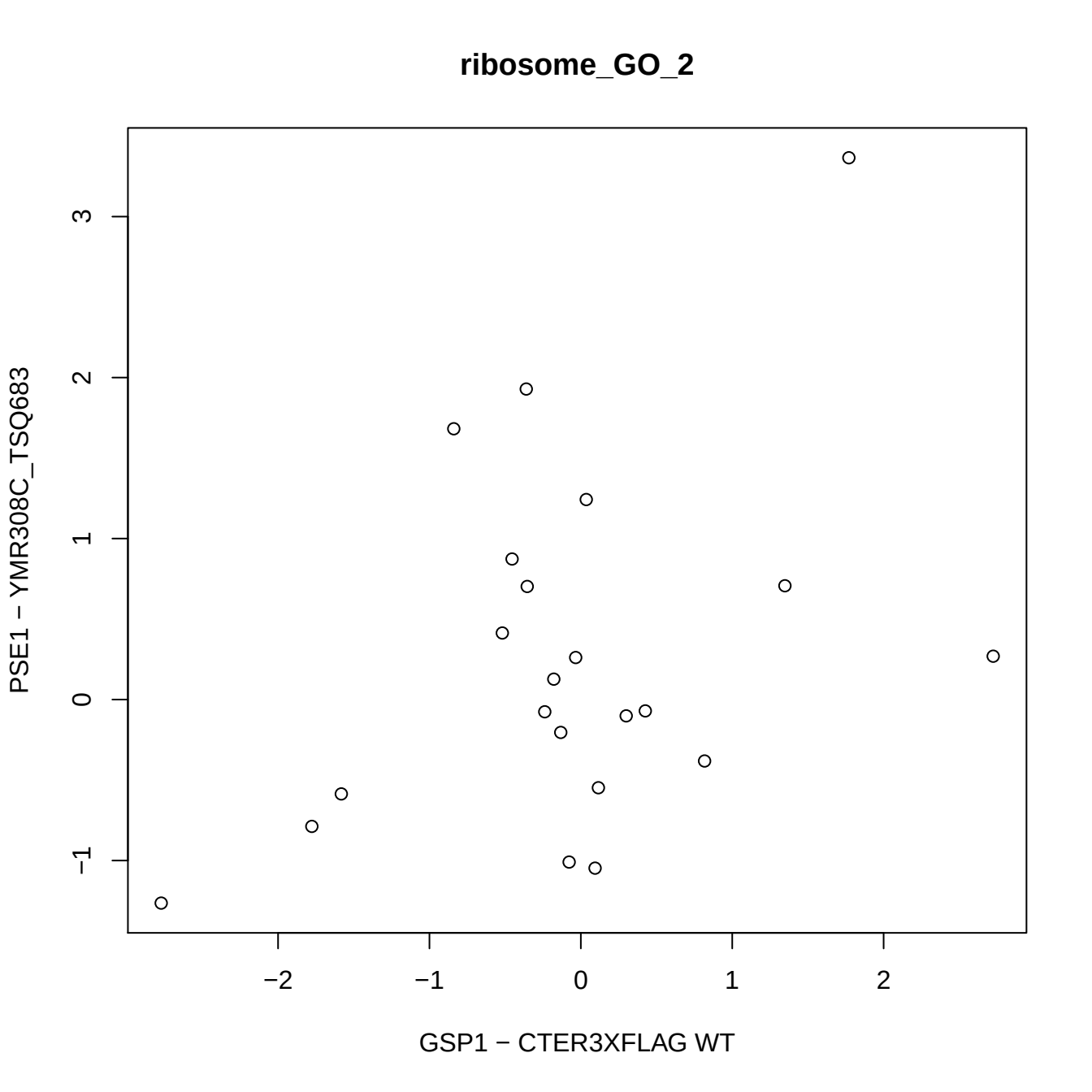
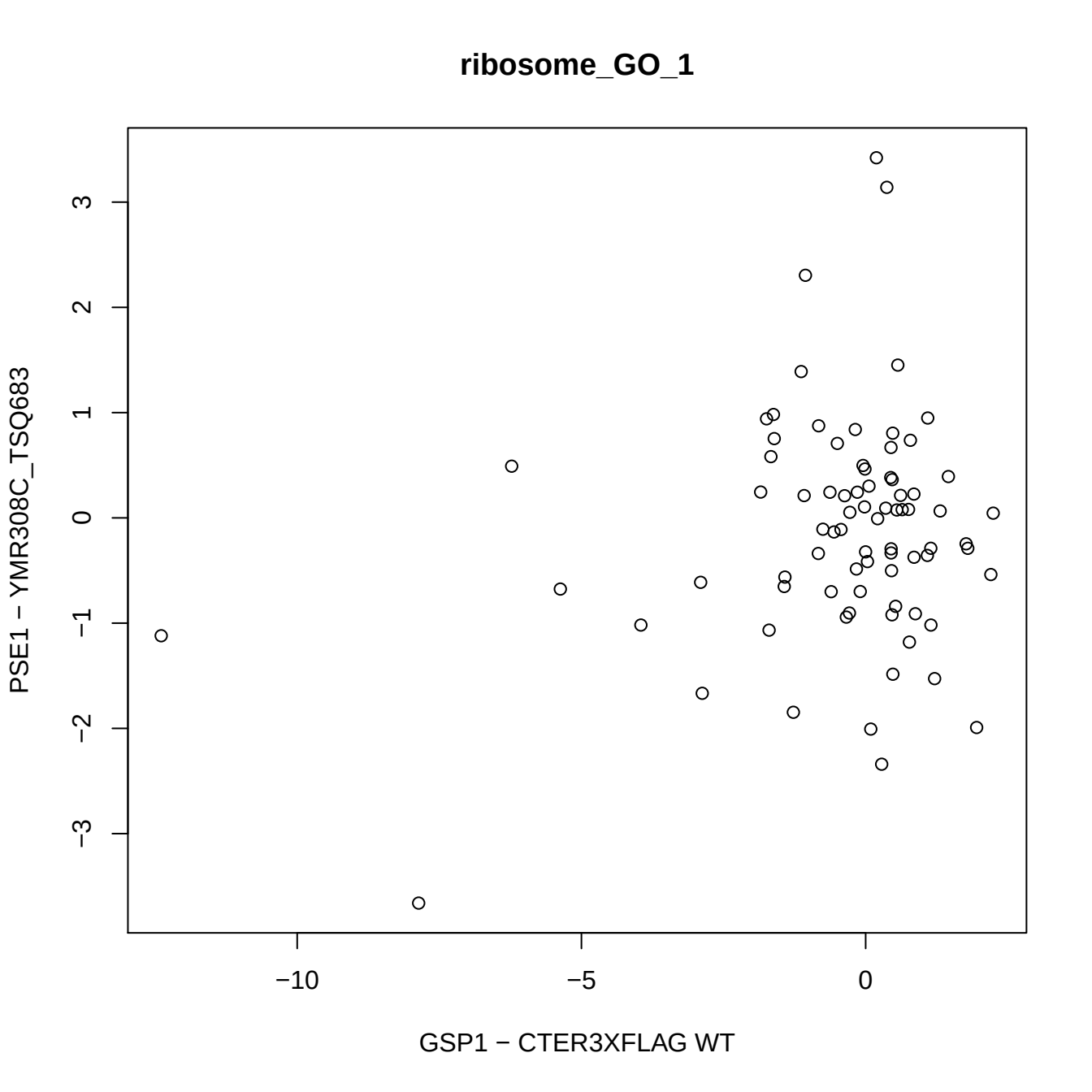
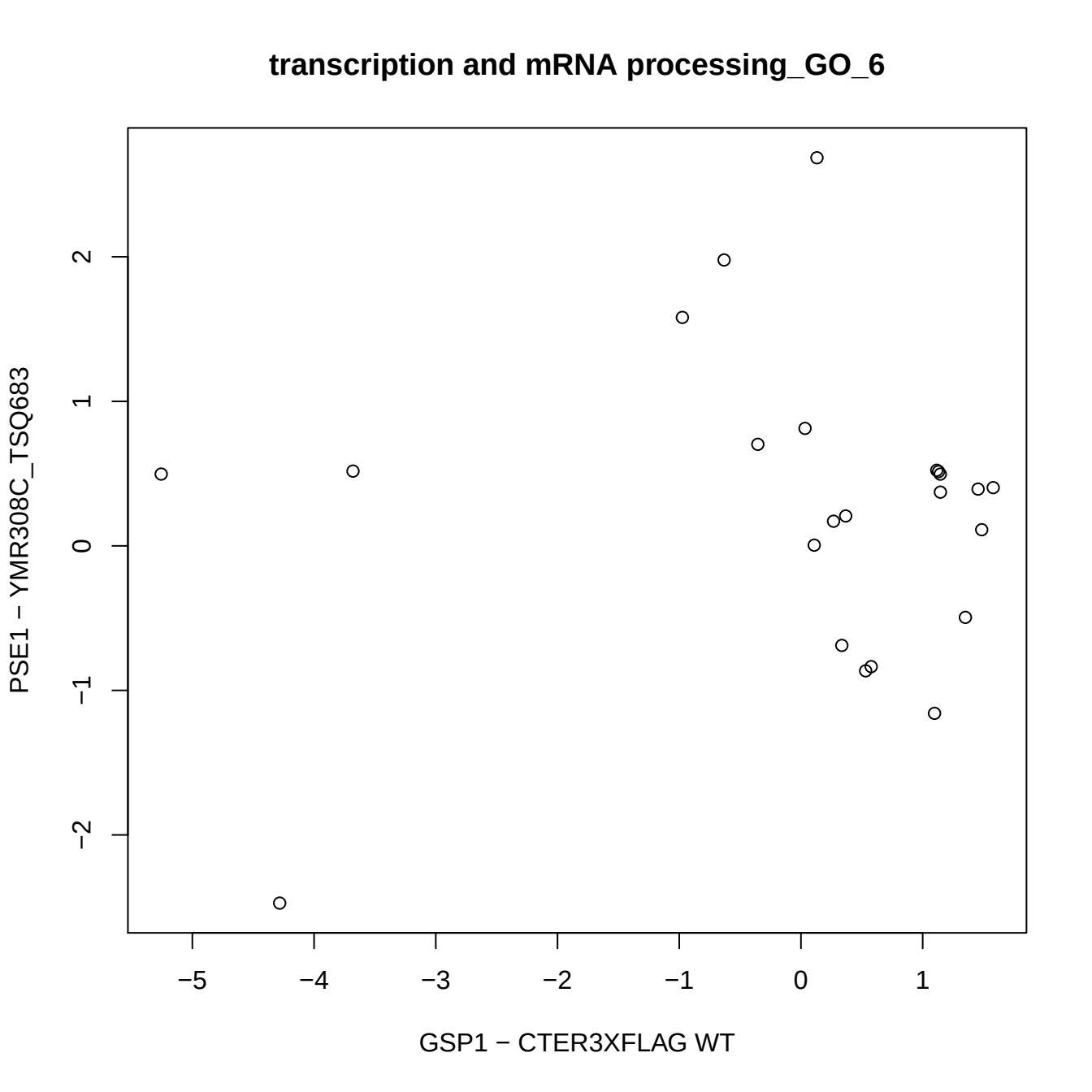
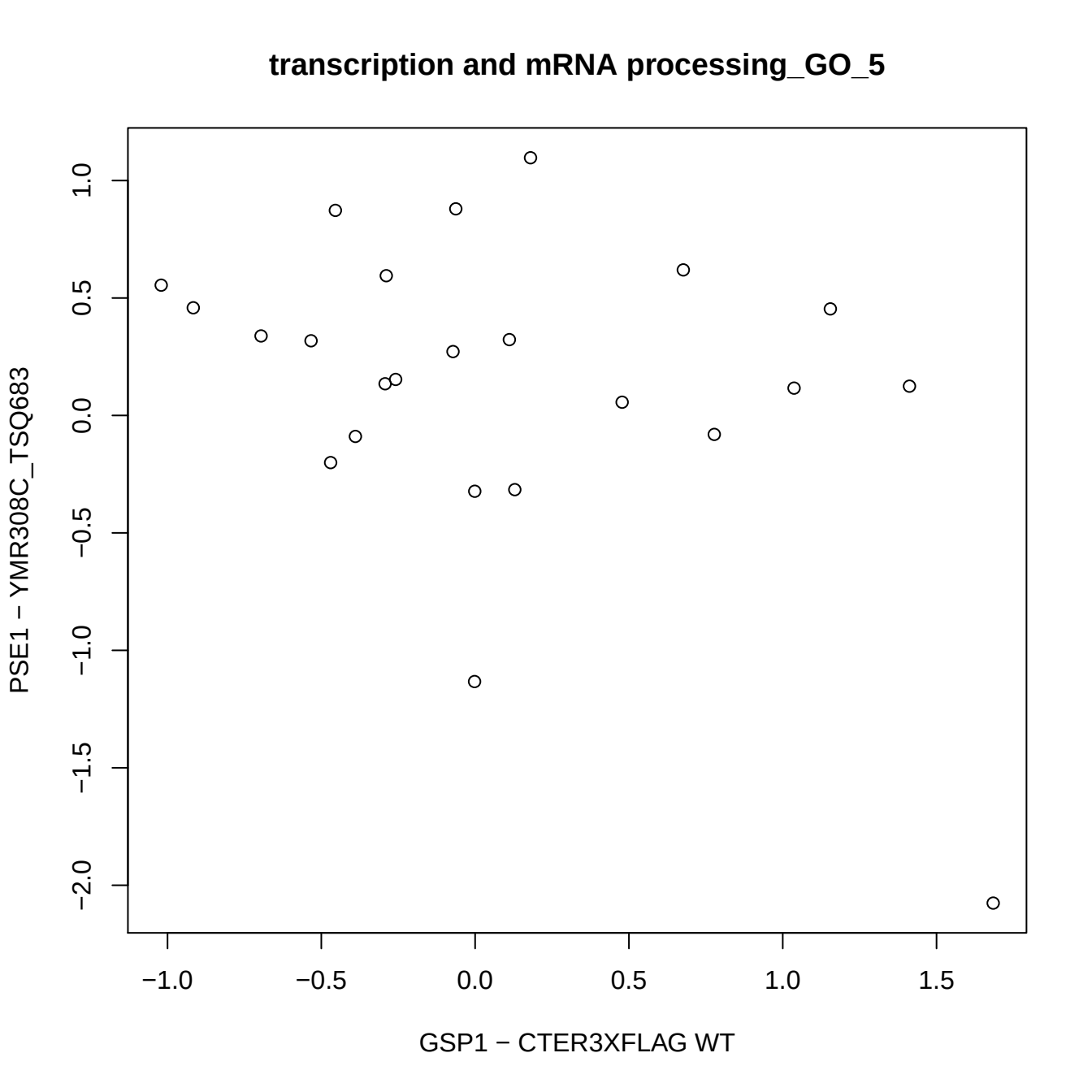
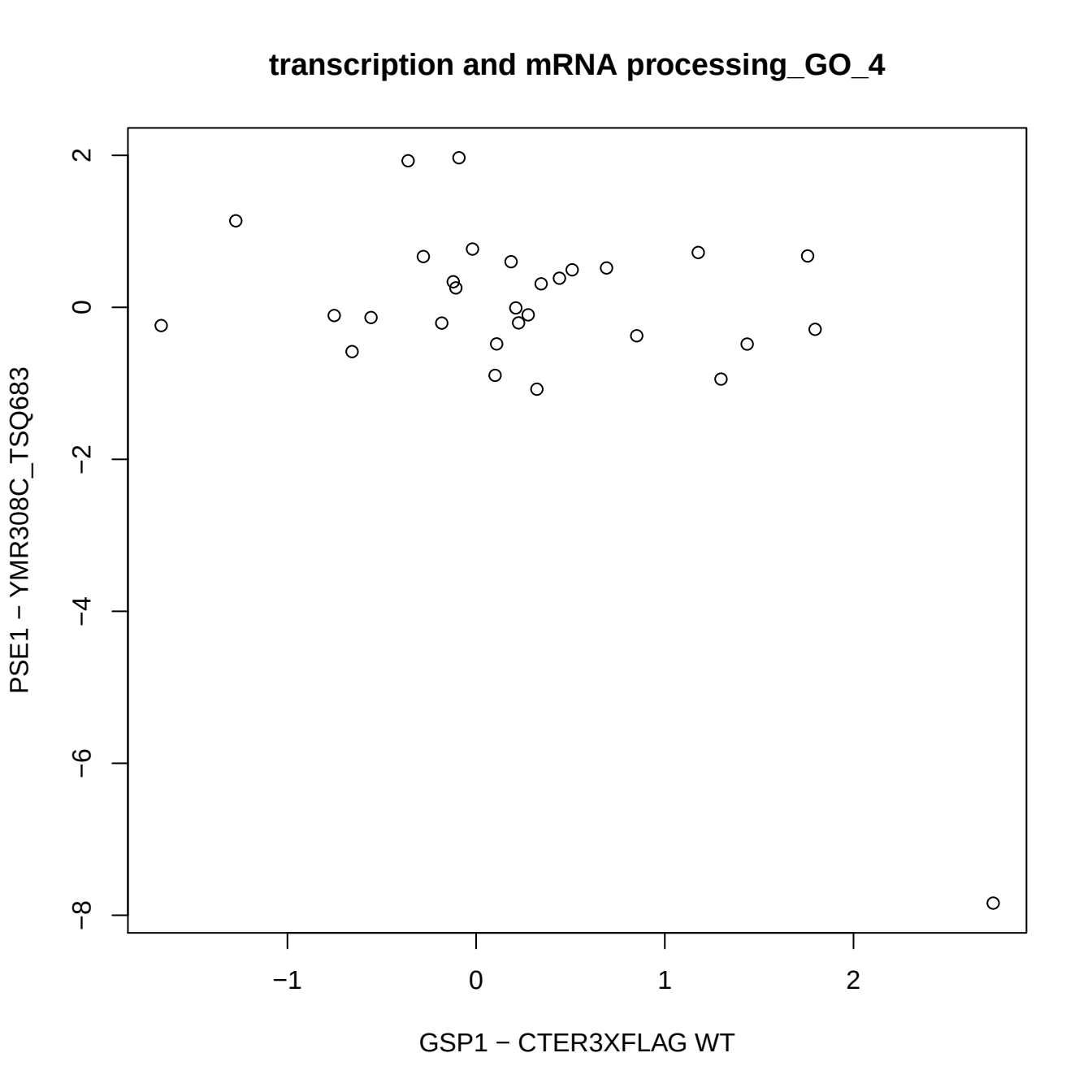
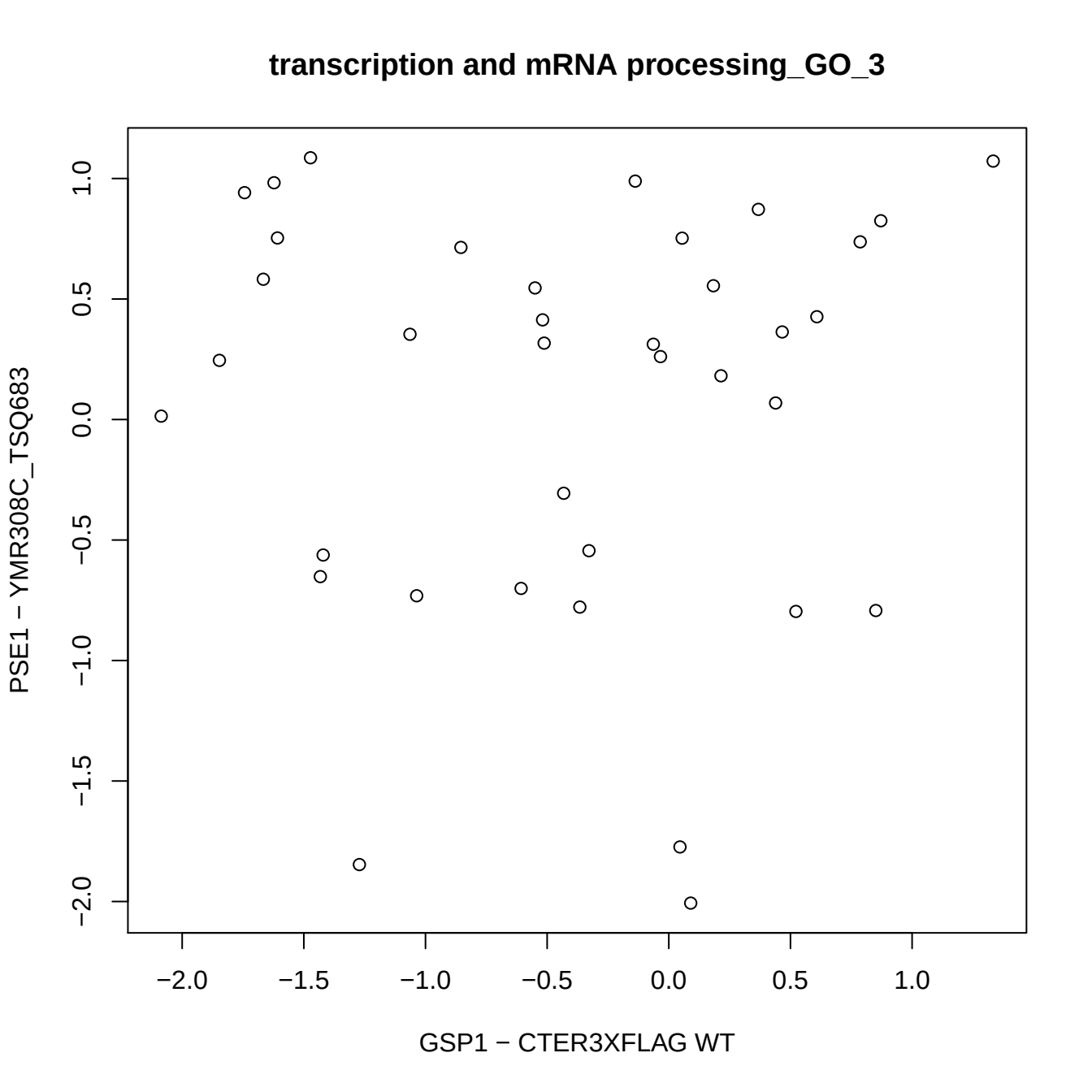

ribosome_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
3
2
1
0
−1
−2
−1
0
1
2

