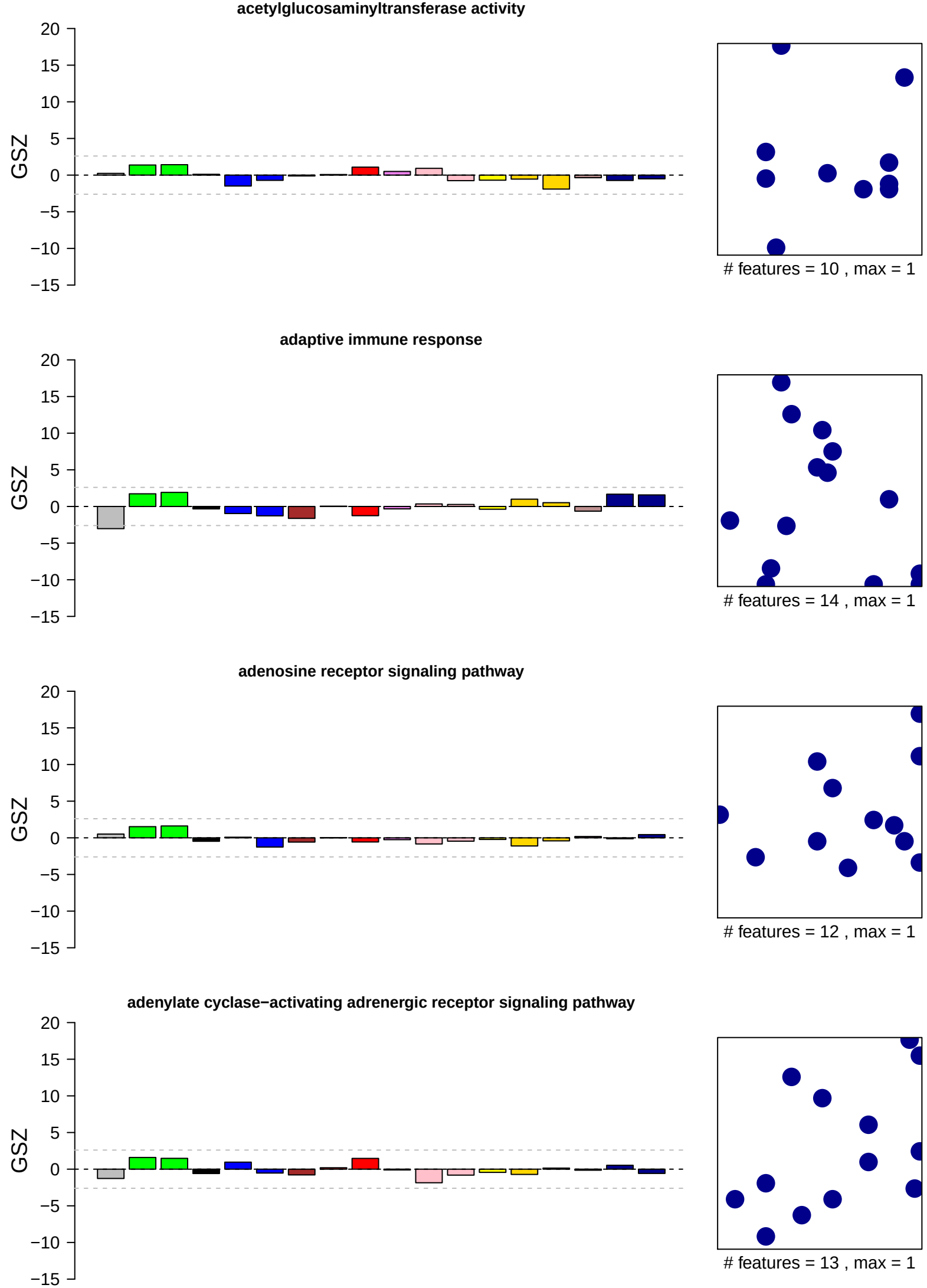
acetylglucosaminyltransferase activity
GSZ
20
15
10
5
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−5
−10
−15
features = 10 , max = 1
adaptive immune response
GSZ
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5
0
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−10
−15
features = 14 , max = 1
adenosine receptor signaling pathway
GSZ
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10
5
0
−5
−10
−15
features = 12 , max = 1
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GSZ
