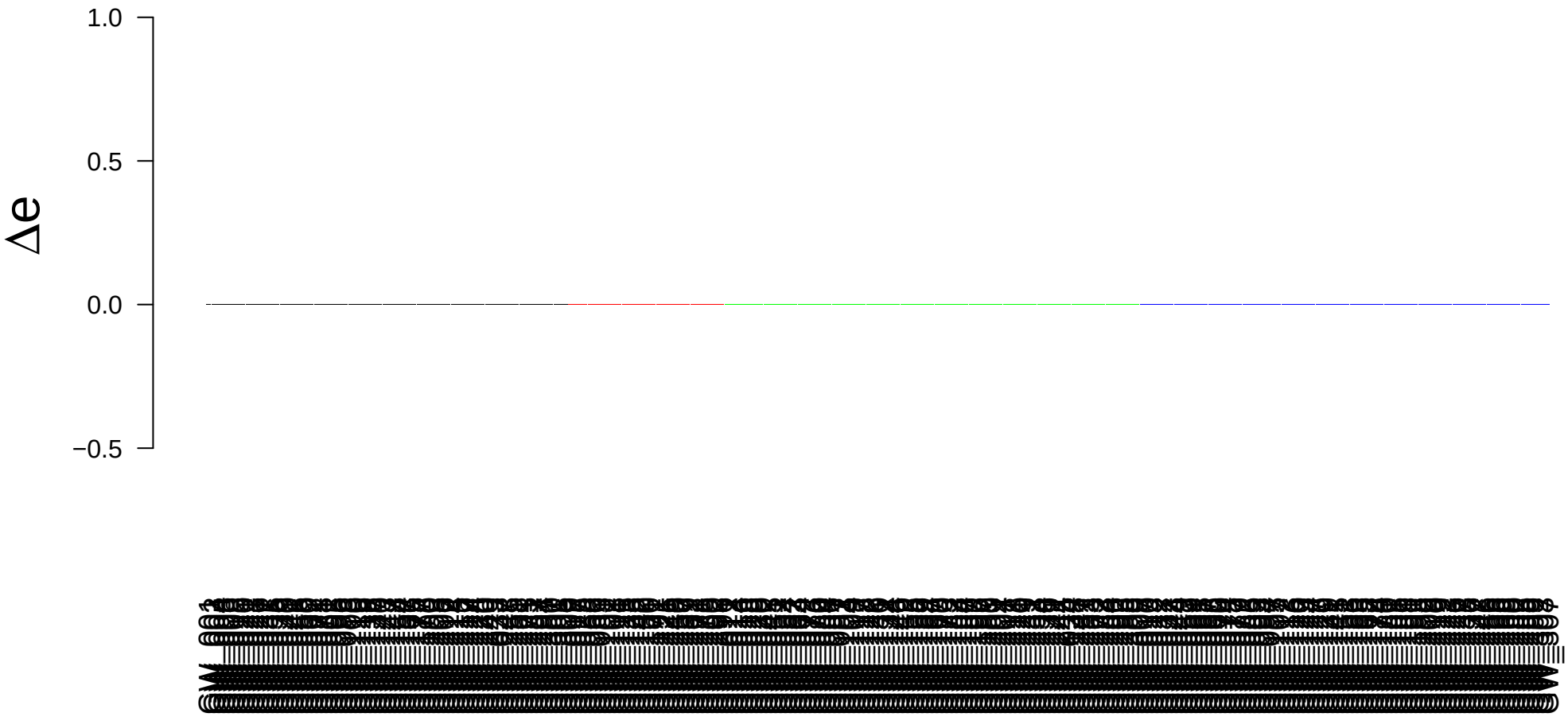
Expression of cyclic nucleotide biosynthetic process in Spot A



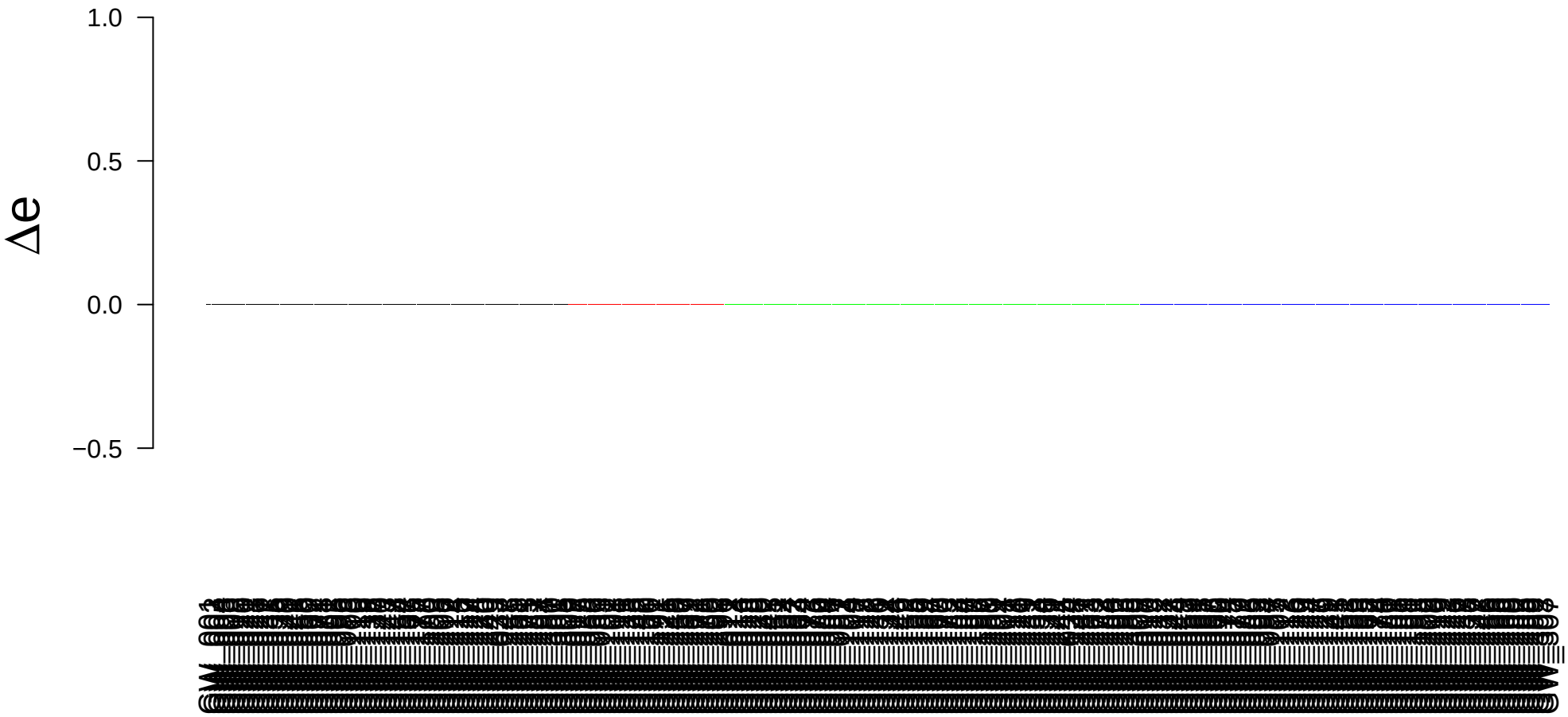
Expression of cyclic nucleotide biosynthetic process in Spot B