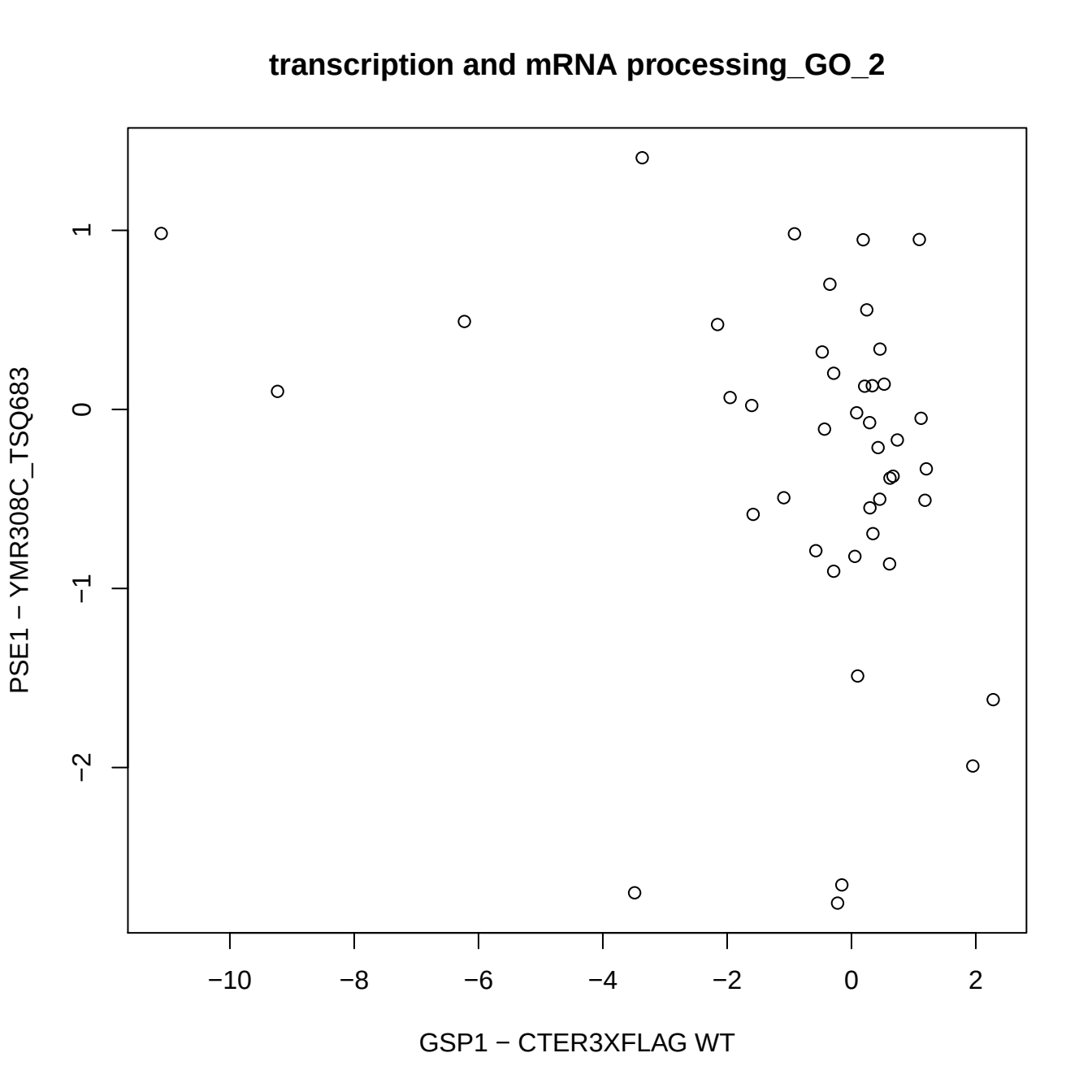

ribosome_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
3
2
1
0
−1
−2
−3
−10
−5
0

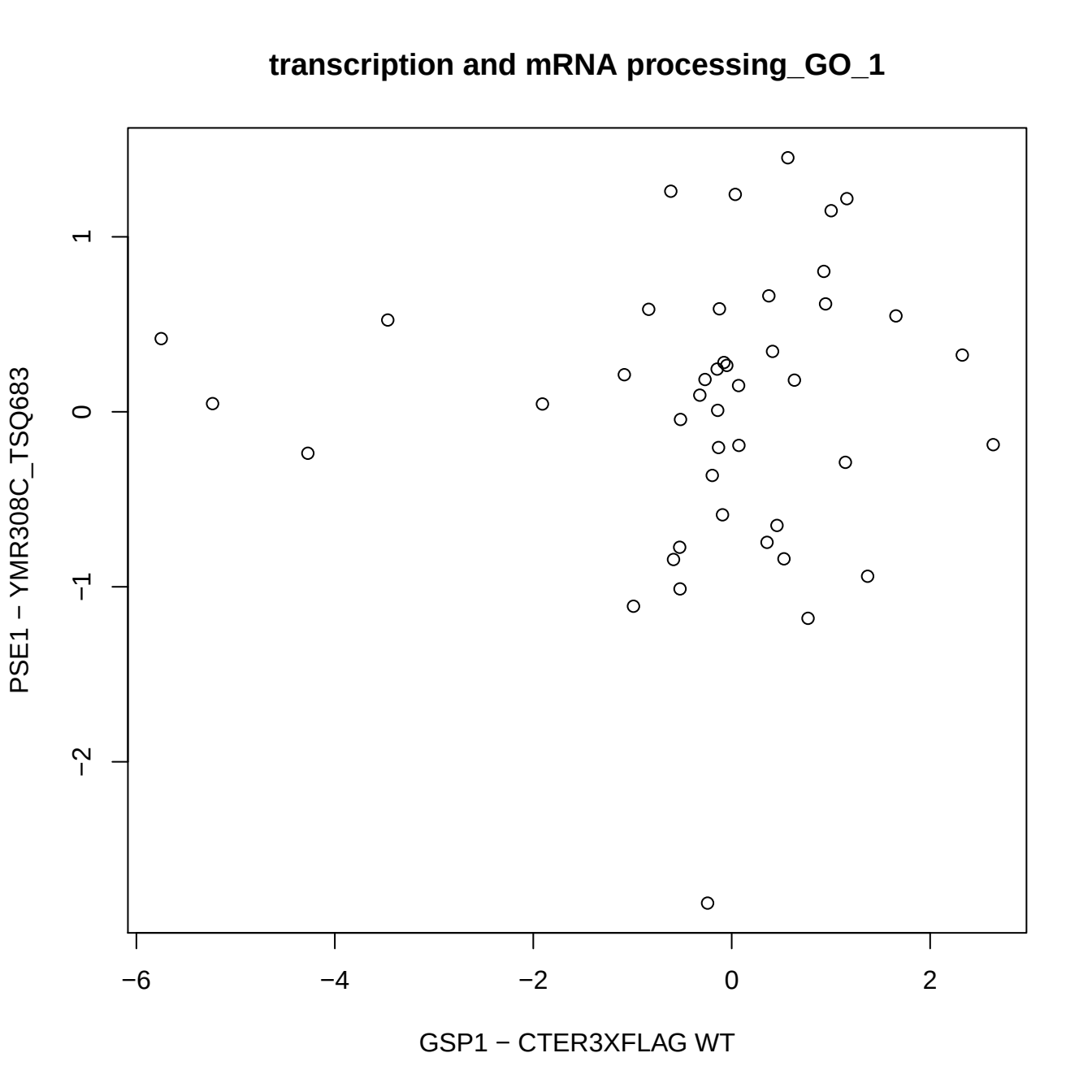

transcription and mRNA processing_GO_6
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
1
0
−1
−2
−5
−4
−3
−2
−1
0
1

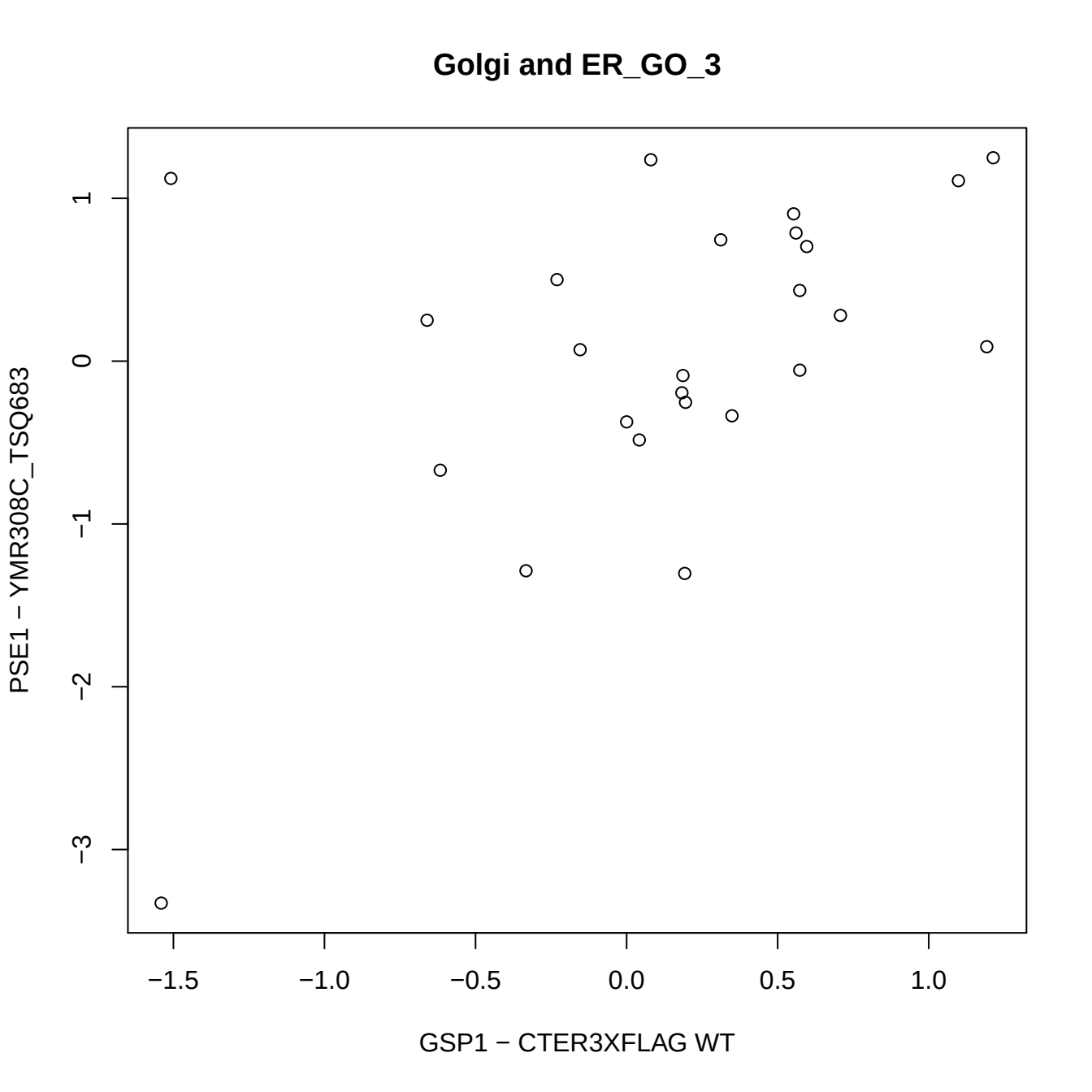

transcription and mRNA processing_GO_5
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1.0
0.5
0.0
−0.5
−1.0
−1.5
−2.0
−1.0
−0.5
0.0
0.5
1.0
1.5

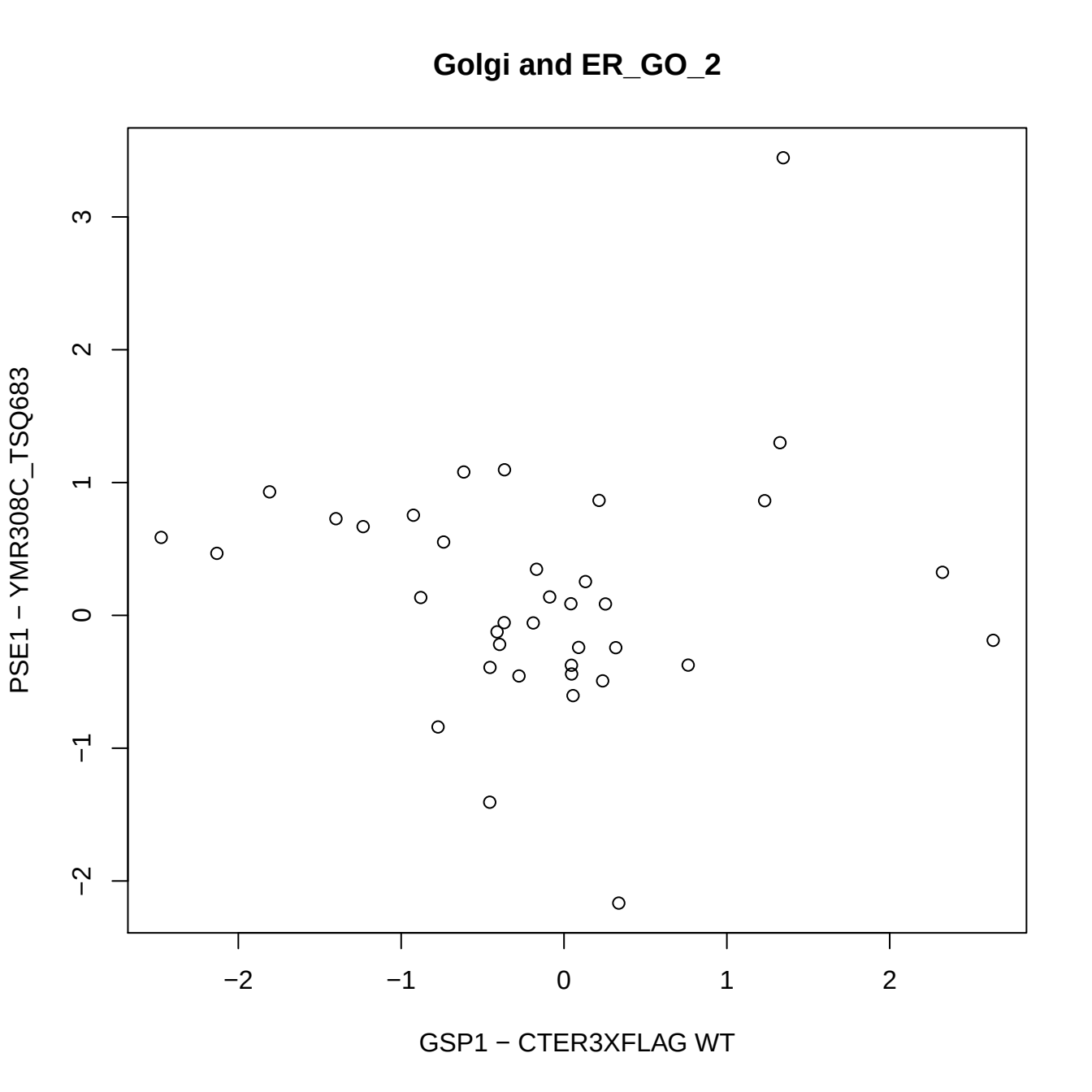

transcription and mRNA processing_GO_4
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
0
−2
−4
−6
−8
−1
0
1
2

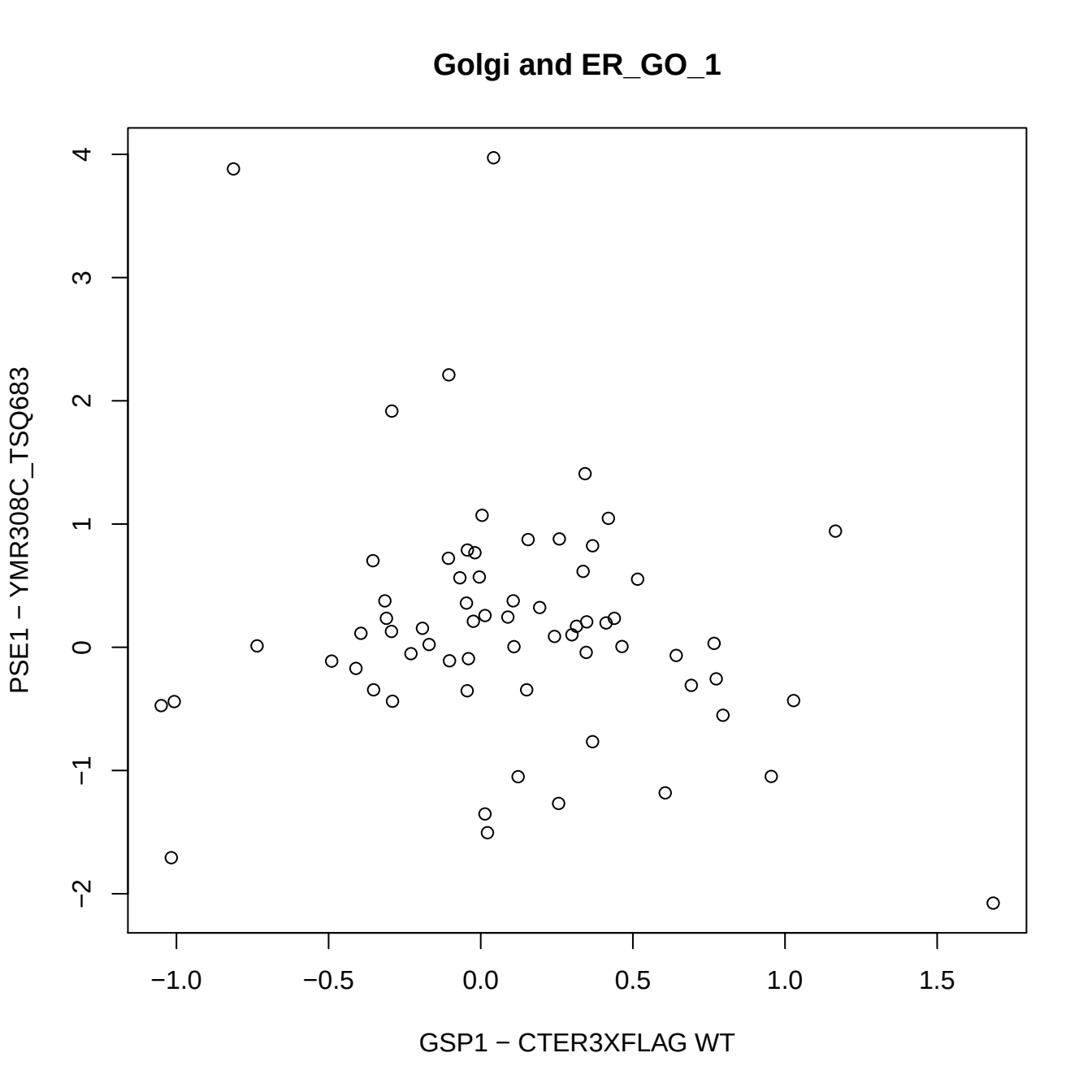

transcription and mRNA processing_GO_3
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1.0
0.5
0.0
−0.5
−1.0
−1.5
−2.0
−2.0
−1.5
−1.0
−0.5
0.0
0.5
1.0

