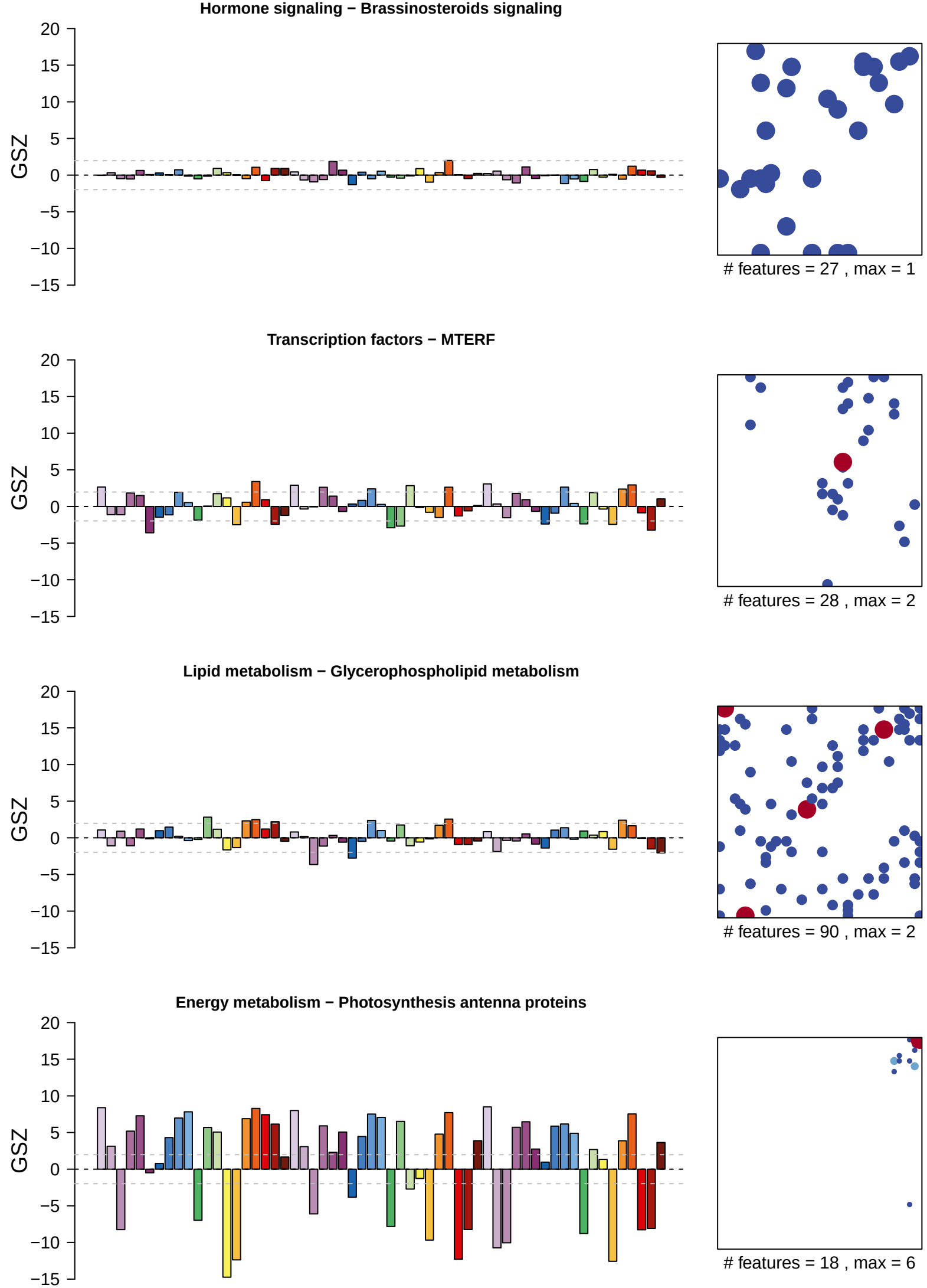
Hormone signaling − Brassinosteroids signaling
GSZ
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15
10
5
0
−5
−10
−15
# features = 27 , max = 1
Transcription factors − MTERF
GSZ
20
15
10
5
0
−5
−10
−15
# features = 28 , max = 2
Lipid metabolism − Glycerophospholipid metabolism
GSZ
20
15
10
5
0
−5
−10
−15
# features = 90 , max = 2
Energy metabolism − Photosynthesis antenna proteins
GSZ
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5