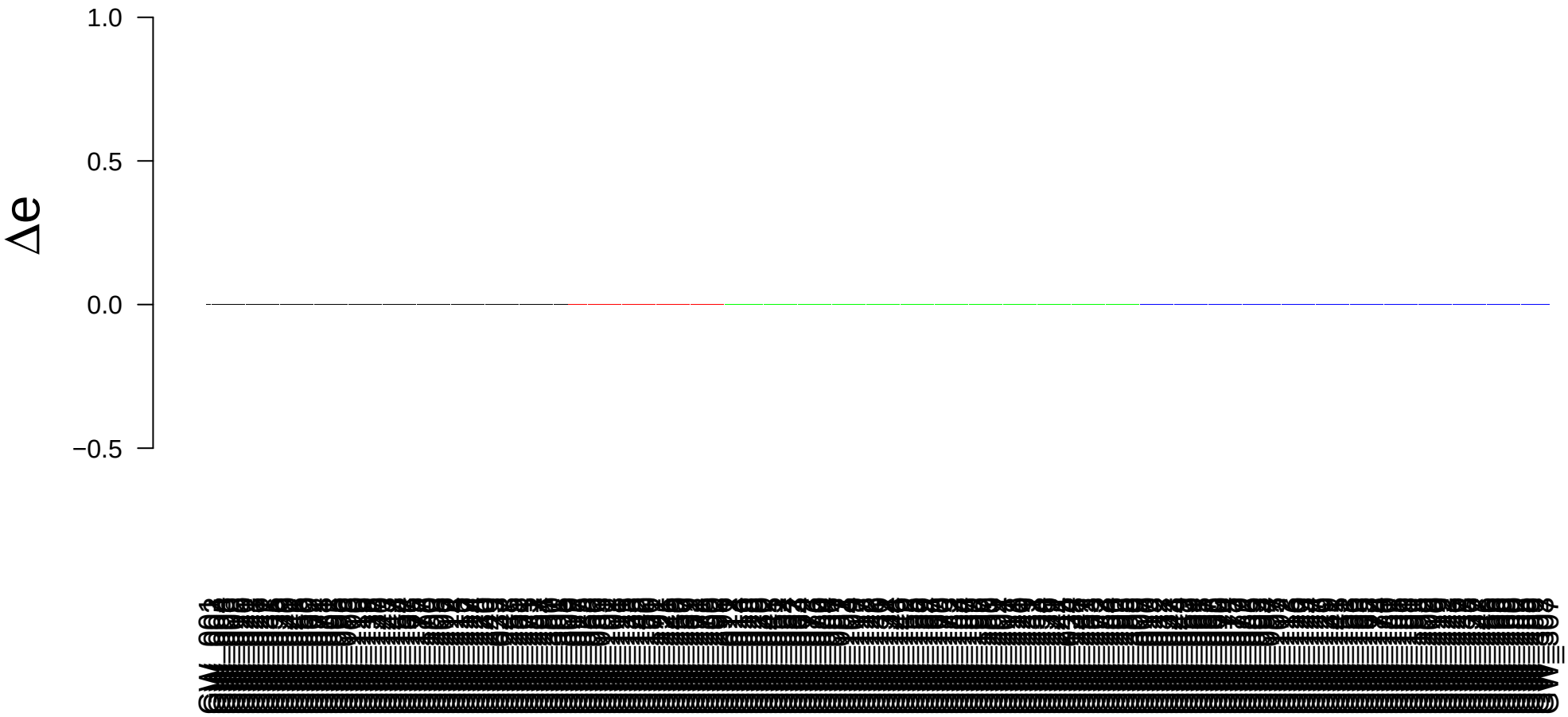
Expression of cyclic nucleotide biosynthetic process in Spot C



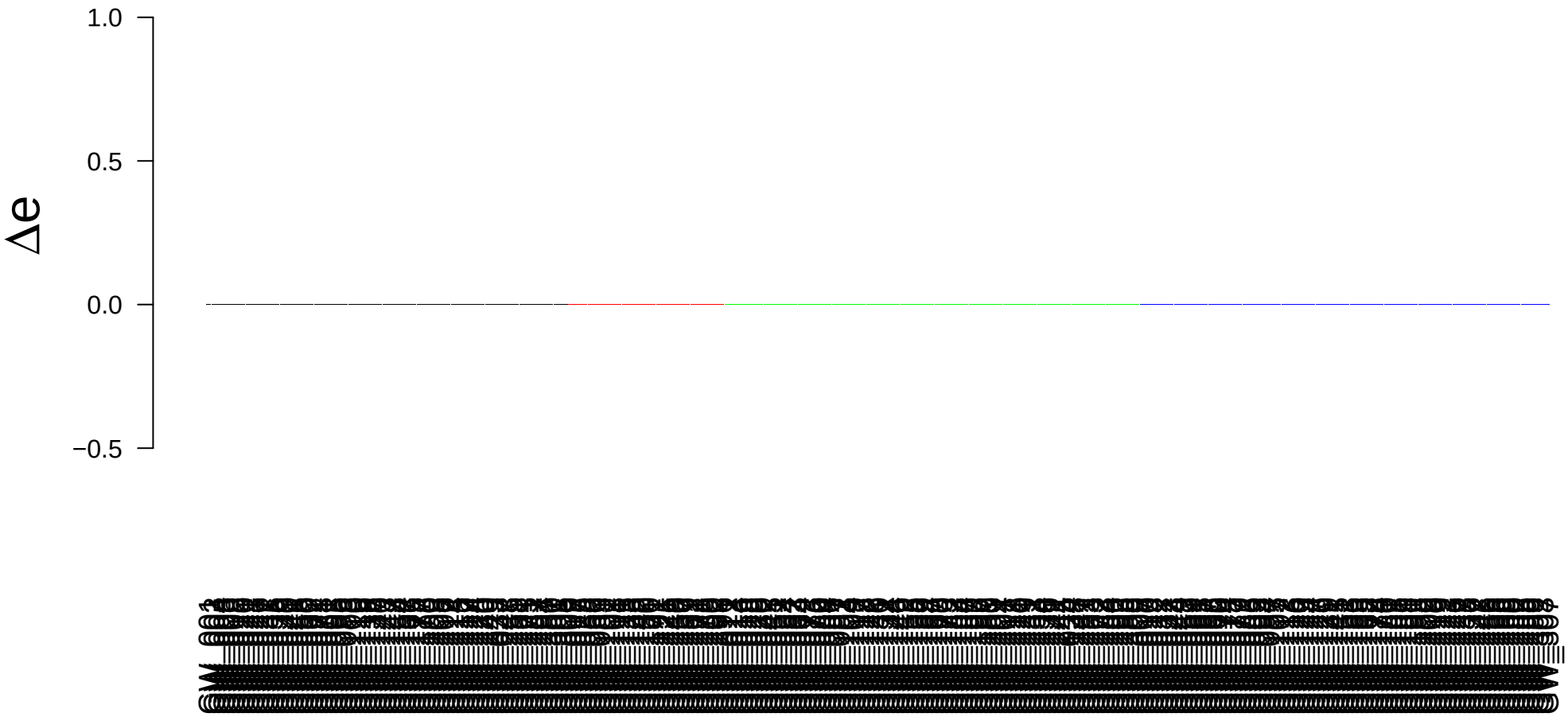
Expression of cyclic nucleotide biosynthetic process in Spot D



