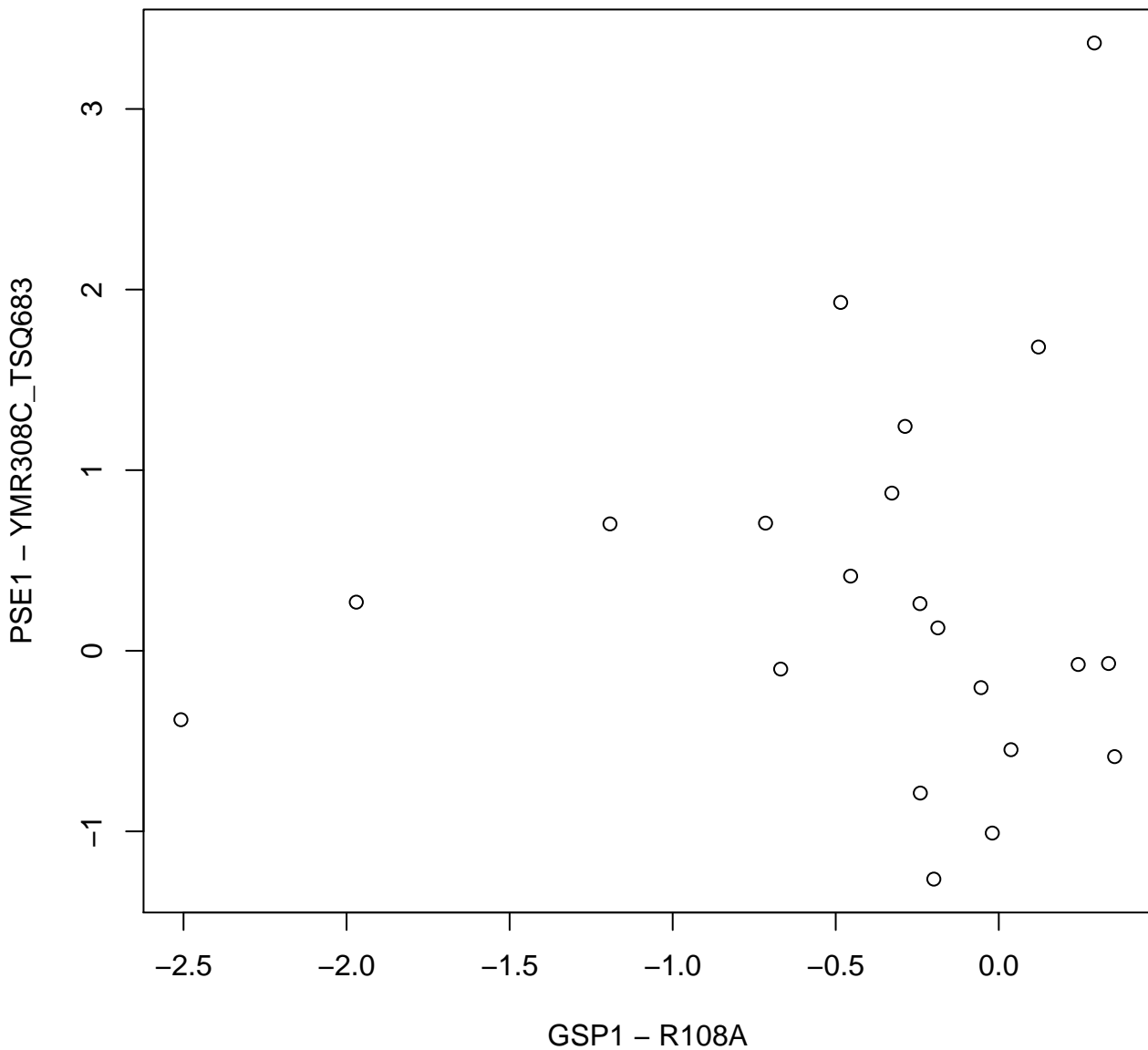
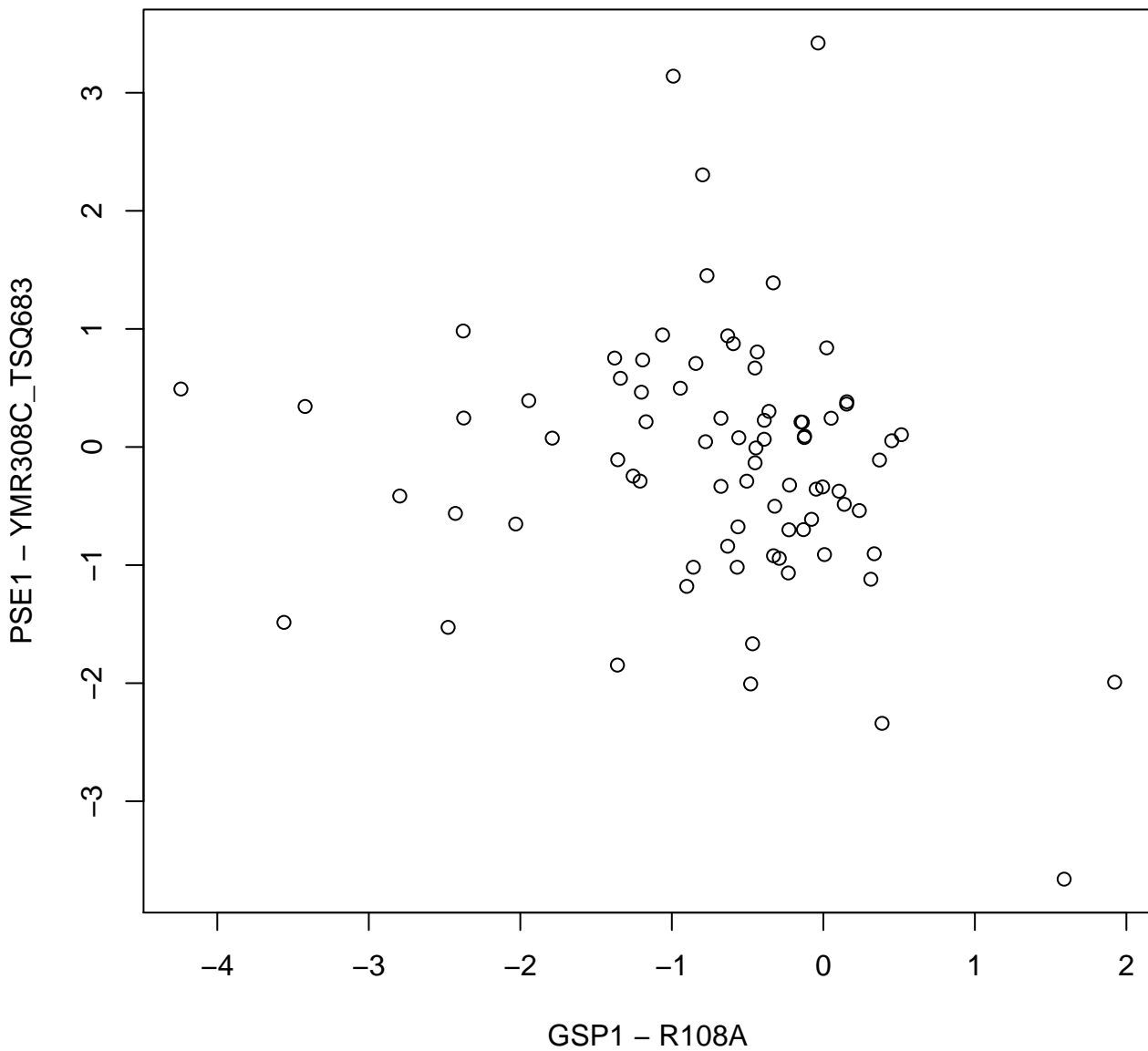


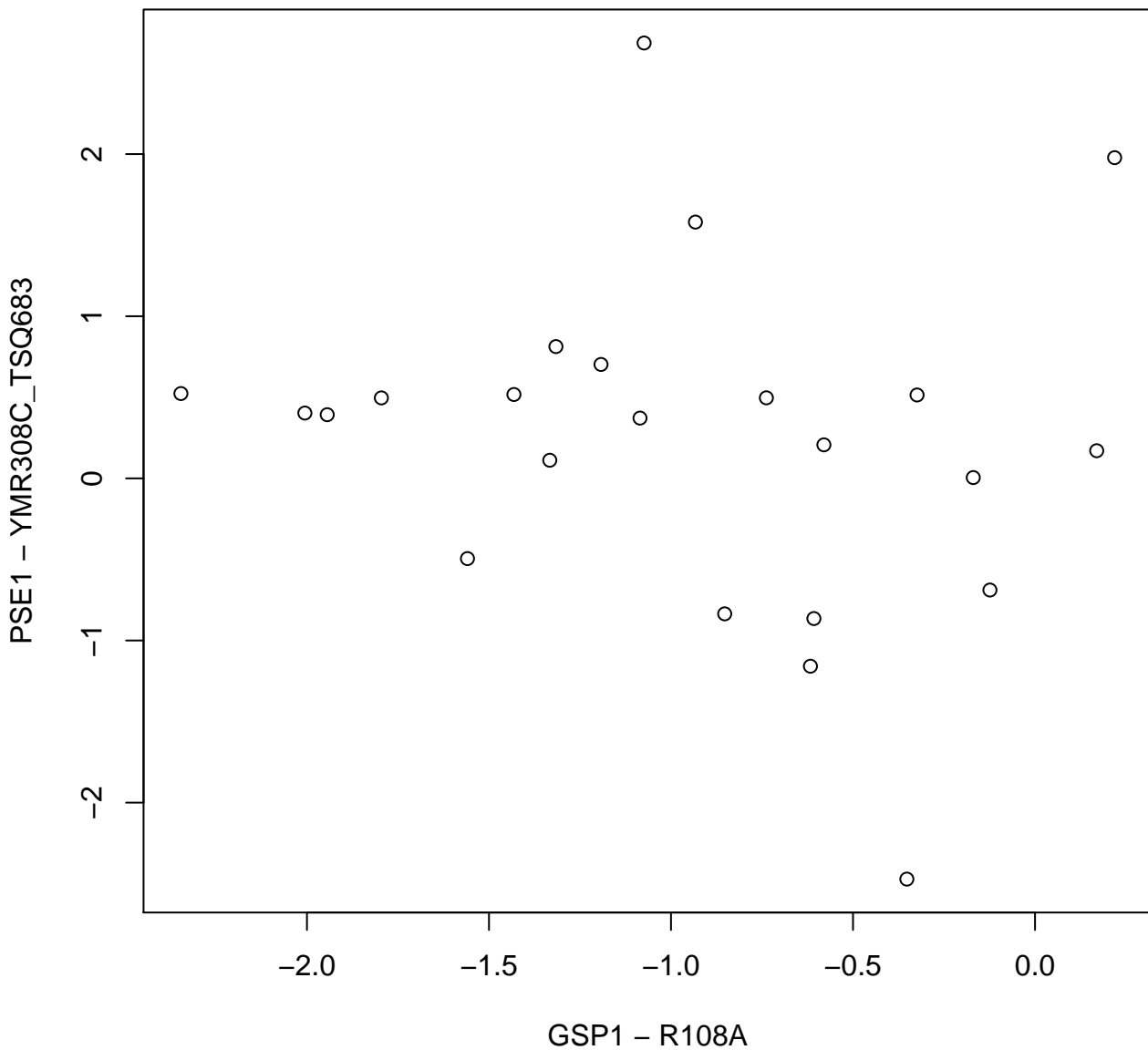
