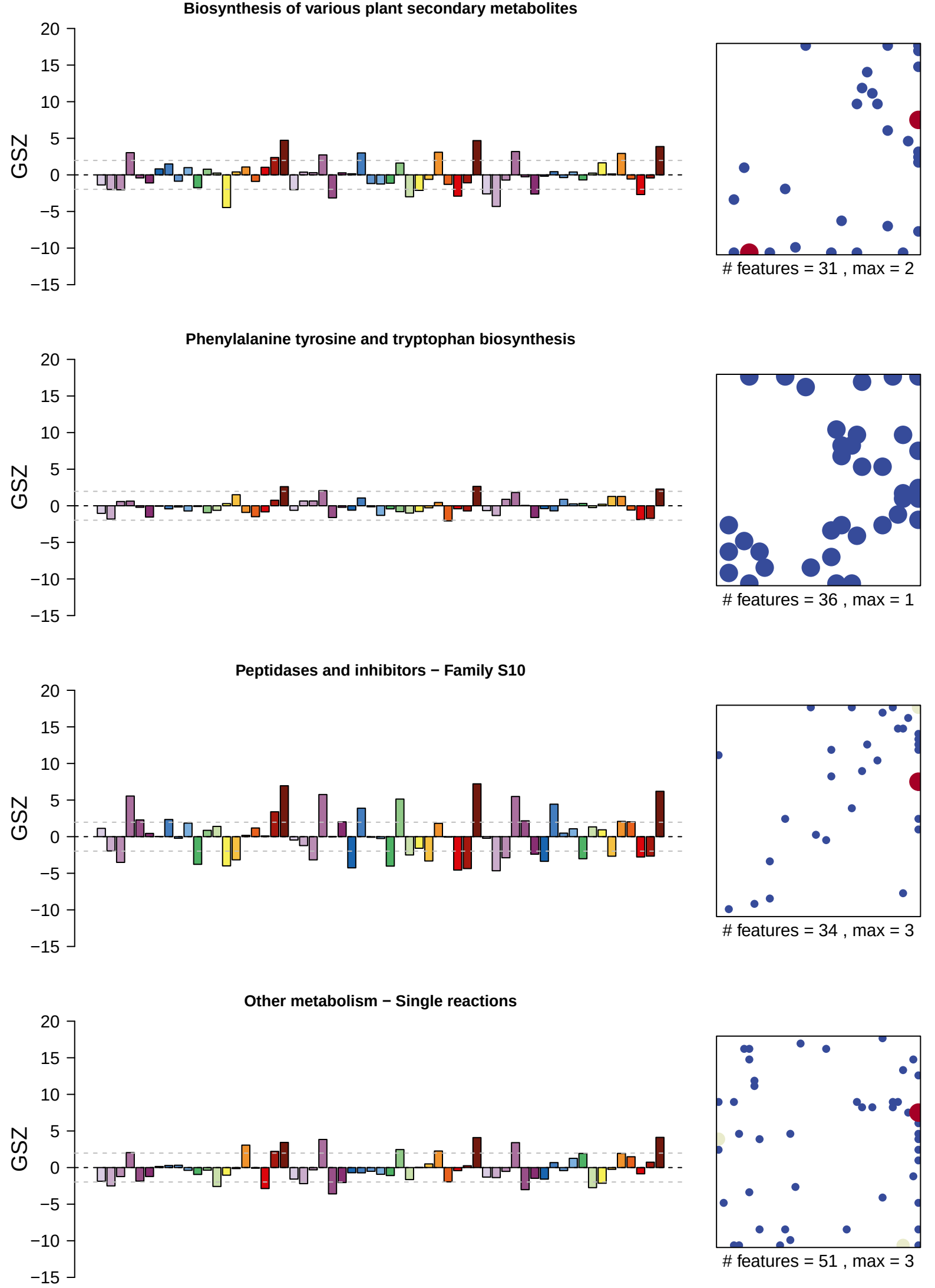
Biosynthesis of various plant secondary metabolites
GSZ
20
15
10
5
0
−5
−10
−15
features = 31 , max = 2
Phenylalanine tyrosine and tryptophan biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 36 , max = 1
Peptidases and inhibitors − Family S10
GSZ
20
15
10
5
0
−5
−10
−15
features = 34 , max = 3