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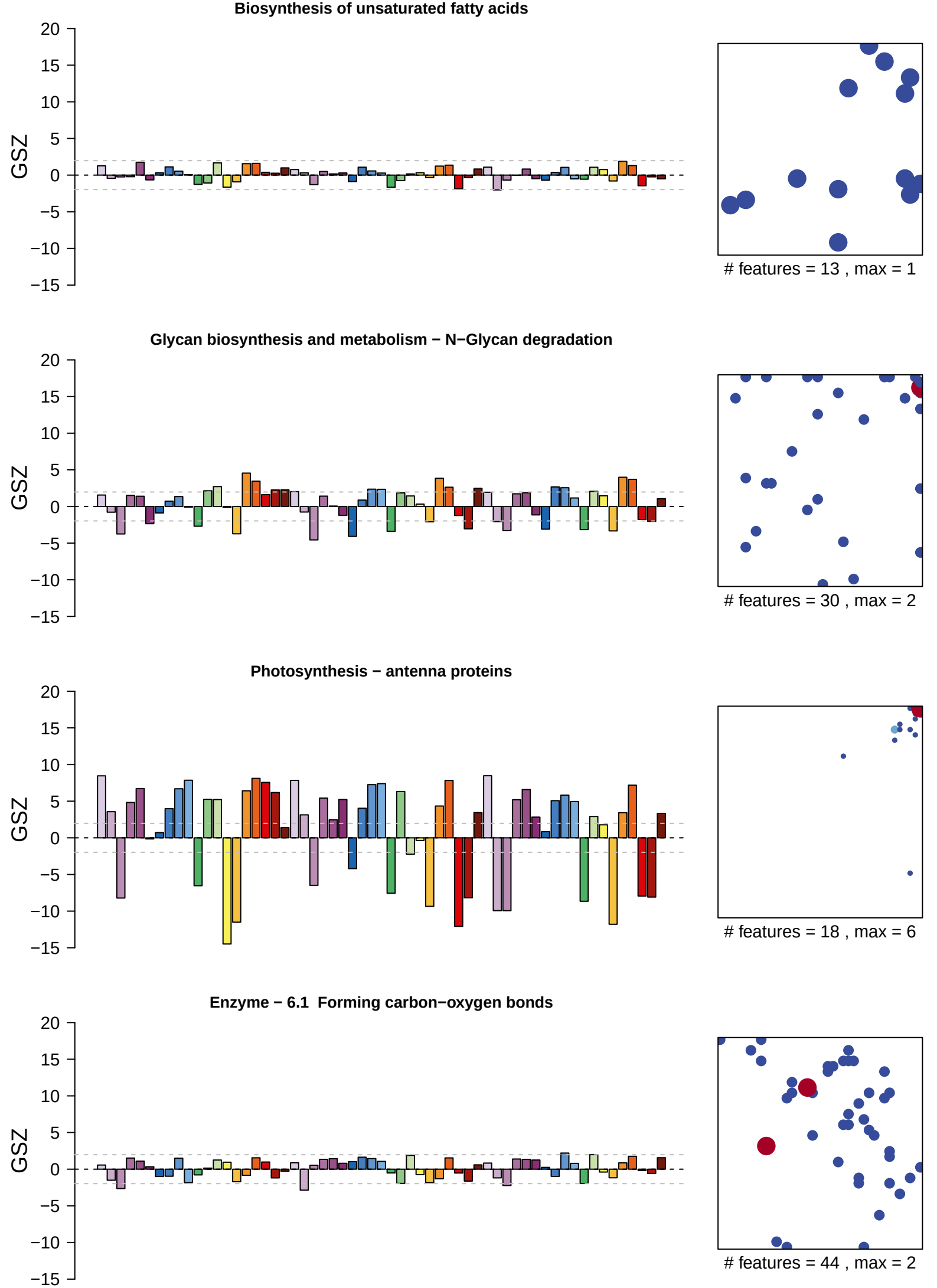
−10
−15
# features = 18 , max = 6


Biosynthesis of unsaturated fatty acids
GSZ
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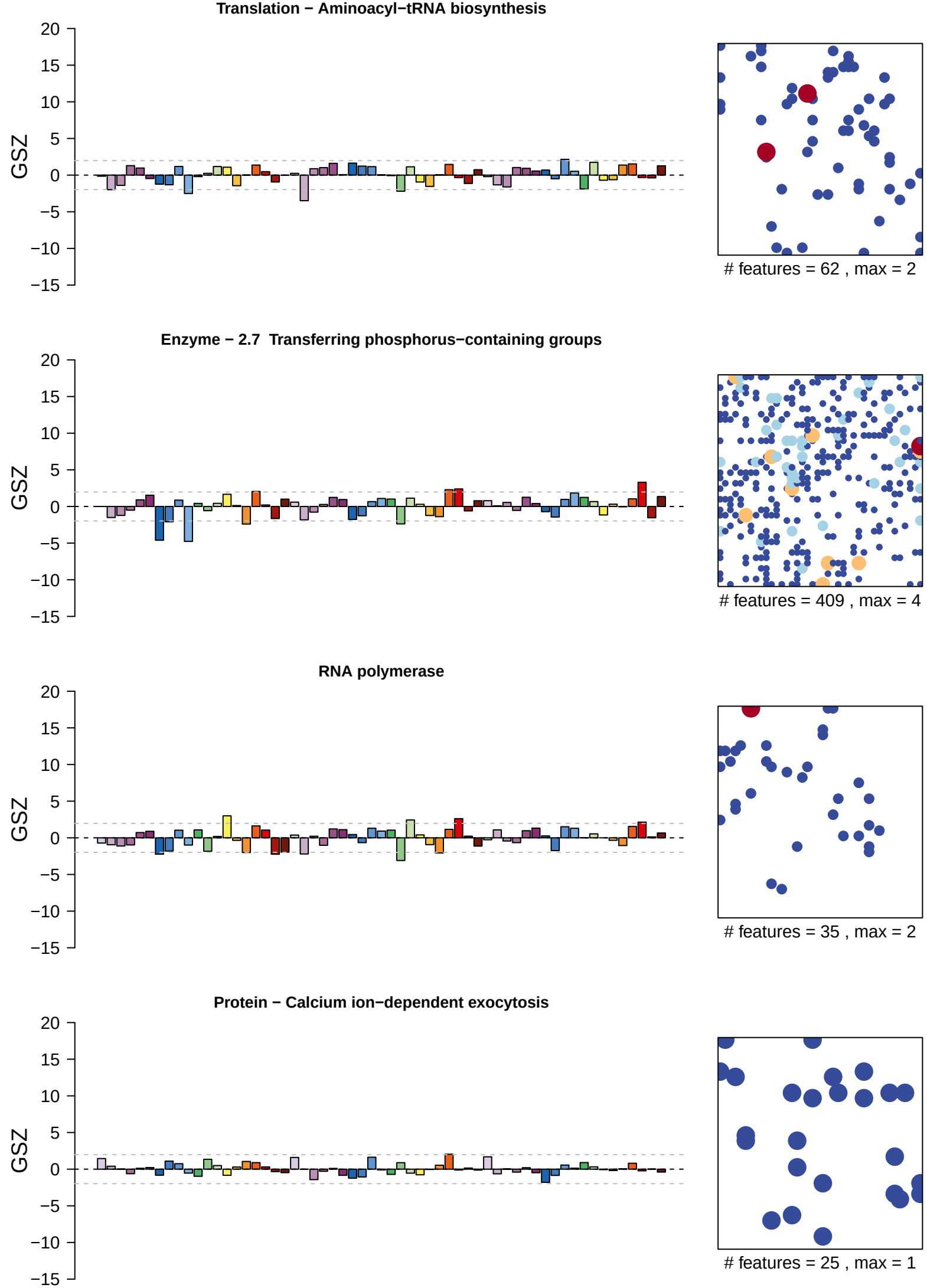
−5
−10
−15
# features = 13 , max = 1
Glycan biosynthesis and metabolism − N−Glycan degradation
GSZ
# features = 30 , max = 2
Photosynthesis − antenna proteins
GSZ
# features = 18 , max = 6
Enzyme − 6.1 Forming carbon−oxygen bonds
GSZ