ribosome_GO_2



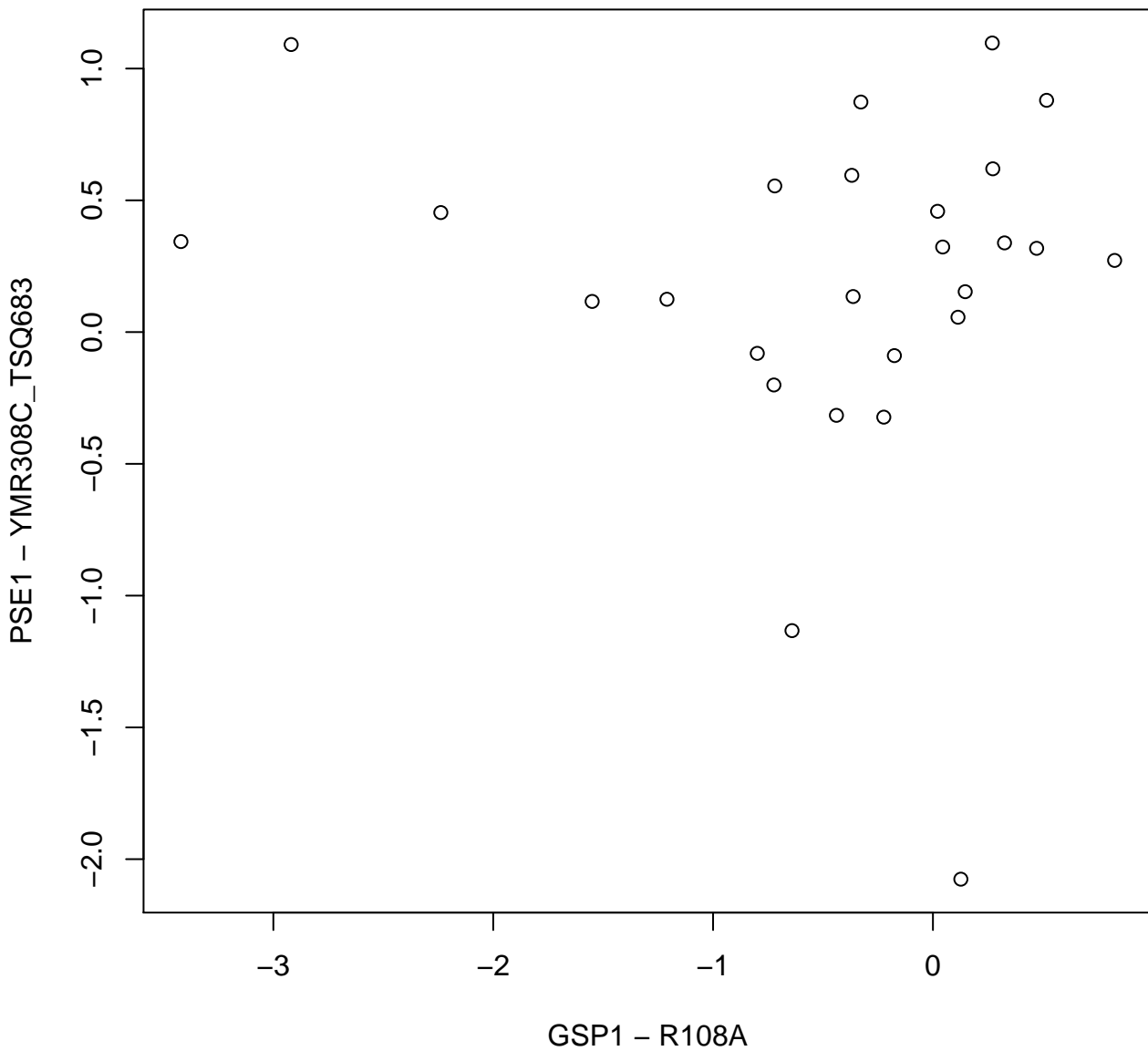
ribosome_GO_1



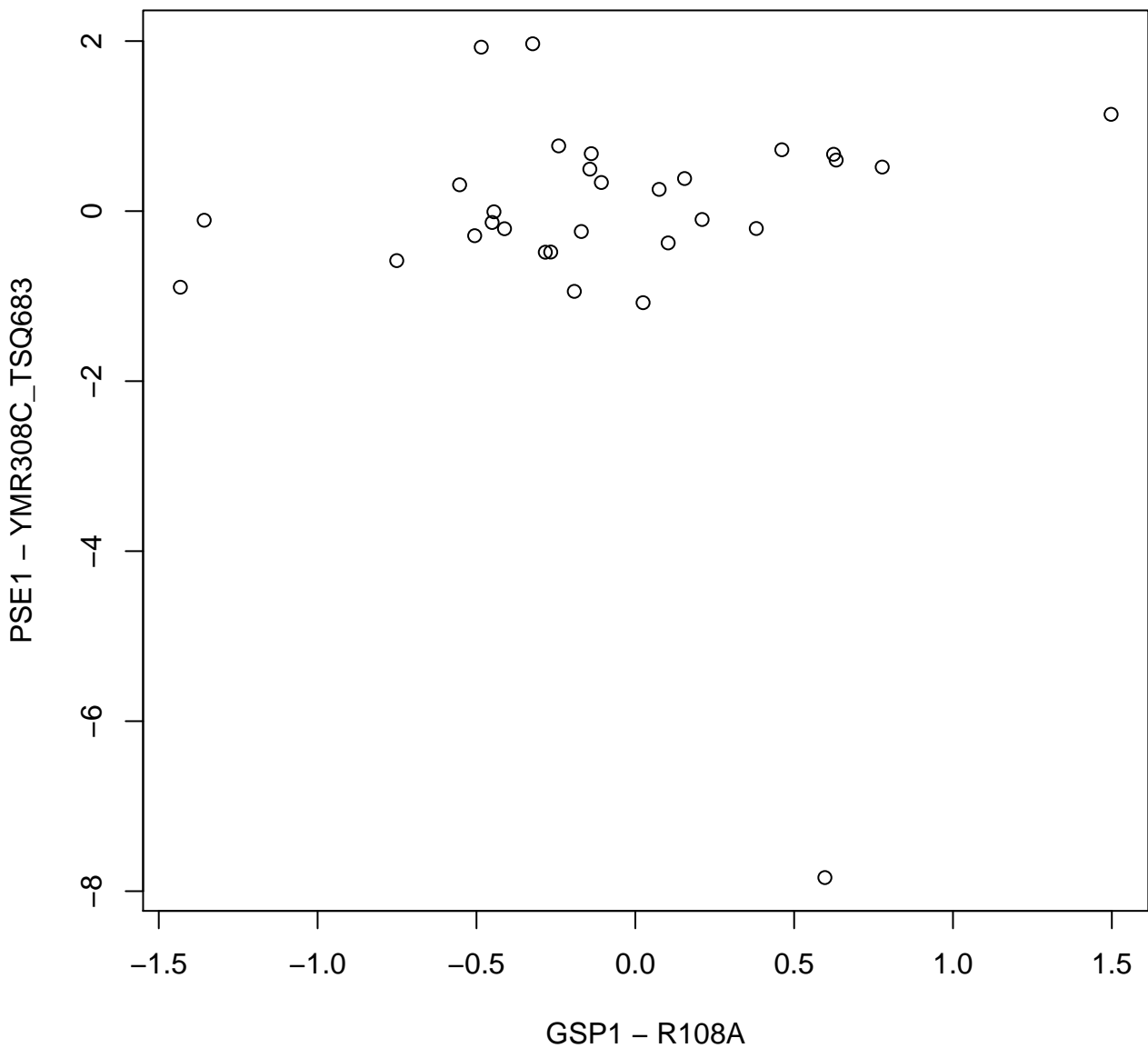
transcription and mRNA processing_GO_6