Other metabolism − Single reactions
GSZ
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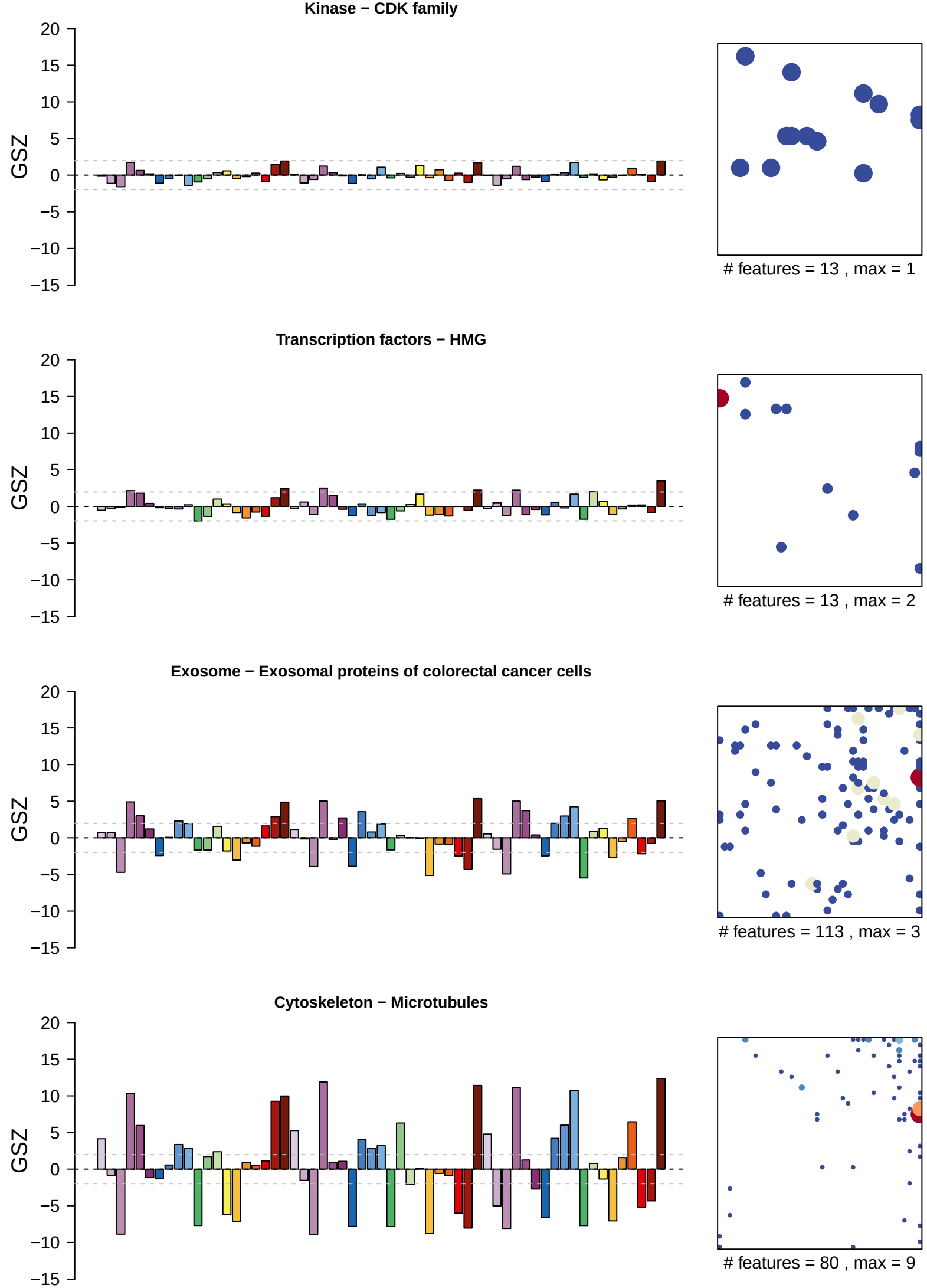
−10
−15
features = 51 , max = 3


Kinase − CDK family
GSZ
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−15
features = 13 , max = 1
Transcription factors − HMG
GSZ
20
15
10
5
0
−5
−10
−15
features = 13 , max = 2