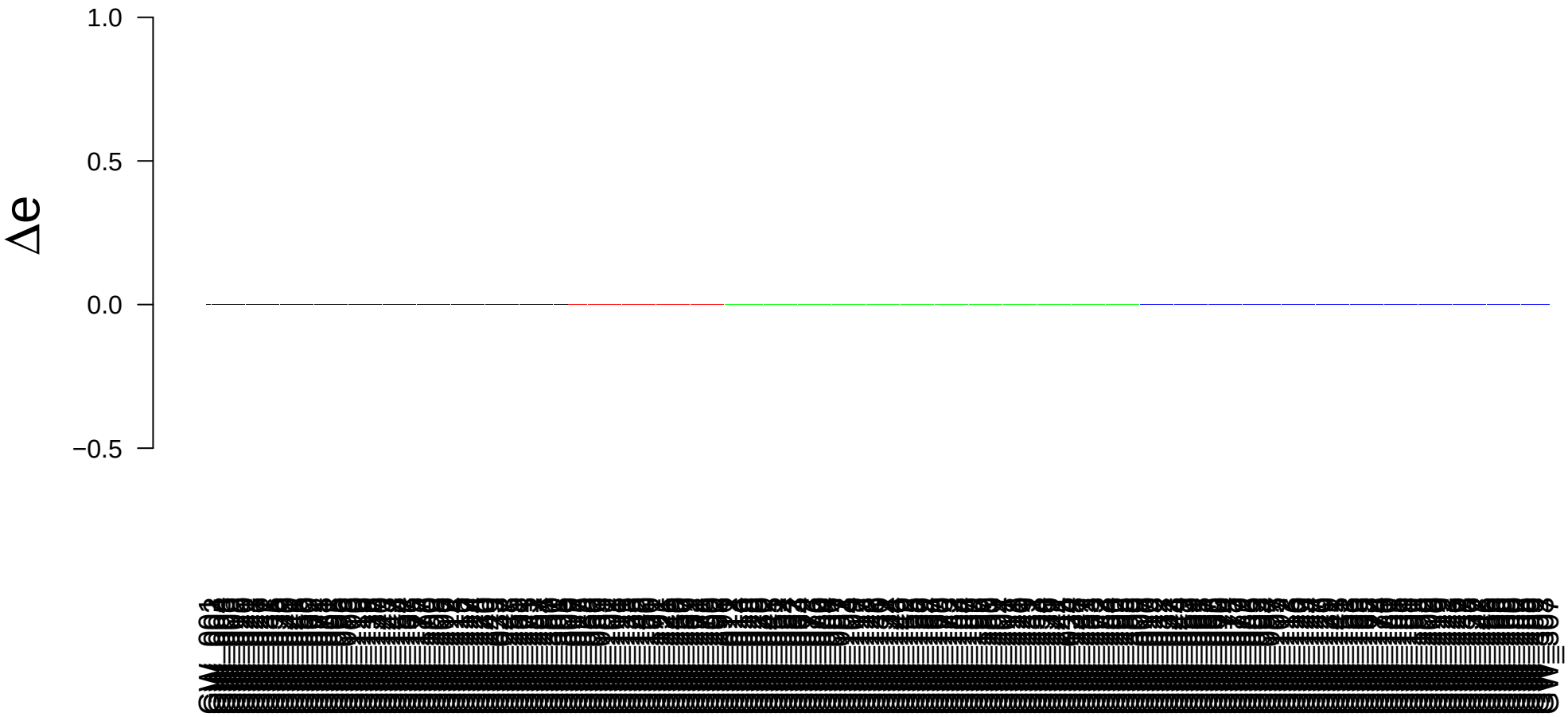
Expression of cyclic nucleotide biosynthetic process in Spot E



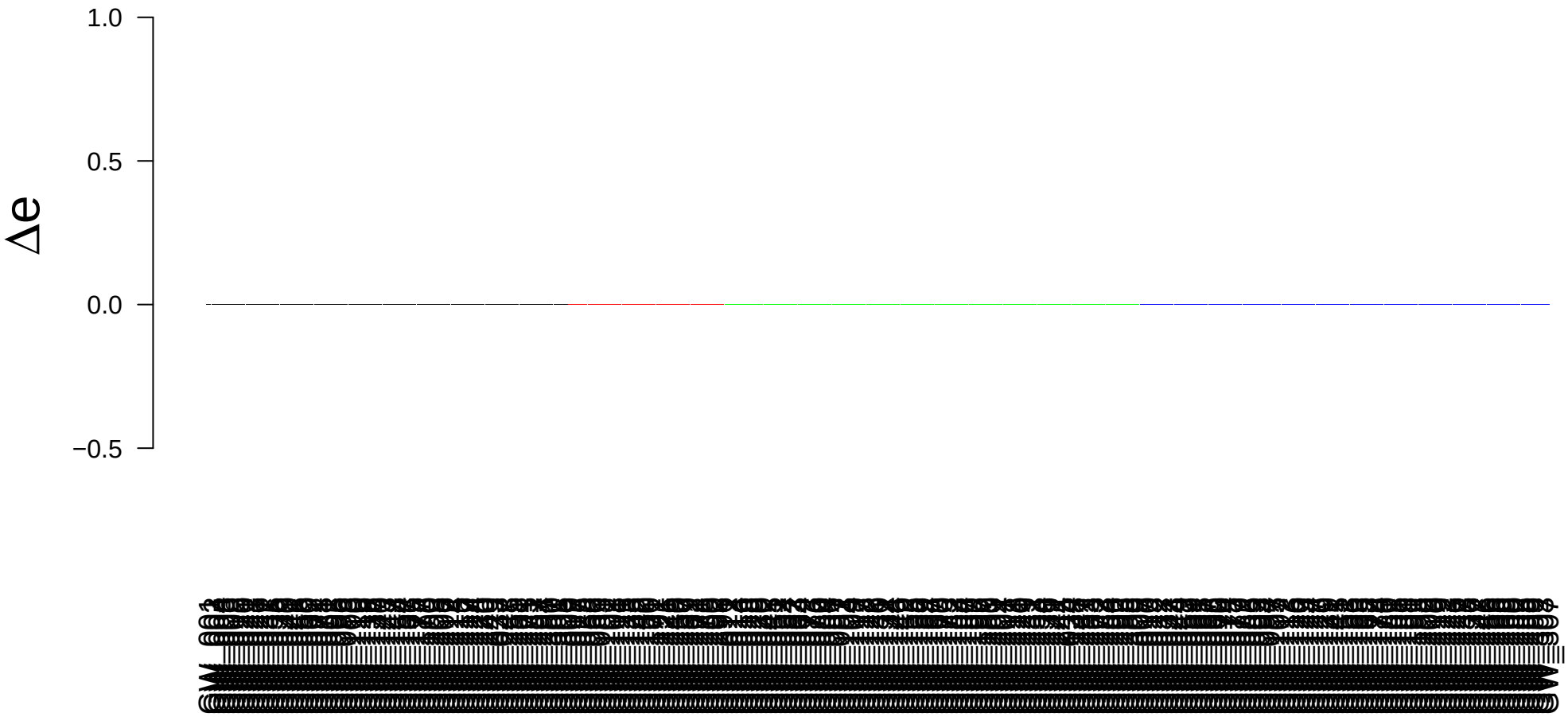
Expression of cyclic nucleotide biosynthetic process in Spot F

