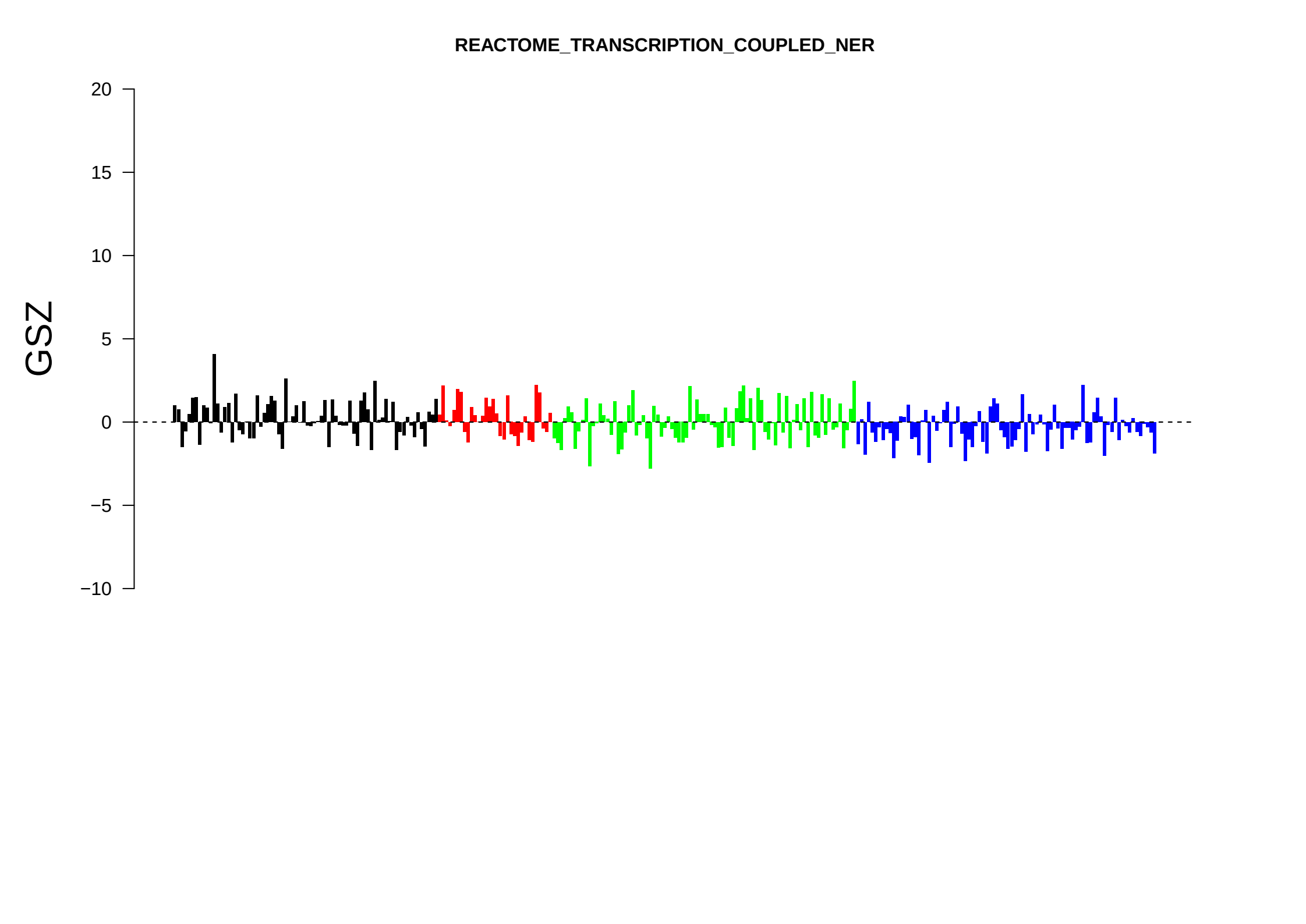
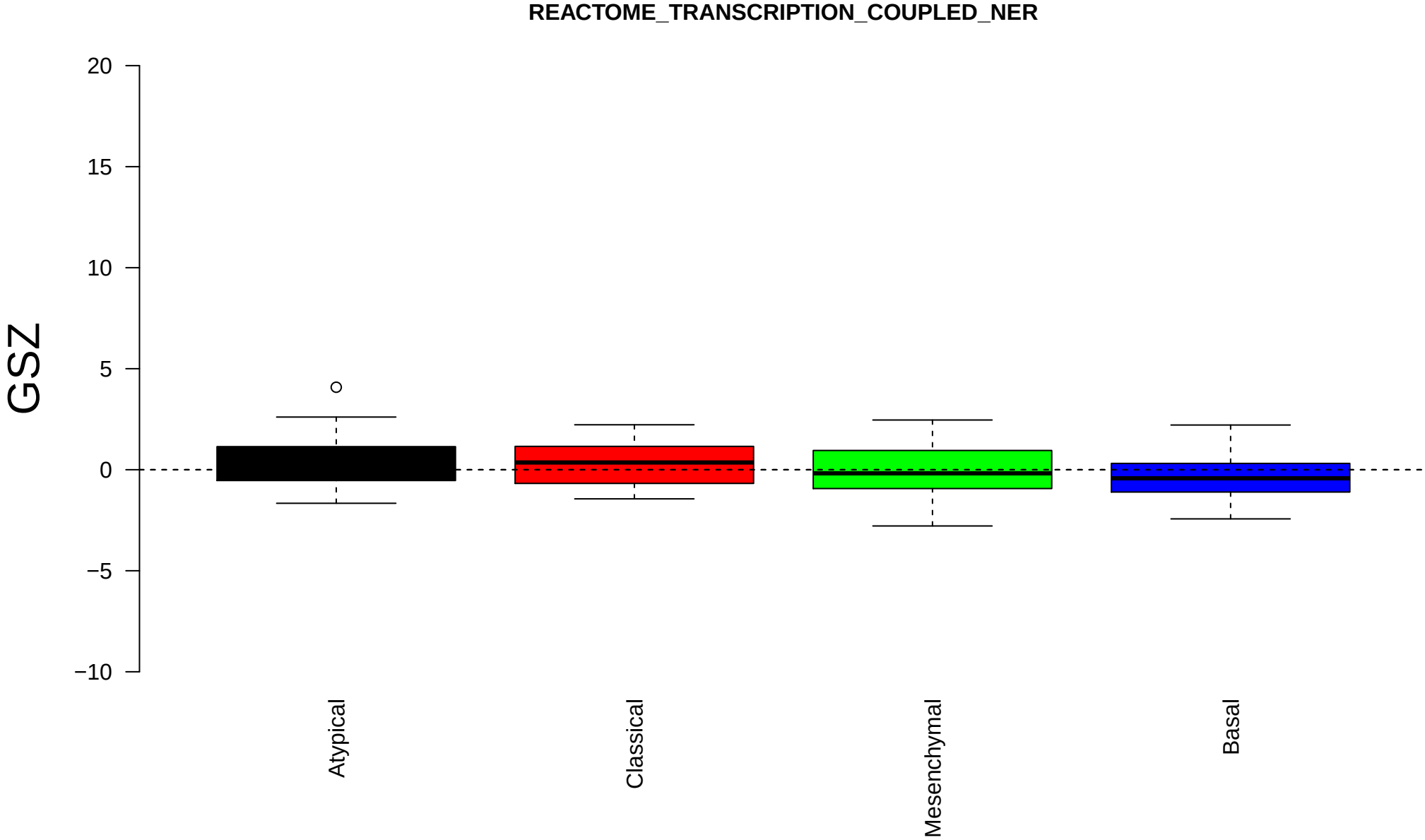
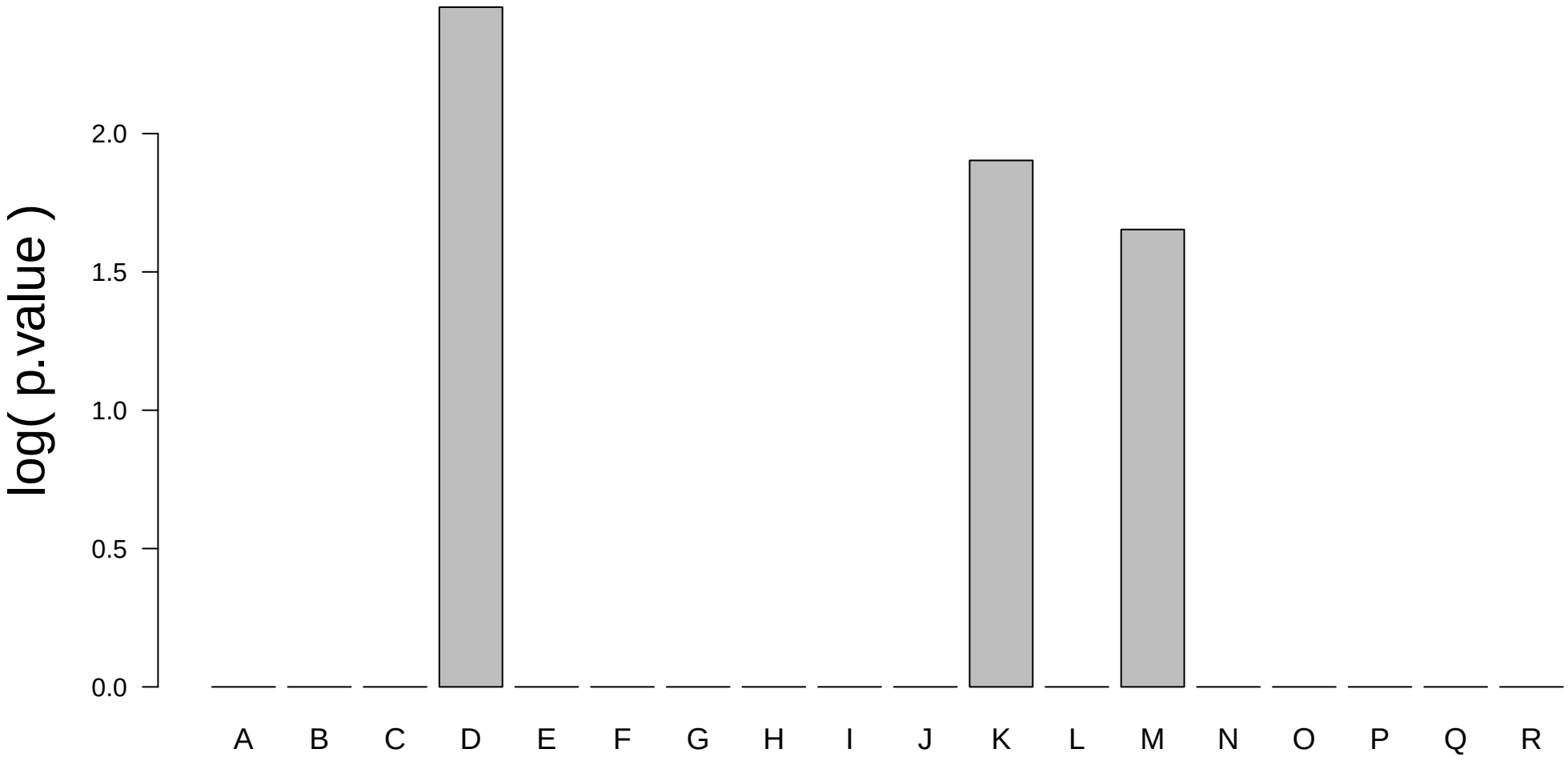