Exosome − Exosomal proteins of colorectal cancer cells
GSZ
20
15
10
5
0
−5
−10
−15
features = 113 , max = 3
Cytoskeleton − Microtubules
GSZ
20
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10
5
0
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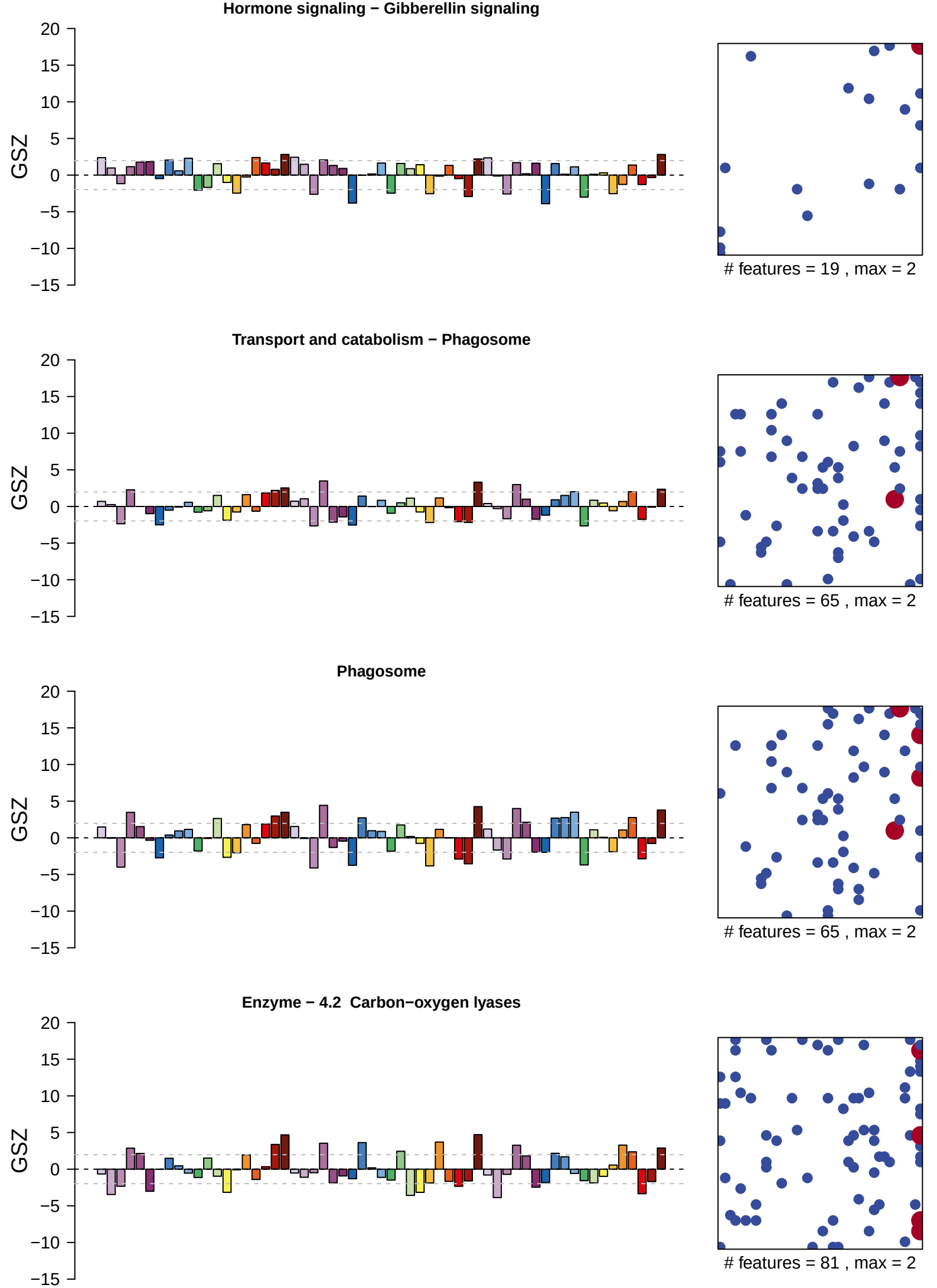
−10
−15
features = 80 , max = 9


Hormone signaling − Gibberellin signaling
GSZ