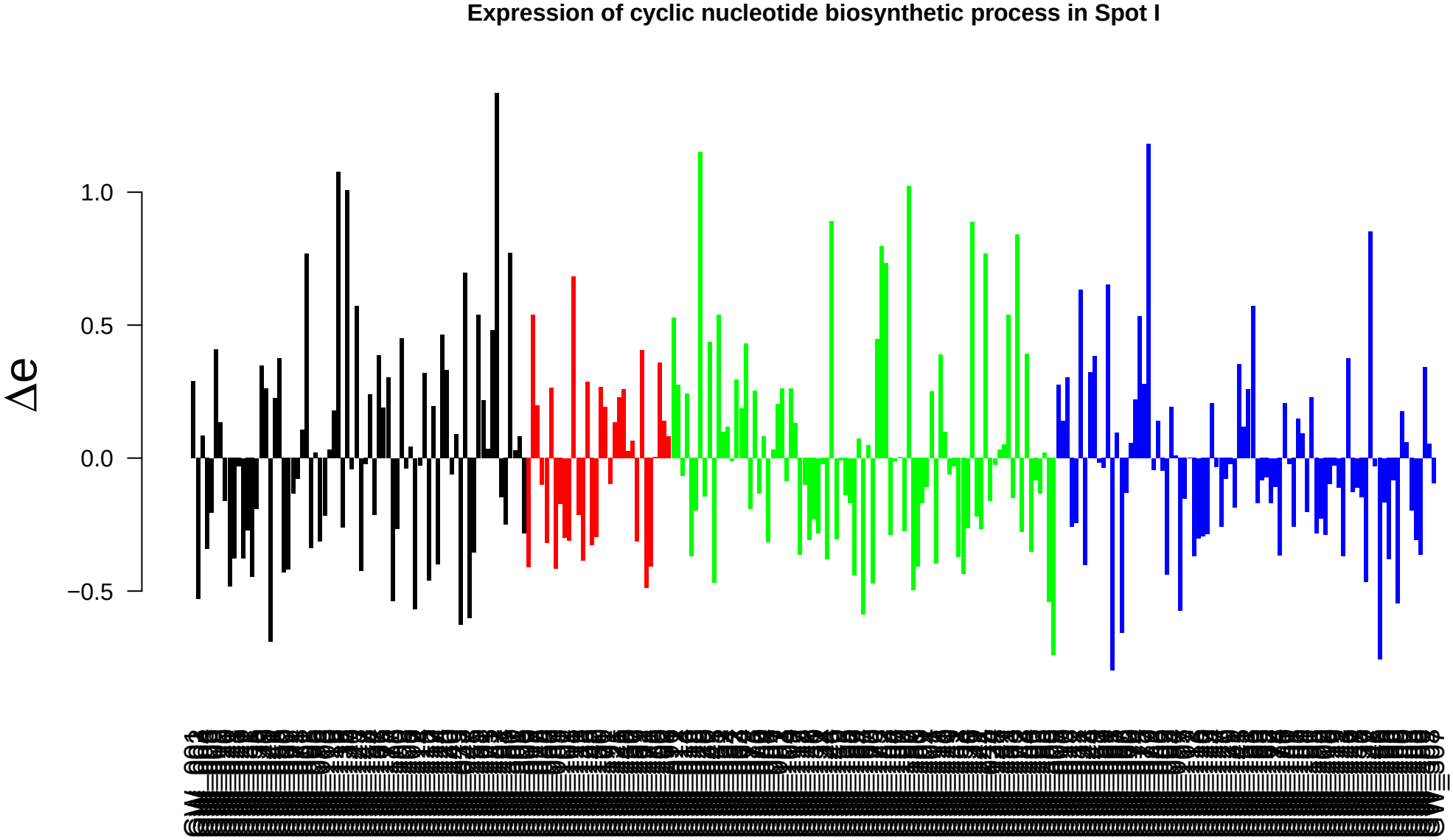


Expression of cyclic nucleotide biosynthetic process in Spot G

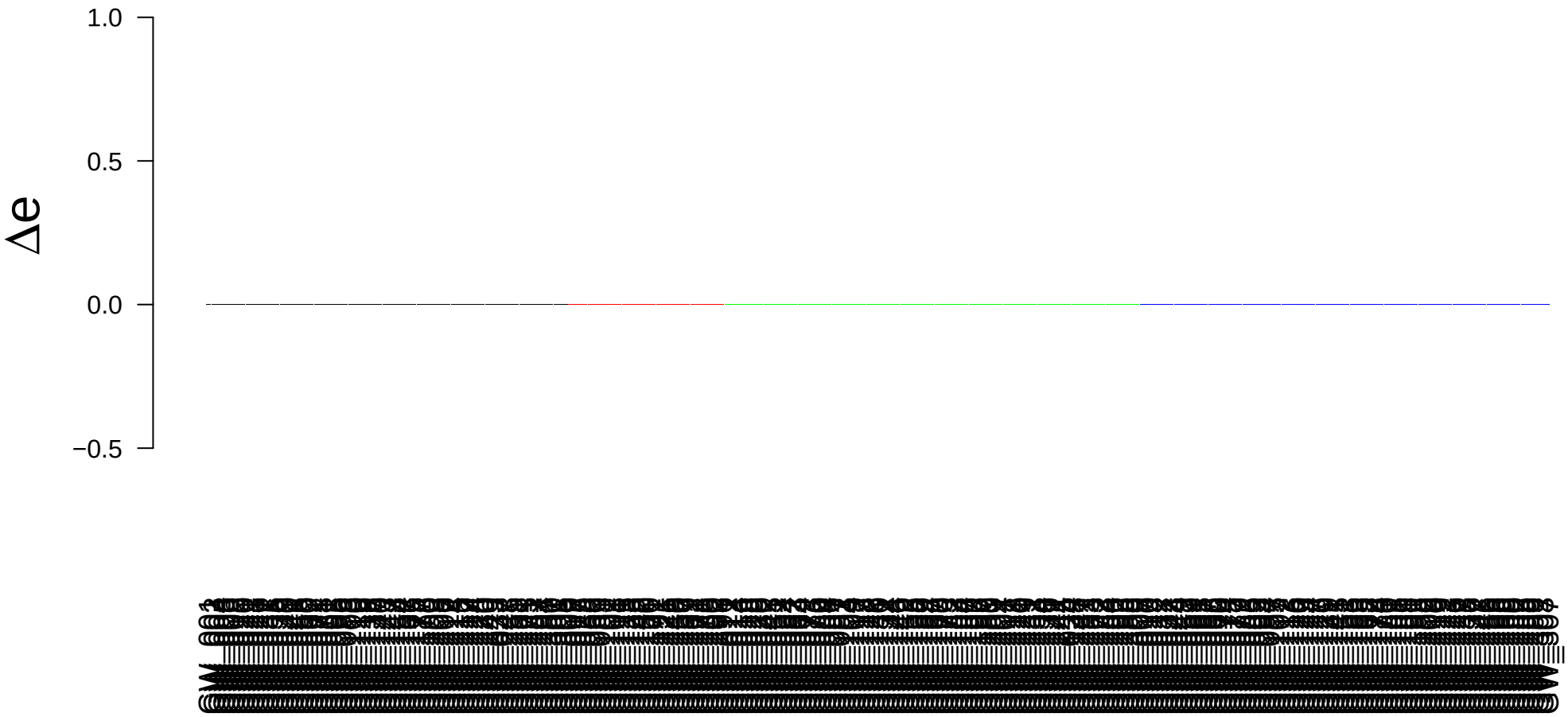