# features = 44 , max = 2


Translation − Aminoacyl−tRNA biosynthesis
GSZ
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−5
−10
−15
# features = 62 , max = 2
Enzyme − 2.7 Transferring phosphorus−containing groups
GSZ
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15
10
5
0
−5
−10
−15
# features = 409 , max = 4
RNA polymerase
GSZ
20
15
10
5
0
−5
−10
−15
# features = 35 , max = 2
Protein − Calcium ion−dependent exocytosis
GSZ
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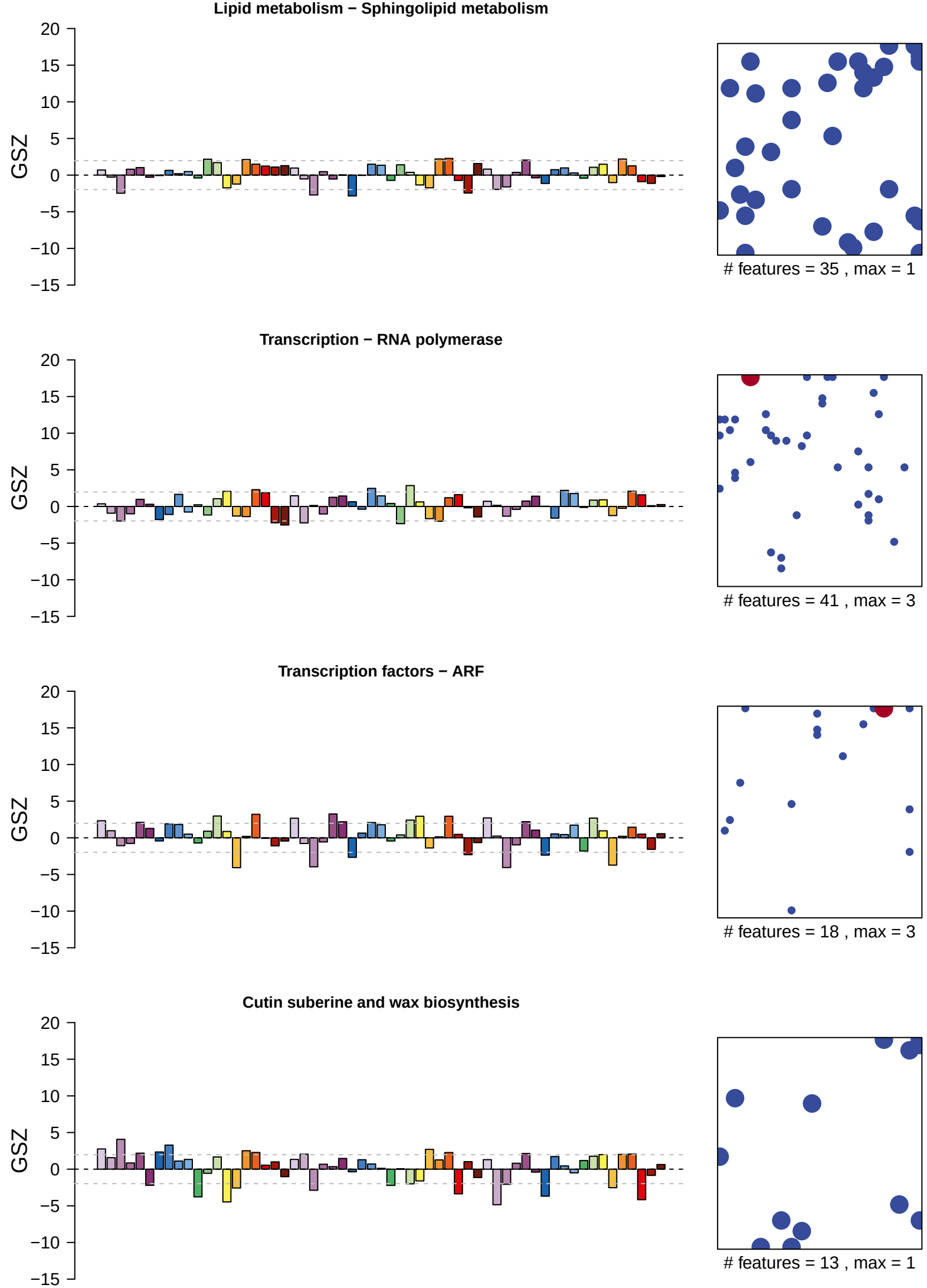