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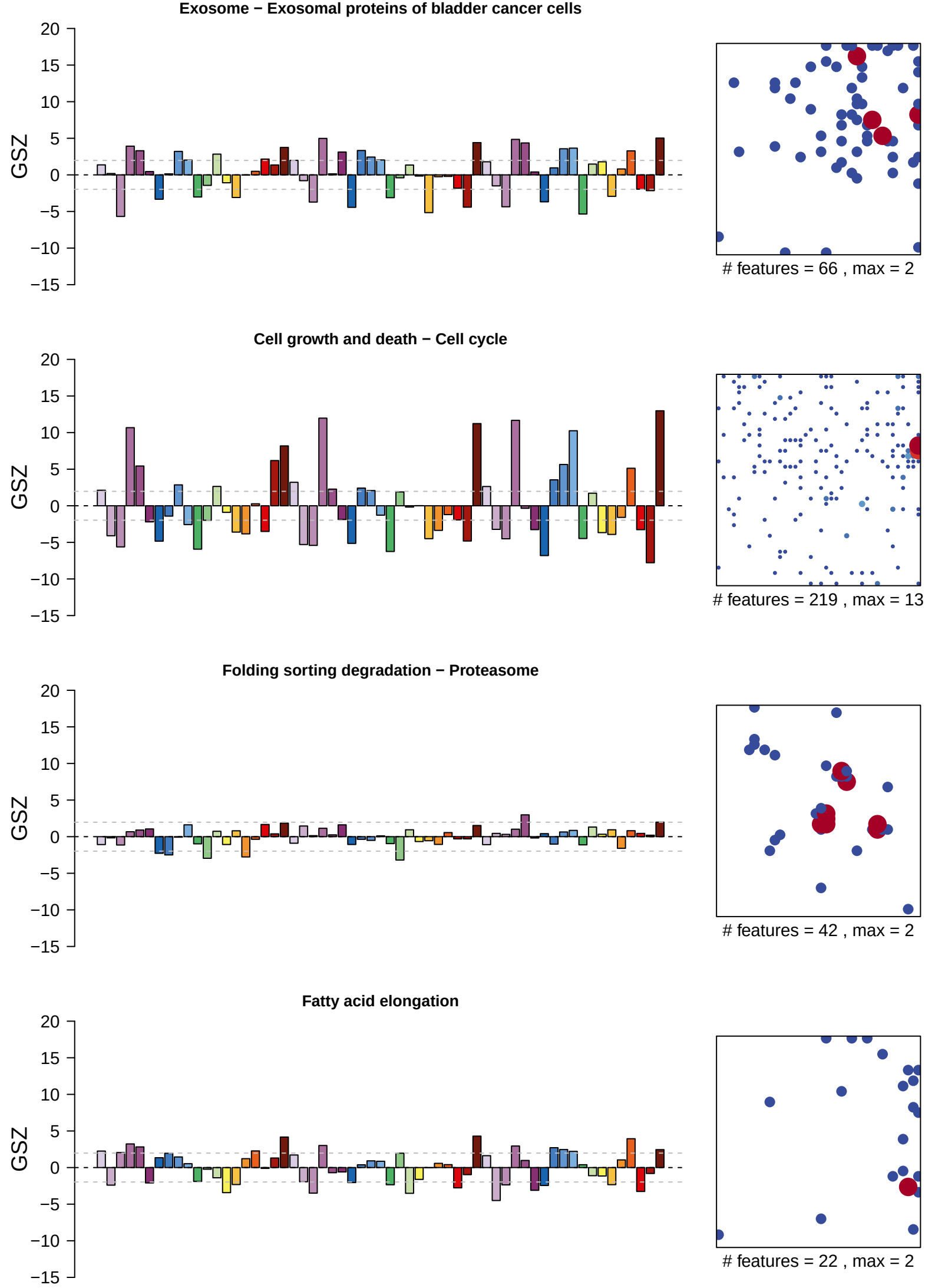
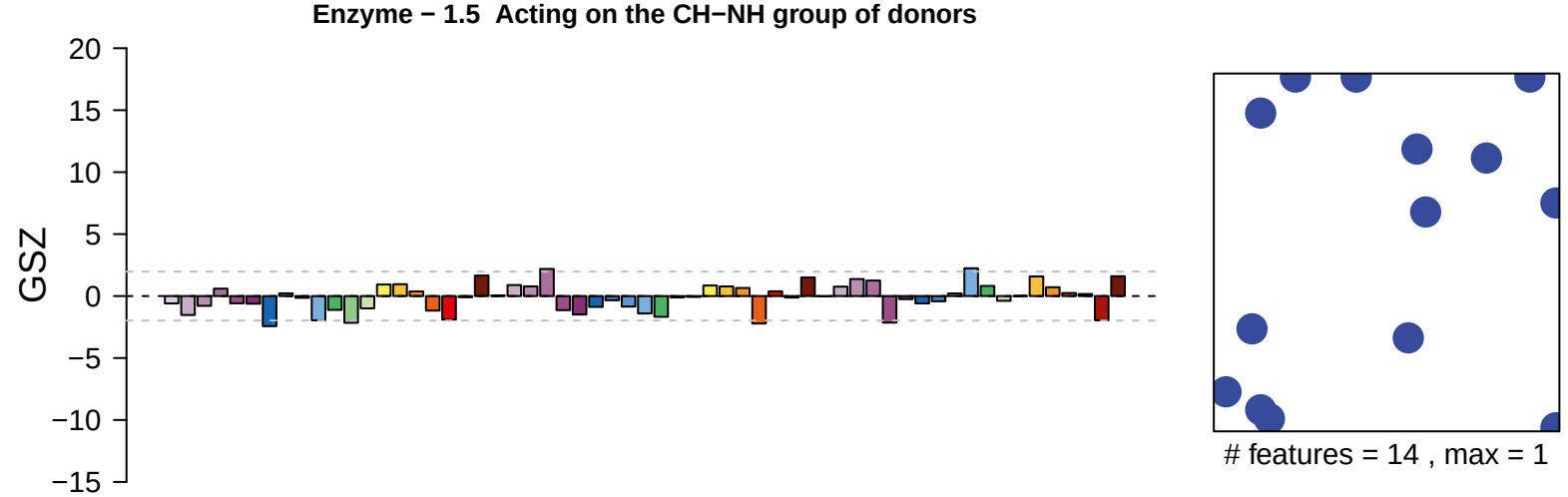
15
10
5
0
−5
−10
−15
features = 19 , max = 2
Transport and catabolism − Phagosome
GSZ
20
15
10
5
0
−5
−10
−15
features = 65 , max = 2
Phagosome
GSZ
20
15
10
5
0
−5
−10
−15
features = 65 , max = 2
Enzyme − 4.2 Carbon−oxygen lyases