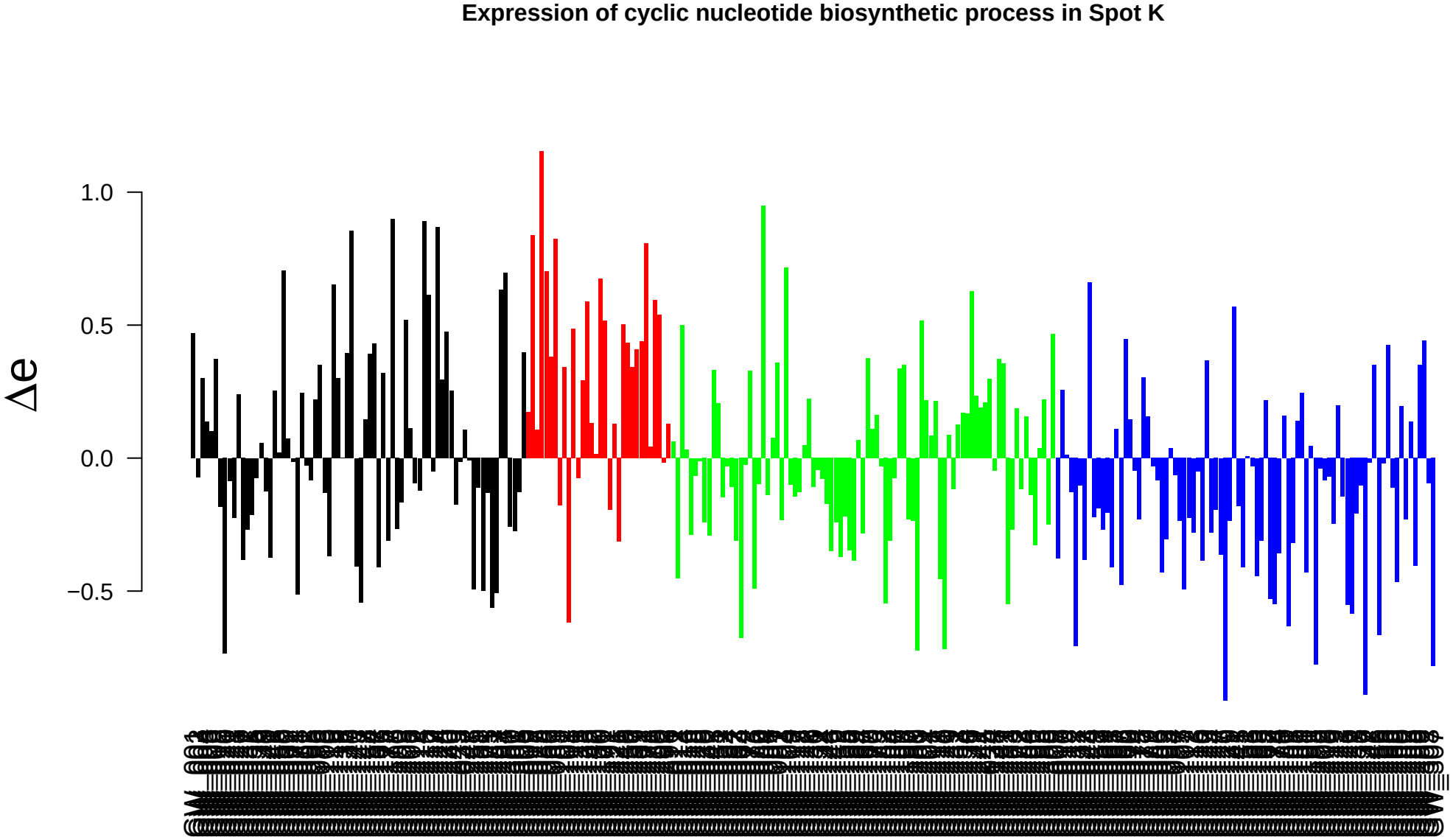
Expression of cyclic nucleotide biosynthetic process in Spot H