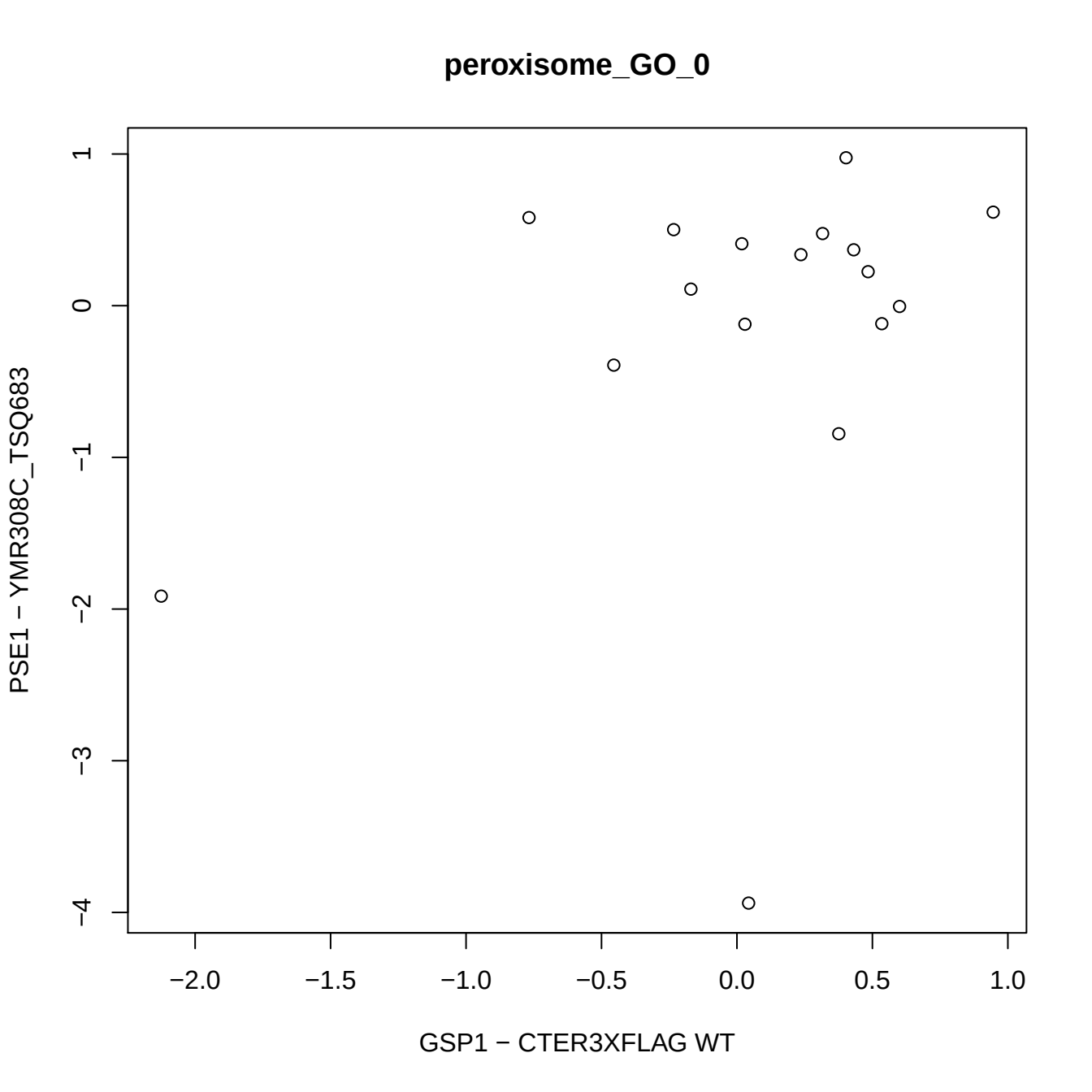

transcription and mRNA processing_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−10
−8
−6
−4
−2
0
2

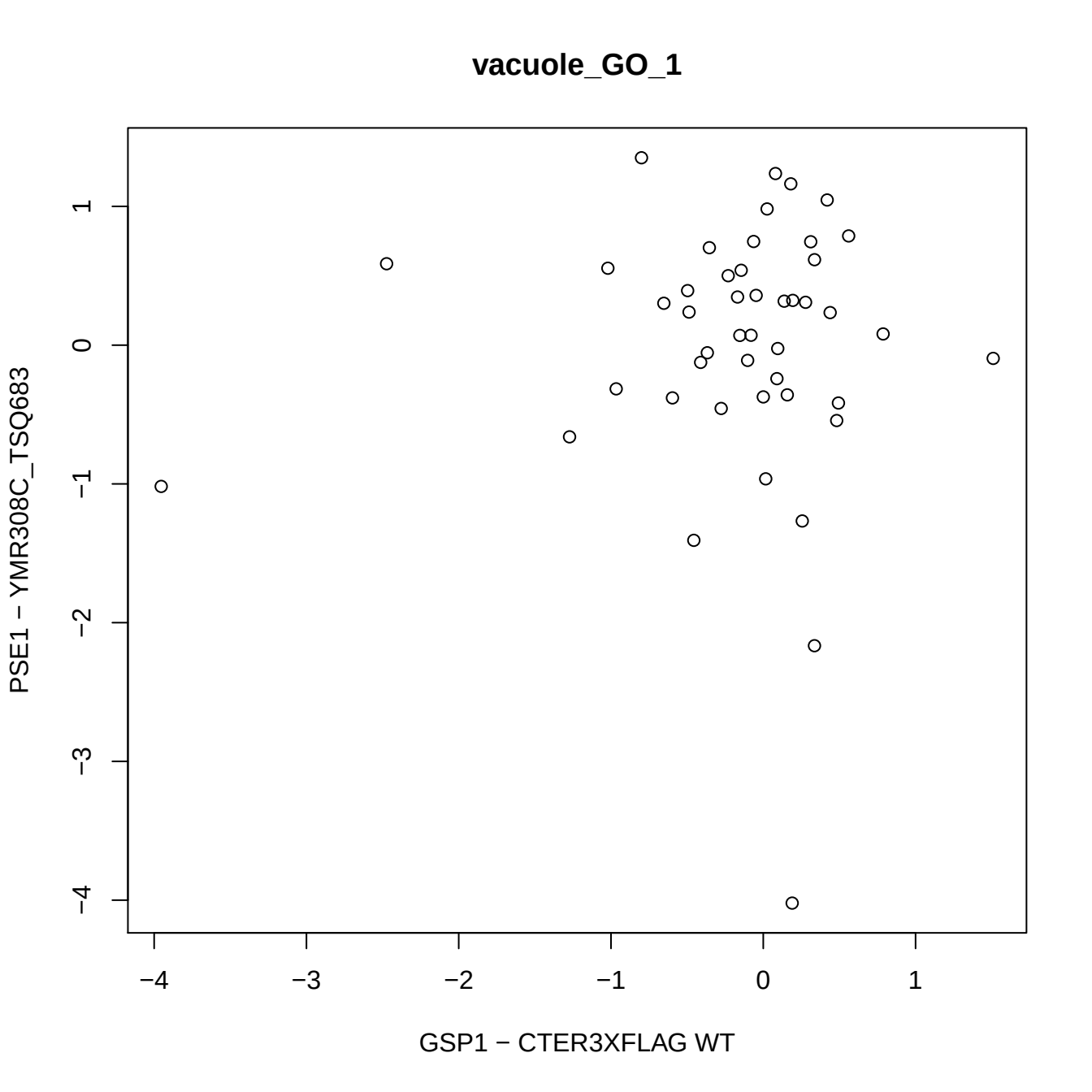

transcription and mRNA processing_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−6
−4
−2
0
2

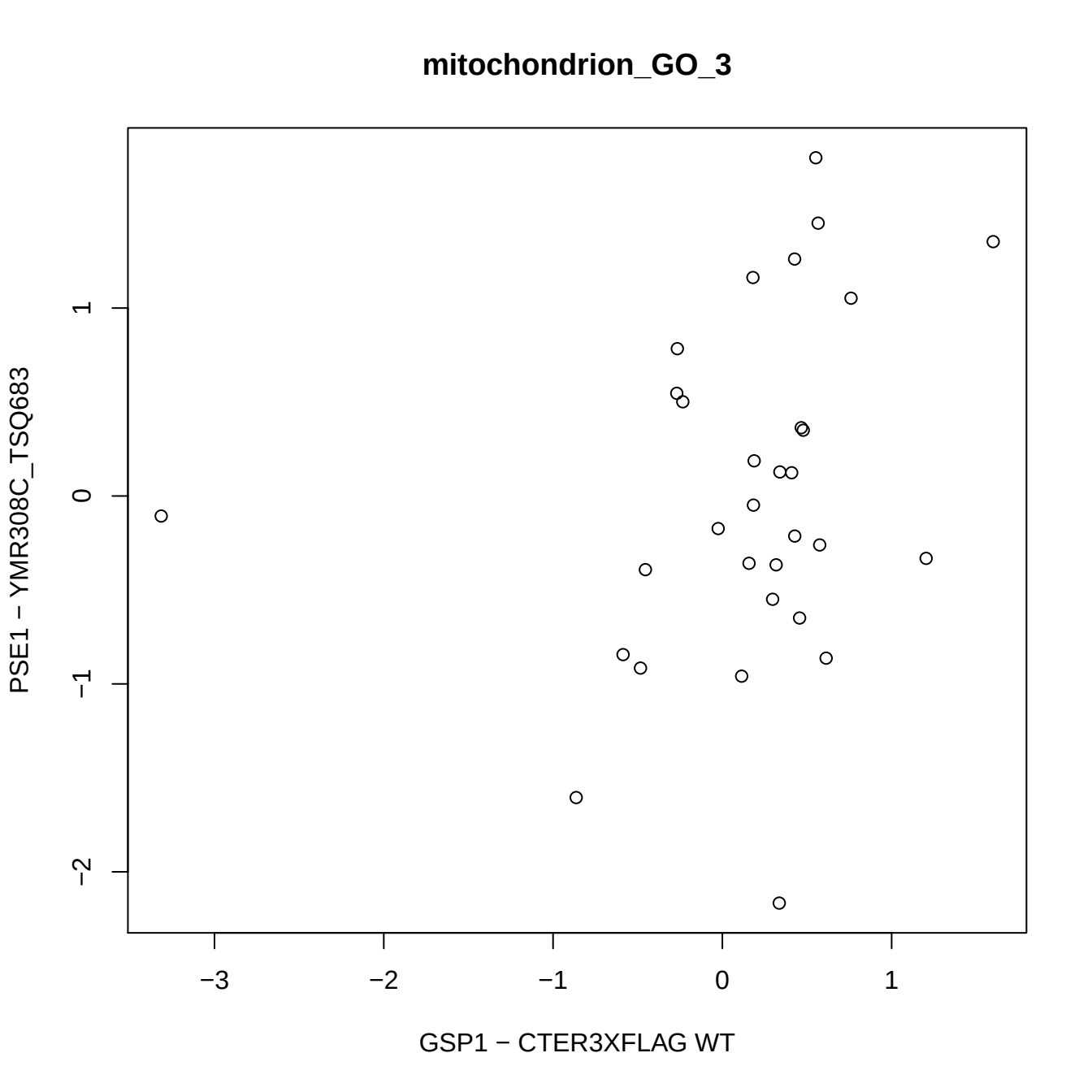

Golgi and ER_GO_3
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−3
−1.5
−1.0
−0.5
0.0
0.5
1.0

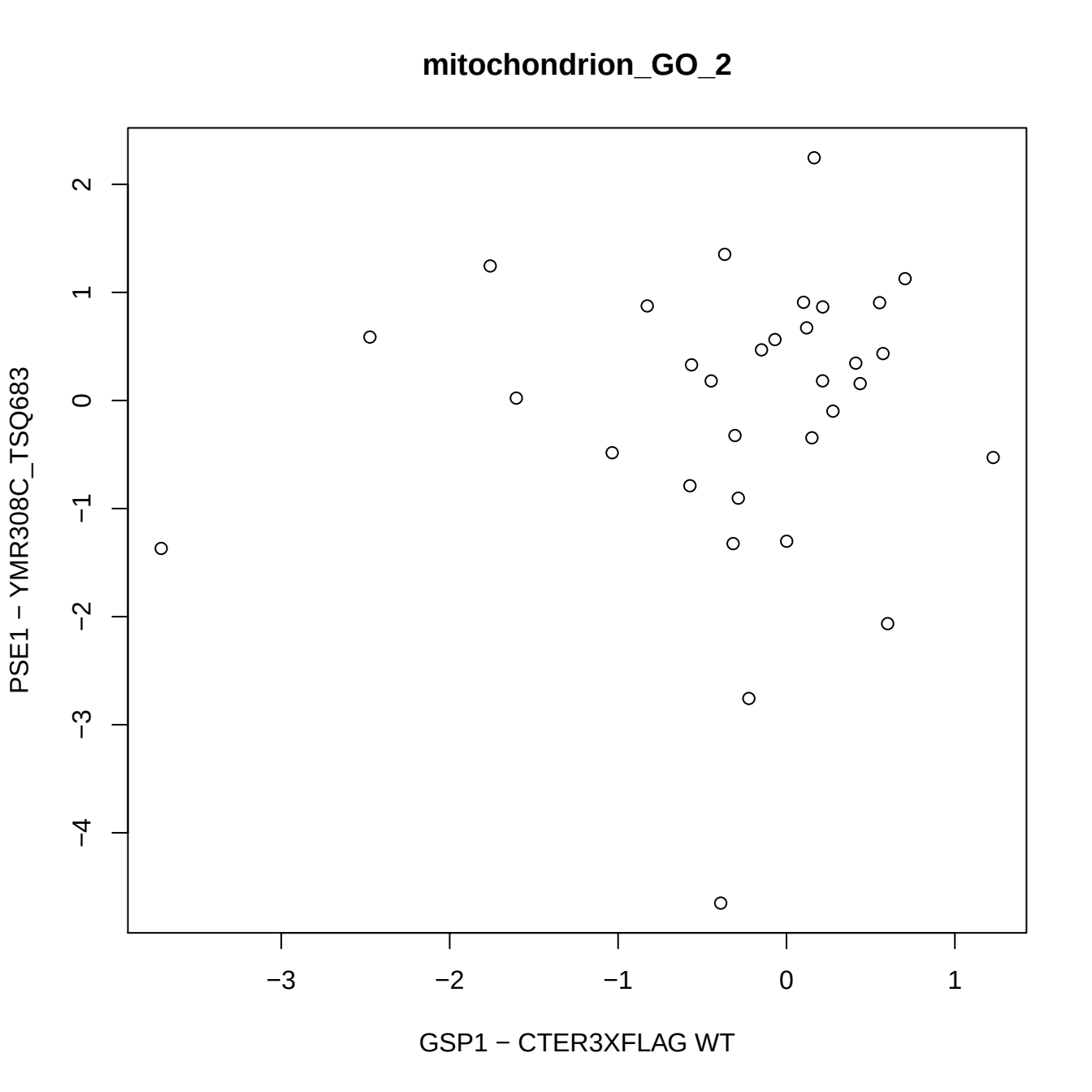

Golgi and ER_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
−2
−1
0
1
2
3
−2
−1
0
1
2

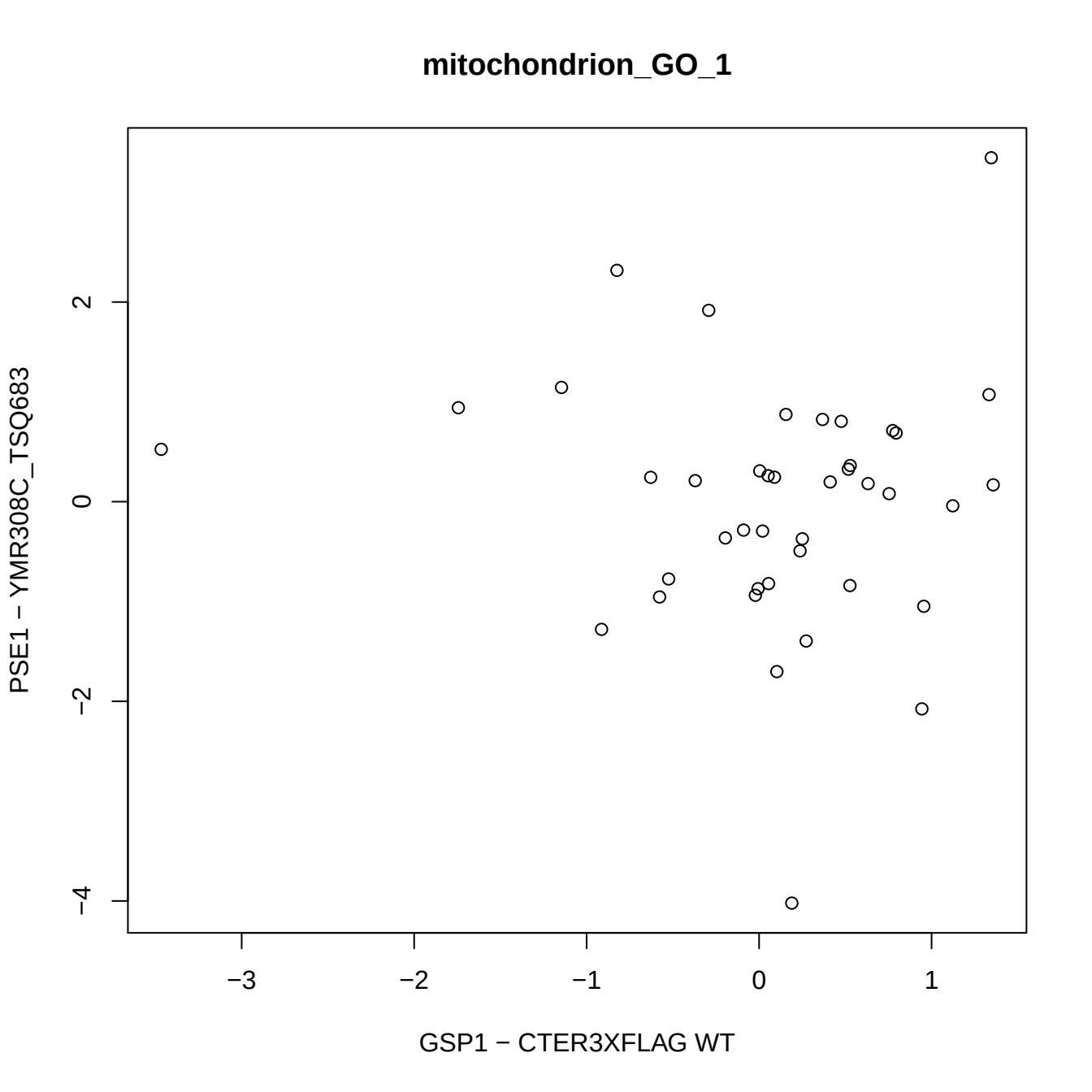

Golgi and ER_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
4
3
2
1
0
−1
−2
−1.0
−0.5
0.0
0.5
1.0
1.5

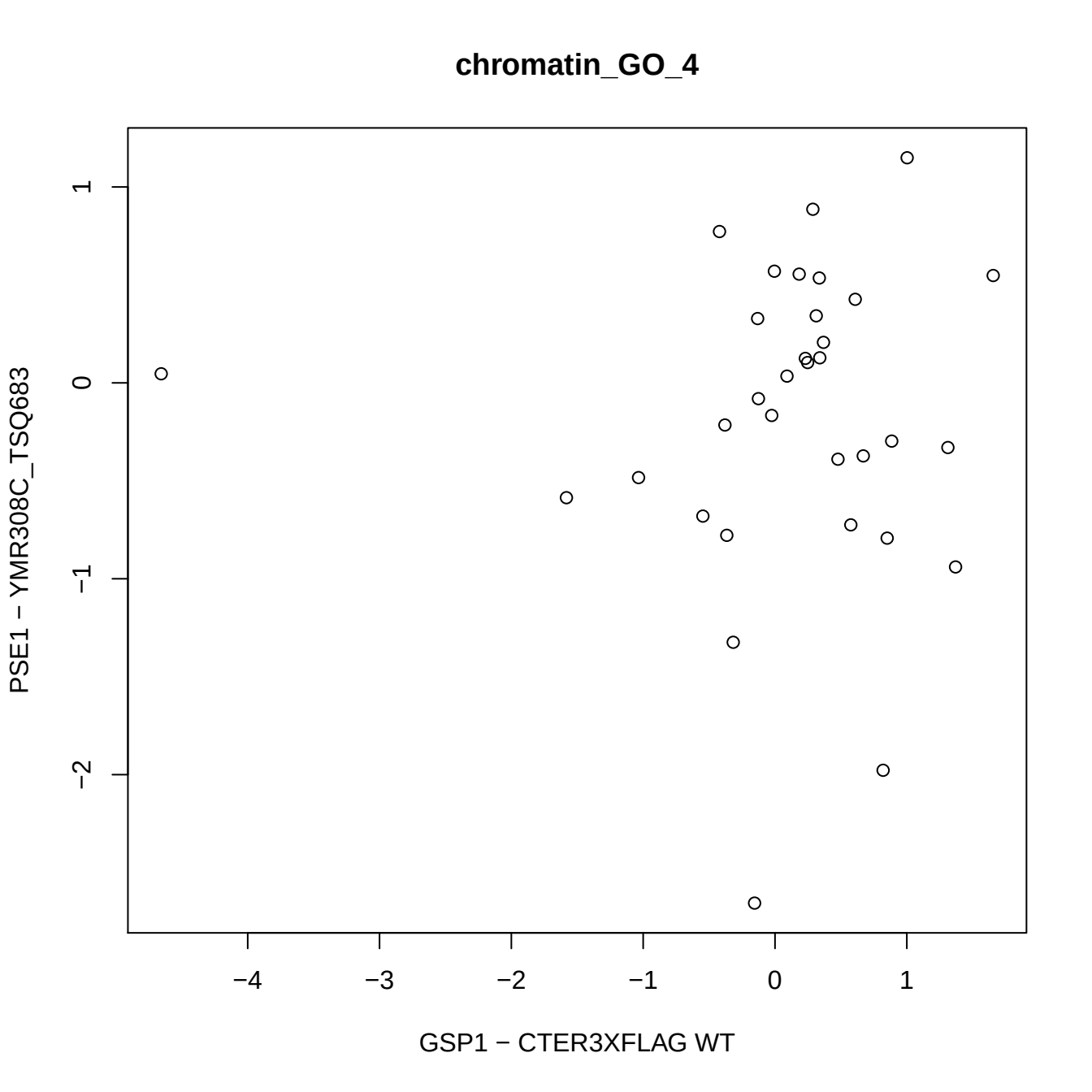

peroxisome_GO_0
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−3
−4
−2.0
−1.5
−1.0
−0.5
0.0
0.5
1.0