GSZ
20
15
10
5
0
−5
−10
−15
features = 81 , max = 2


Exosome − Exosomal proteins of bladder cancer cells
GSZ
20
15
10
5
0
−5
−10
−15
features = 66 , max = 2
Cell growth and death − Cell cycle
GSZ
features = 219 , max = 13
Folding sorting degradation − Proteasome
GSZ
features = 42 , max = 2
Fatty acid elongation
GSZ
features = 22 , max = 2


Enzyme − 1.5 Acting on the CH−NH group of donors
GSZ
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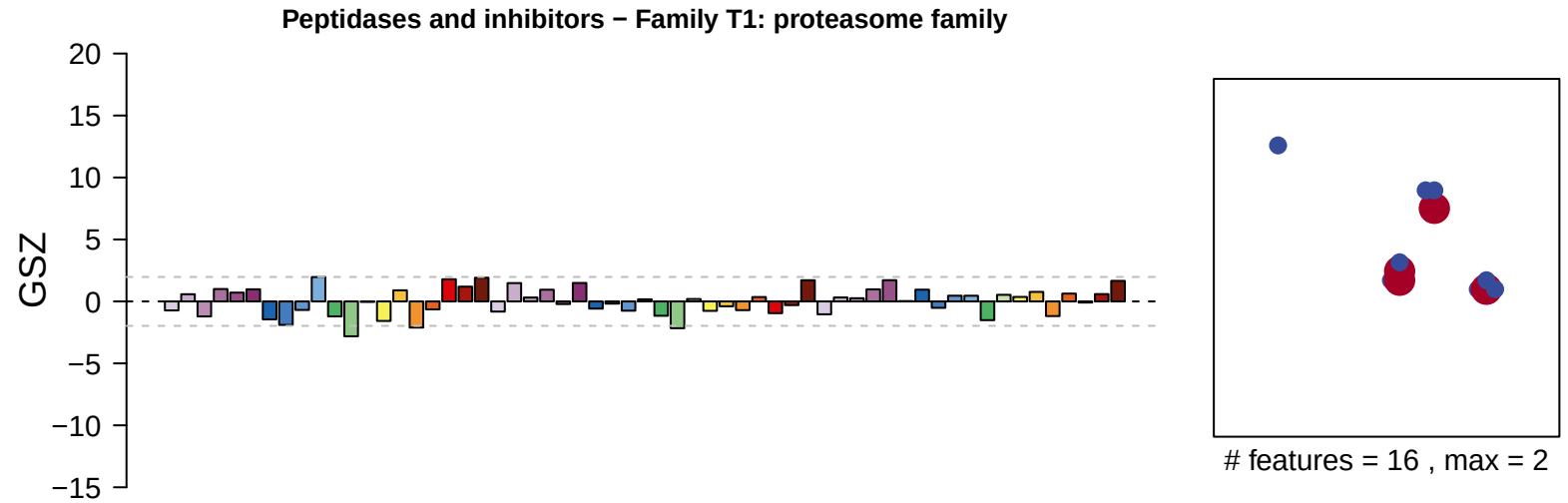
10
5
0
−5
−10
−15
features = 14 , max = 1


Peptidases and inhibitors − Family T1: proteasome family
GSZ
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15
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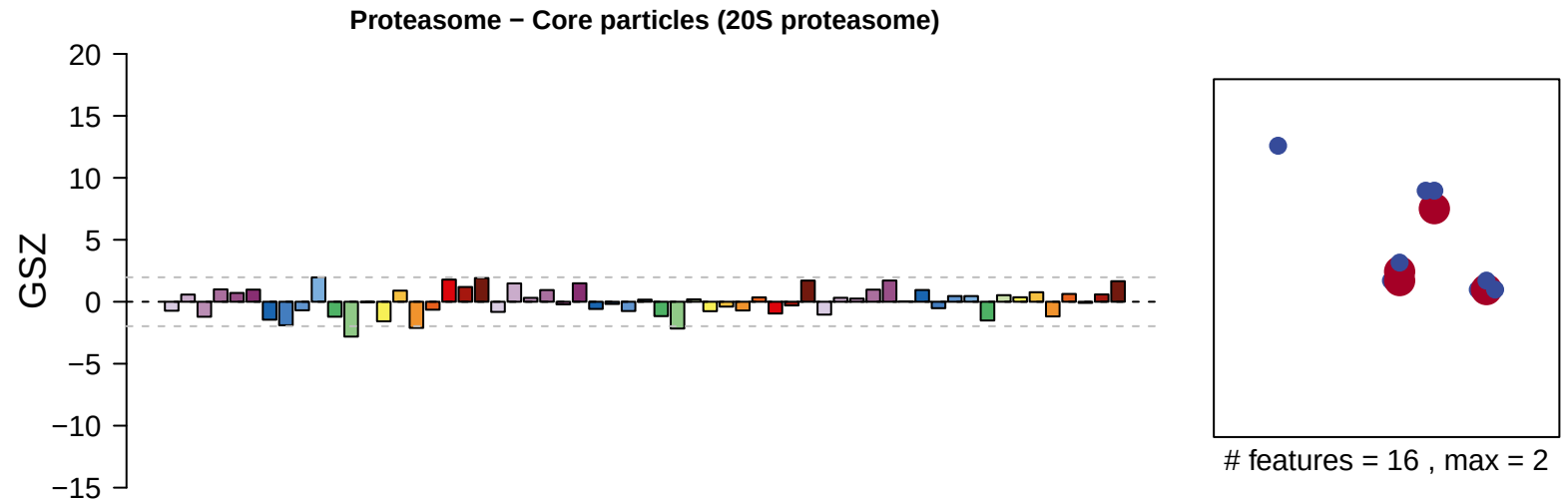