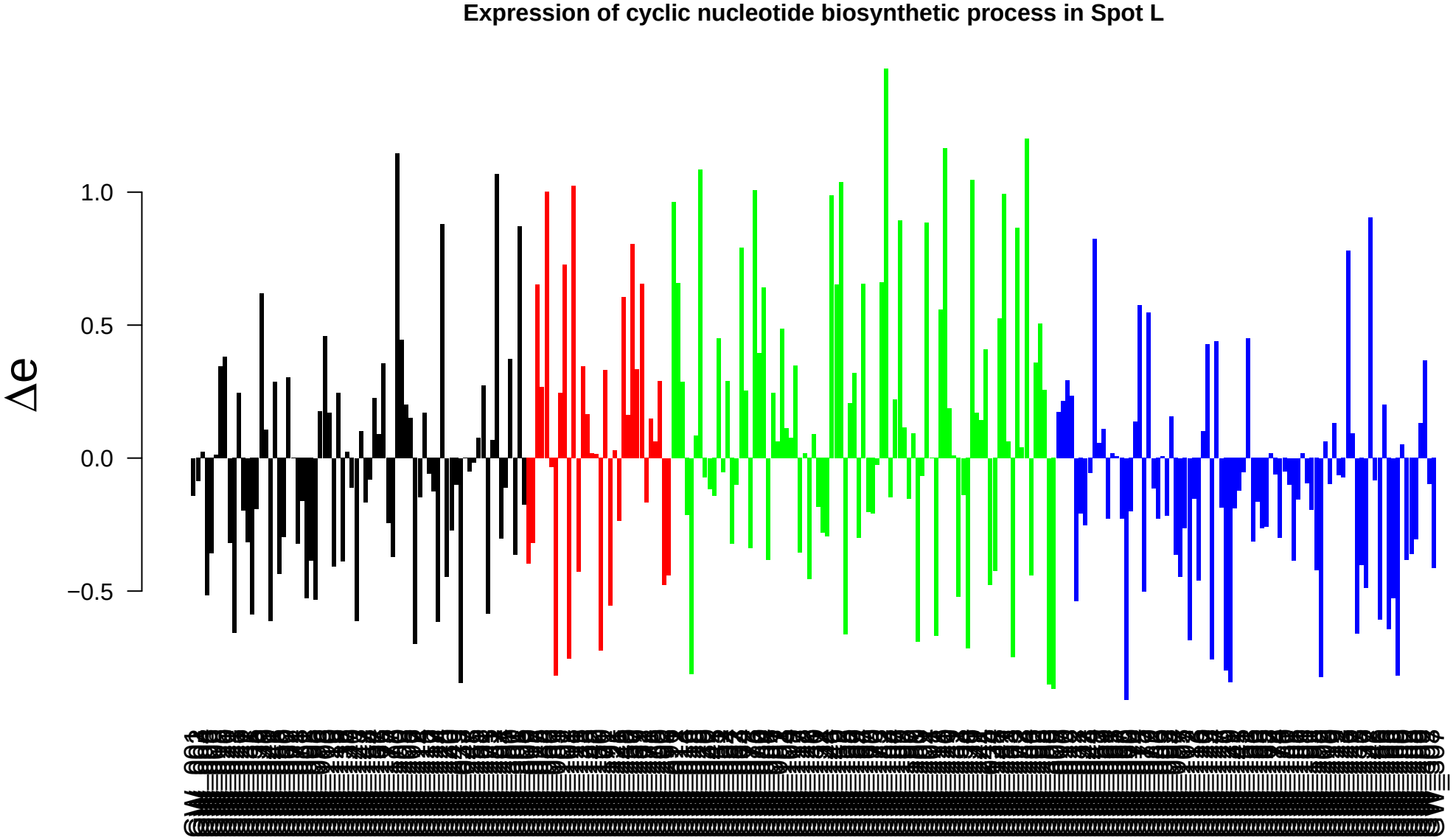



Expression of cyclic nucleotide biosynthetic process in Spot I
Δe
1.0
0.5
0.0
−0.5

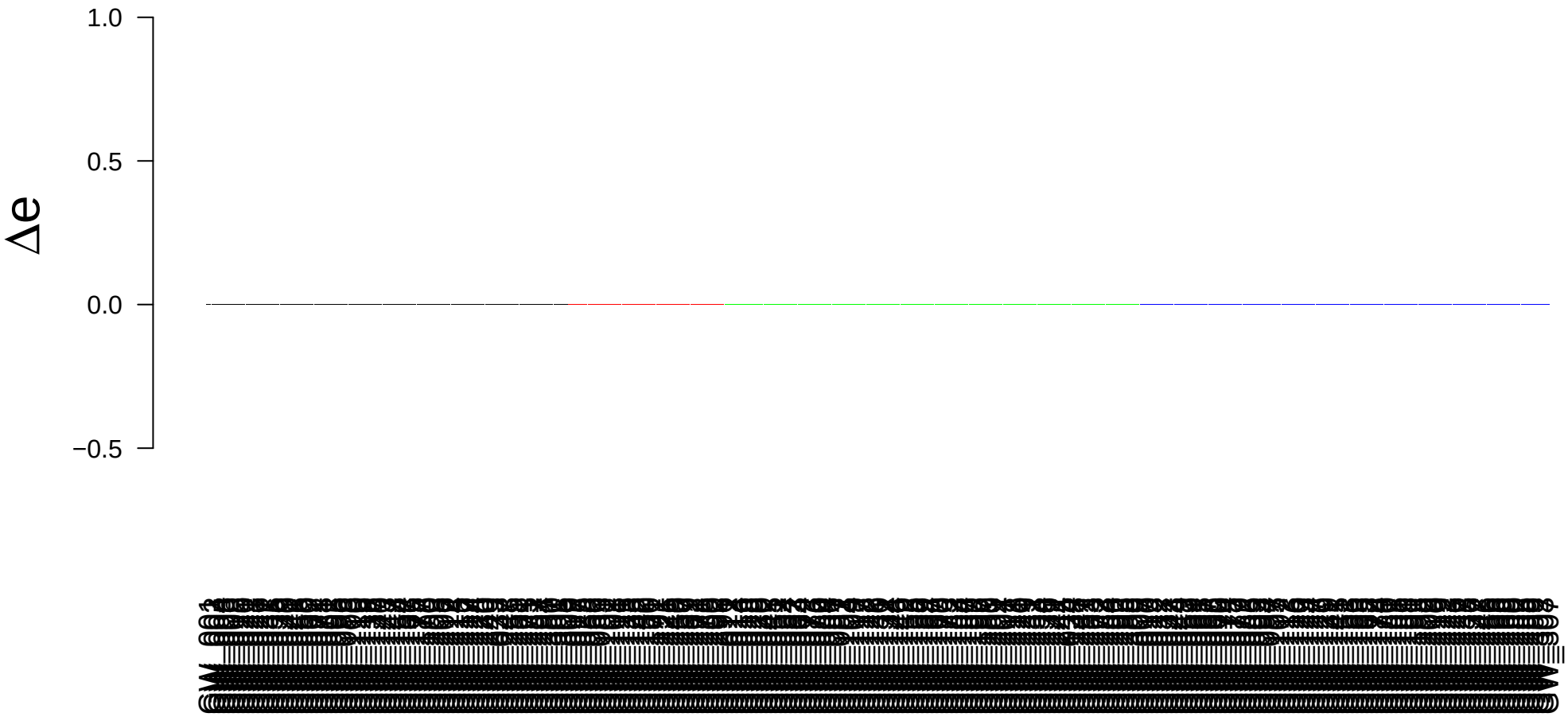
Expression of cyclic nucleotide biosynthetic process in Spot J



