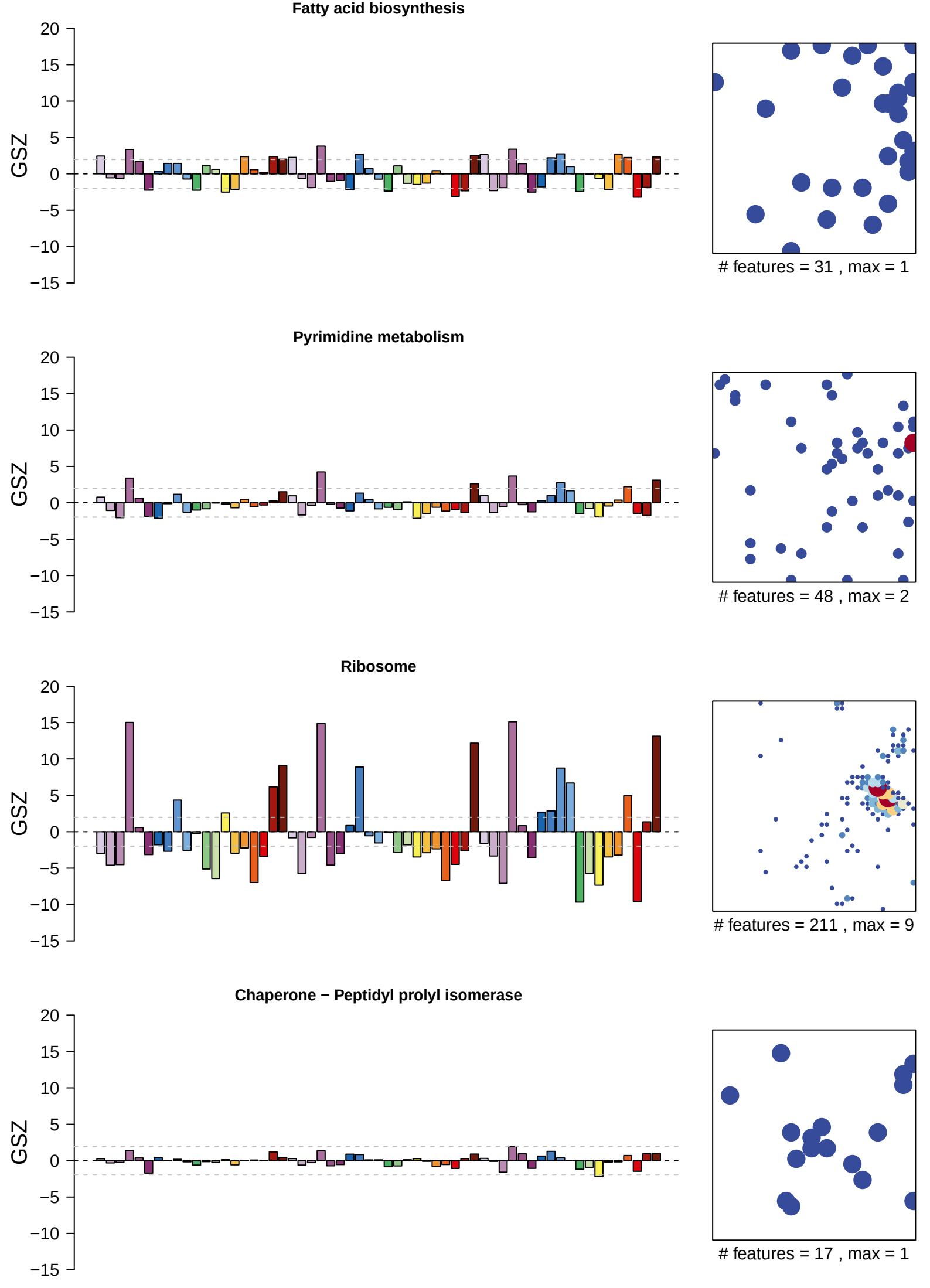
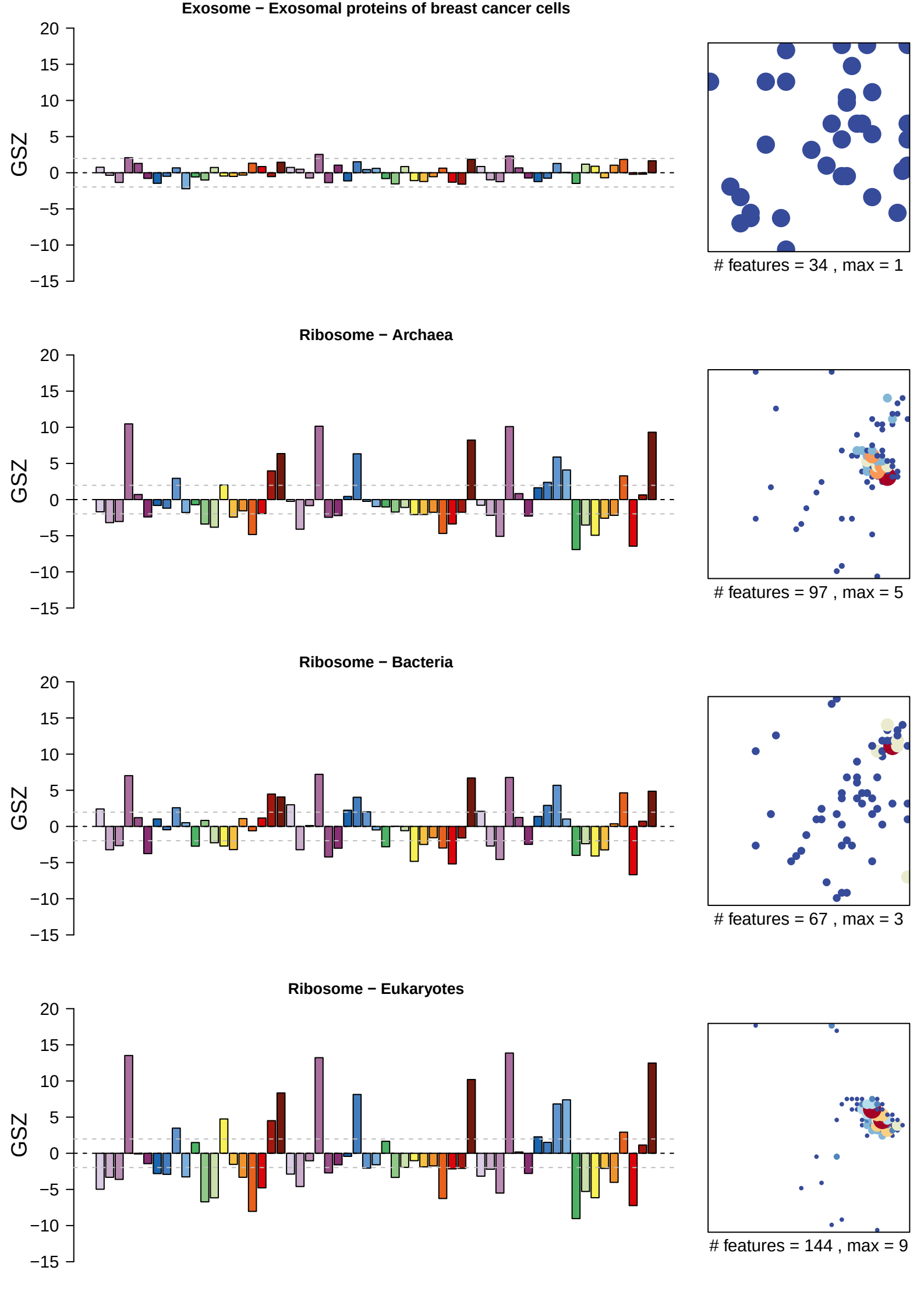
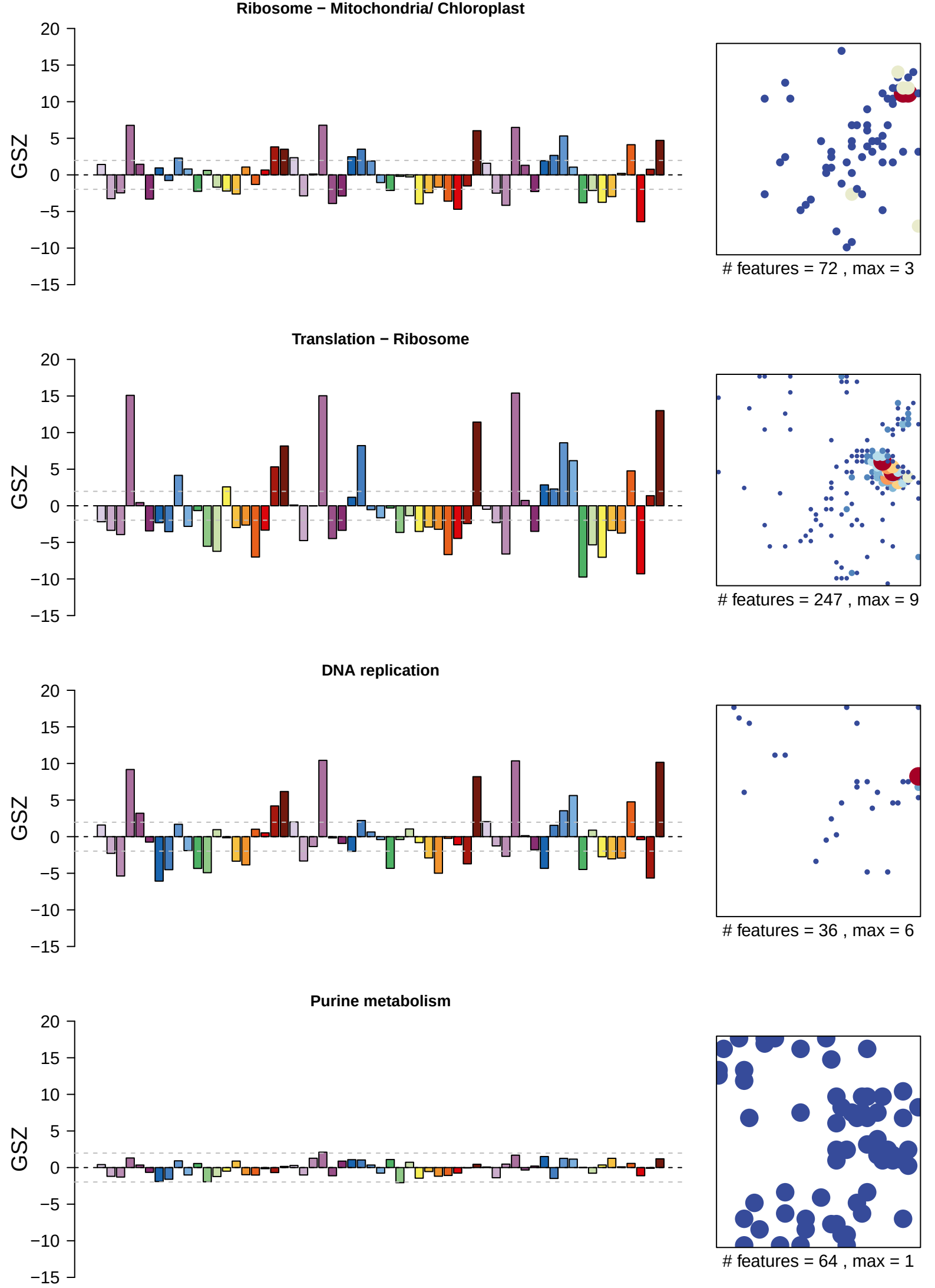
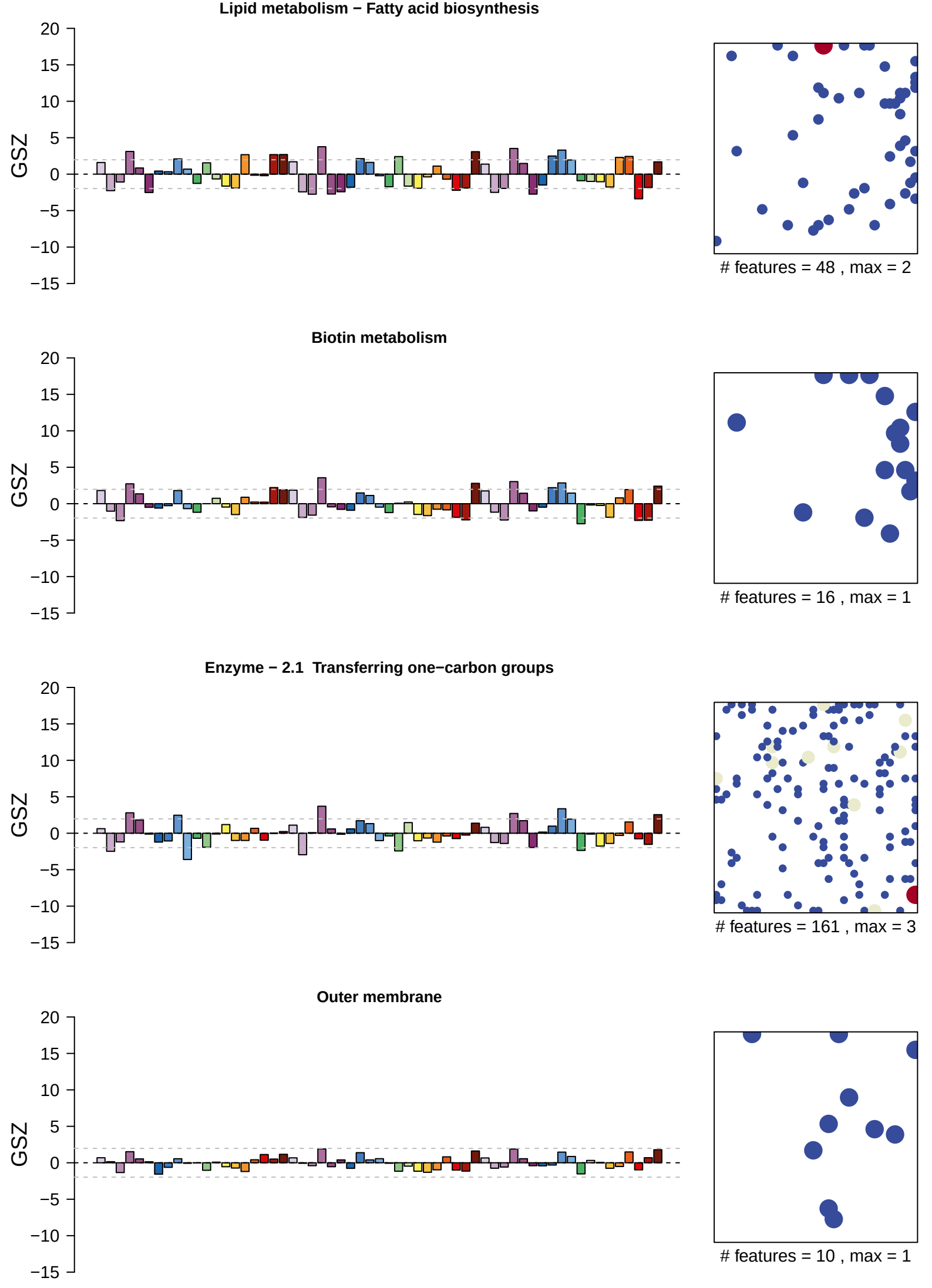
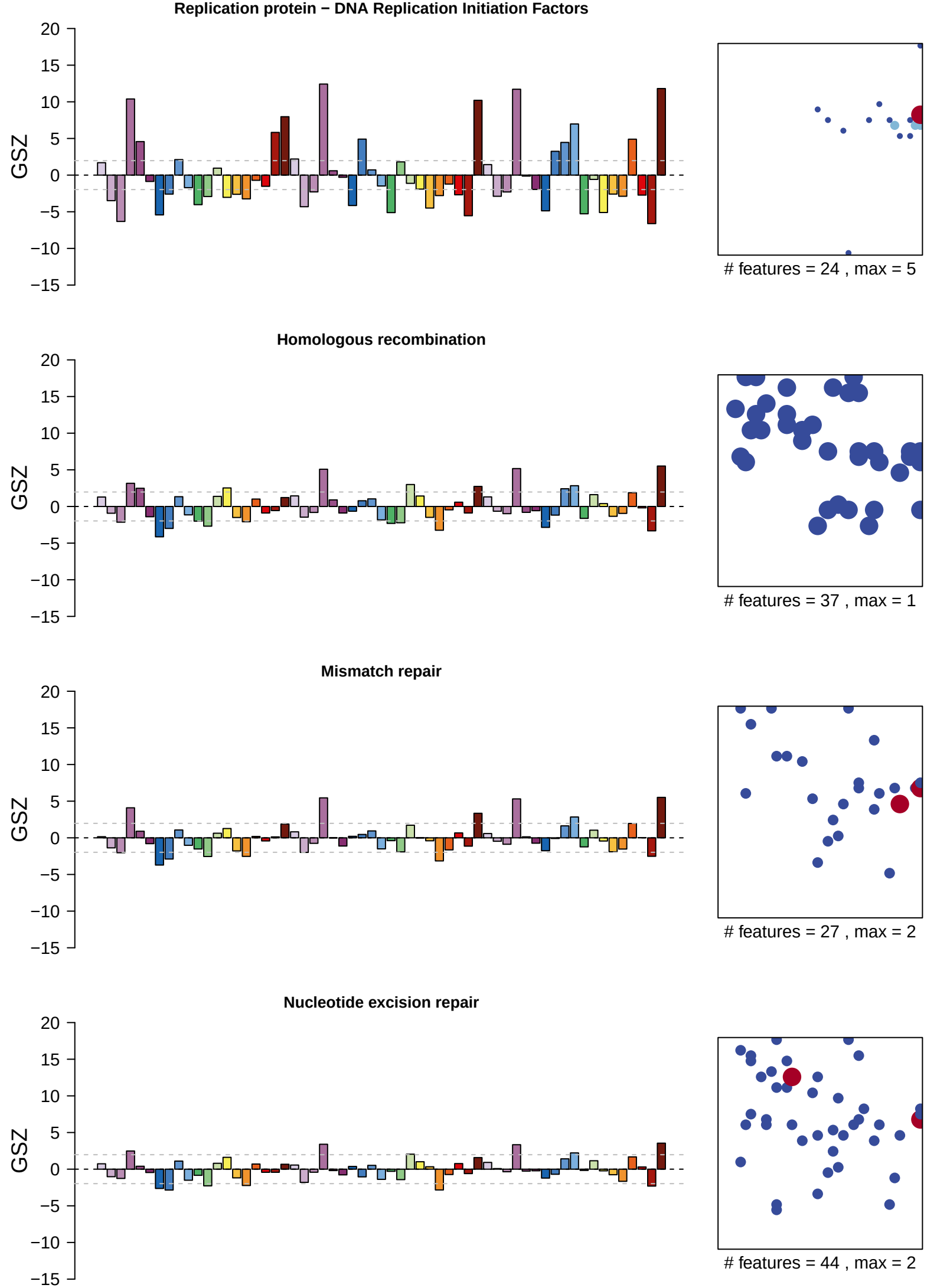

Fatty acid biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 31 , max = 1
Pyrimidine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 48 , max = 2