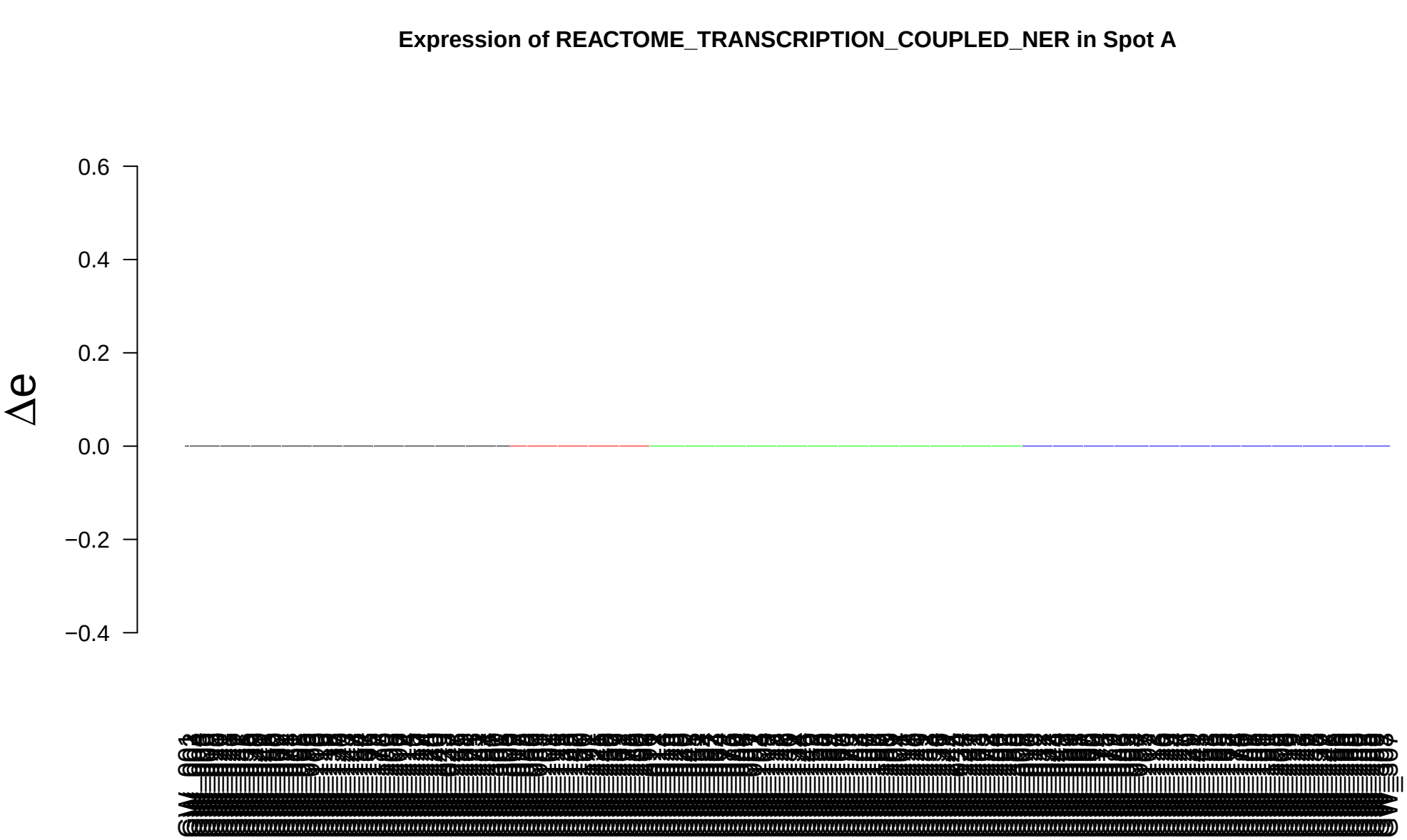
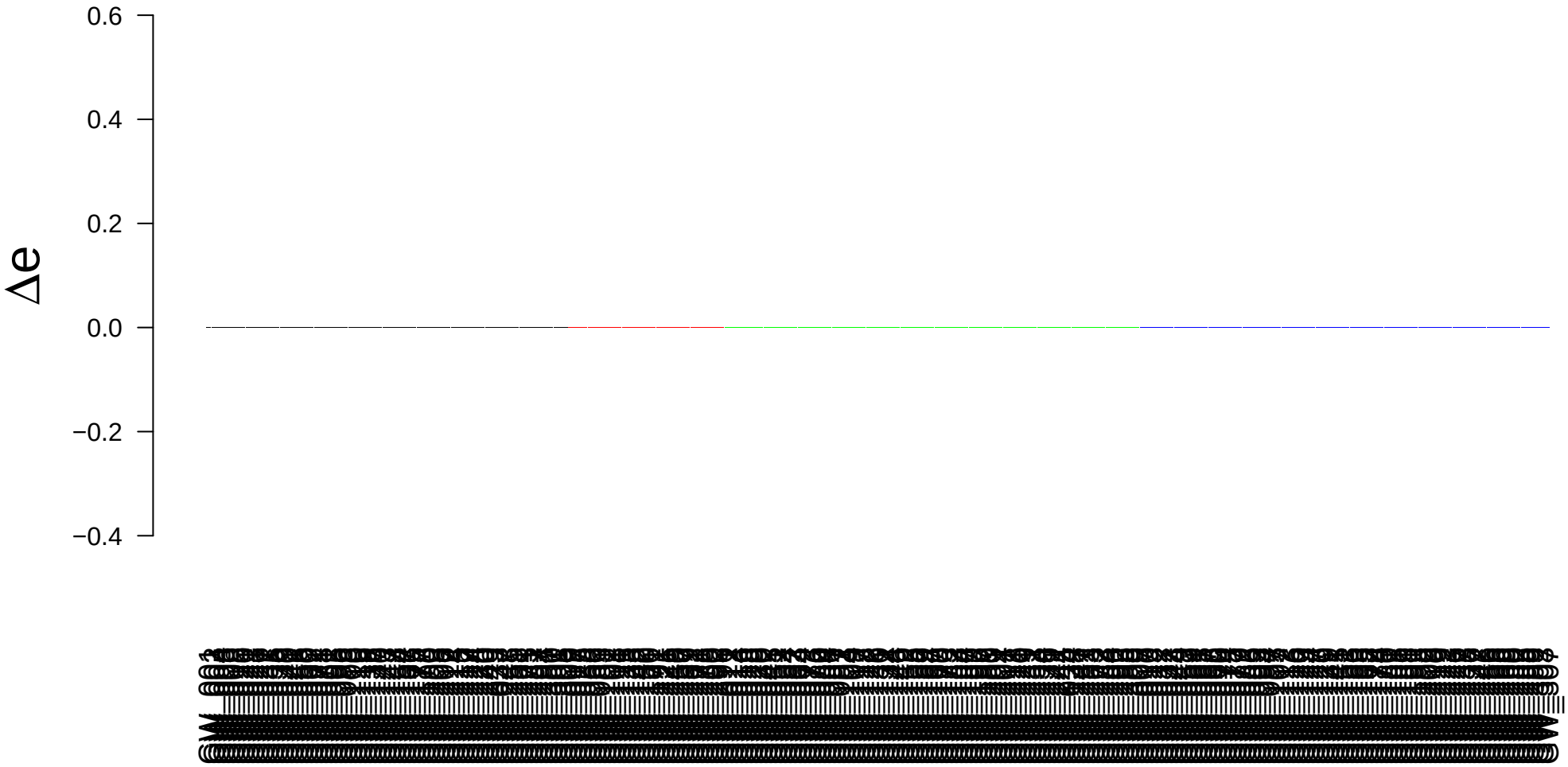
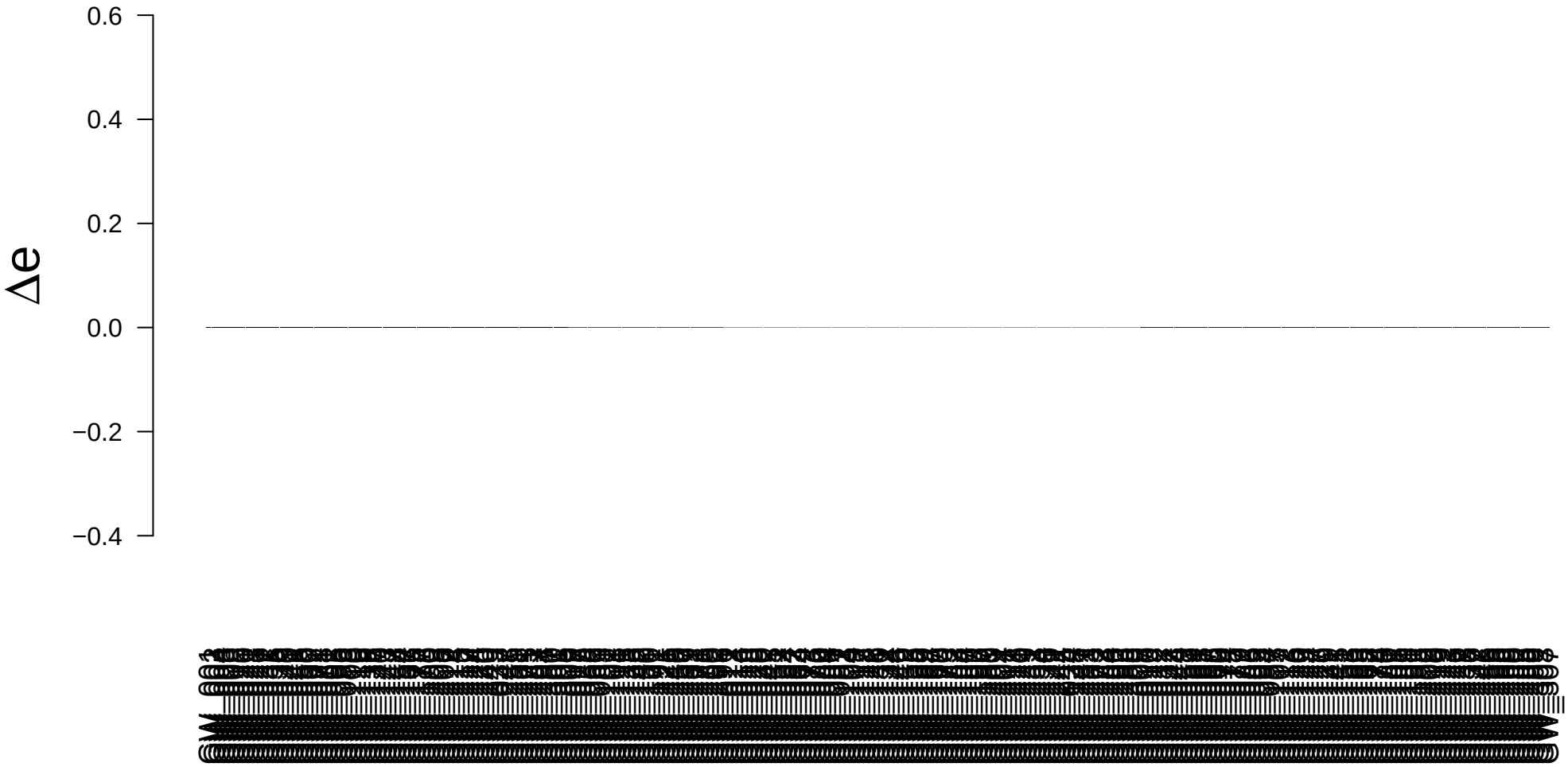