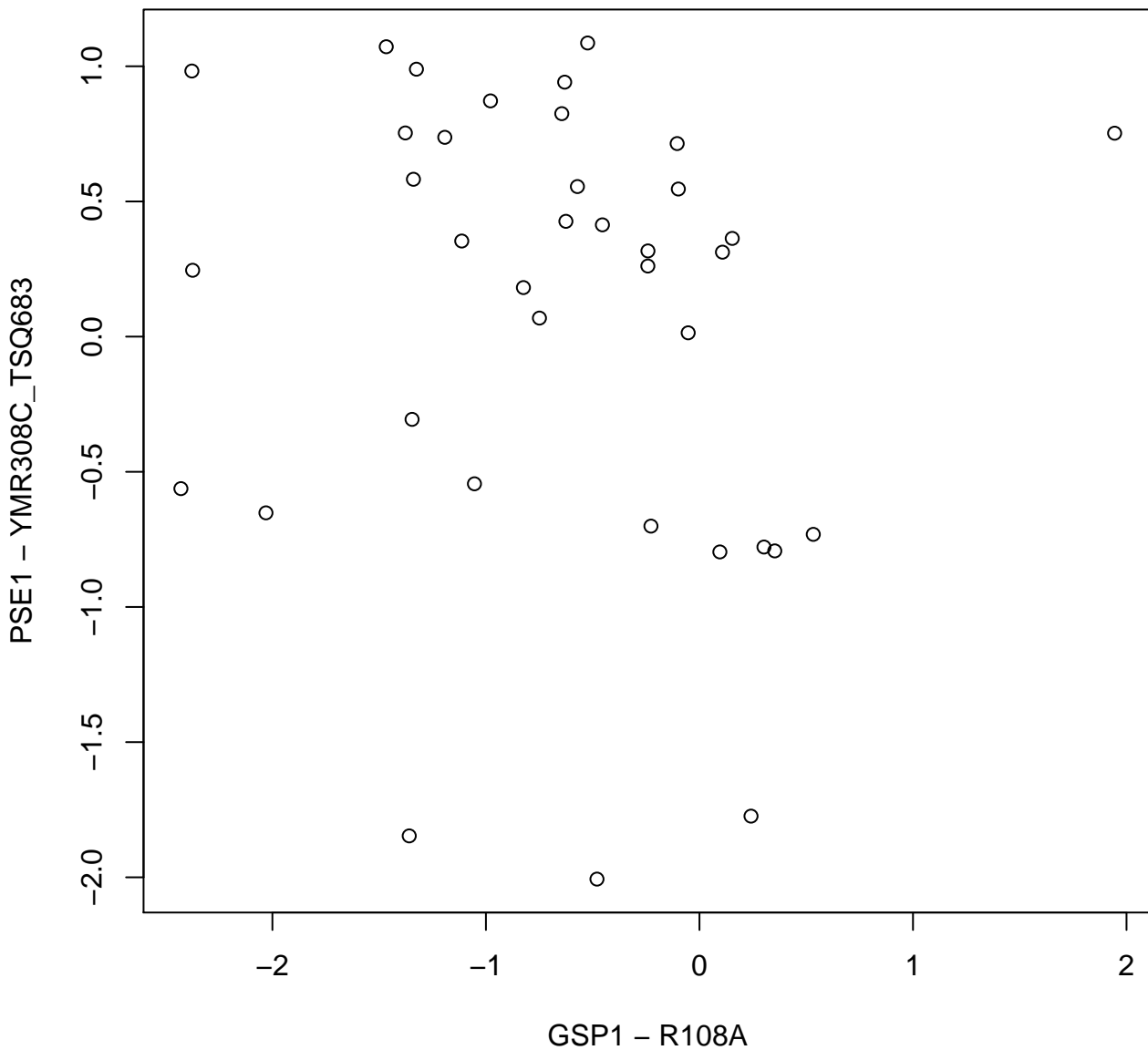
transcription and mRNA processing_GO_5



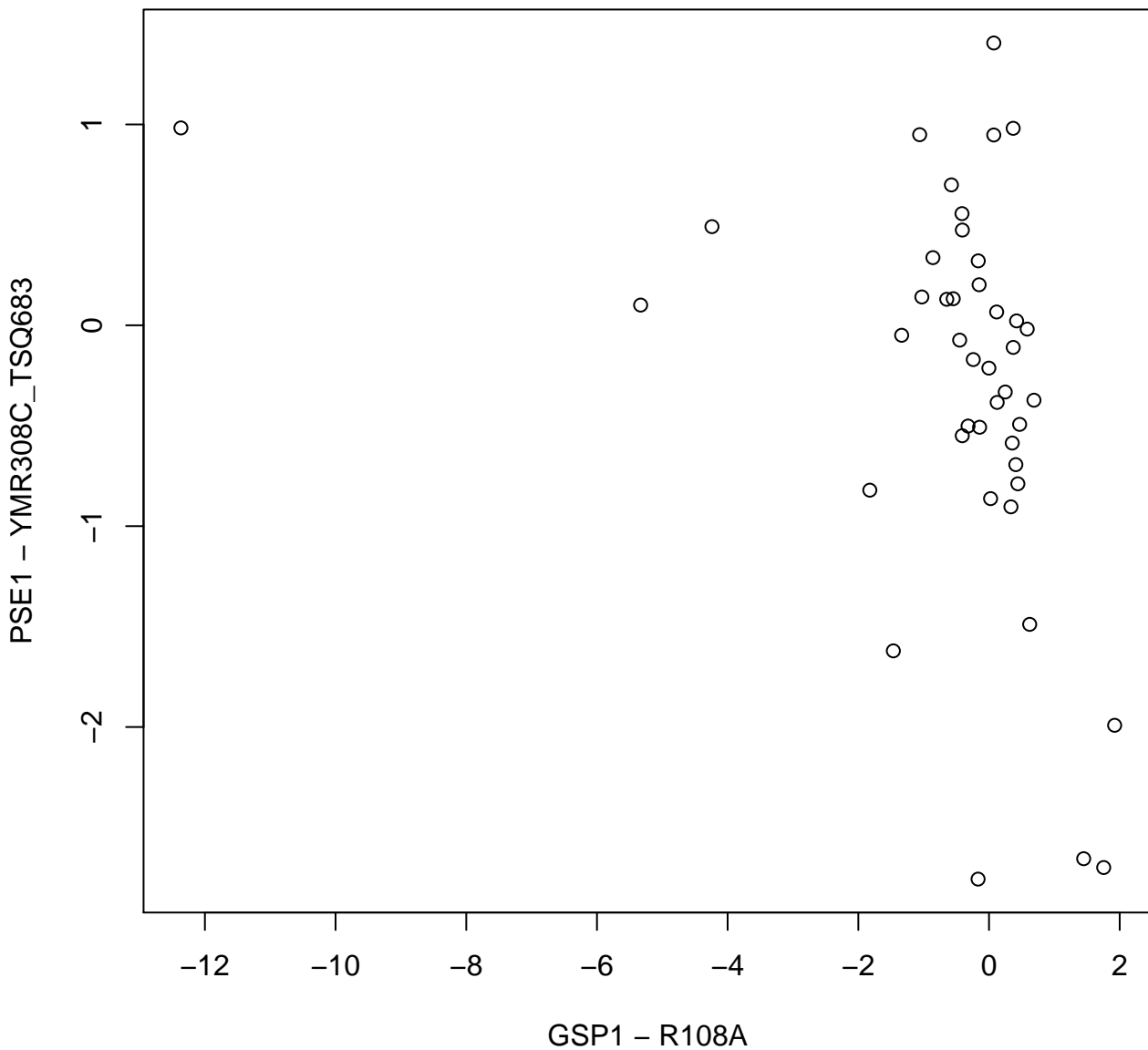