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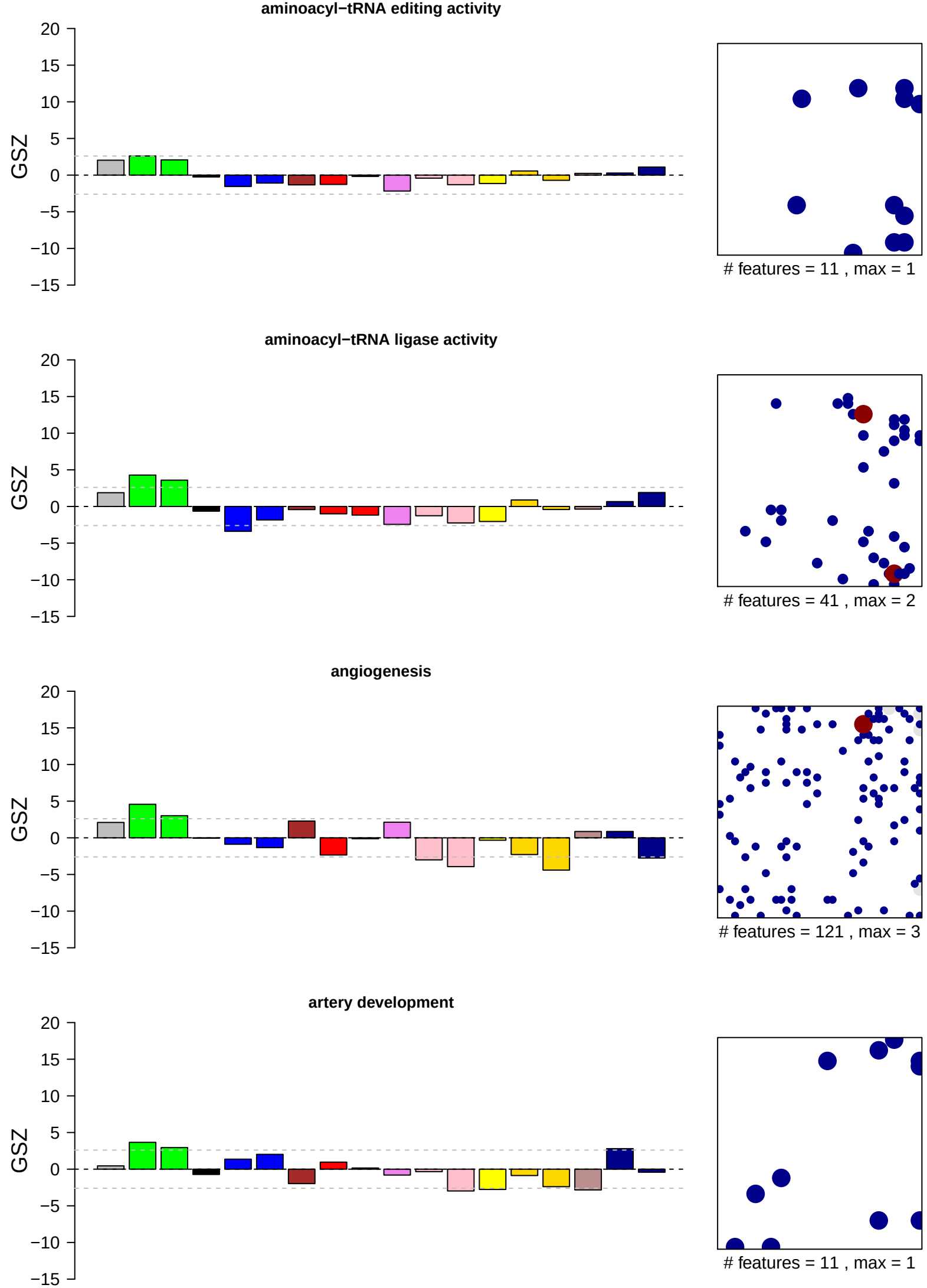
features = 13 , max = 1


aminoacyl−tRNA editing activity
GSZ
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−10
−15
features = 11 , max = 1
aminoacyl−tRNA ligase activity
GSZ
20
15
10
5
0
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−10
−15
features = 41 , max = 2
angiogenesis
GSZ
20
15
10
5
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−5
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−15
features = 121 , max = 3
artery development
GSZ
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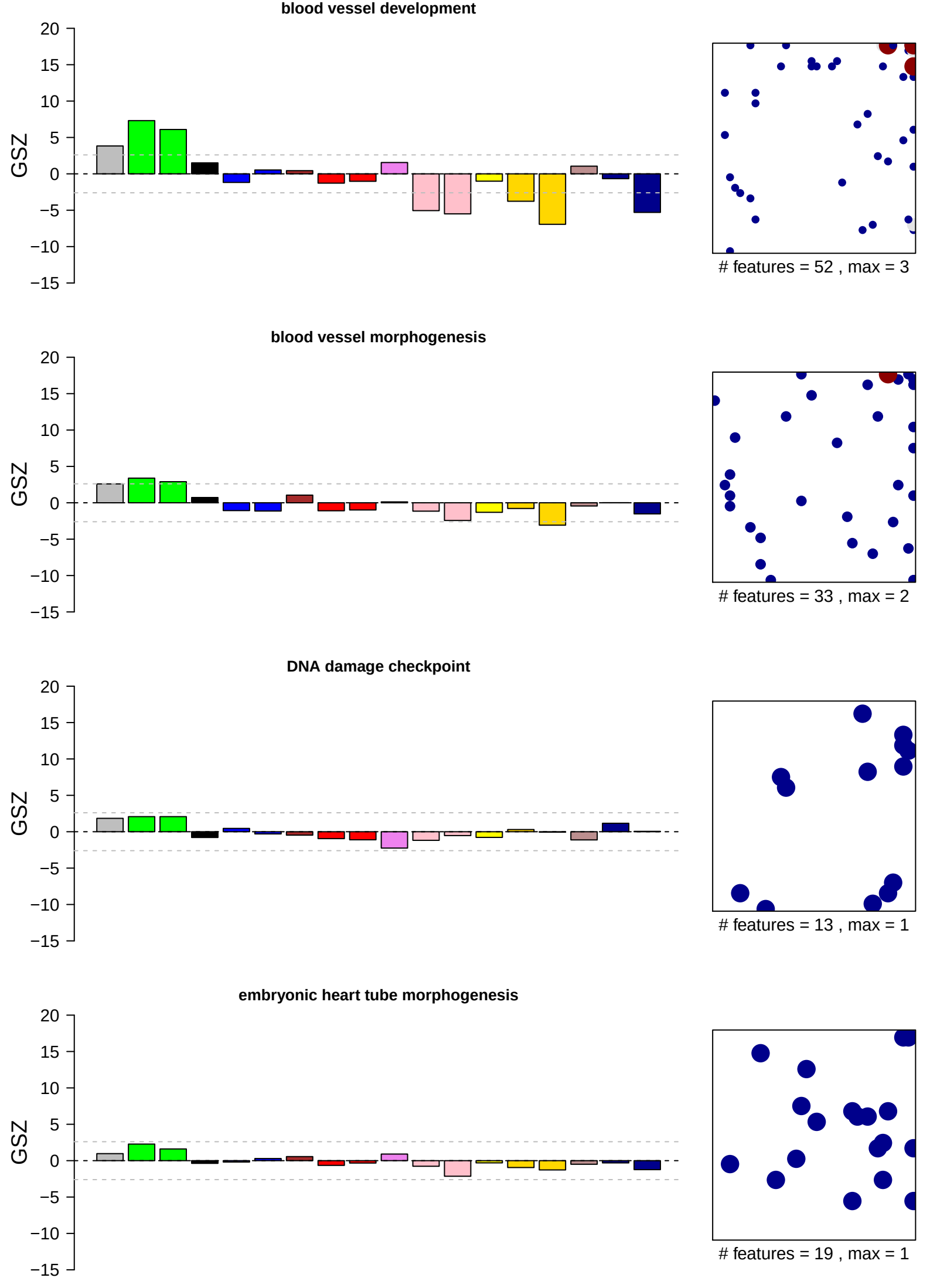