REACTOME_TRANSCRIPTION_COUPLED_NER
GSZ
20
15
10
5
0
−5
−10

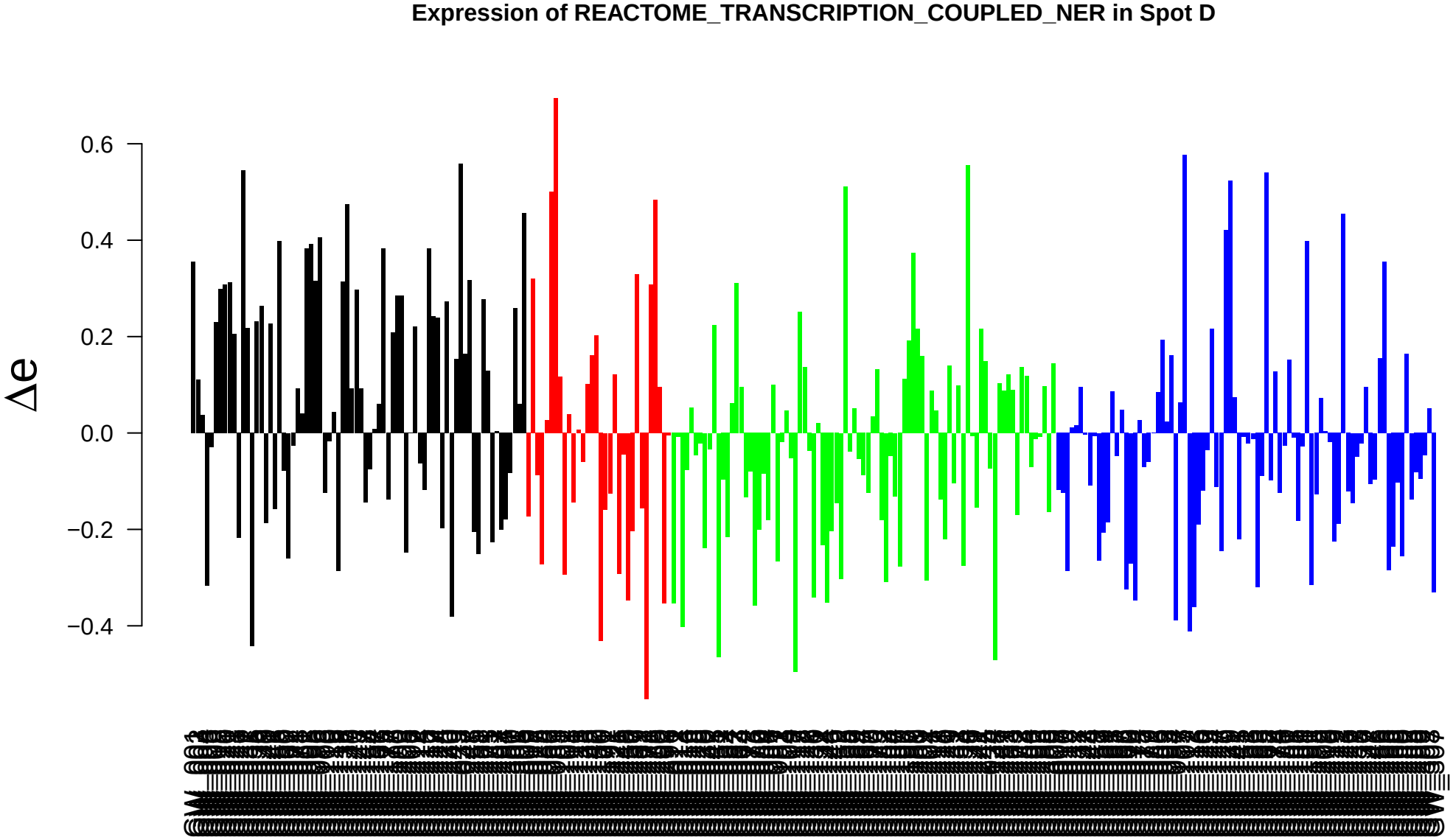

REACTOME_TRANSCRIPTION_COUPLED_NER
GSZ
20
15
10
5
0
−5
−10
Atypical
Classical
Mesenchymal
Basal