transcription and mRNA processing_GO_4



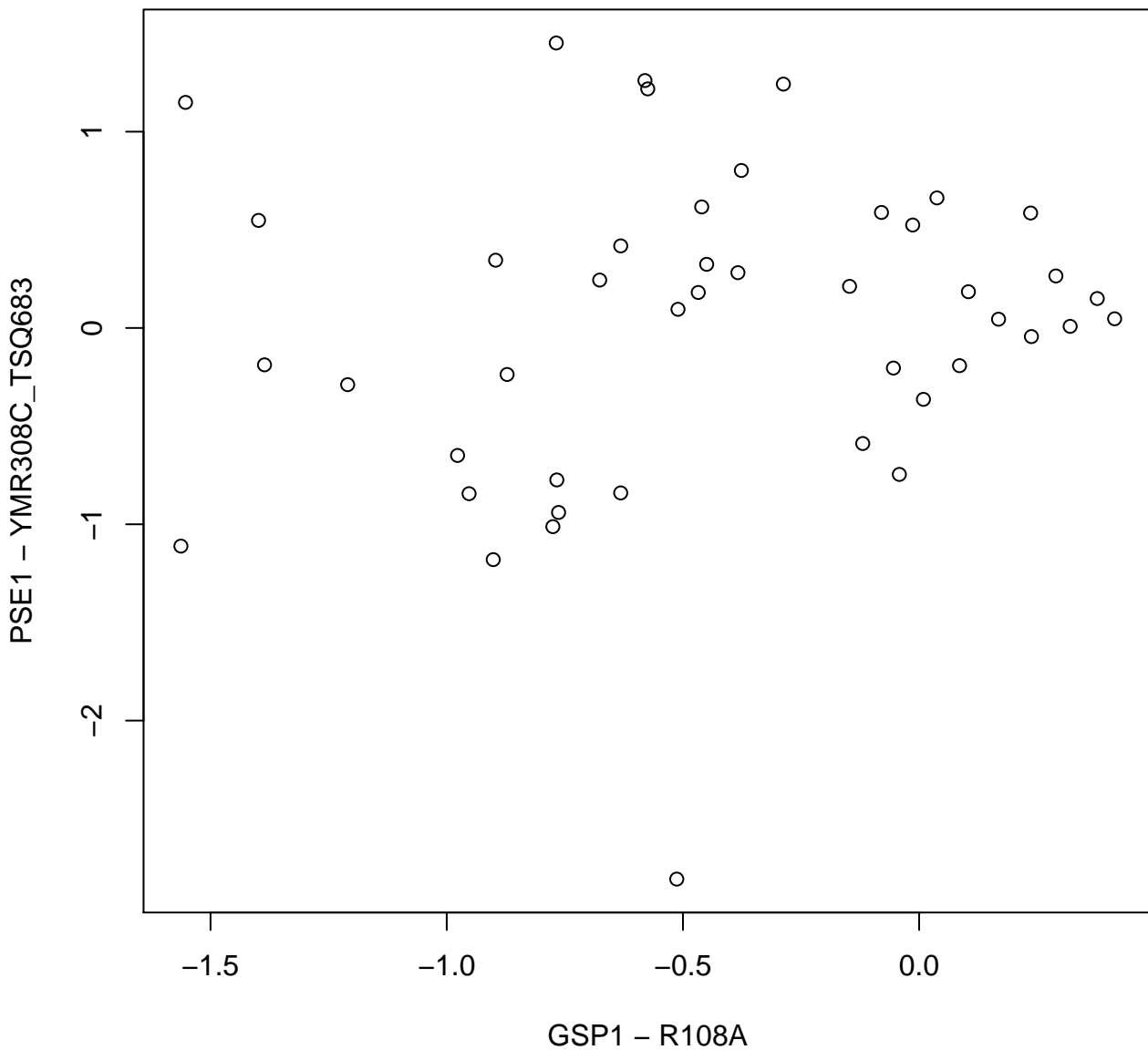
transcription and mRNA processing_GO_3