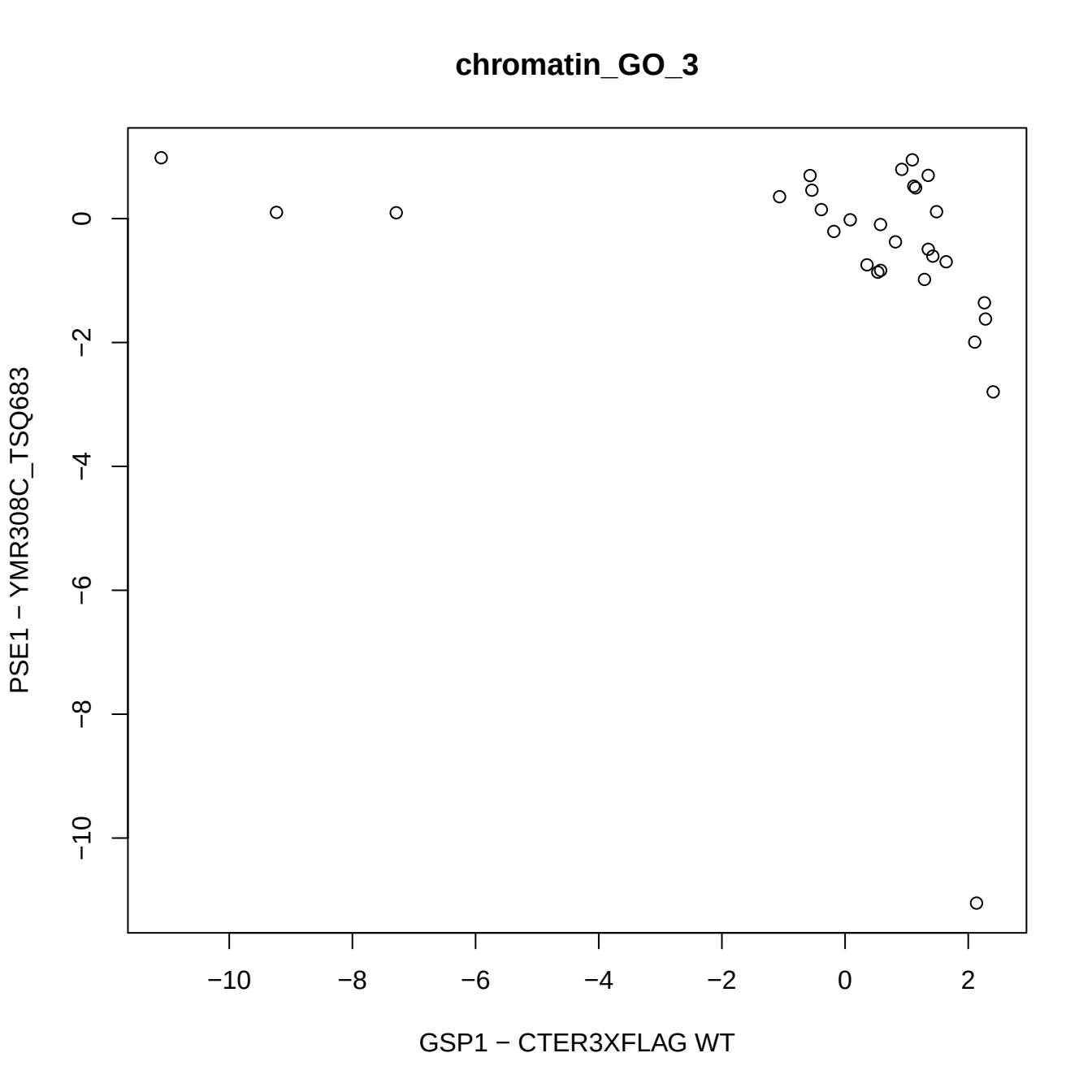

vacuole_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
−4
−3
−2
−1
0
1

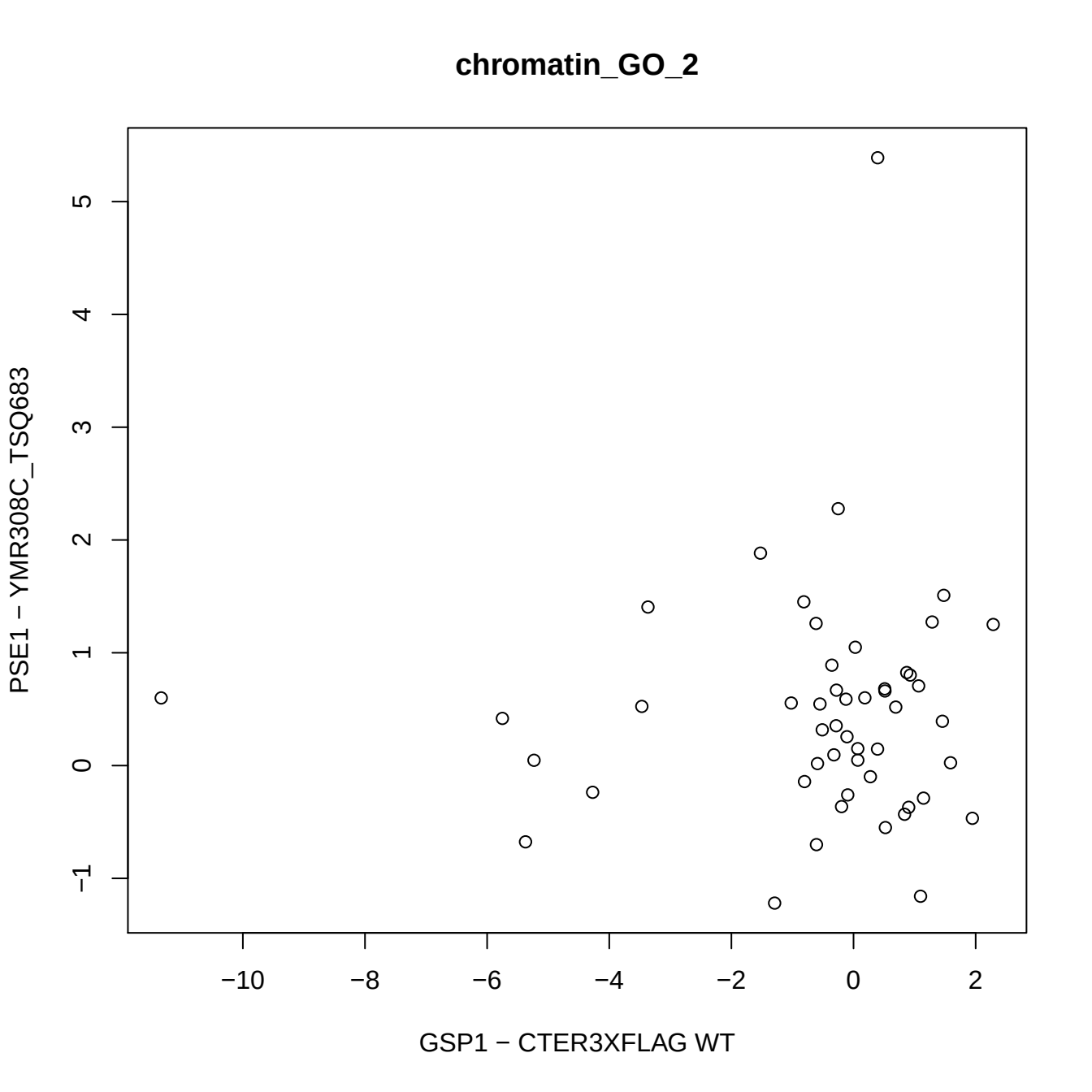

mitochondrion_GO_3
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
−3
−2
−1
0
1
−2
−1
0
1

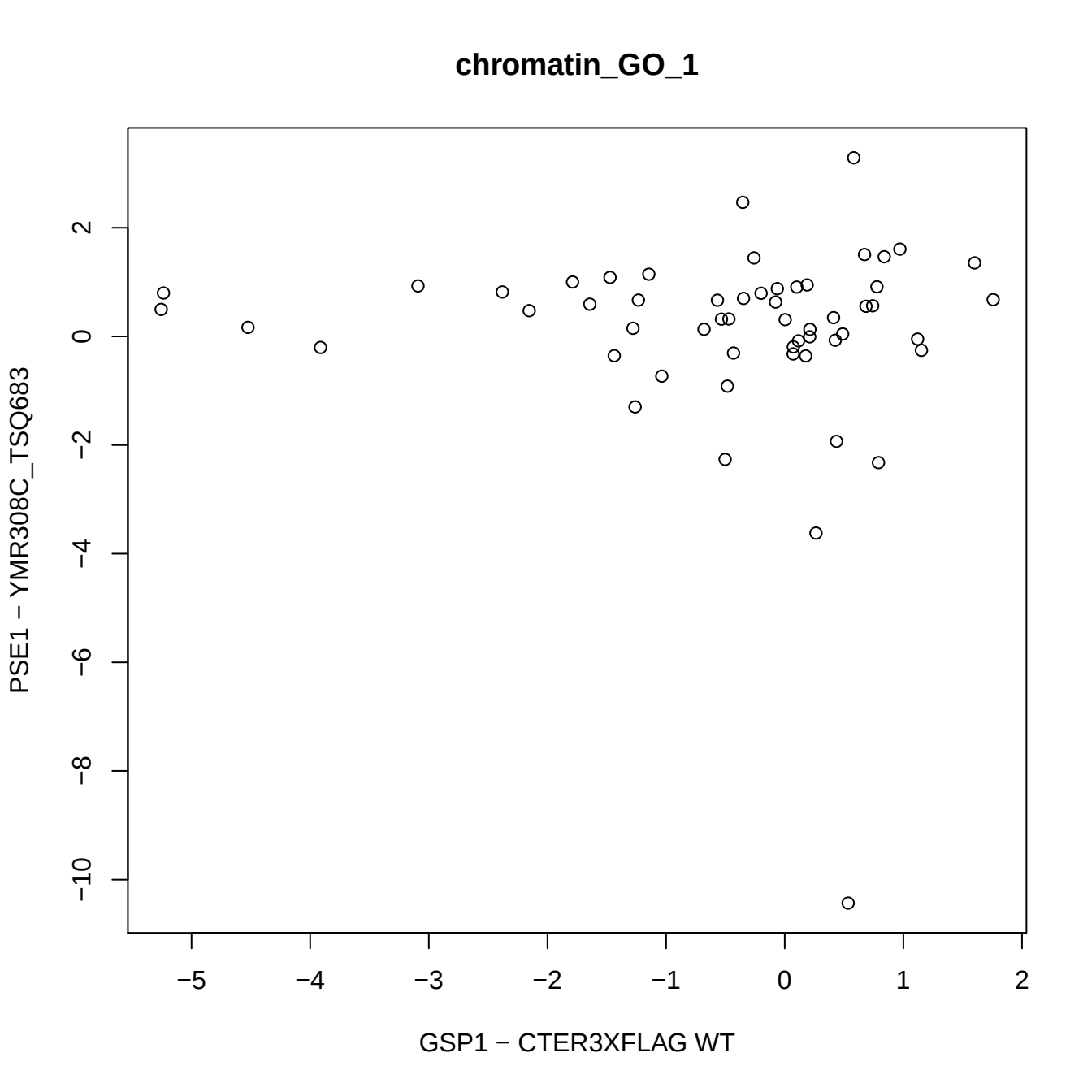

mitochondrion_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
1
0
−1
−2
−3
−4
−3
−2
−1
0
1

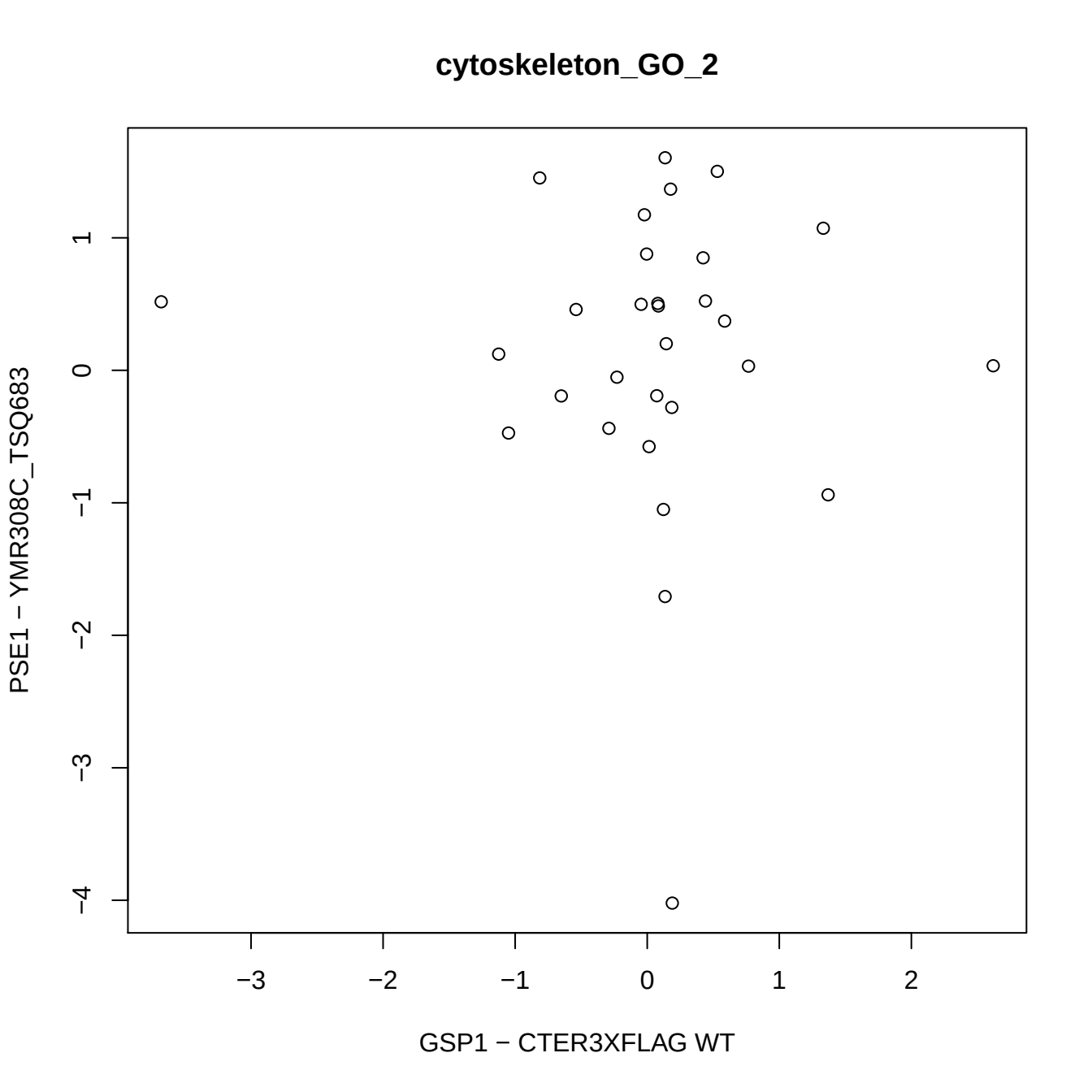

mitochondrion_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
0
−2
−4
−3
−2
−1
0
1

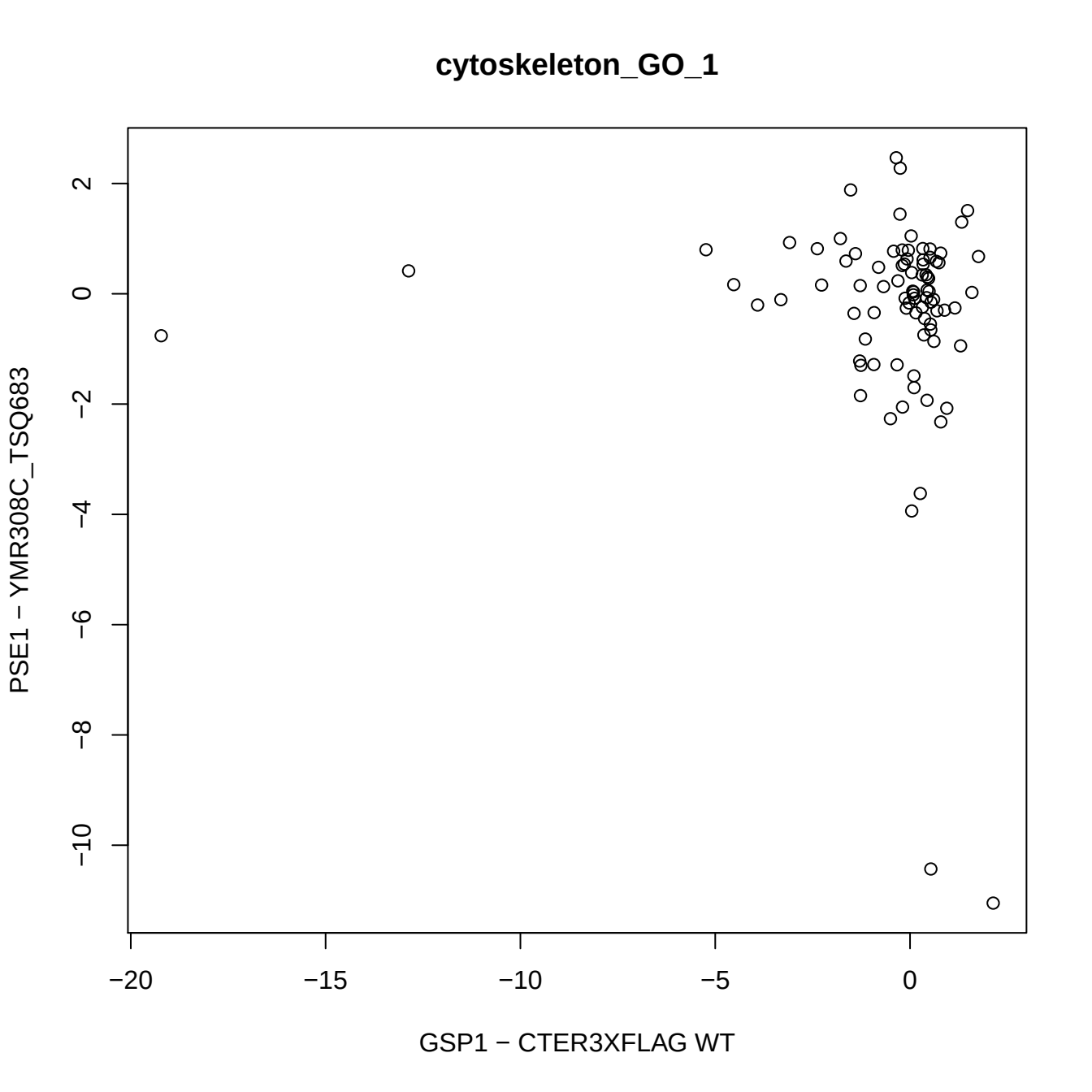

chromatin_GO_4
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−4
−3
−2
−1
0
1