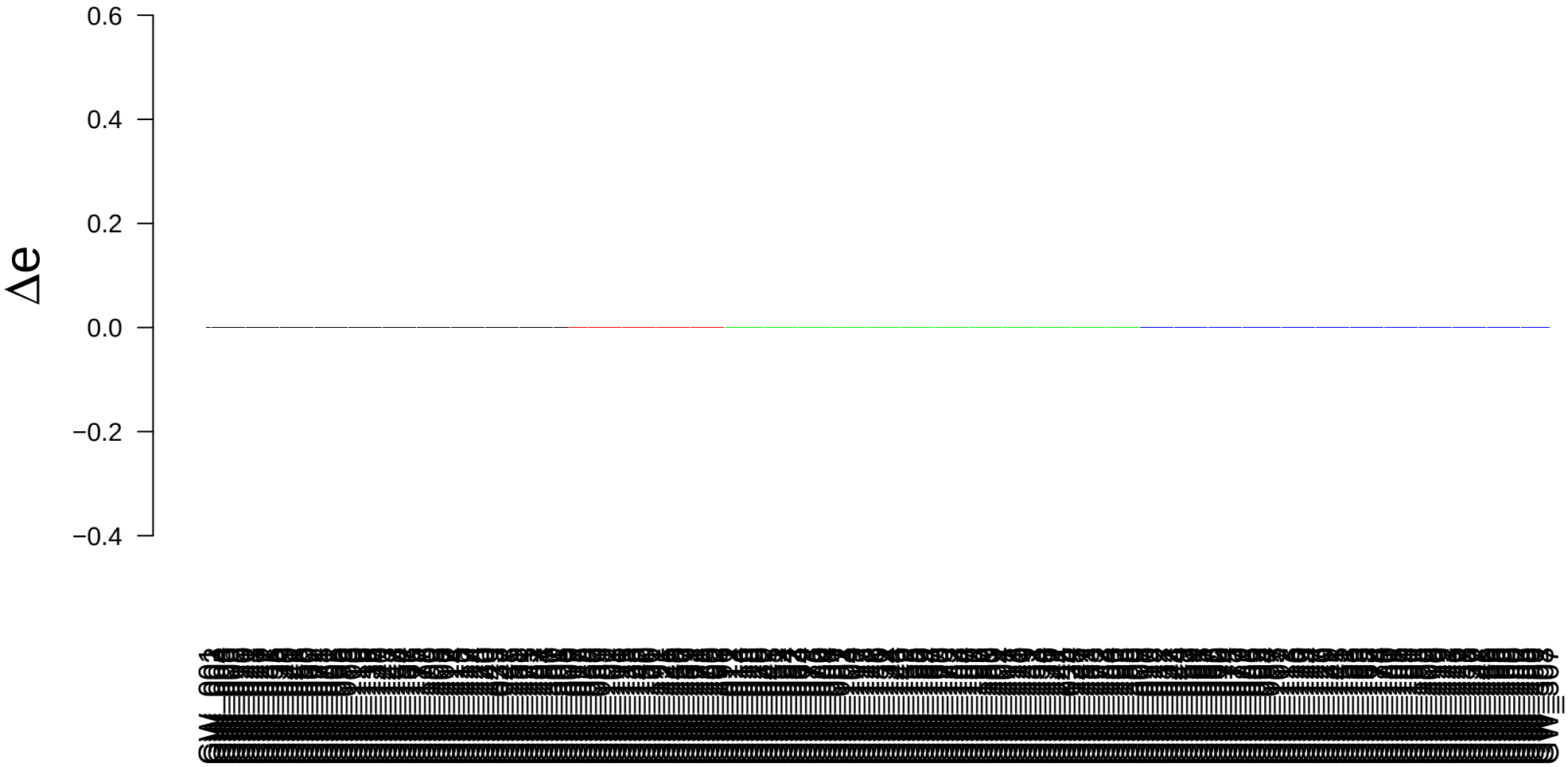

Enrichment in spots: REACTOME_TRANSCRIPTION_COUPLED_NER
log(p.value)
2.0
1.5
1.0
0.5
0.0
A
B
C
D
E
F
G
H
I
J
K
L
M
N
O
P
Q
R


Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot A
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

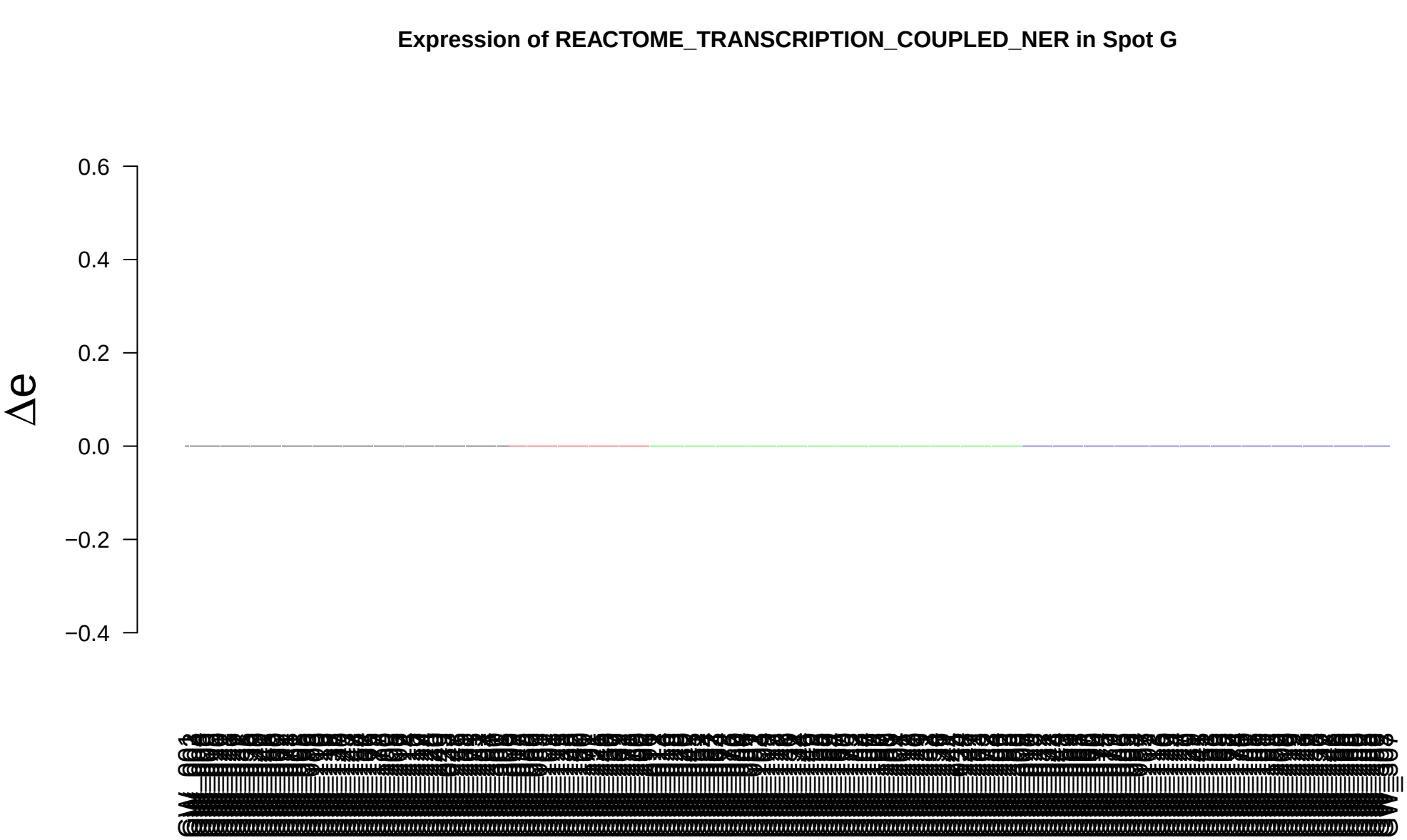

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot B
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