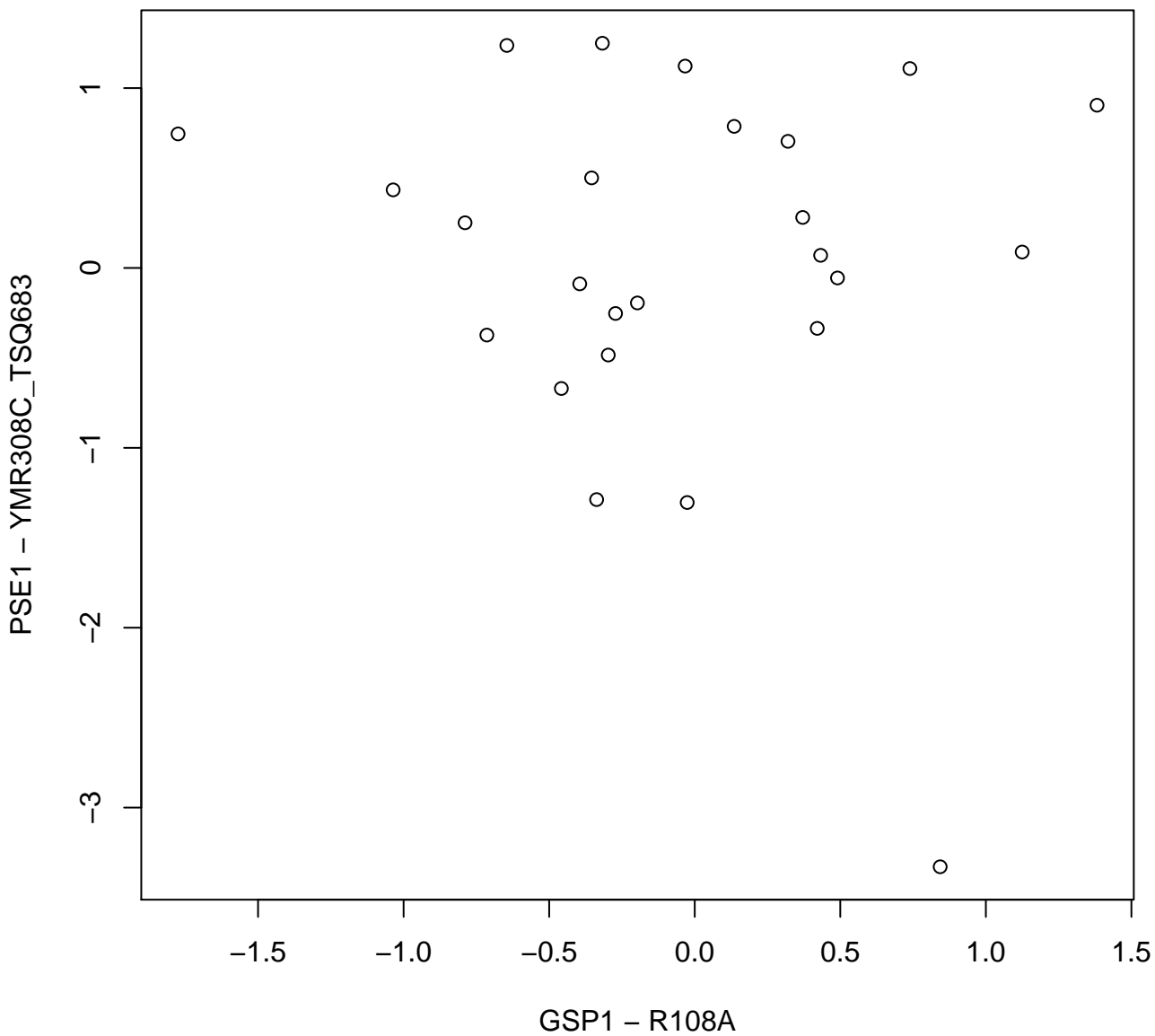
transcription and mRNA processing_GO_2



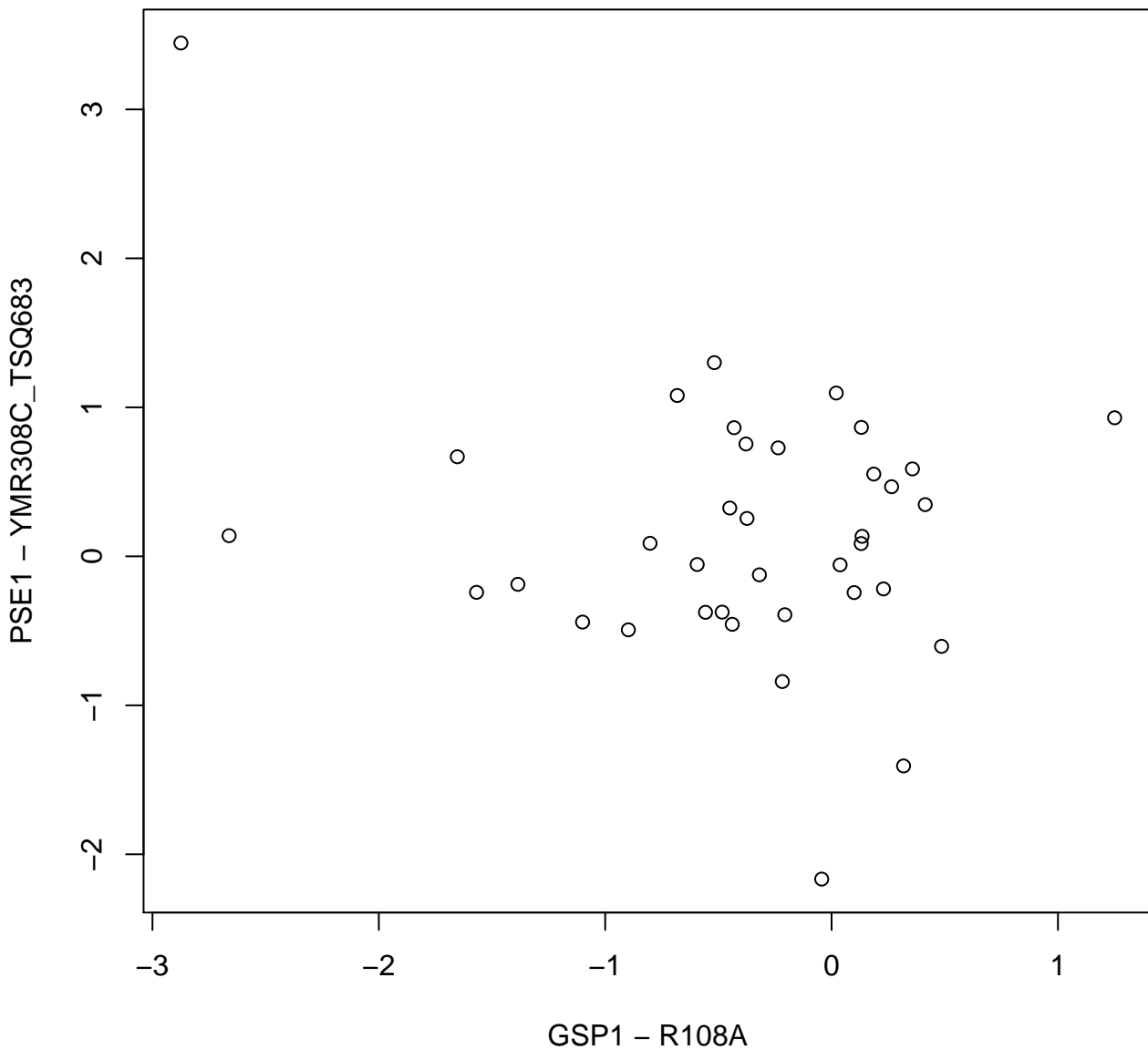