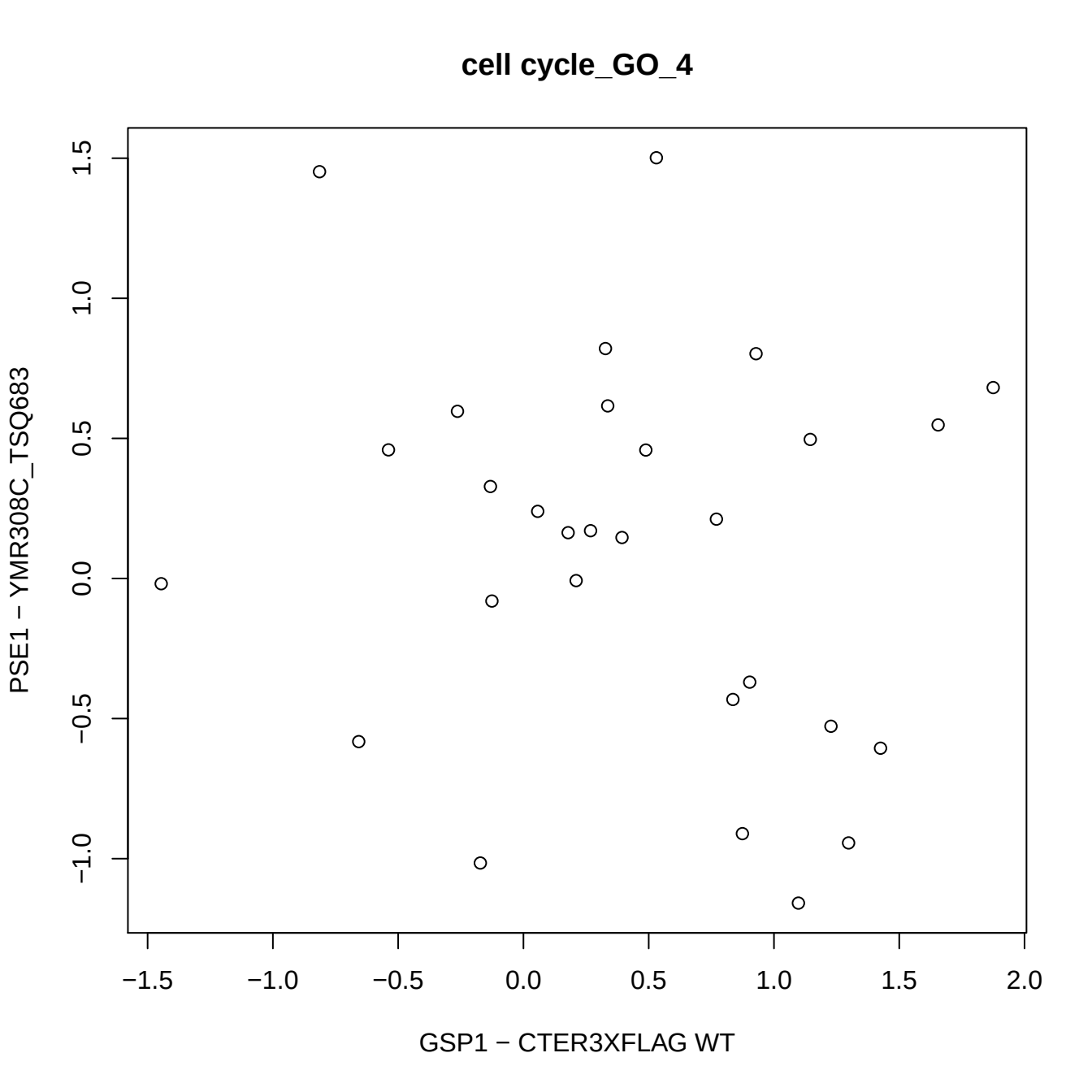

chromatin_GO_3
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
0
−2
−4
−6
−8
−10
−10
−8
−6
−4
−2
0
2

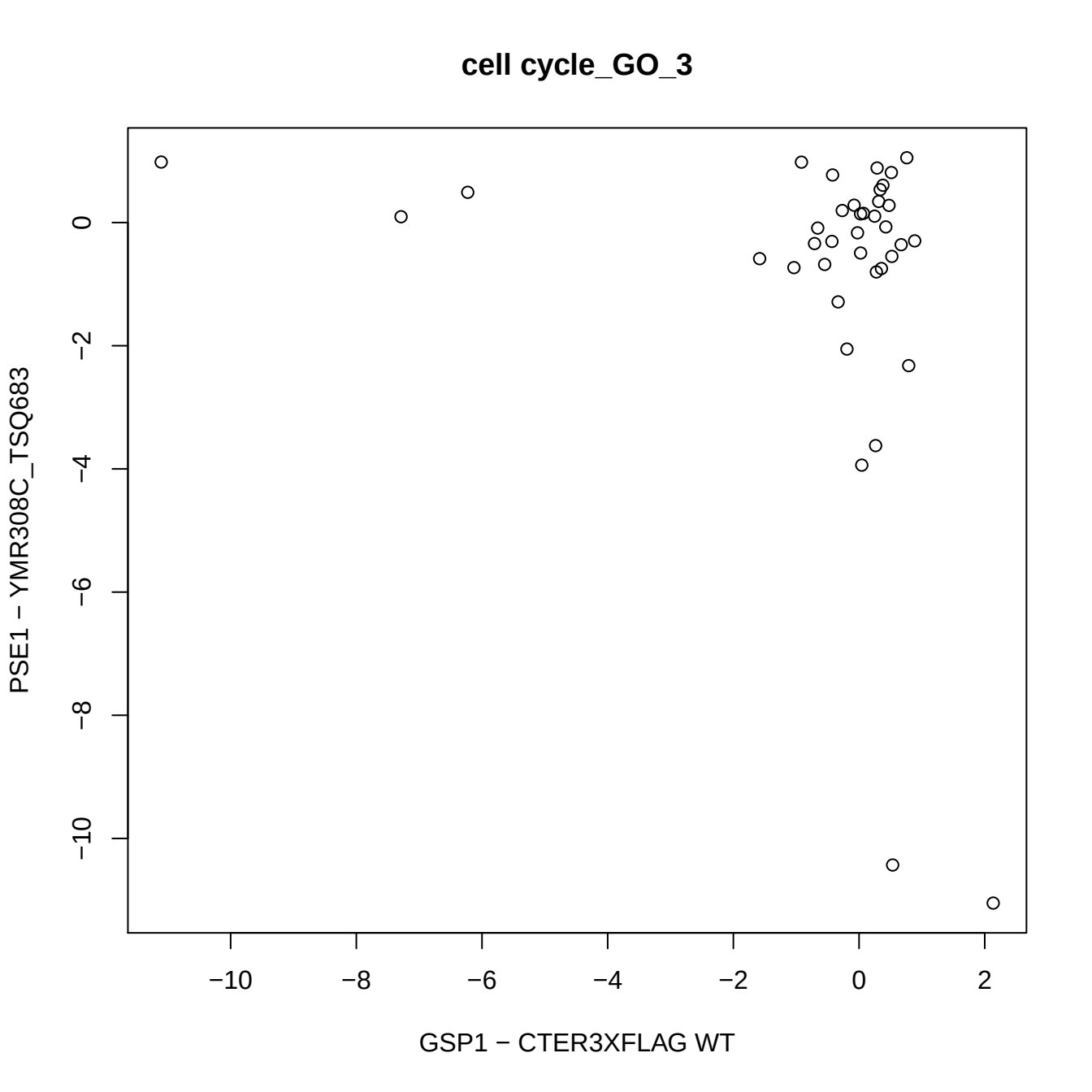

chromatin_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
−1
0
1
2
3
4
5
−10
−8
−6
−4
−2
0
2

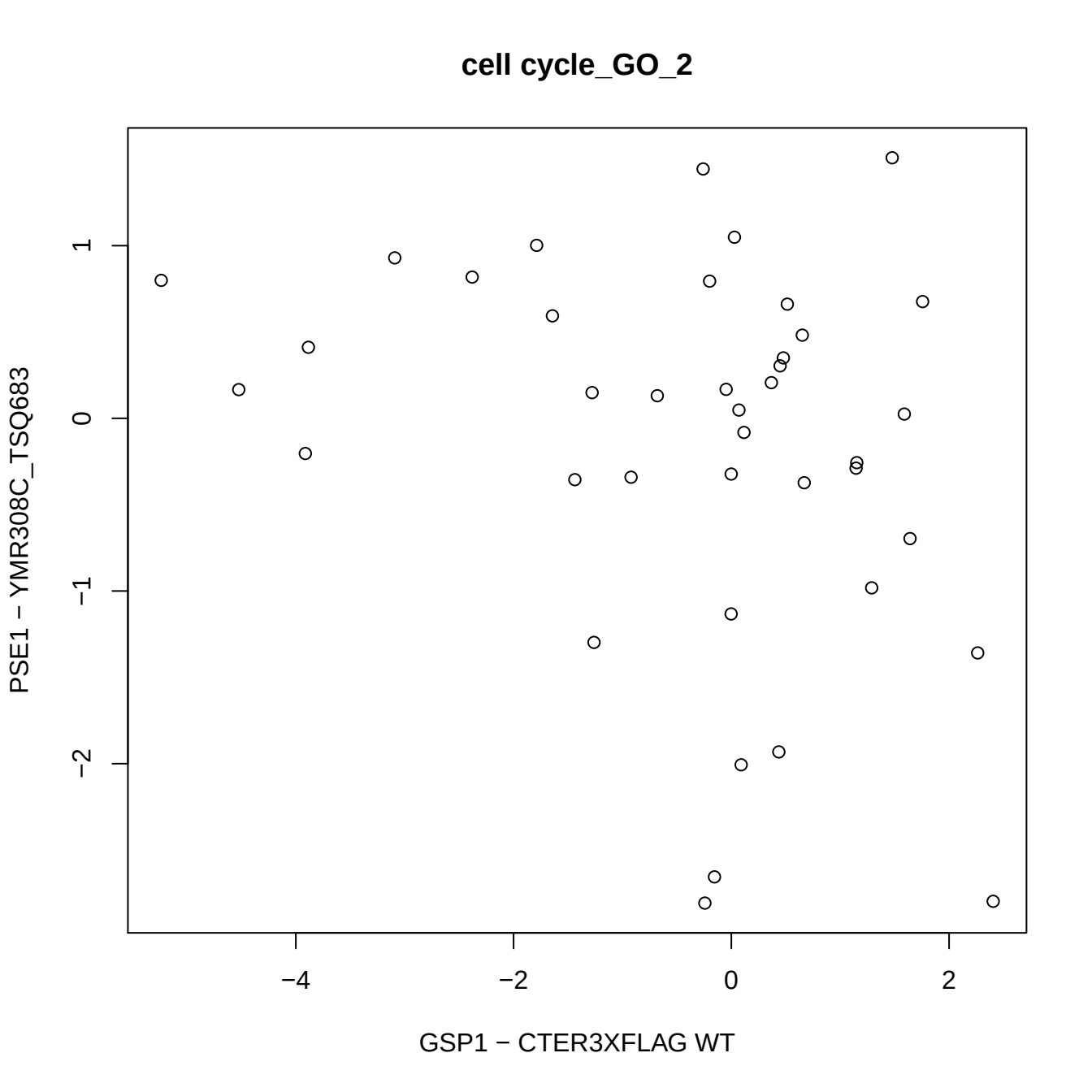

chromatin_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
0
−2
−4
−6
−8
−10
−5
−4
−3
−2
−1
0
1
2

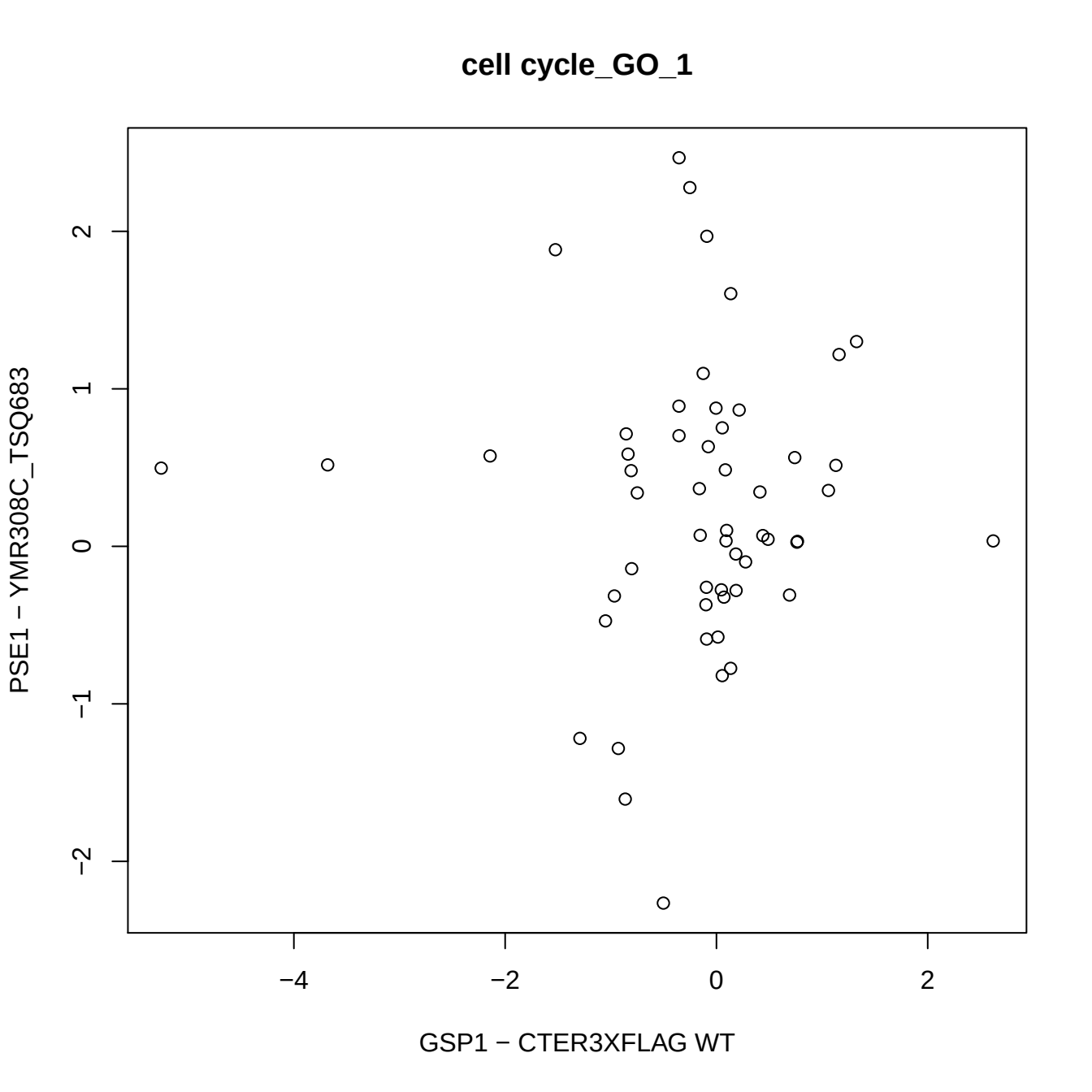

cytoskeleton_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−3
−4
−3
−2
−1
0
1
2

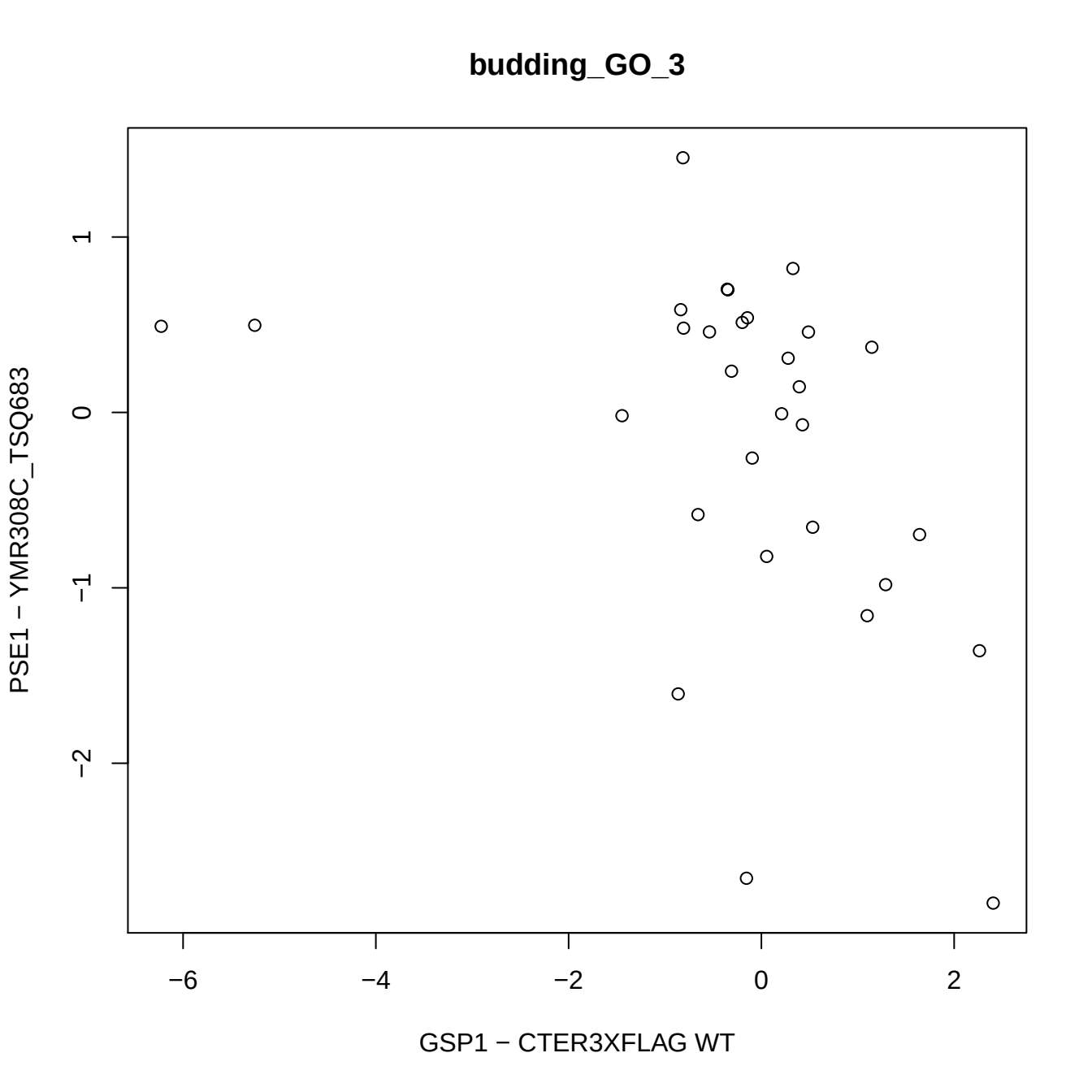

cytoskeleton_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
0
−2
−4
−6
−8
−10
−20
−15
−10
−5
0

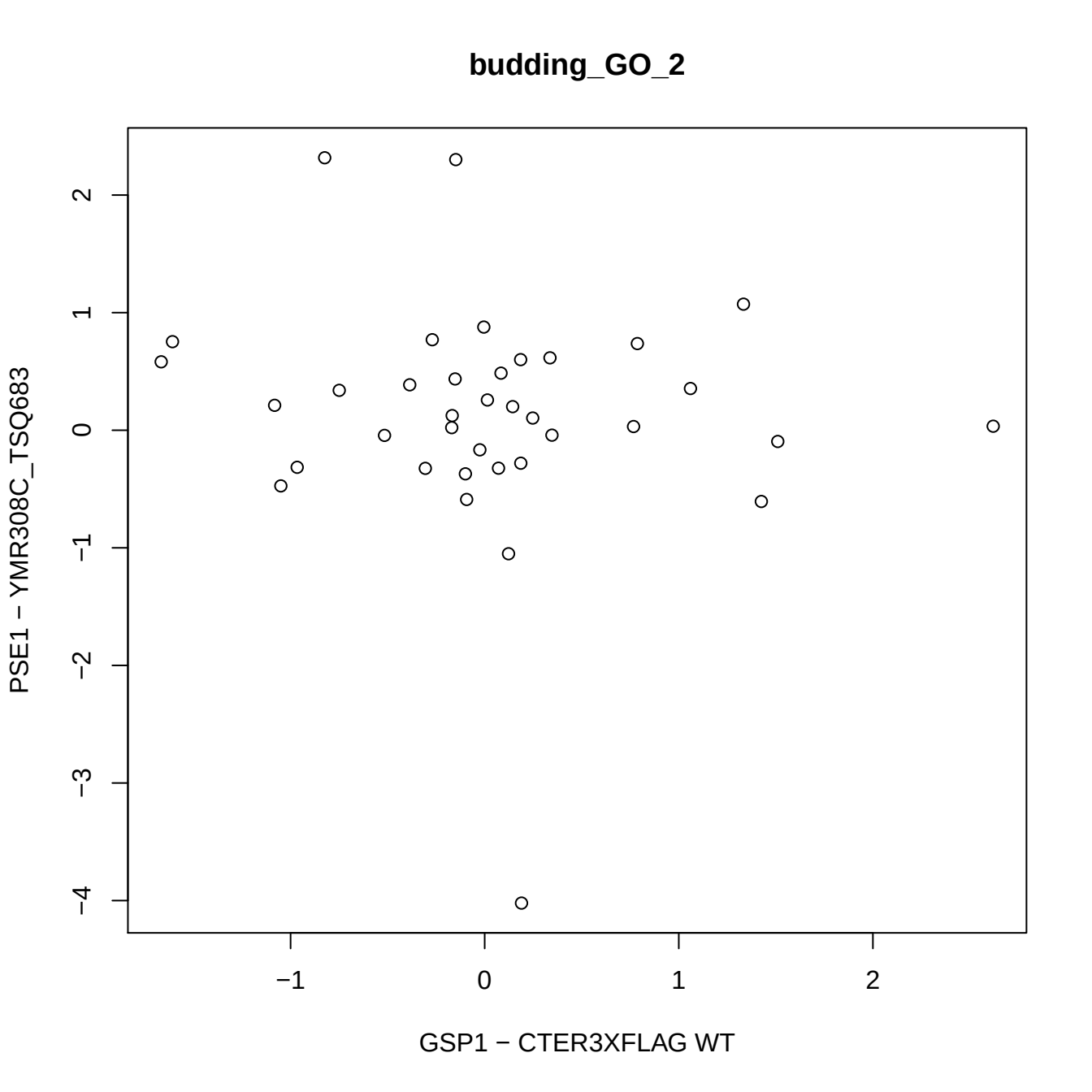

cell cycle_GO_4
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5
−1.0
−0.5
0.0
0.5
1.0
1.5
2.0