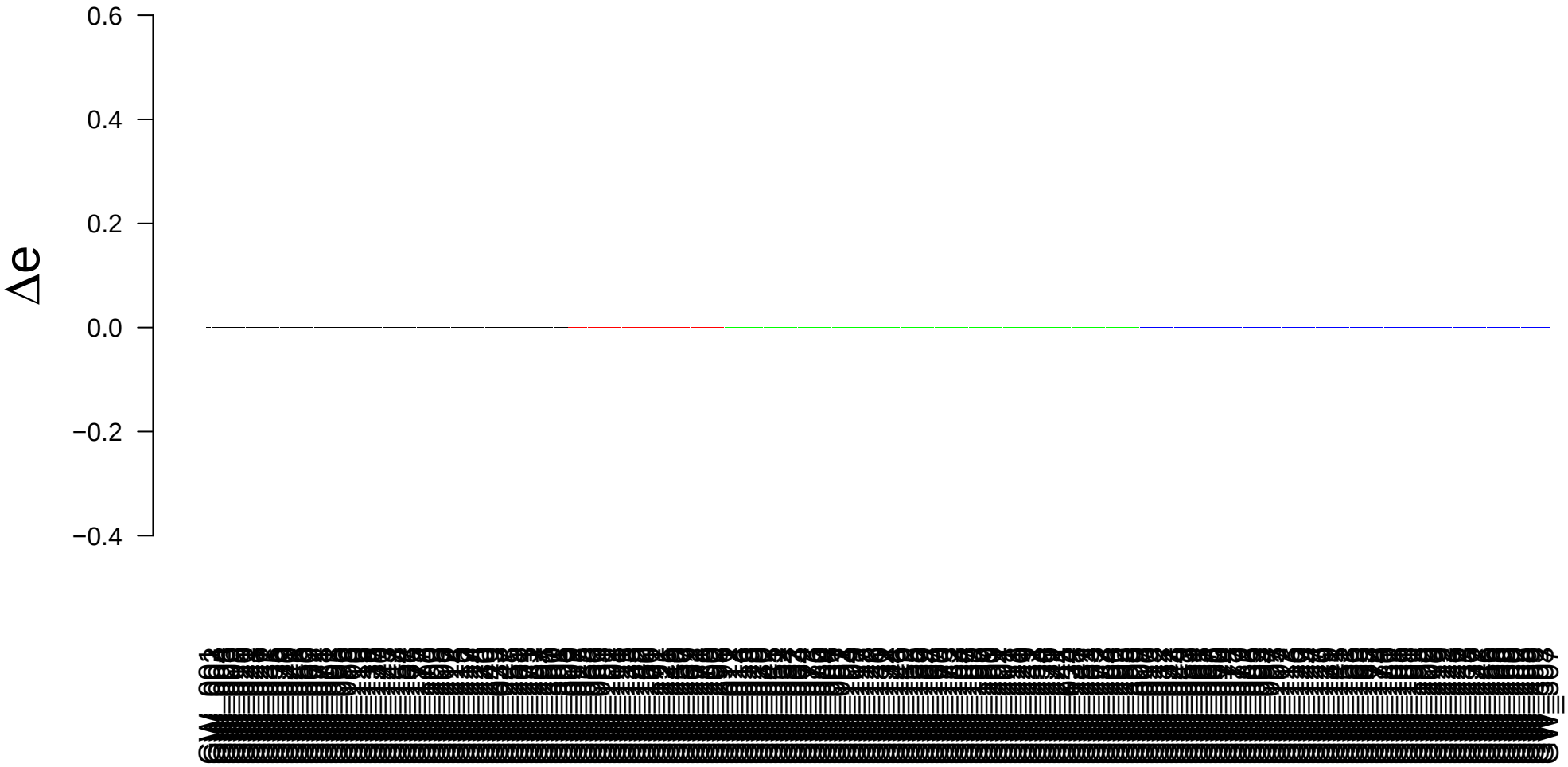

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot C
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

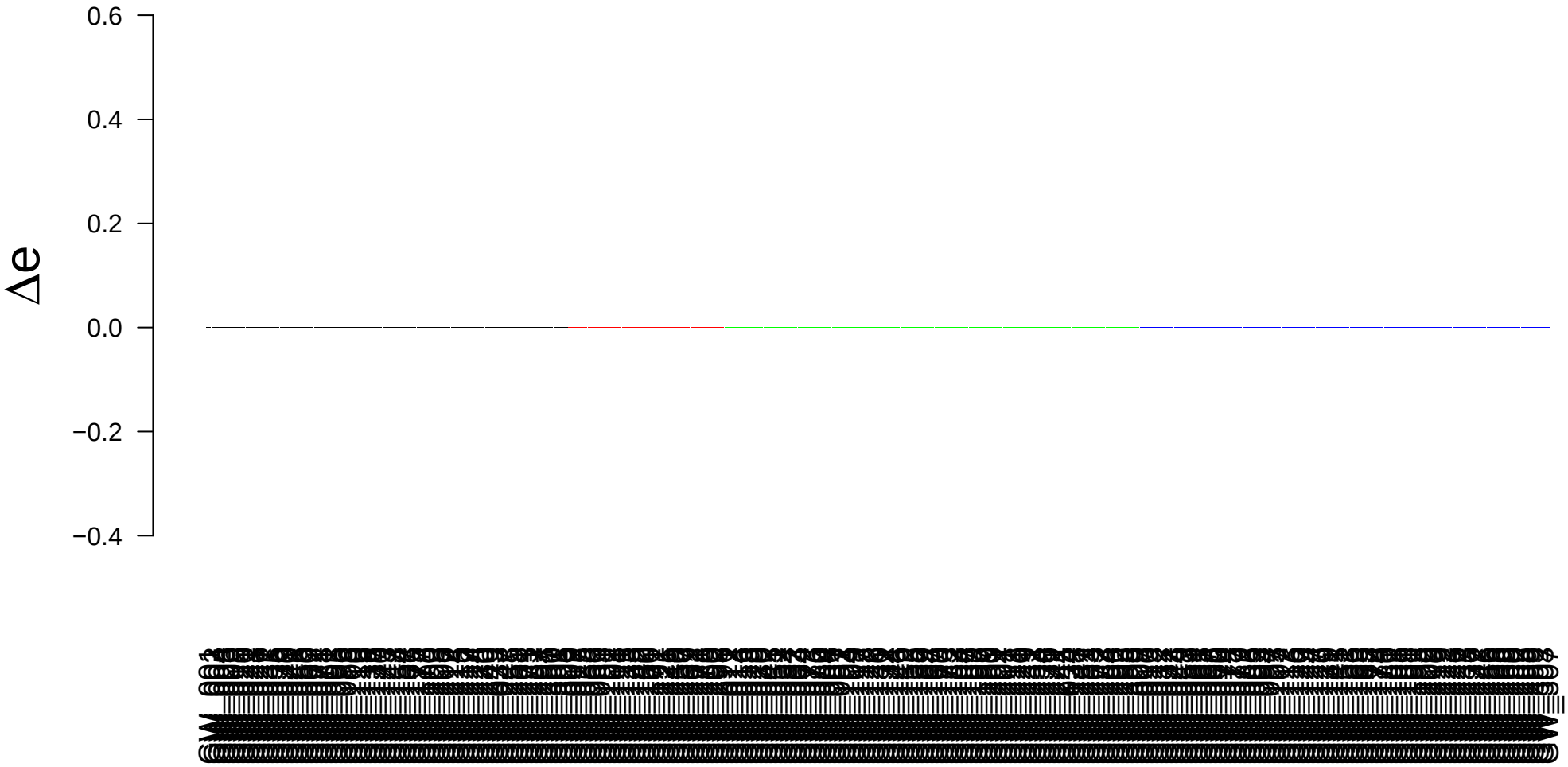

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot D
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

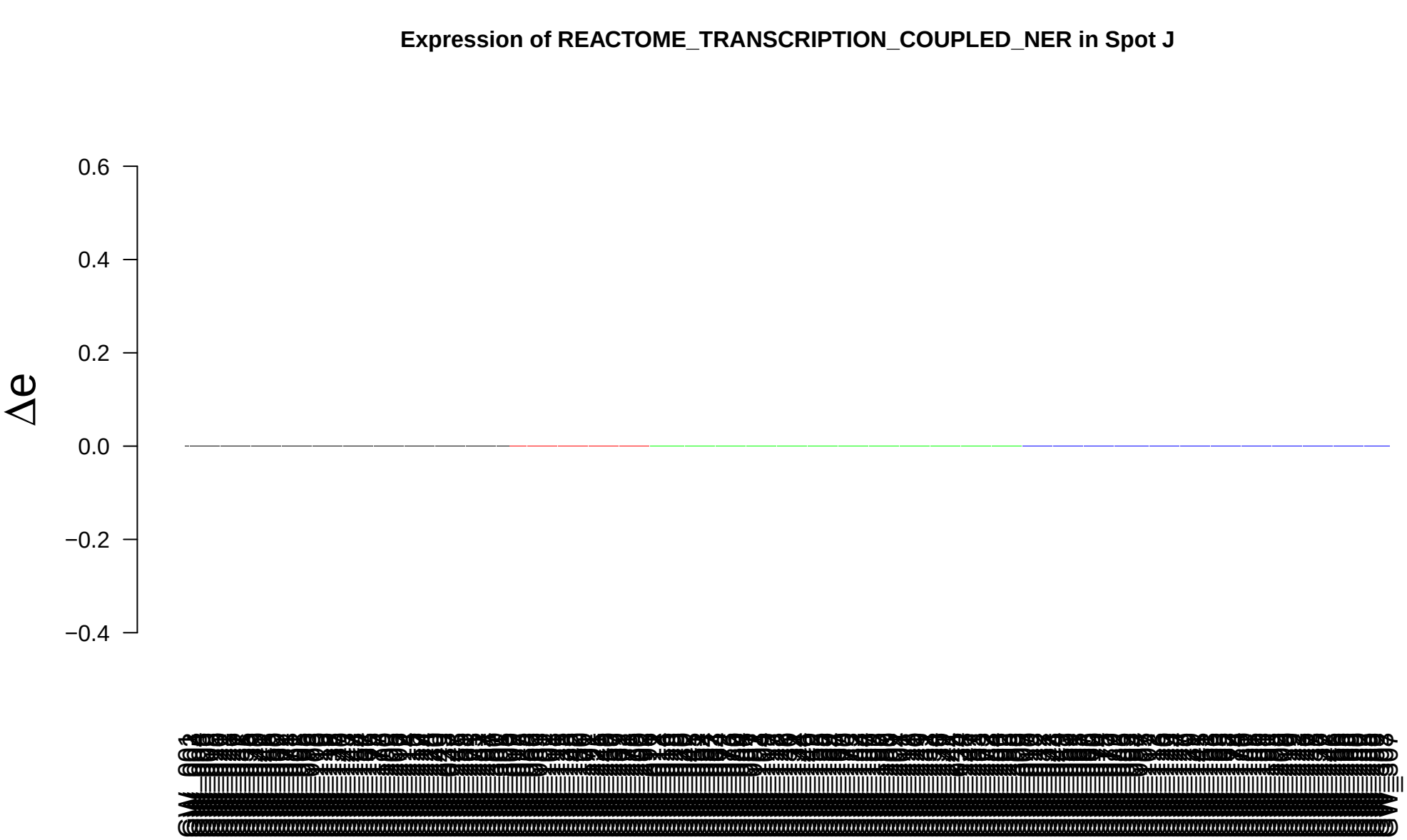

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot E
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