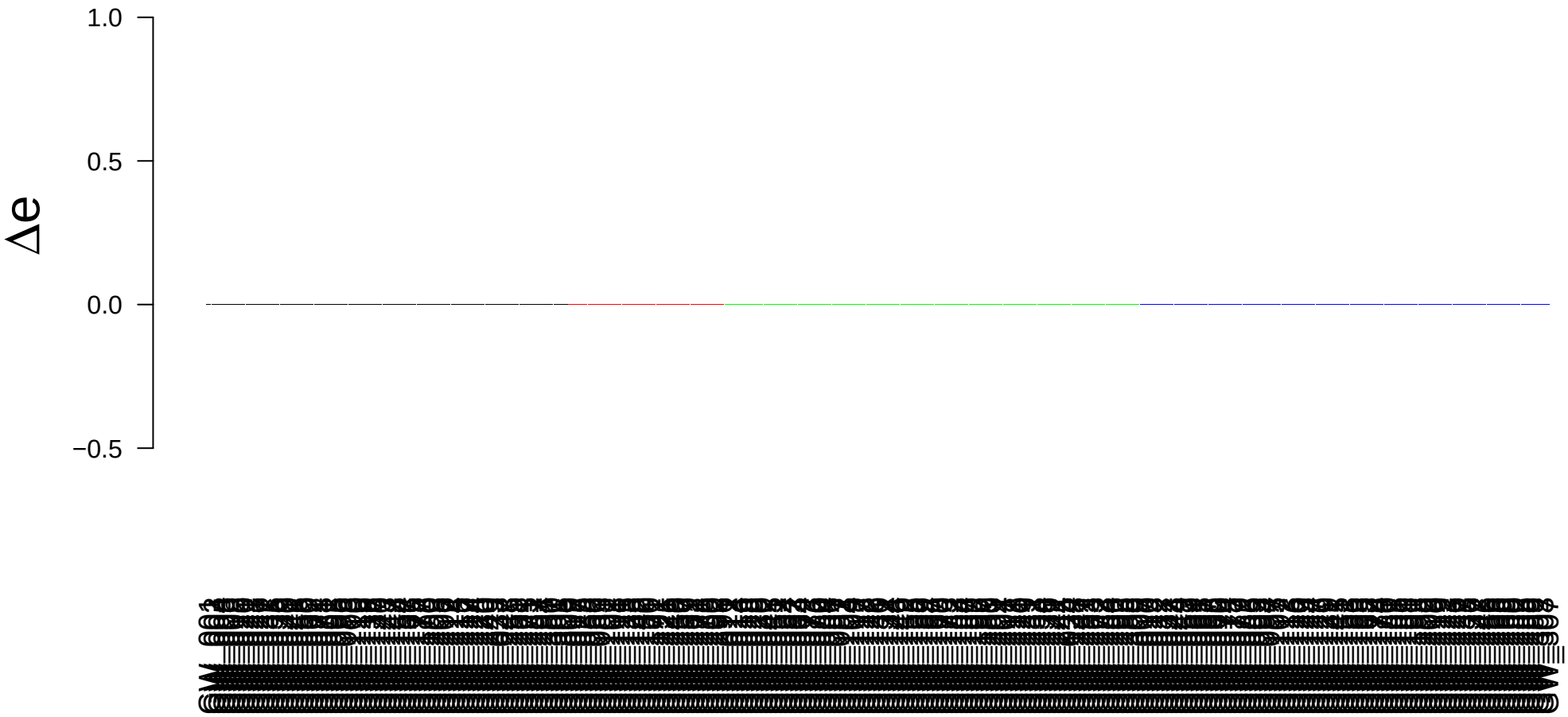

Expression of cyclic nucleotide biosynthetic process in Spot K
Δe
1.0
0.5
0.0
−0.5


Expression of cyclic nucleotide biosynthetic process in Spot L
Δe
1.0
0.5
0.0
−0.5

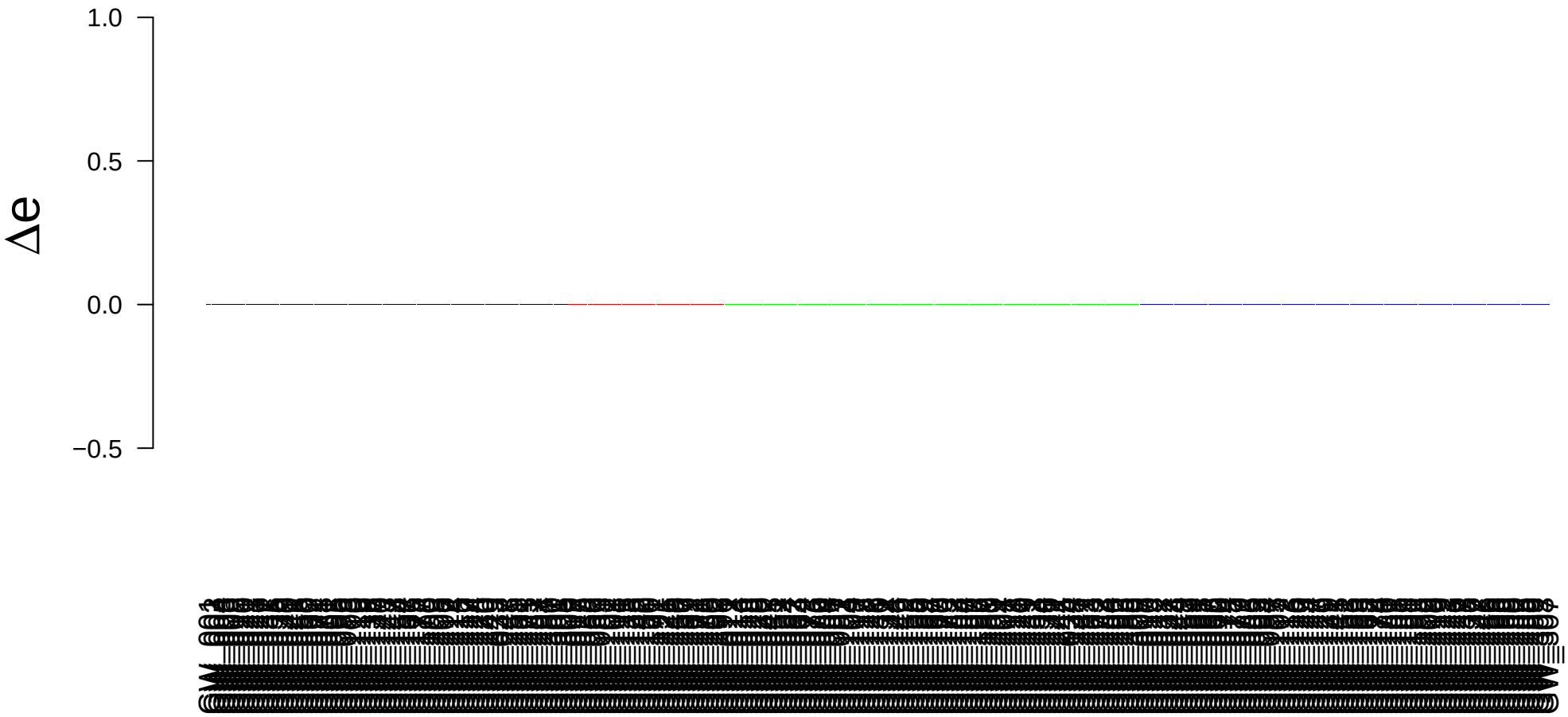
Expression of cyclic nucleotide biosynthetic process in Spot M