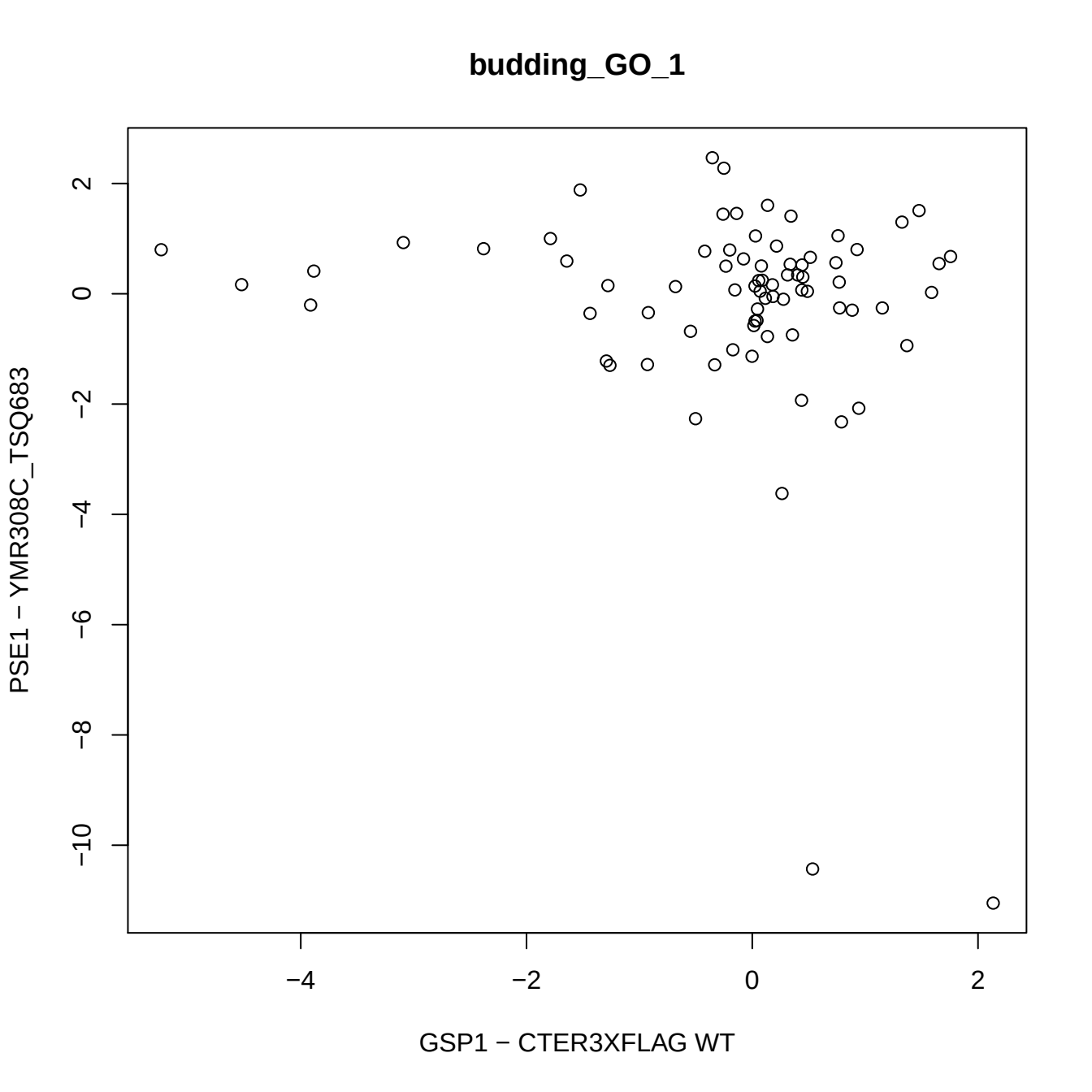

cell cycle_GO_3
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
0
−2
−4
−6
−8
−10
−10
−8
−6
−4
−2
0
2

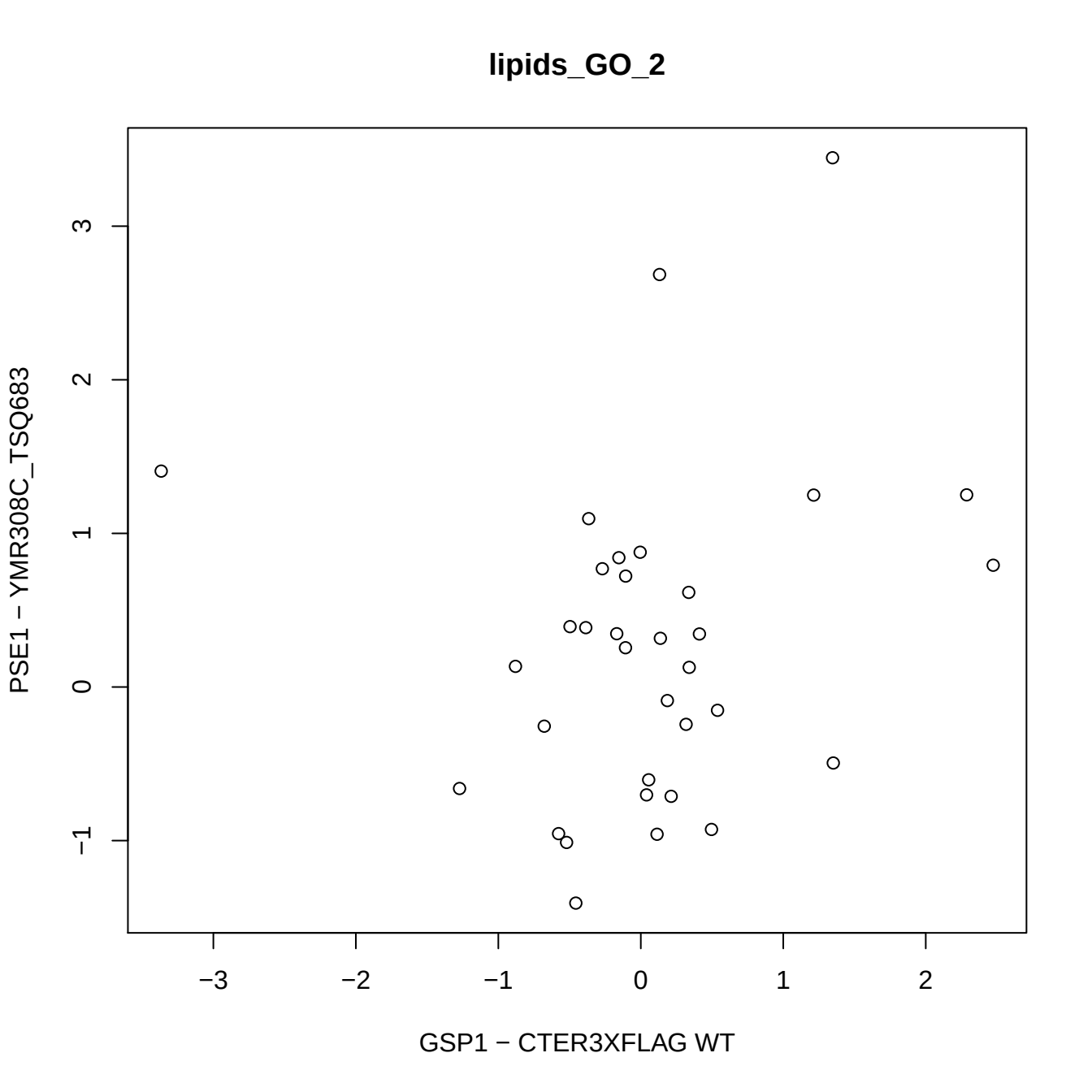

cell cycle_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−4
−2
0
2

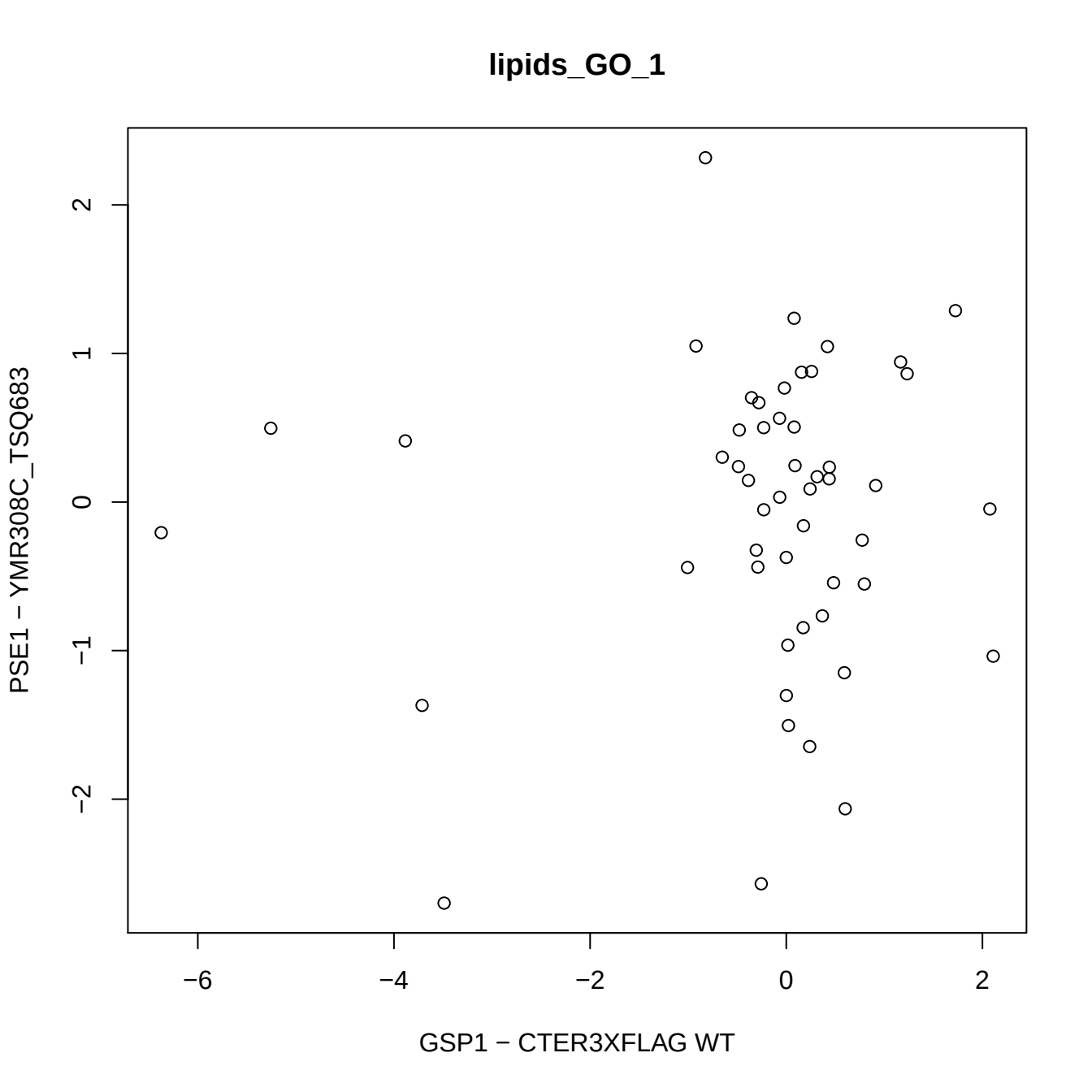

cell cycle_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
1
0
−1
−2
−4
−2
0
2

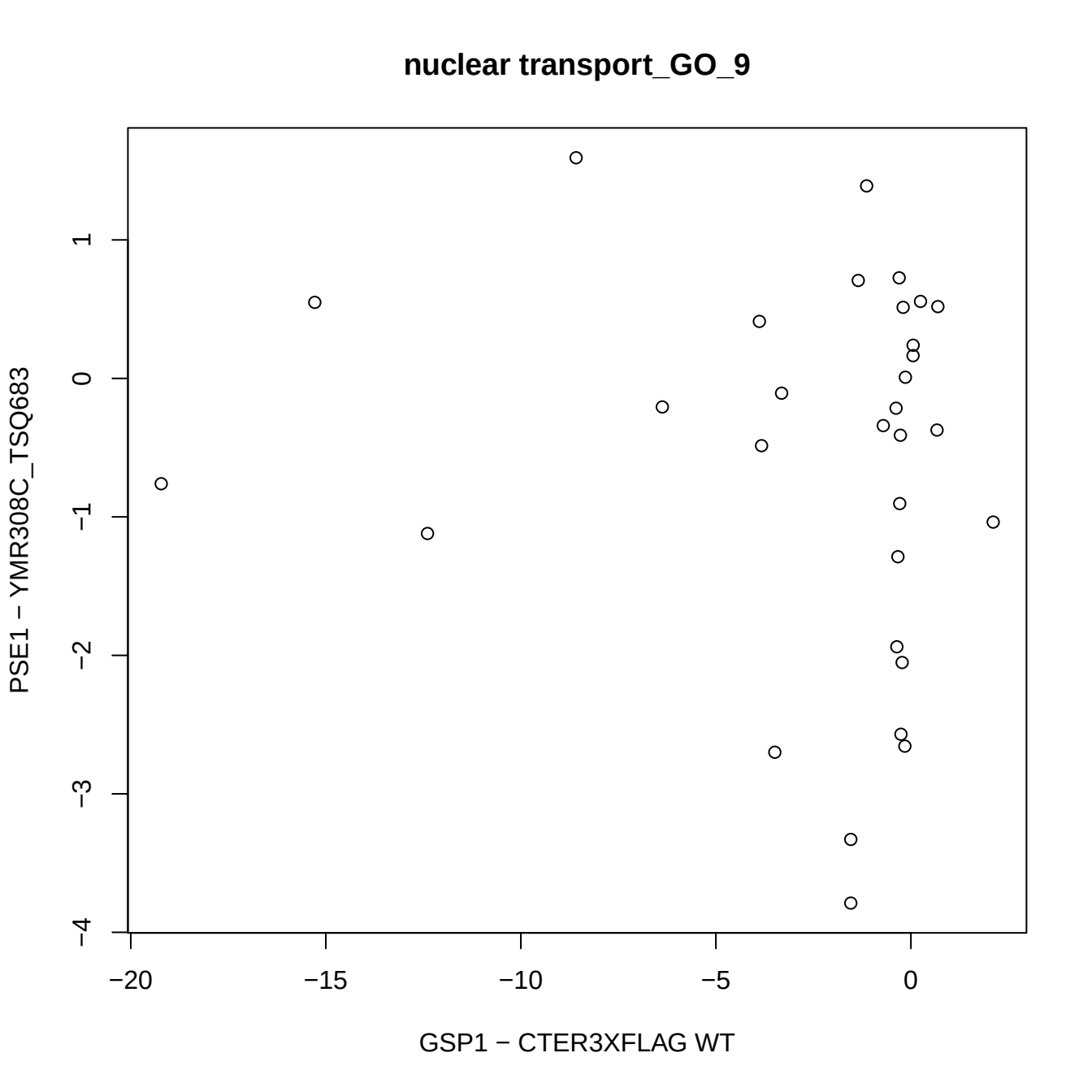

budding_GO_3
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−6
−4
−2
0
2

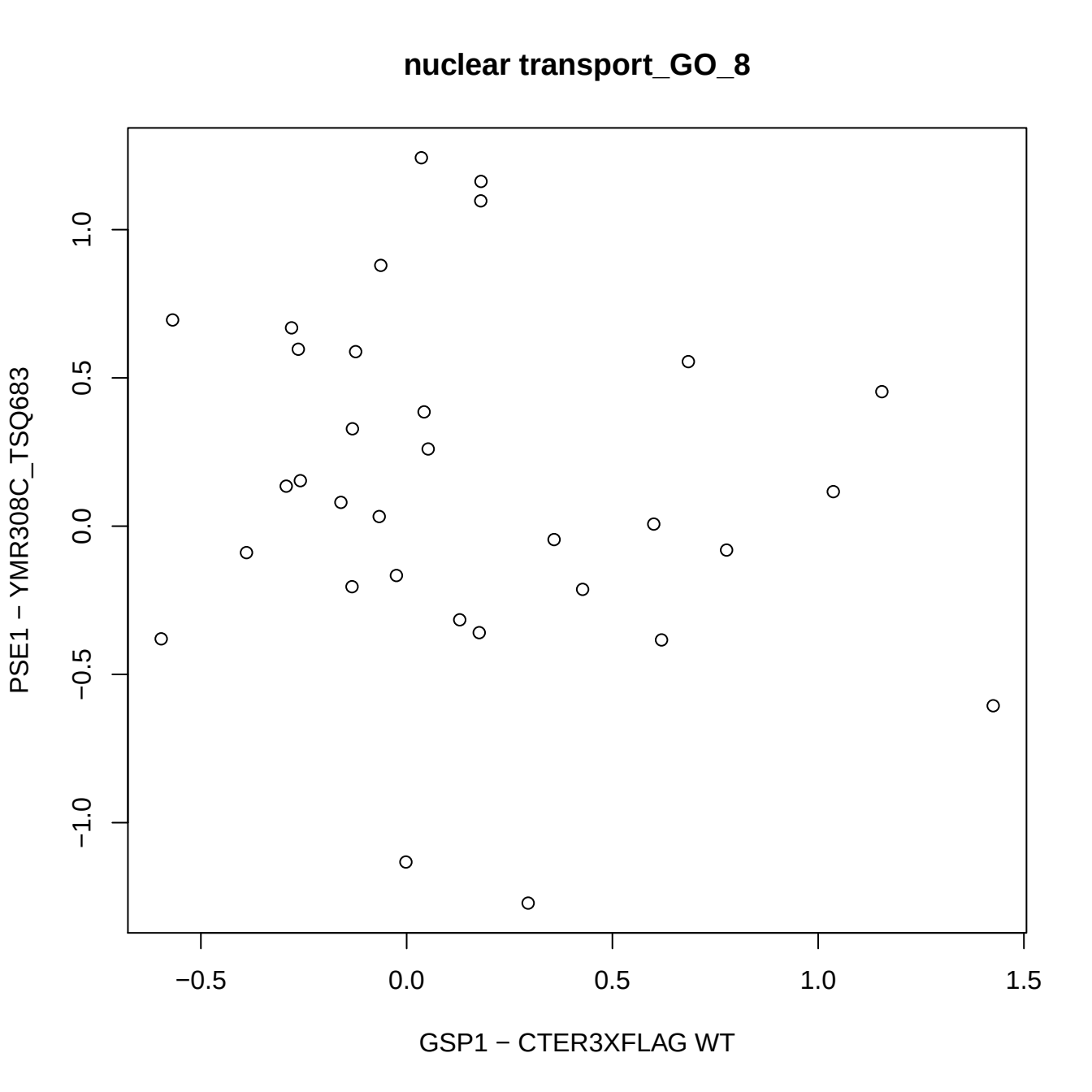

budding_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
1
0
−1
−2
−3
−4
−1
0
1
2