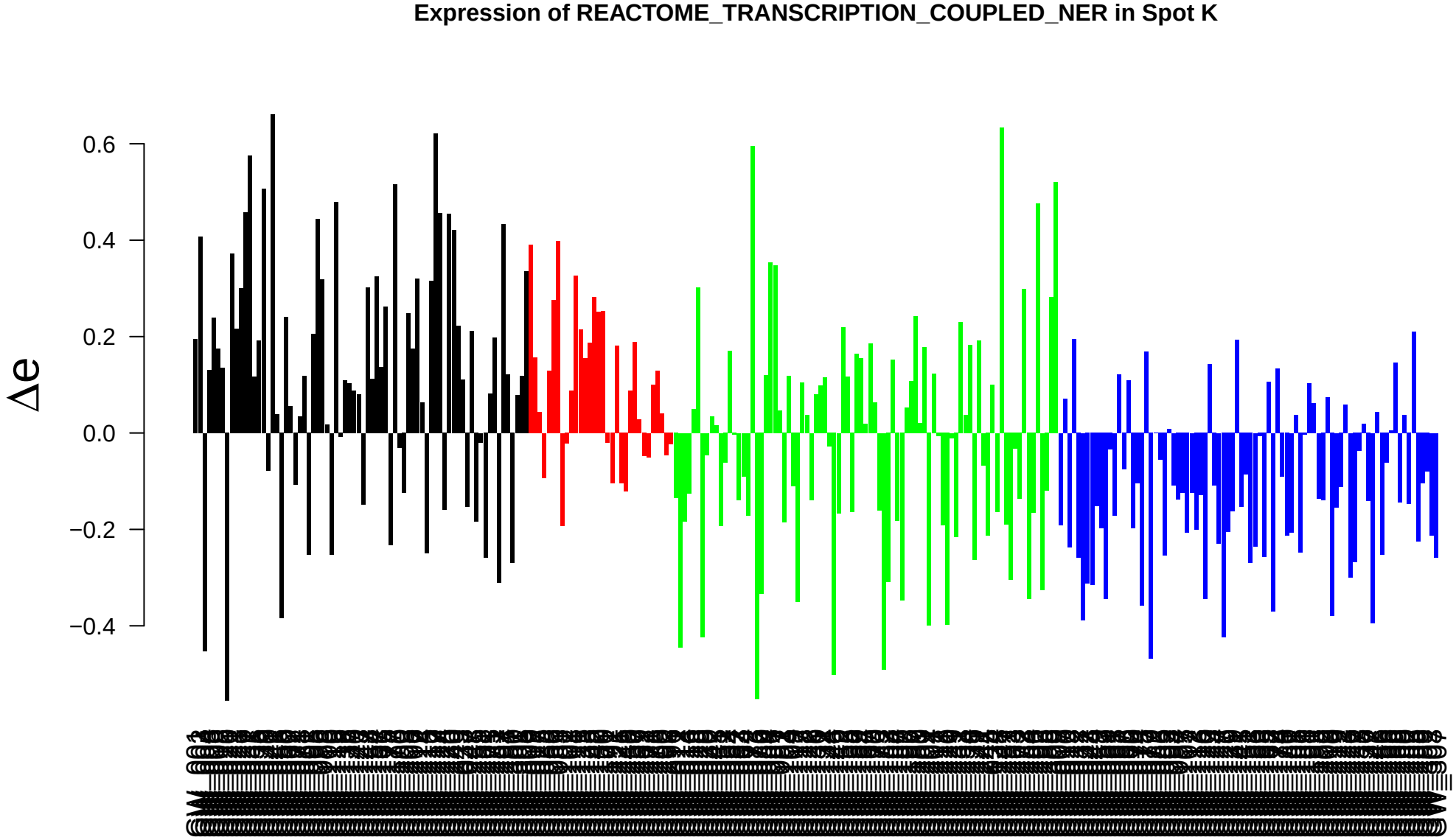
Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot F
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

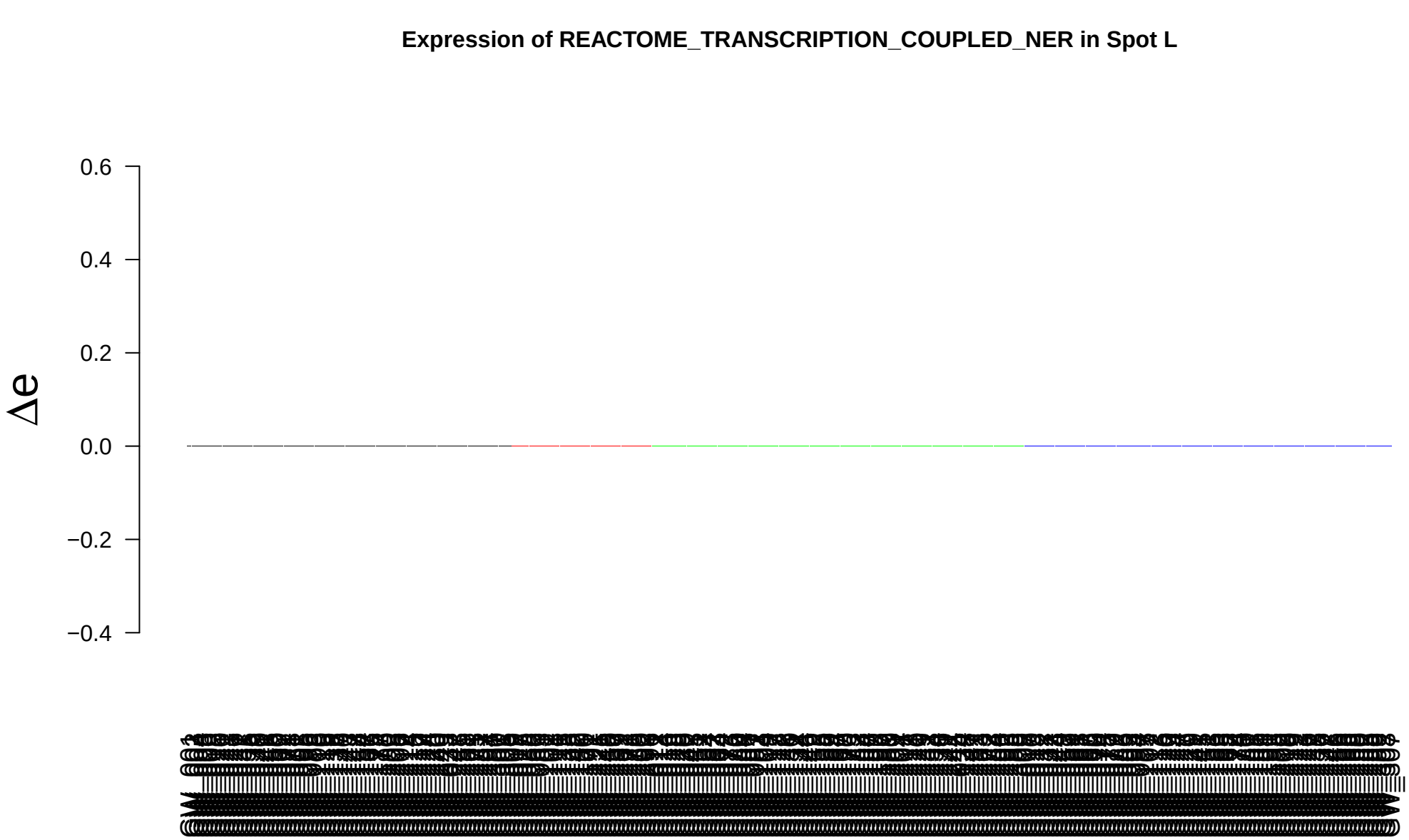

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot G
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