−10
−15
# features = 25 , max = 1


Lipid metabolism − Sphingolipid metabolism
GSZ
20
15
10
5
0
−5
−10
−15
# features = 35 , max = 1
Transcription − RNA polymerase
GSZ
20
15
10
5
0
−5
−10
−15
# features = 41 , max = 3
Transcription factors − ARF
GSZ
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15
10
5
0
−5
−10
−15
# features = 18 , max = 3
Cutin suberine and wax biosynthesis
GSZ
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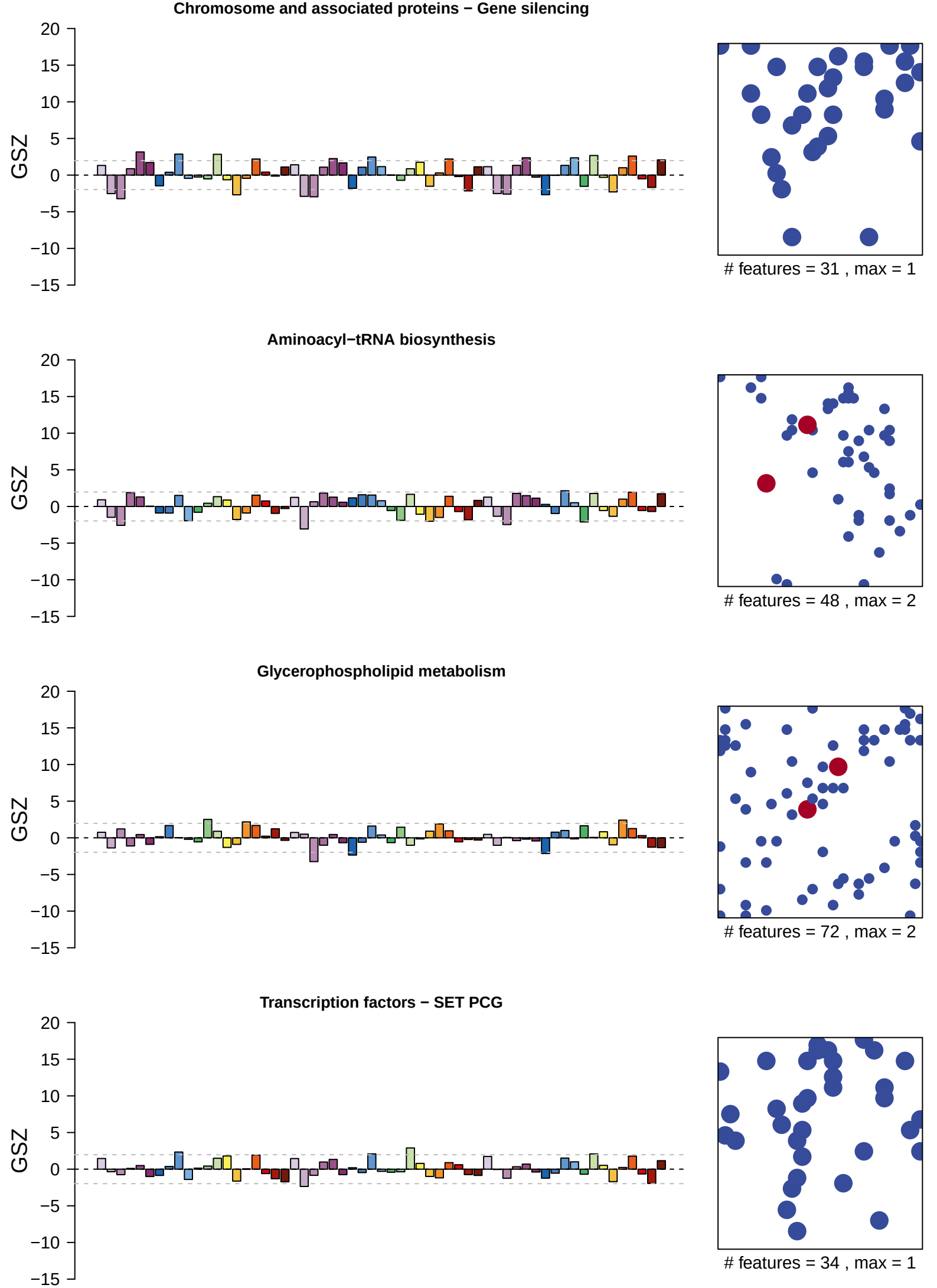
−10
−15
# features = 13 , max = 1


Chromosome and associated proteins − Gene silencing
GSZ
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Aminoacyl−tRNA biosynthesis
GSZ
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5
0
−5
−10
−15
# features = 48 , max = 2
Glycerophospholipid metabolism
GSZ
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5
0
−5
−10
−15
# features = 72 , max = 2
Transcription factors − SET PCG
GSZ
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15
10
5
0
−5
−10
−15
# features = 34 , max = 1