−10
−15
features = 11 , max = 1


blood vessel development
GSZ
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5
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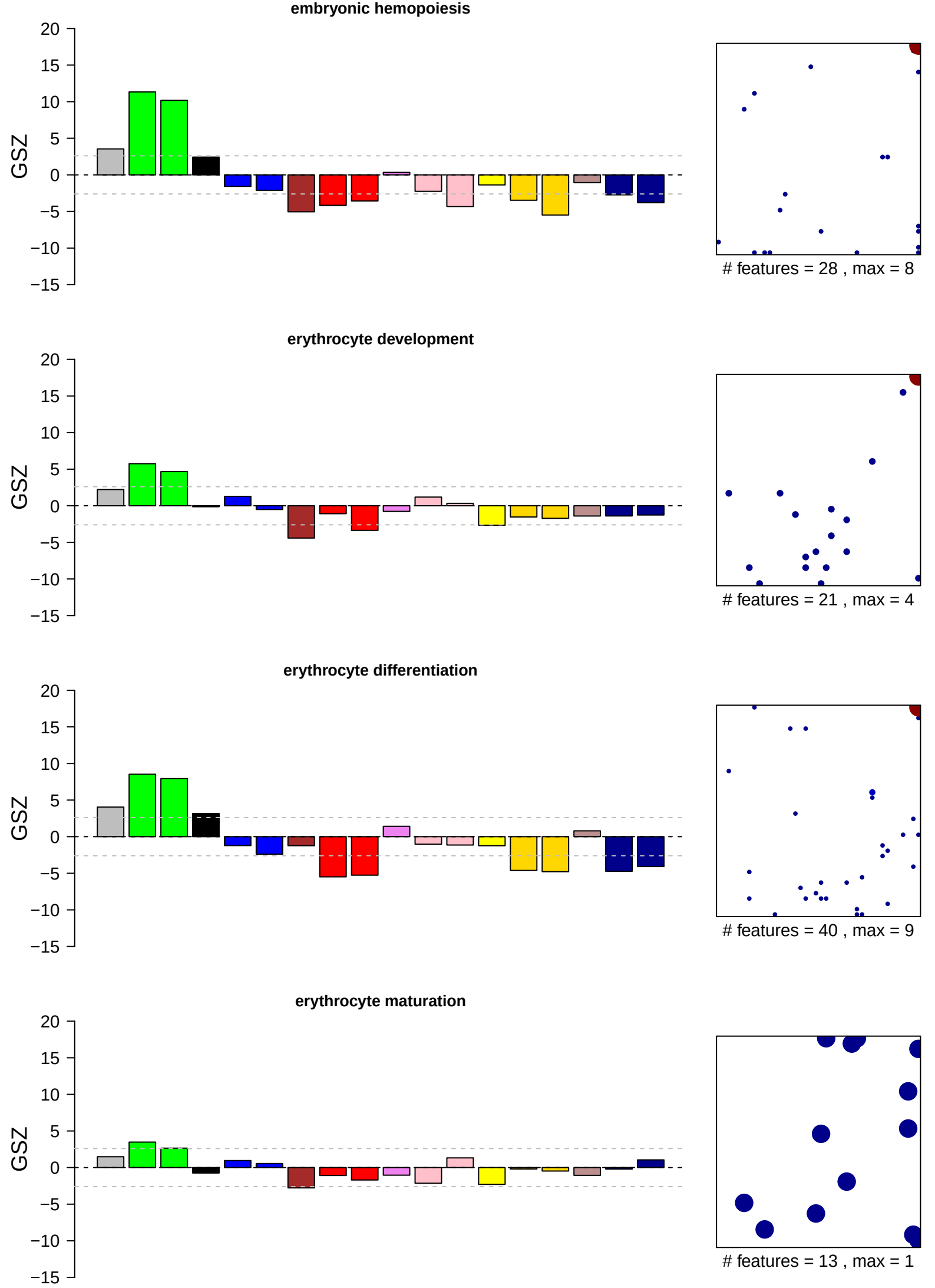
−10
−15
features = 52 , max = 3
blood vessel morphogenesis
GSZ
features = 33 , max = 2
DNA damage checkpoint
GSZ
features = 13 , max = 1
embryonic heart tube morphogenesis
GSZ
features = 19 , max = 1


embryonic hemopoiesis
GSZ
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features = 28 , max = 8
erythrocyte development