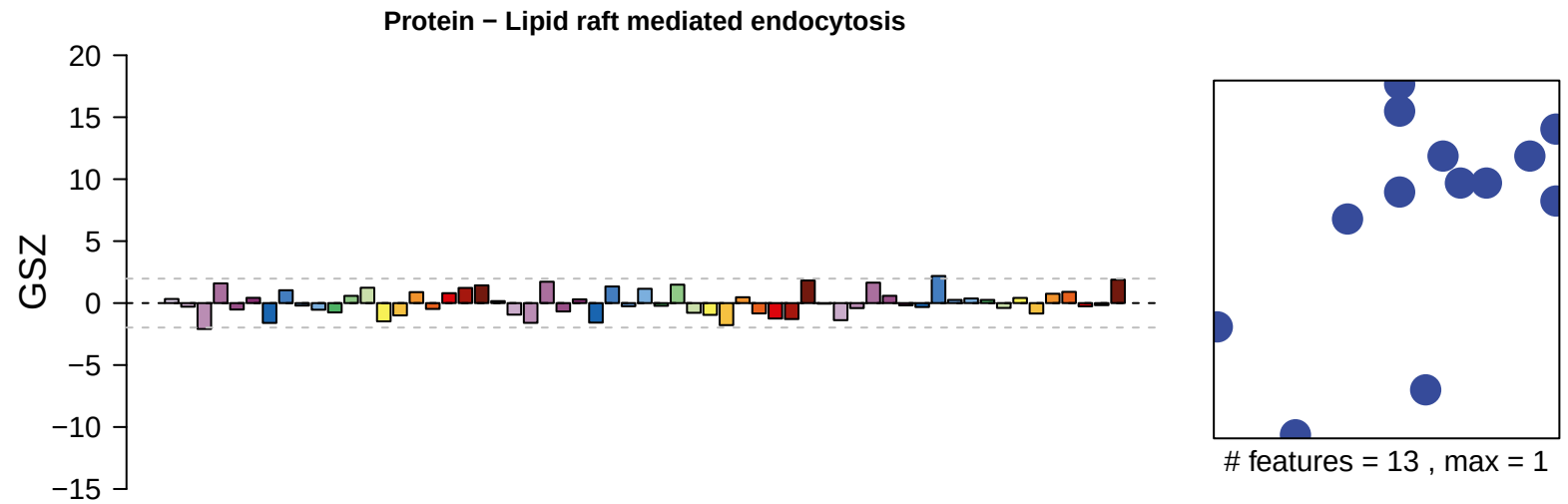
5
0
−5
−10
−15
features = 16 , max = 2


Proteasome − Core particles (20S proteasome)
GSZ
20
15
10
5
0
−5
−10
−15
features = 16 , max = 2


Protein − Lipid raft mediated endocytosis
GSZ
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5
0
−5
−10
−15
features = 13 , max = 1