transcription and mRNA processing_GO_1



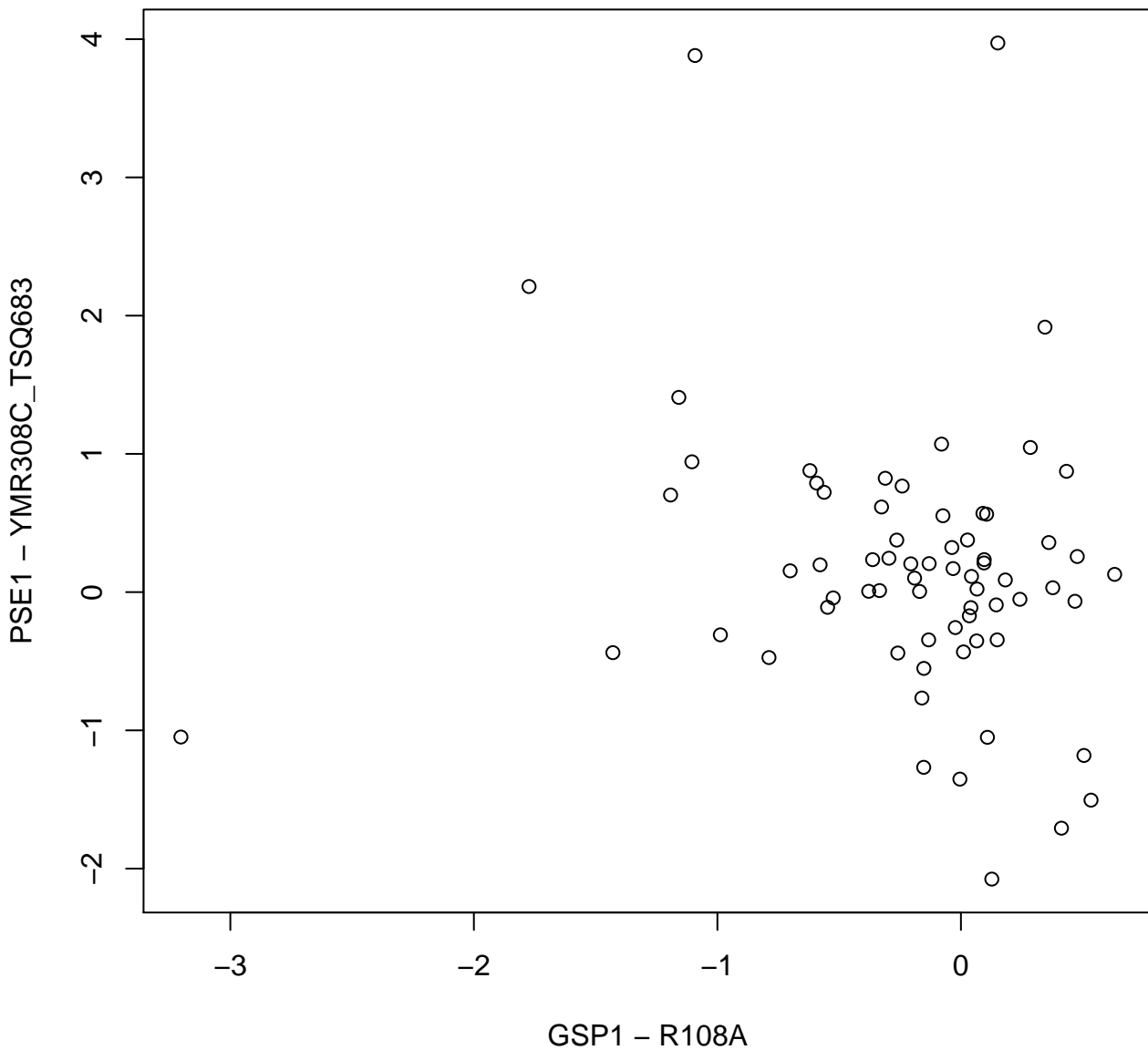
Golgi and ER_GO_3



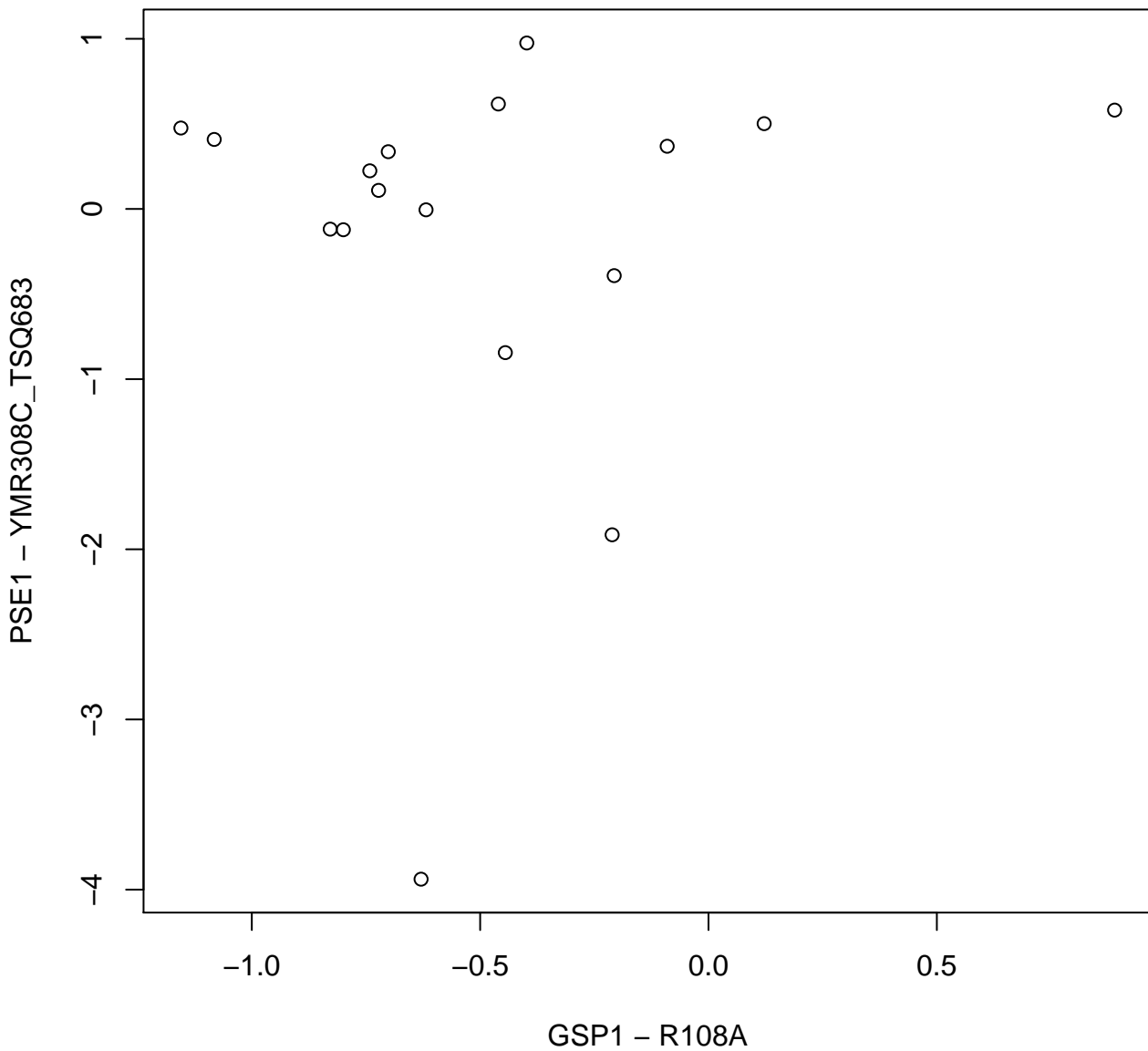