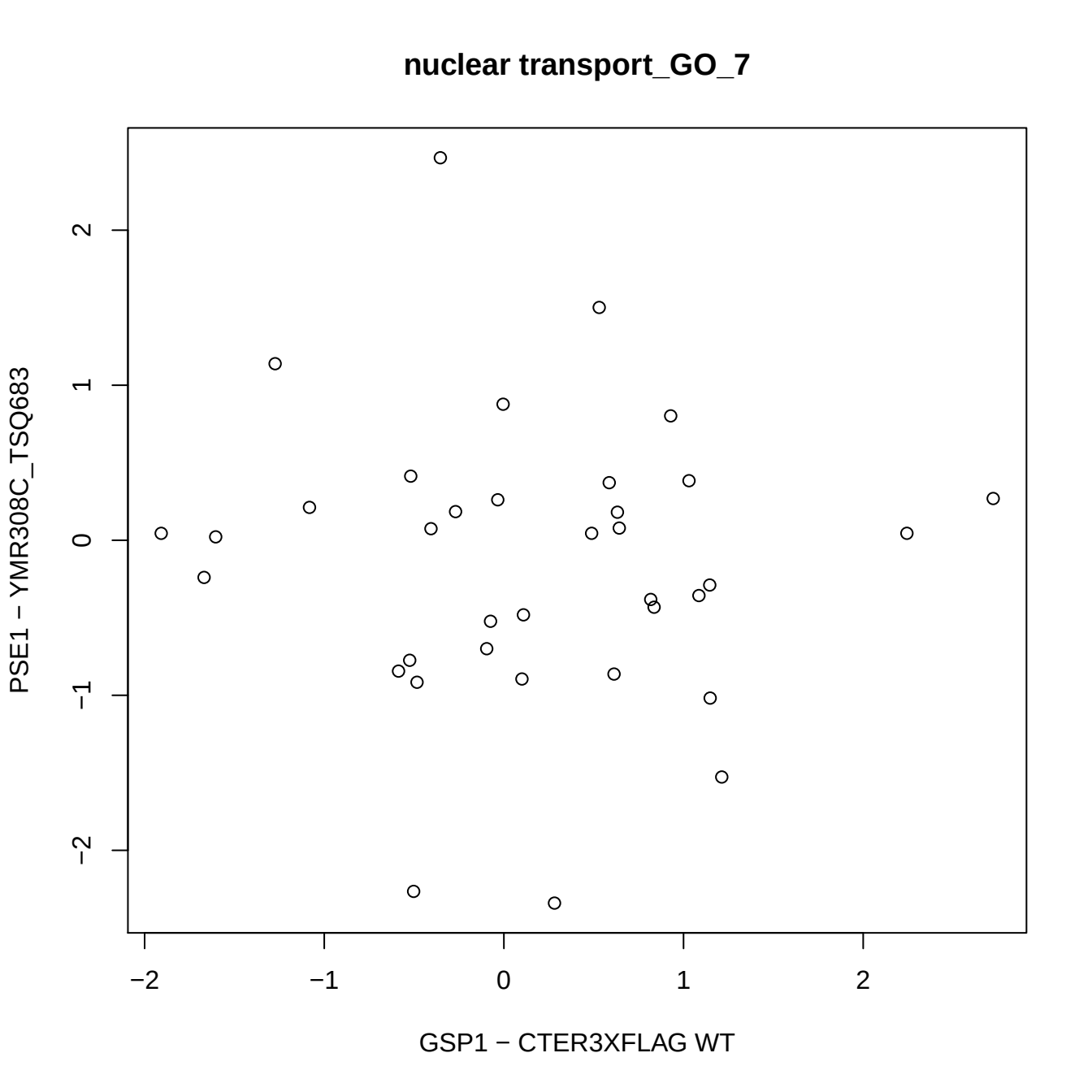

budding_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
0
−2
−4
−6
−8
−10
−4
−2
0
2

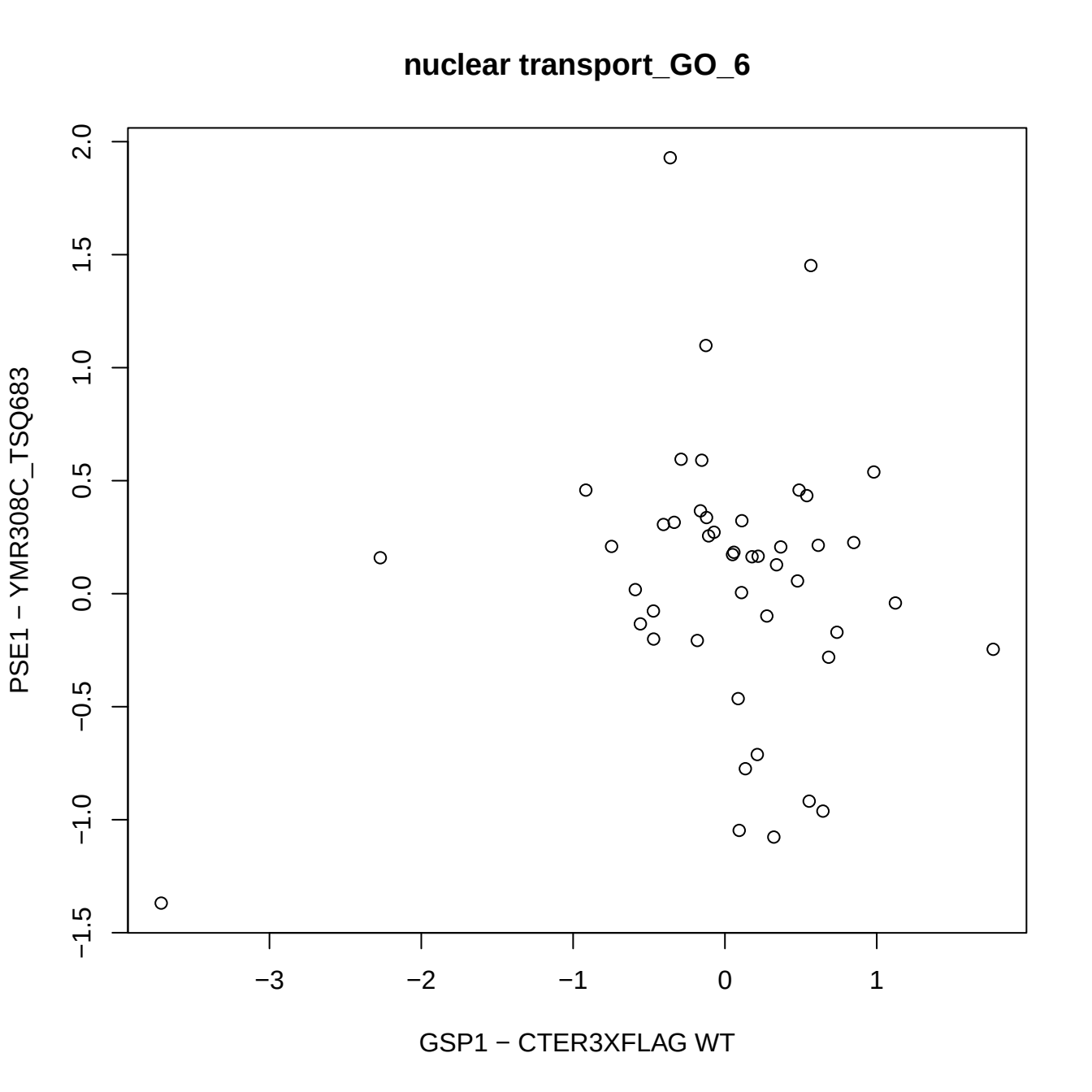

lipids_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
3
2
1
0
−1
−3
−2
−1
0
1
2

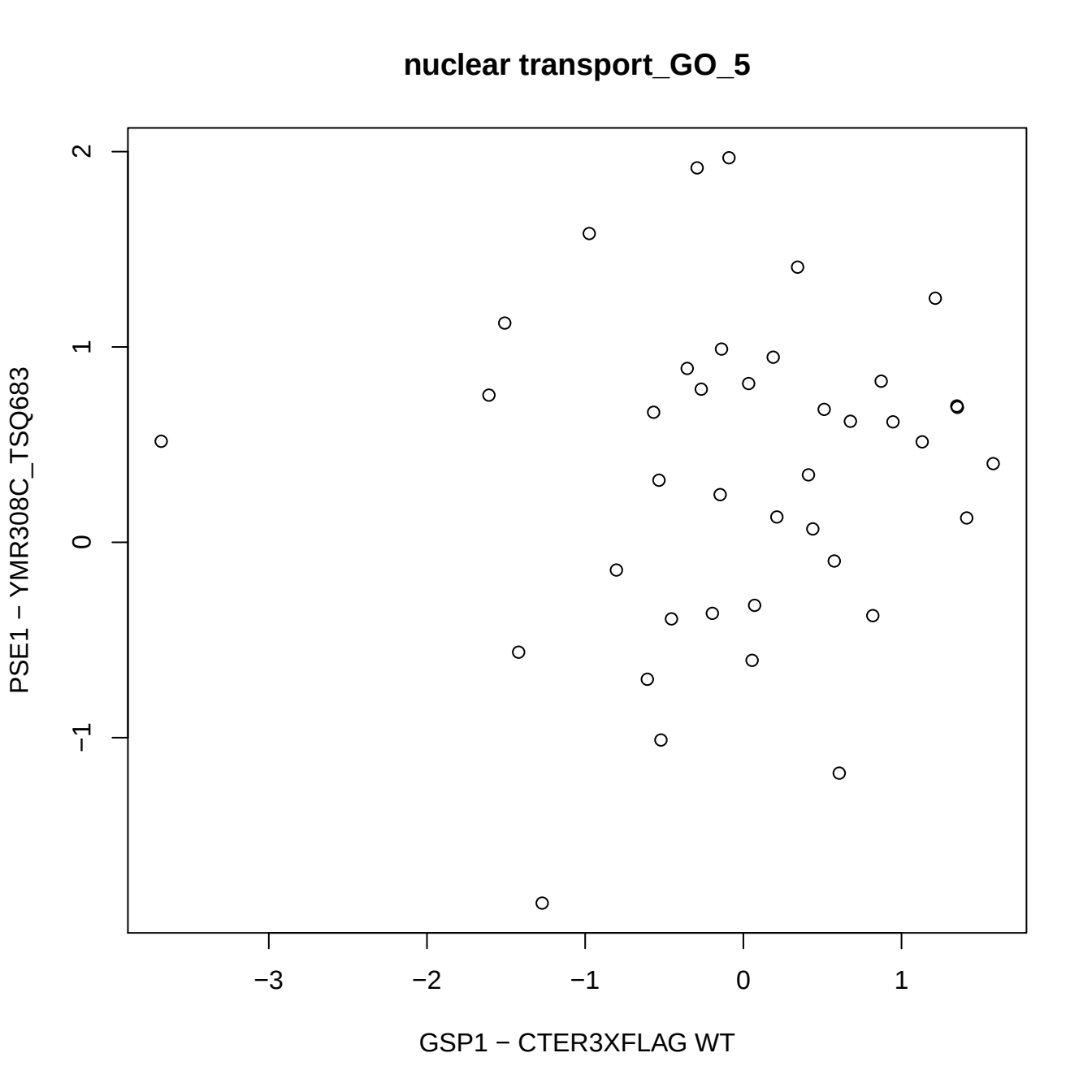

lipids_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
1
0
−1
−2
−6
−4
−2
0
2

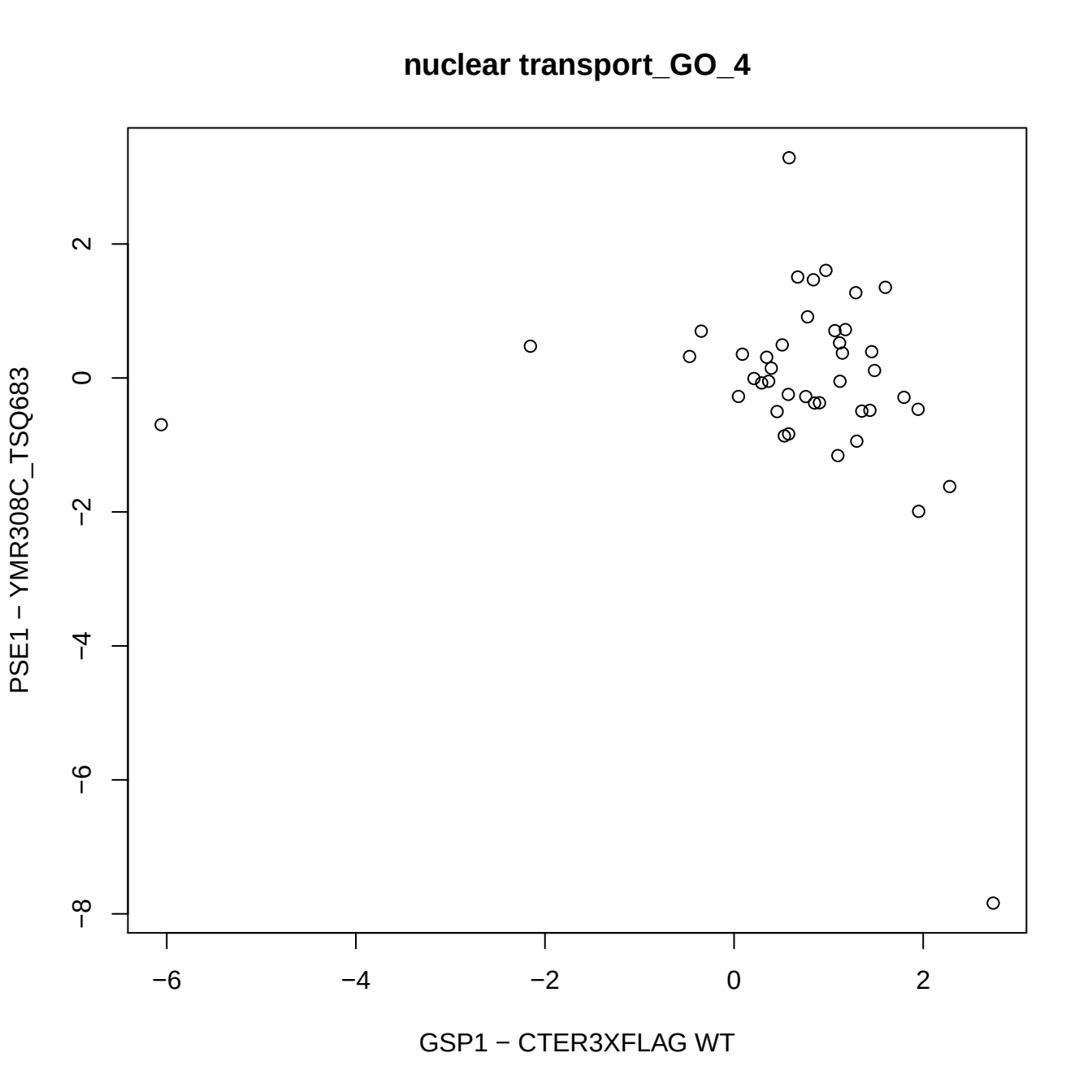

nuclear transport_GO_9
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−3
−4
−20
−15
−10
−5
0

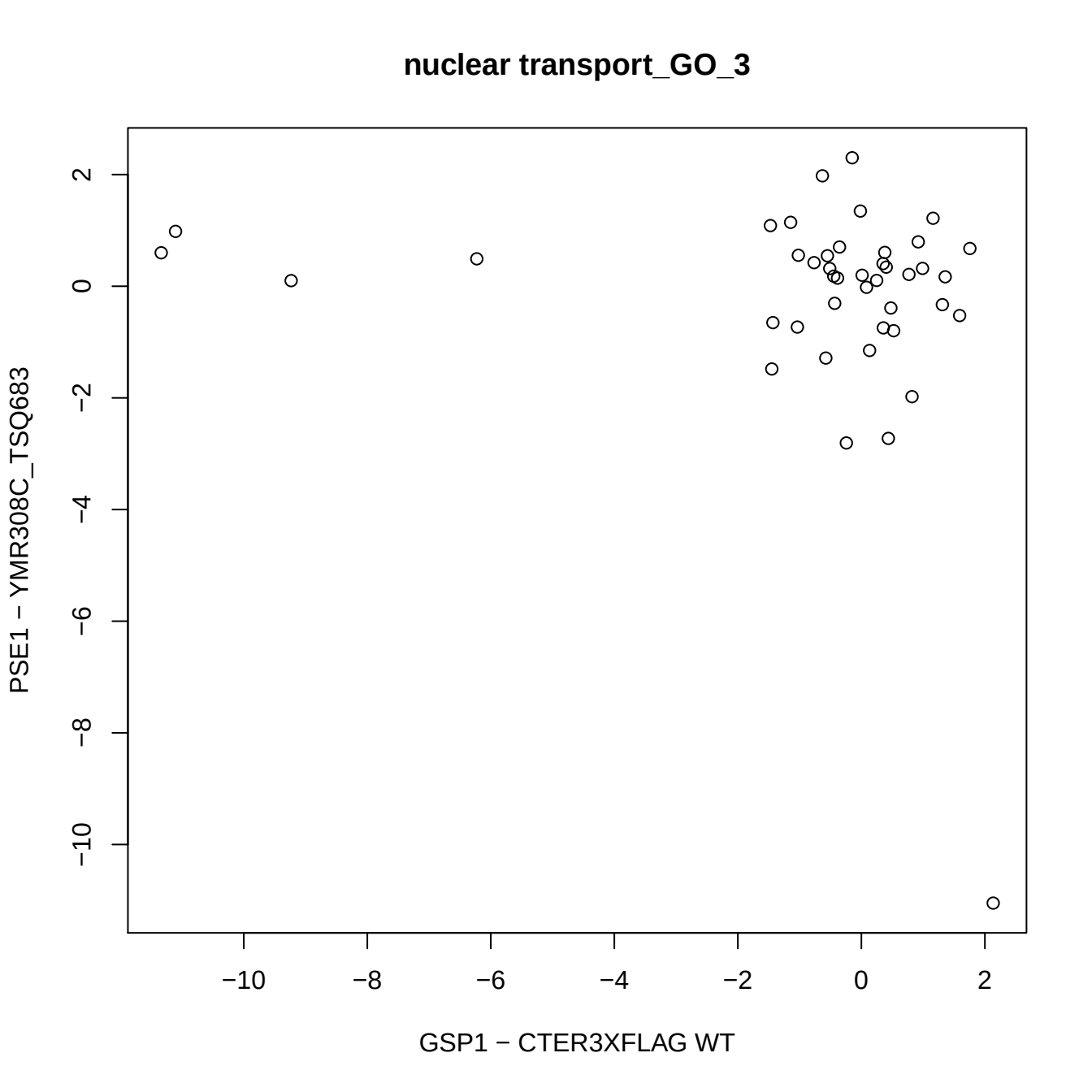

nuclear transport_GO_8
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
−1.0
−0.5
0.0
0.5
1.0
−0.5
0.0
0.5
1.0
1.5

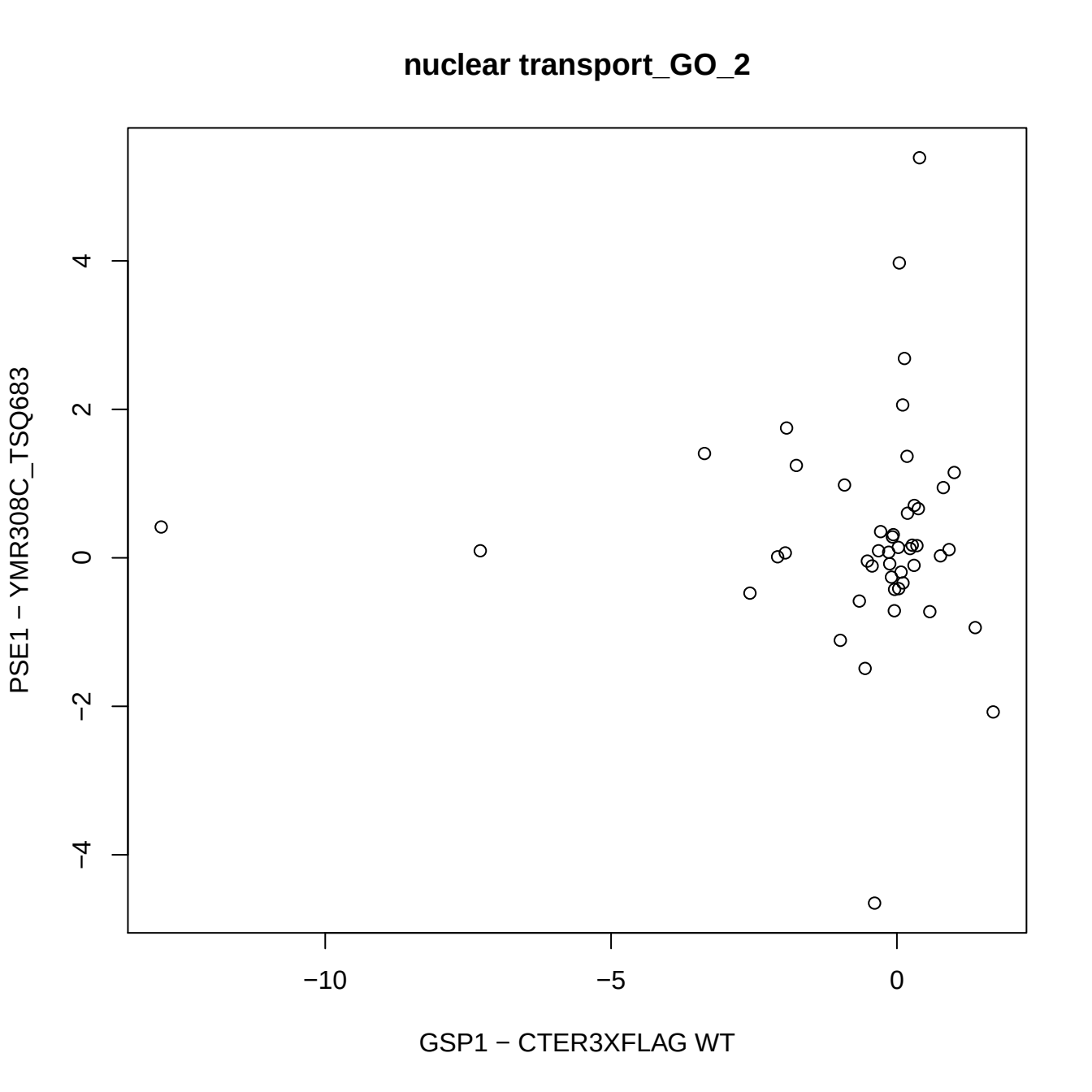

nuclear transport_GO_7
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
−2
−1
0
1
2