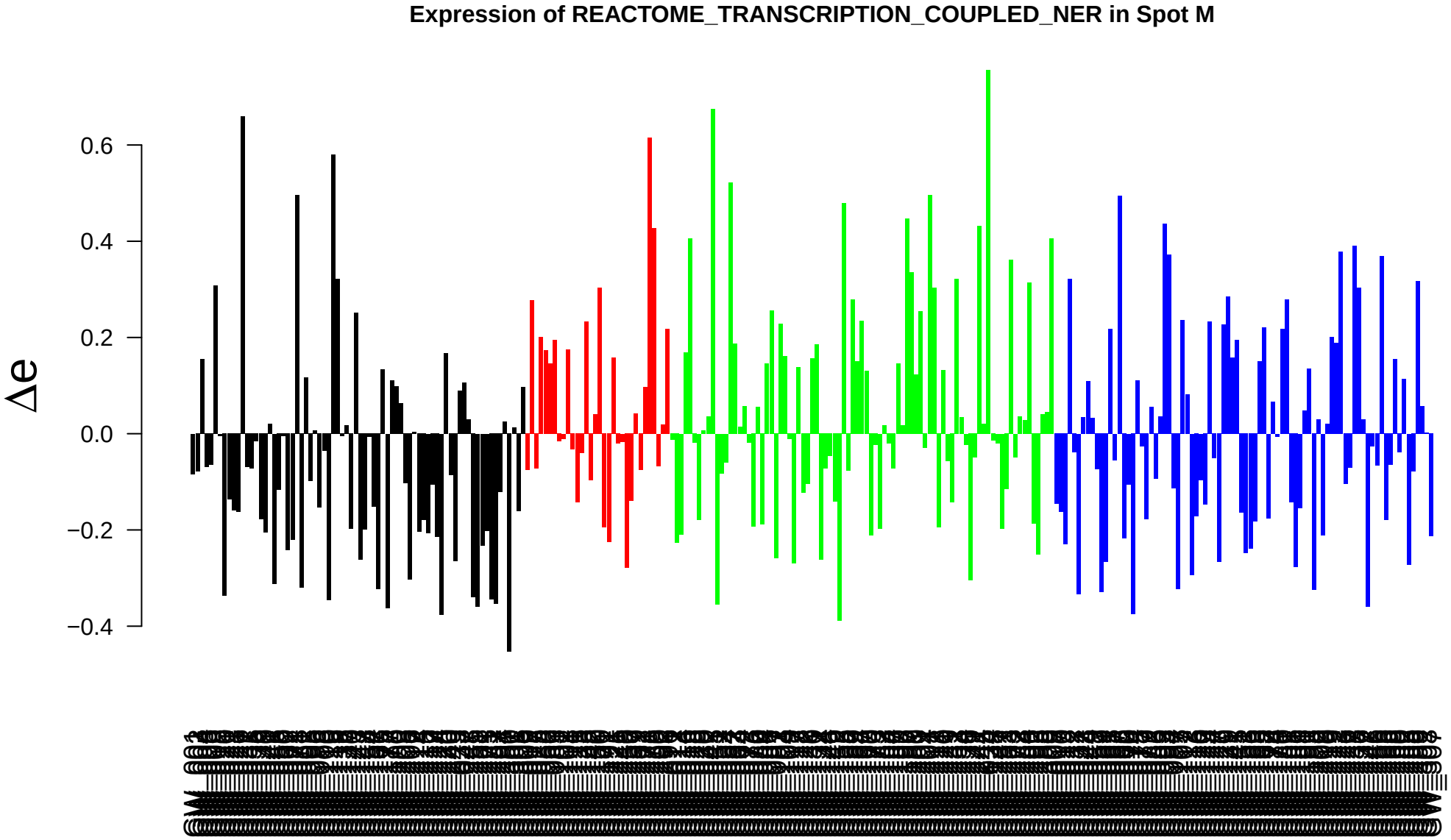

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot H
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

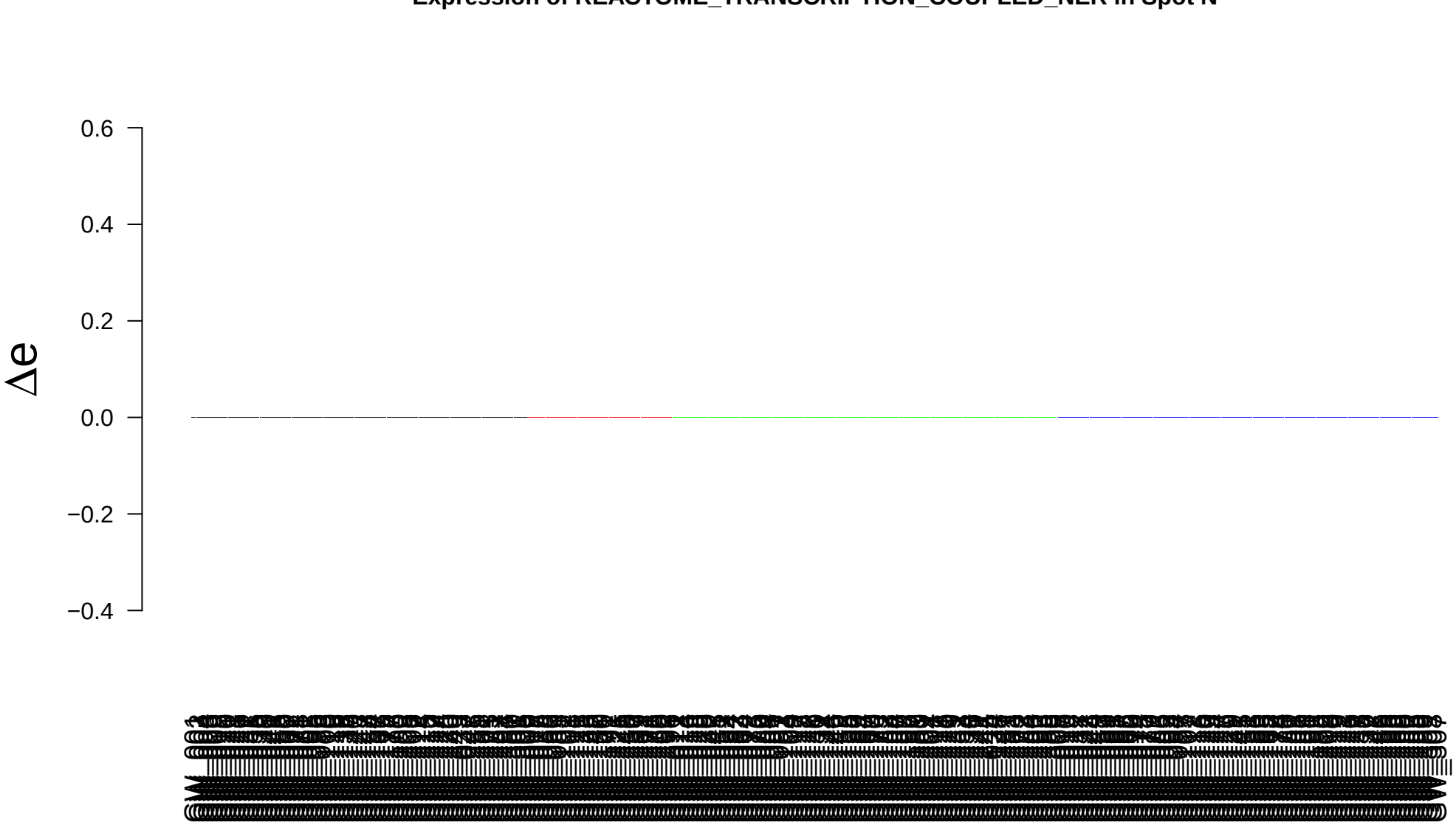

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot I
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

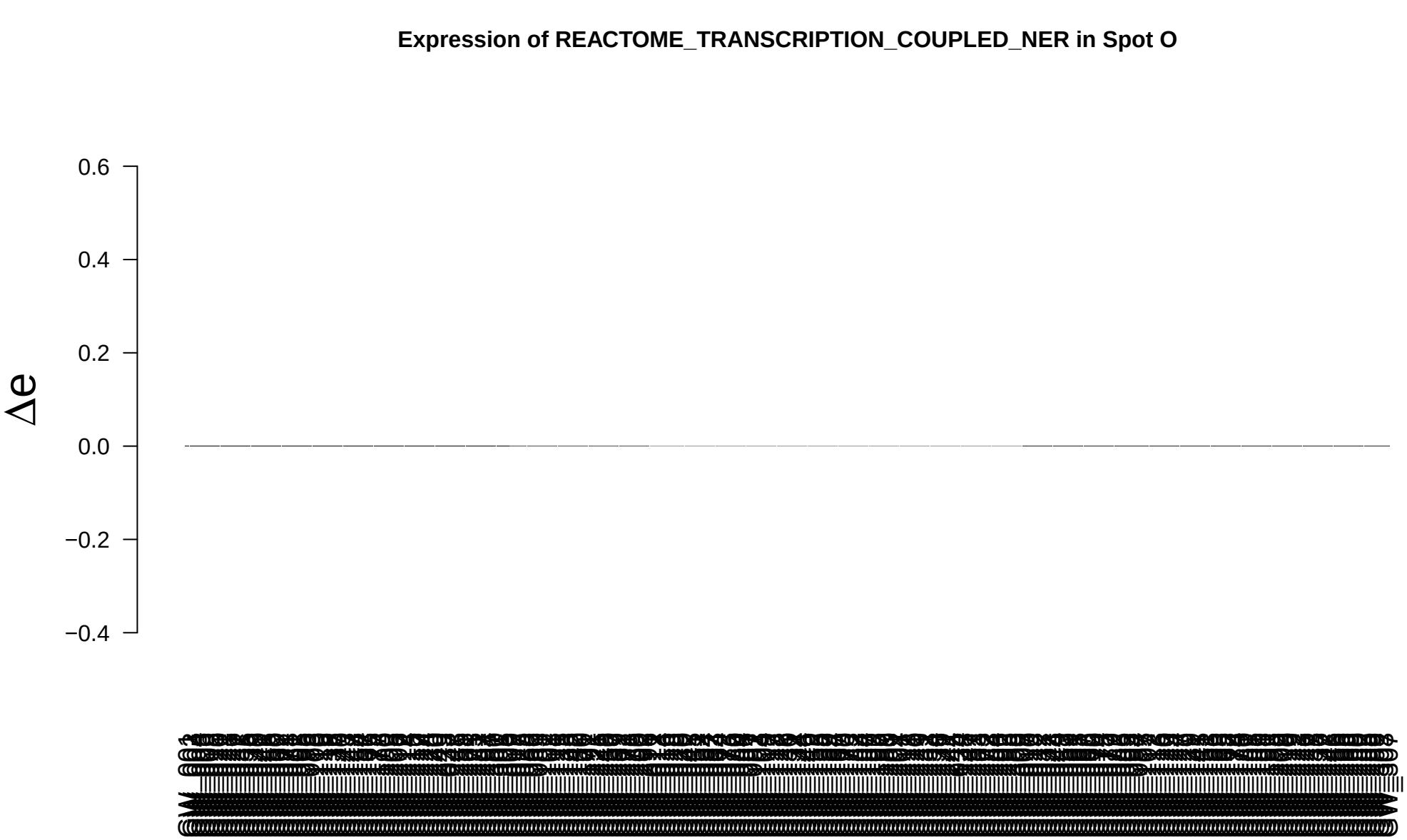

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot J
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4


Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot K
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

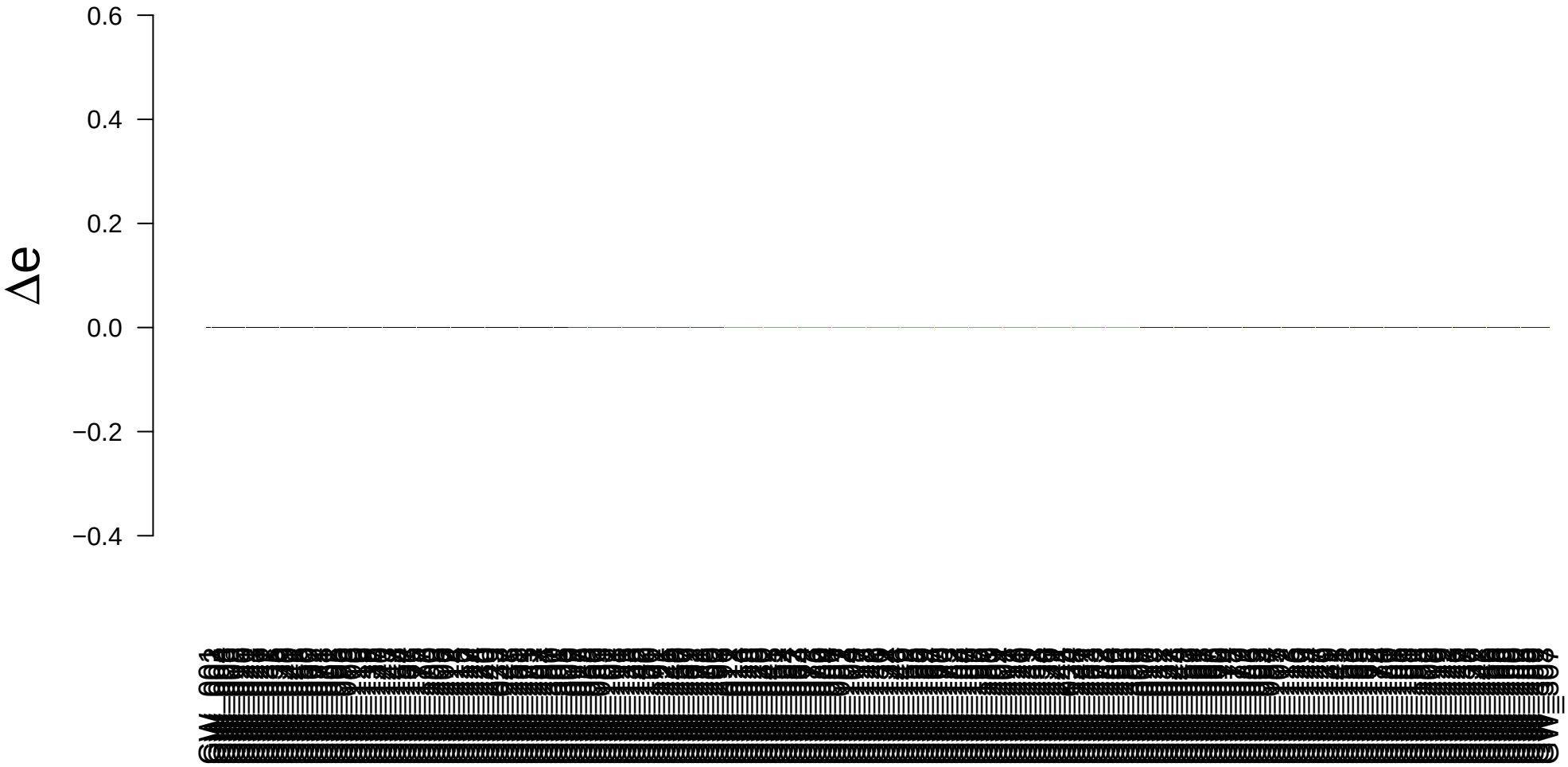

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot L
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

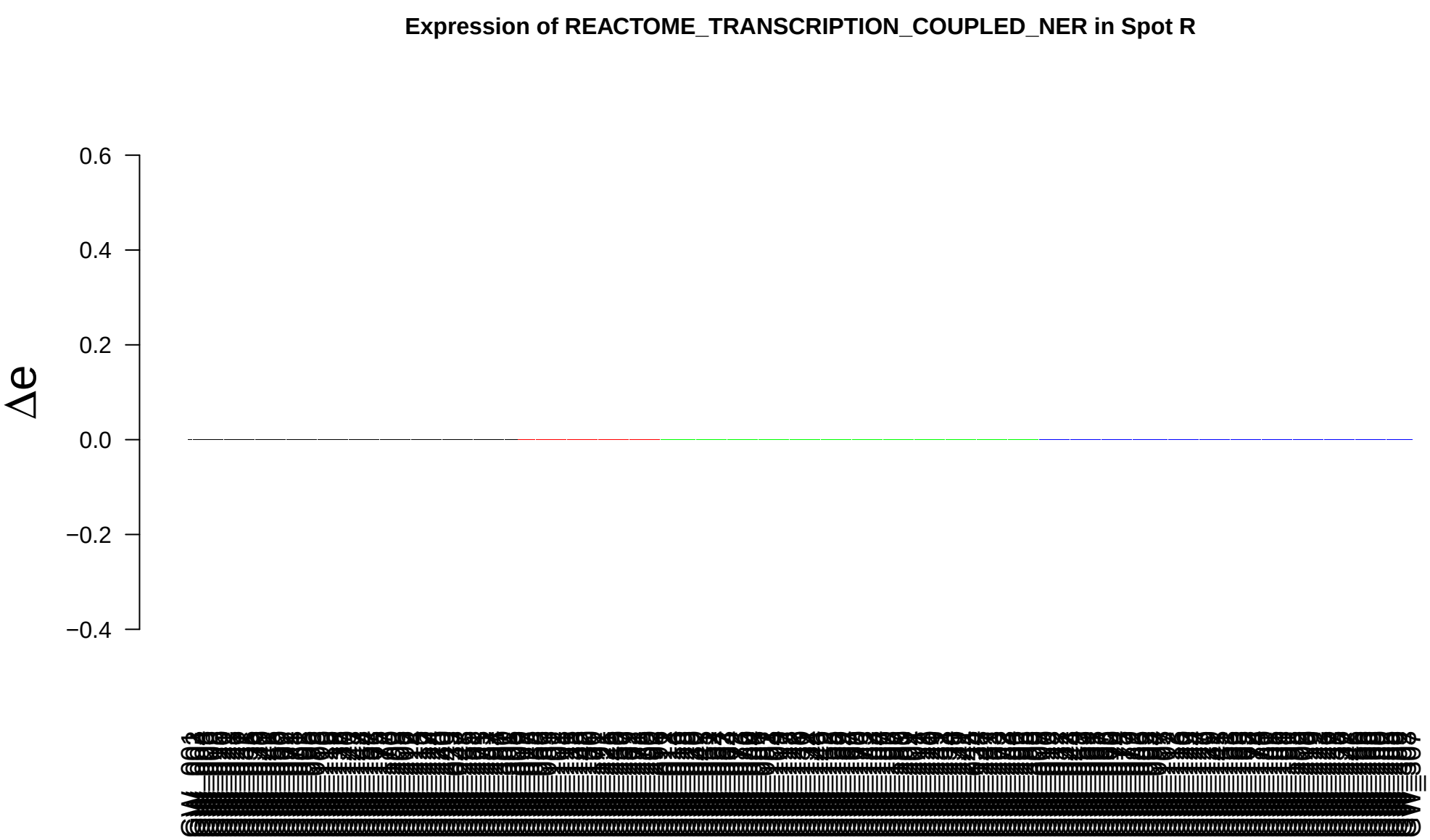

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot M
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4


Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot N
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4


Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot O
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4

Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot P
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4


Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot Q
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4


Expression of REACTOME_TRANSCRIPTION_COUPLED_NER in Spot R
Δe
0.6
0.4
0.2
0.0
−0.2
−0.4