Golgi and ER_GO_2



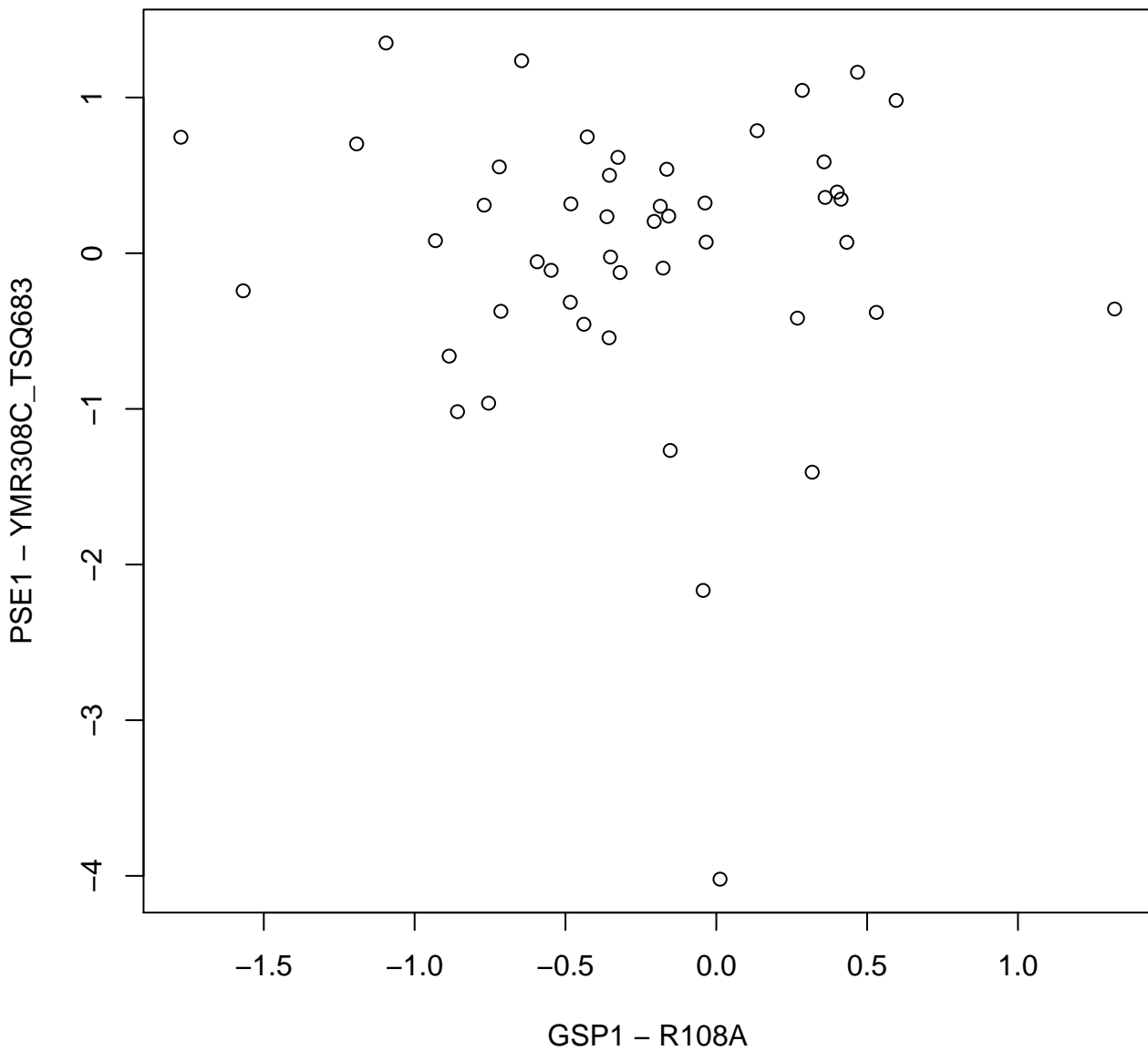
Golgi and ER_GO_1



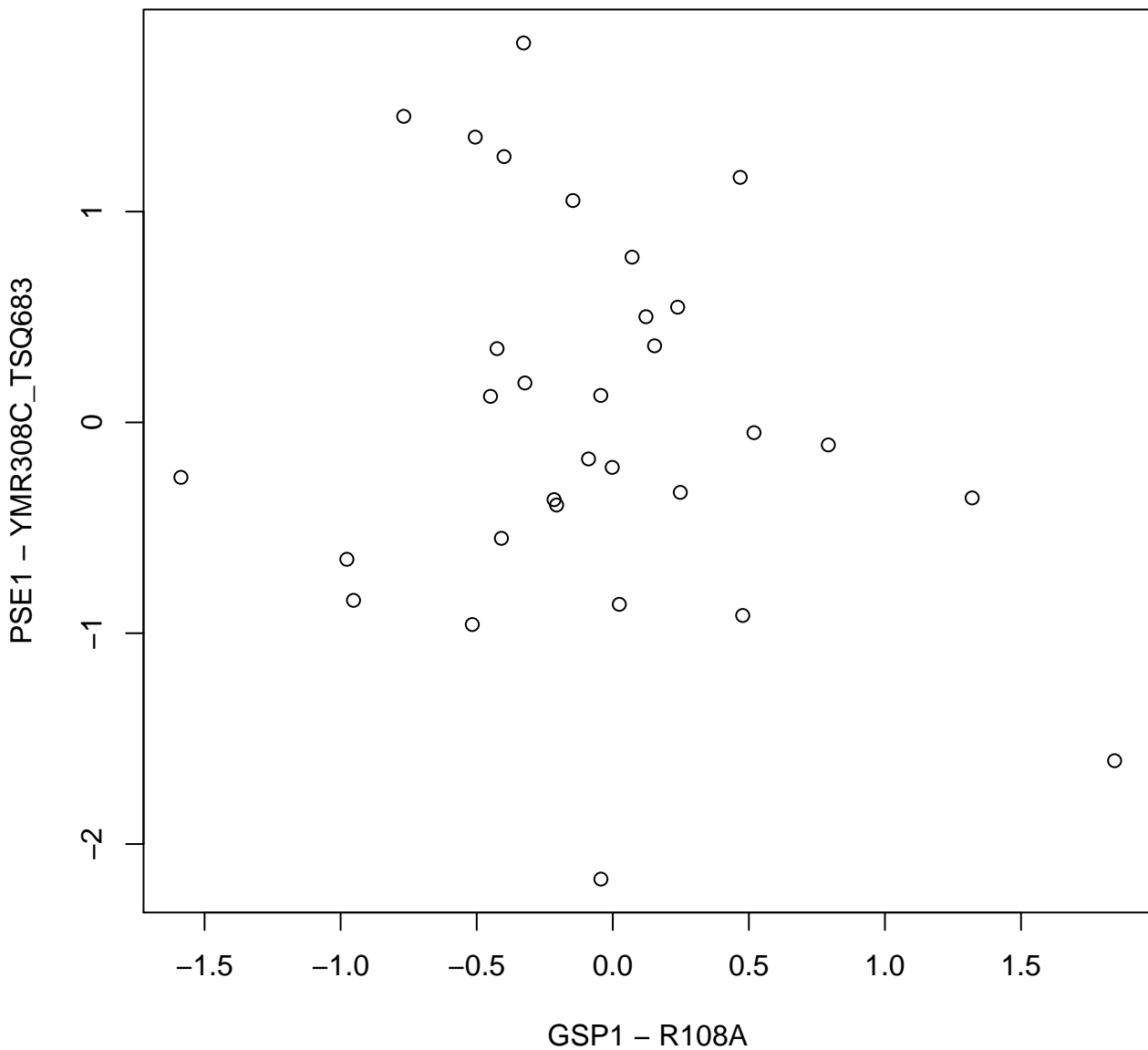