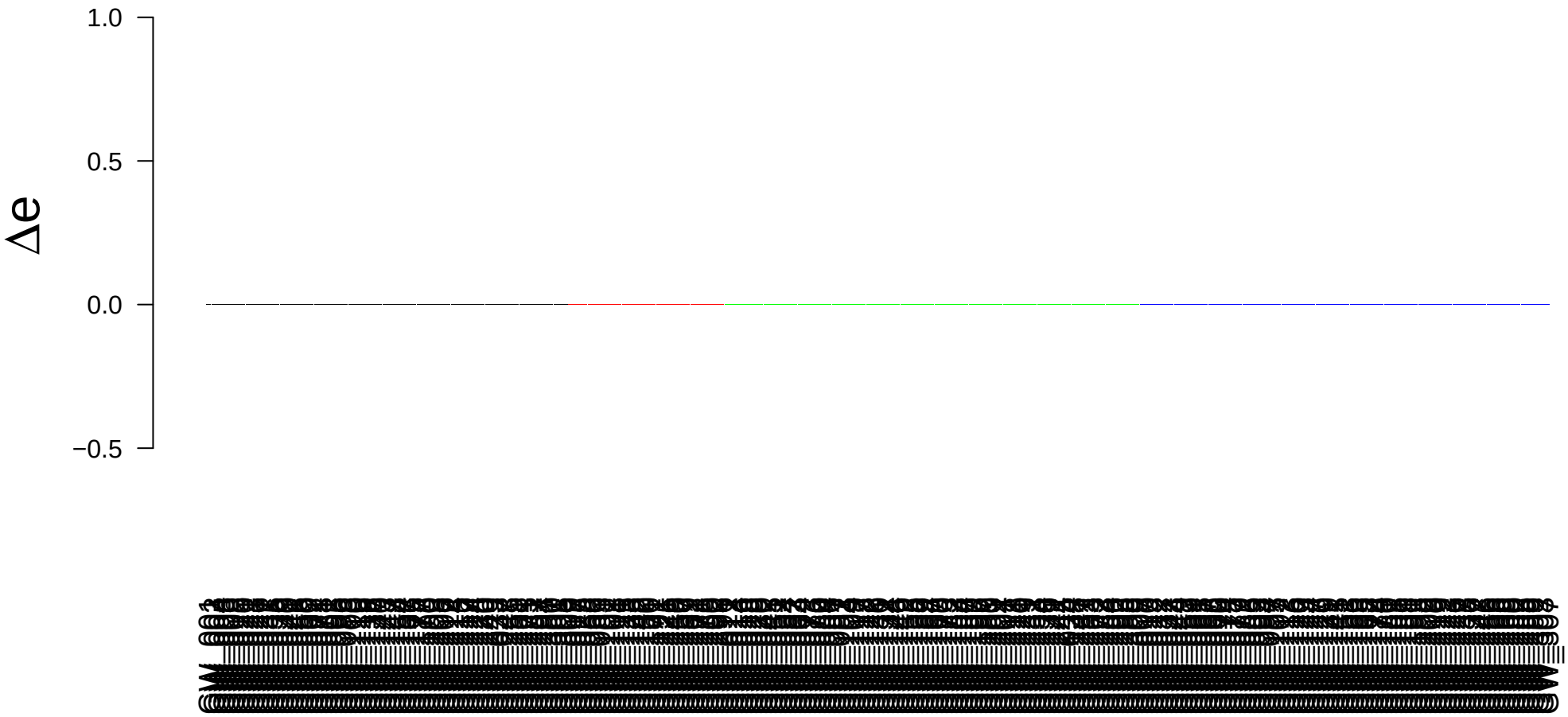
Expression of cyclic nucleotide biosynthetic process in Spot N



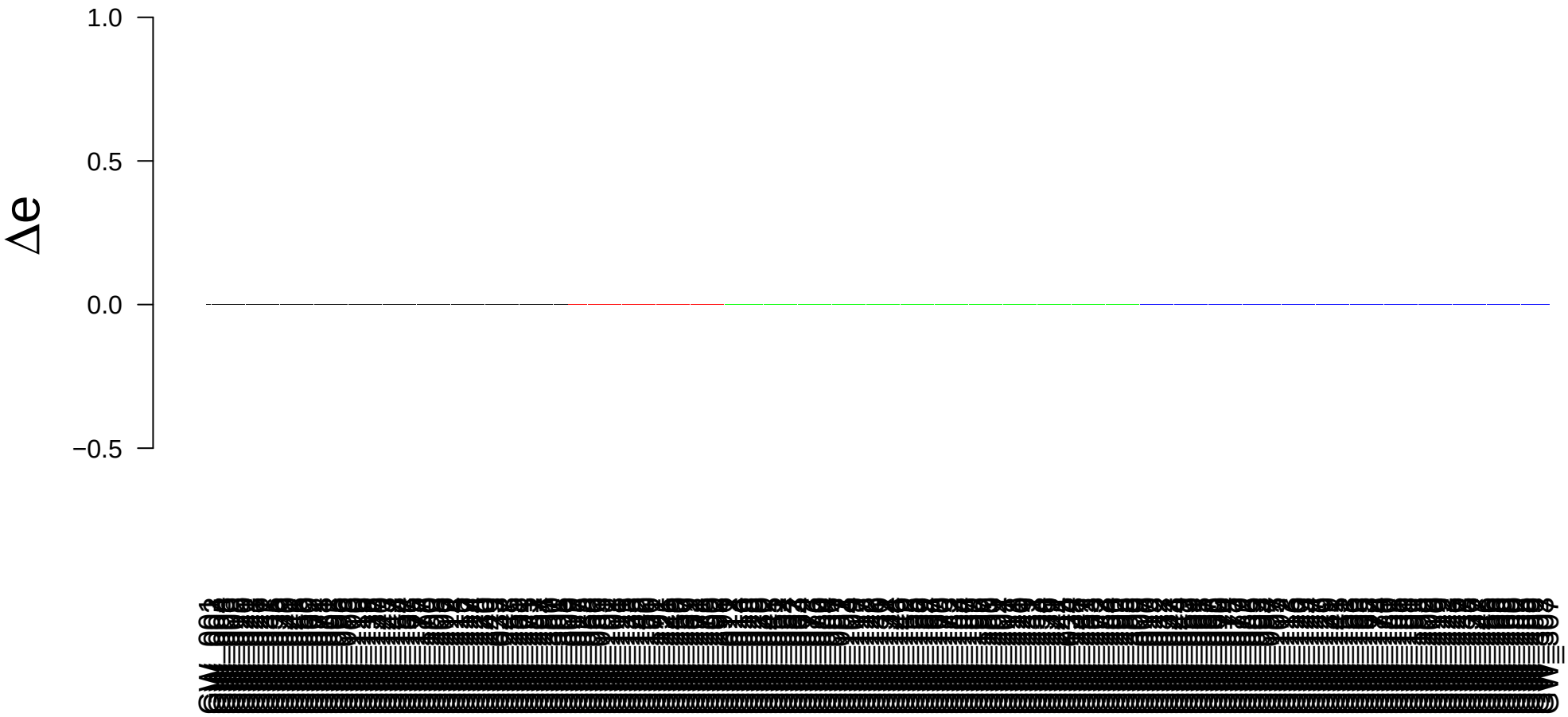
Expression of cyclic nucleotide biosynthetic process in Spot O



Expression of cyclic nucleotide biosynthetic process in Spot P



Expression of cyclic nucleotide biosynthetic process in Spot Q



Expression of cyclic nucleotide biosynthetic process in Spot R