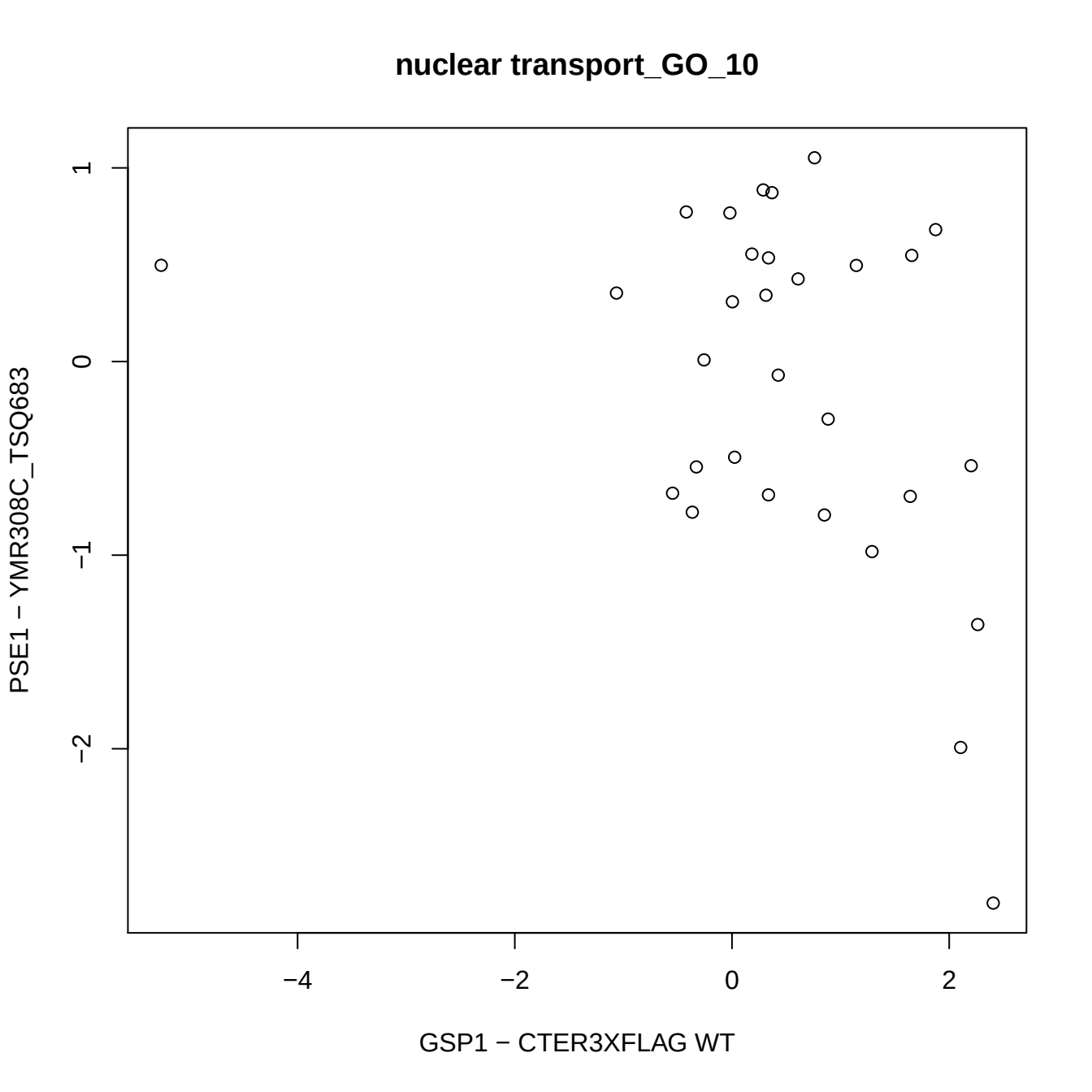

nuclear transport_GO_6
PSE1 – YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2.0
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5
−3
−2
−1
0
1

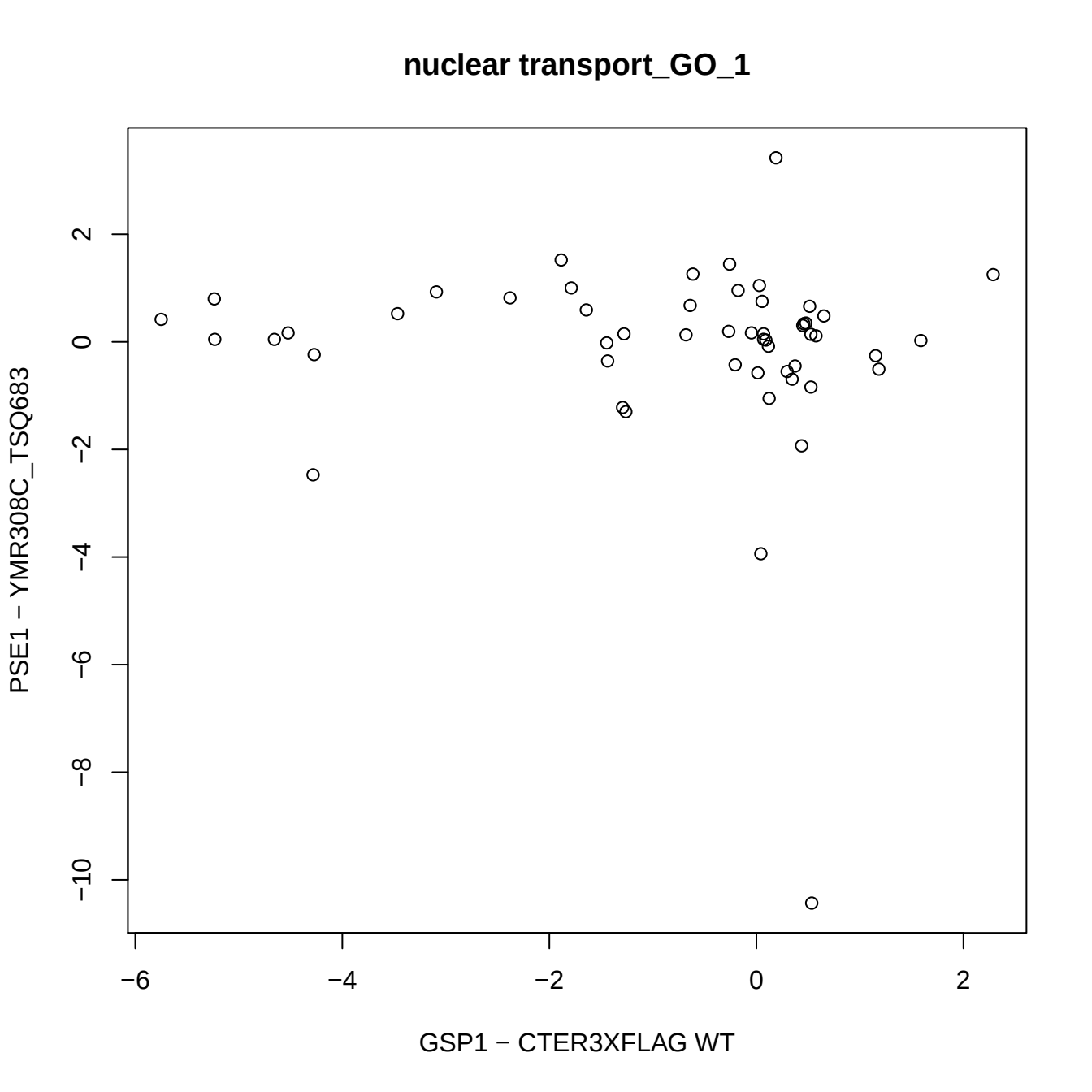

nuclear transport_GO_5
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
1
0
−1
−3
−2
−1
0
1

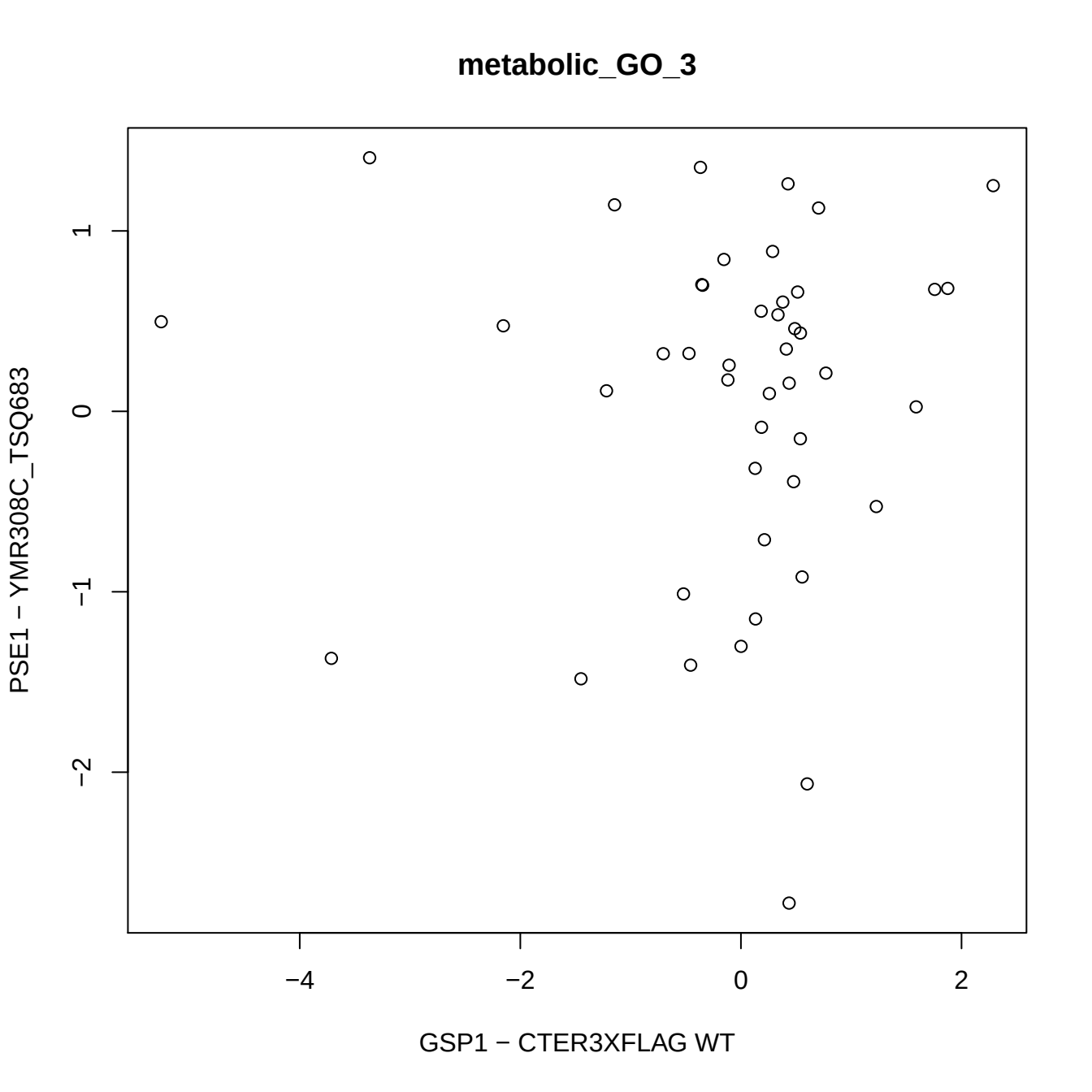

nuclear transport_GO_4
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
0
−2
−4
−6
−8
−6
−4
−2
0
2

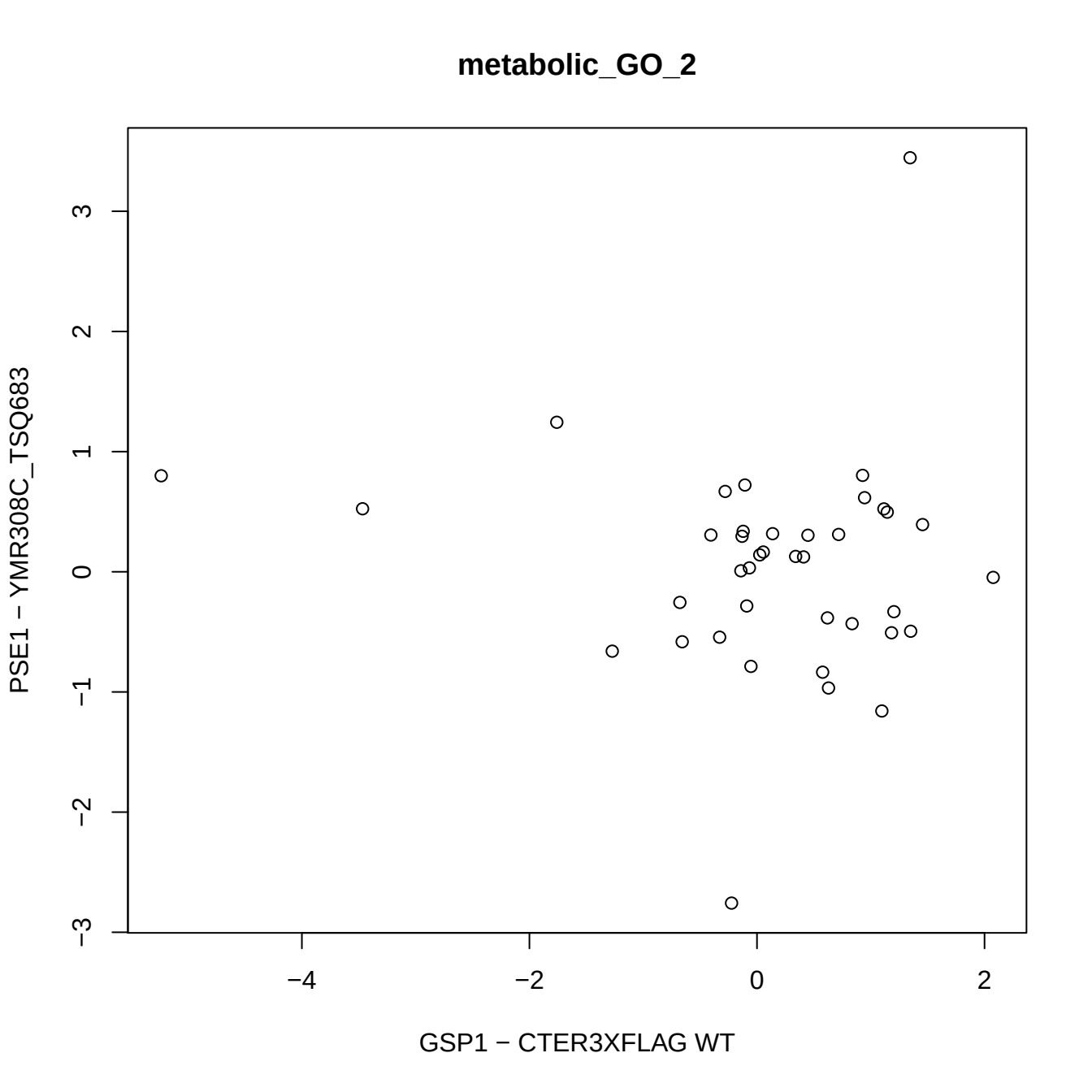

nuclear transport_GO_3
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
0
−2
−4
−6
−8
−10
−10
−8
−6
−4
−2
0
2

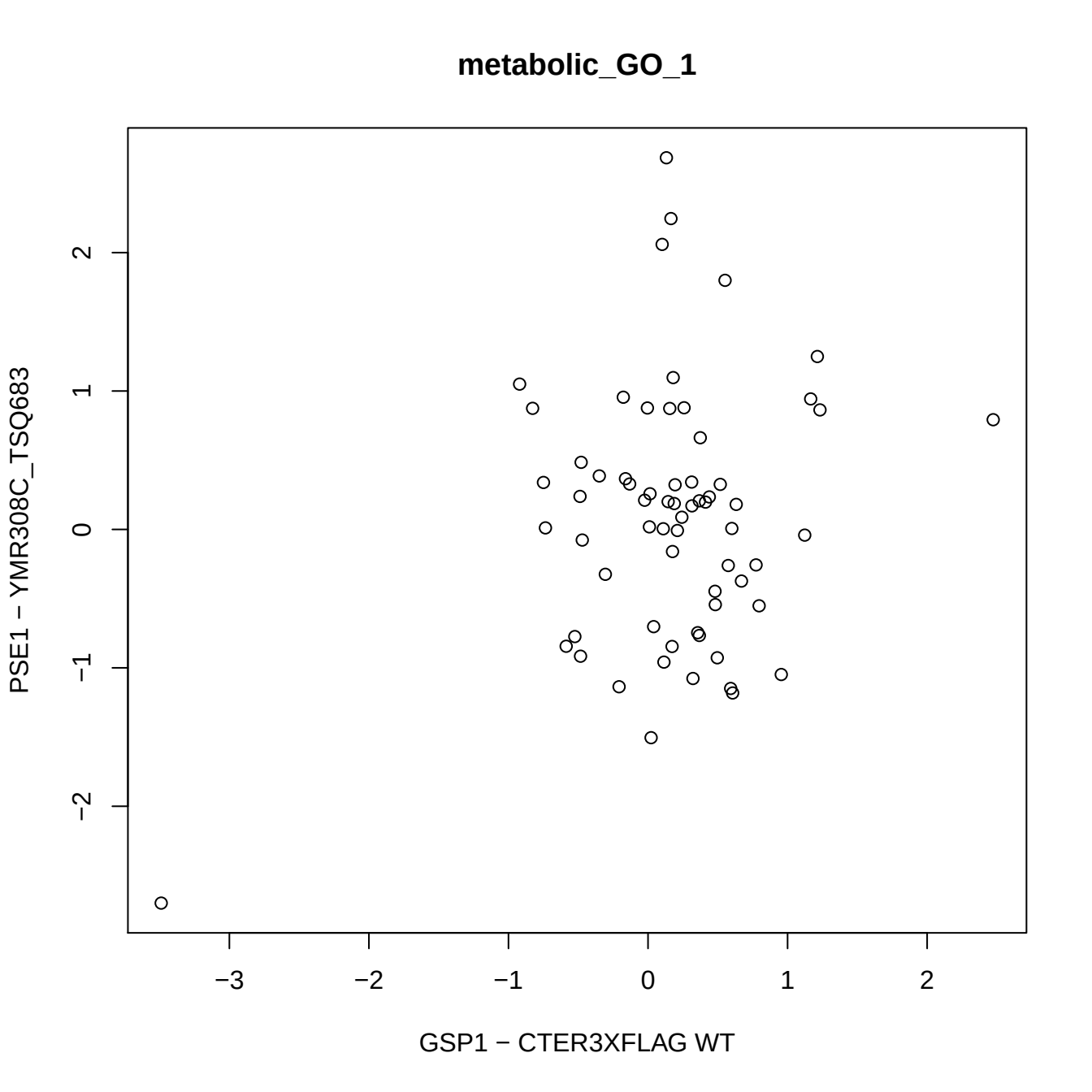

nuclear transport_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
4
2
0
−2
−4
−10
−5
0


nuclear transport_GO_10
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−4
−2
0
2


nuclear transport_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
2
0
−2
−4
−6
−8
−10
−6
−4
−2
0
2


metabolic_GO_3
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
1
0
−1
−2
−4
−2
0
2


metabolic_GO_2
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
3
2
1
0
−1
−2
−3
−4
−2
0
2


metabolic_GO_1
PSE1 − YMR308C_TSQ683
GSP1 − CTER3XFLAG WT
−3
−2
−1
0
1
2
−2
−1
0
1
2