GSZ
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5
0
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−10
−15
features = 21 , max = 4
erythrocyte differentiation
GSZ
20
15
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0
−5
−10
−15
features = 40 , max = 9
erythrocyte maturation
GSZ
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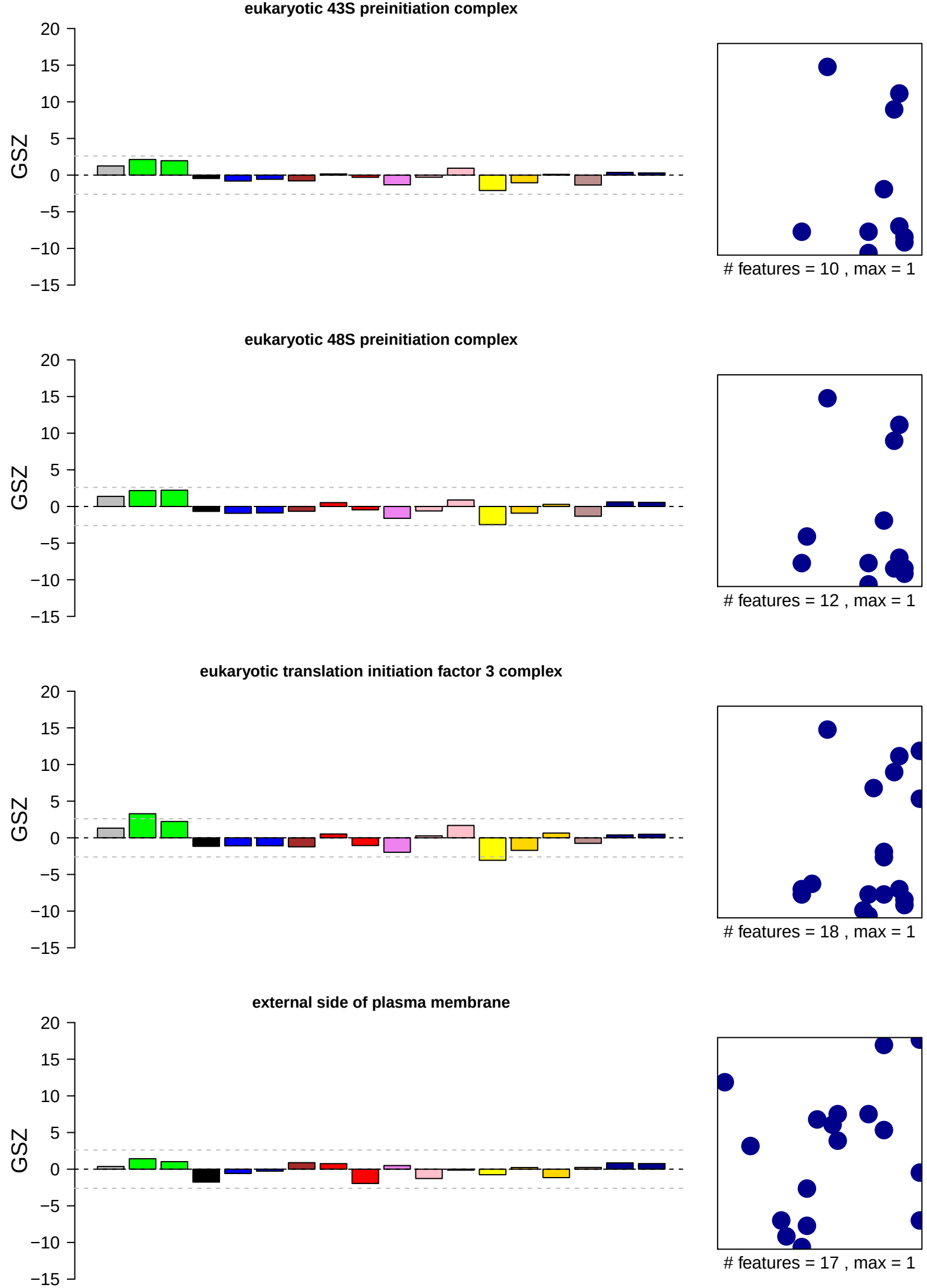
−15
features = 13 , max = 1


eukaryotic 43S preinitiation complex
GSZ
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15
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5
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features = 10 , max = 1
eukaryotic 48S preinitiation complex
GSZ
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15
10
5
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−15
features = 12 , max = 1
eukaryotic translation initiation factor 3 complex
GSZ
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15
10
5
0
−5
−10
−15
features = 18 , max = 1
external side of plasma membrane
GSZ
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5
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−10
−15
features = 17 , max = 1