Ribosome
GSZ
20
15
10
5
0
−5
−10
−15
features = 211 , max = 9
Chaperone − Peptidyl prolyl isomerase
GSZ
20
15
10
5
0
−5
−10
−15
features = 17 , max = 1


Exosome − Exosomal proteins of breast cancer cells
GSZ
20
15
10
5
0
−5
−10
−15
features = 34 , max = 1
Ribosome − Archaea
GSZ
20
15
10
5
0
−5
−10
−15
features = 97 , max = 5
Ribosome − Bacteria
GSZ
20
15
10
5
0
−5
−10
−15
features = 67 , max = 3
Ribosome − Eukaryotes
GSZ
20
15
10
5
0
−5
−10
−15
features = 144 , max = 9


Ribosome − Mitochondria/ Chloroplast
GSZ
20
15
10
5
0
−5
−10
−15
features = 72 , max = 3
Translation − Ribosome
GSZ
20
15
10
5
0
−5
−10
−15
features = 247 , max = 9
DNA replication
GSZ
20
15
10
5
0
−5
−10
−15
features = 36 , max = 6
Purine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 64 , max = 1


Lipid metabolism − Fatty acid biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 48 , max = 2
Biotin metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 16 , max = 1
Enzyme − 2.1 Transferring one−carbon groups
GSZ
20
15
10
5
0
−5
−10
−15
features = 161 , max = 3
Outer membrane
GSZ
20
15
10
5
0
−5
−10
−15
features = 10 , max = 1


Replication protein − DNA Replication Initiation Factors
GSZ
20
15
10
5
0
−5
−10
−15
features = 24 , max = 5
Homologous recombination
GSZ
20
15
10
5
0
−5
−10
−15
features = 37 , max = 1
Mismatch repair
GSZ
20
15
10
5
0
−5
−10
−15
features = 27 , max = 2
Nucleotide excision repair
GSZ
20
15
10
5
0
−5
−10
−15
features = 44 , max = 2