peroxisome_GO_0



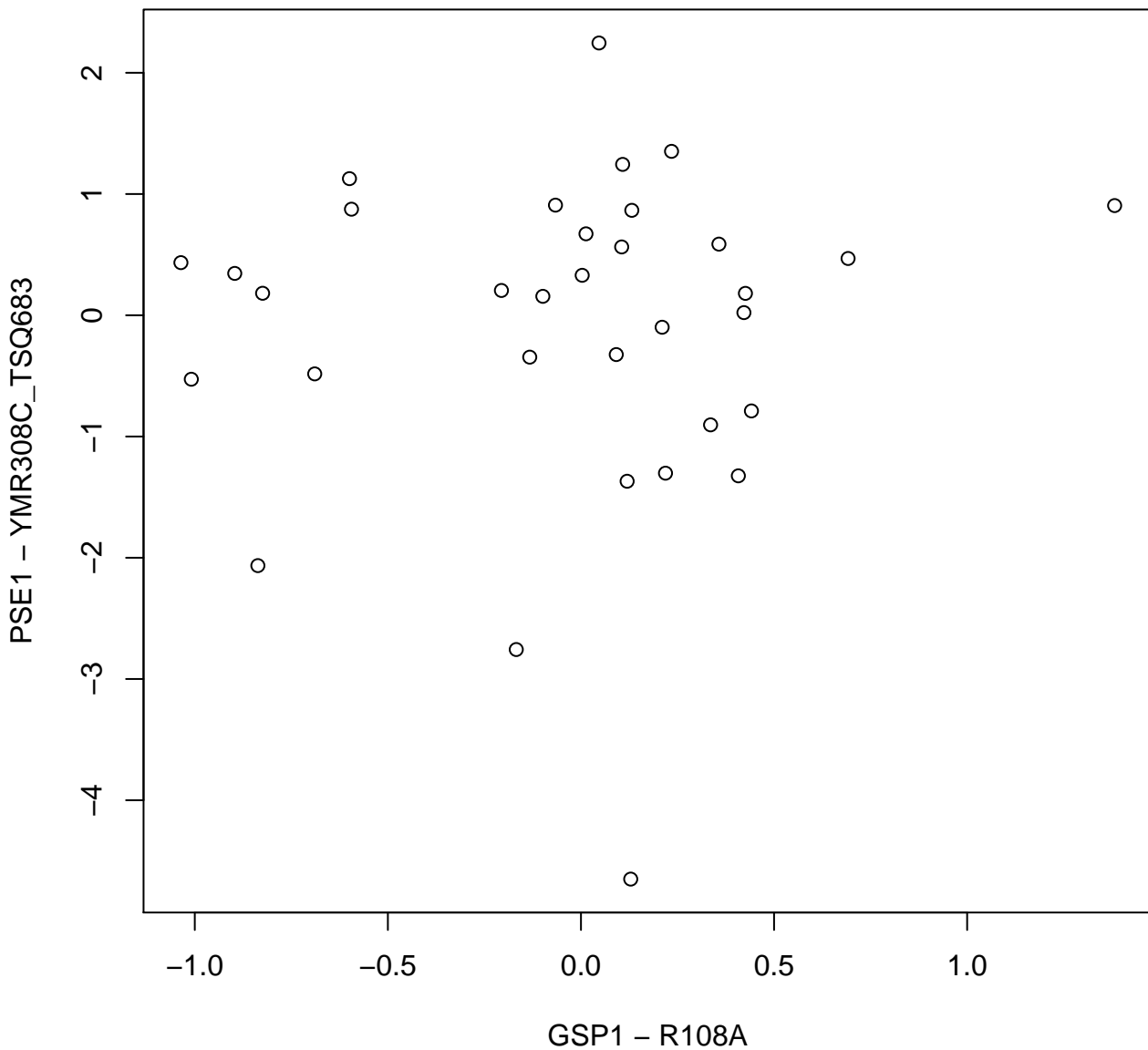
vacuole_GO_1



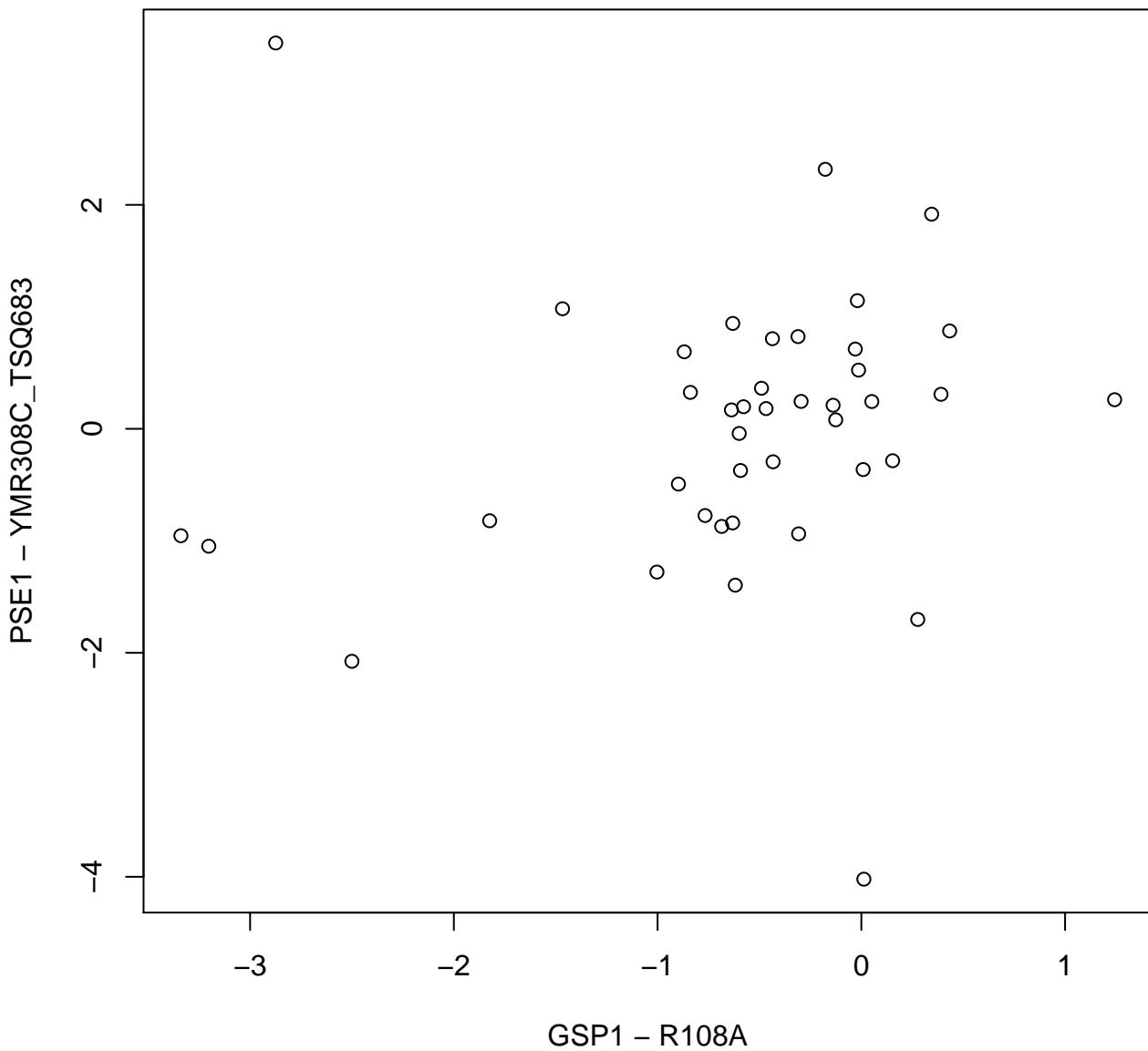
mitochondrion_GO_3



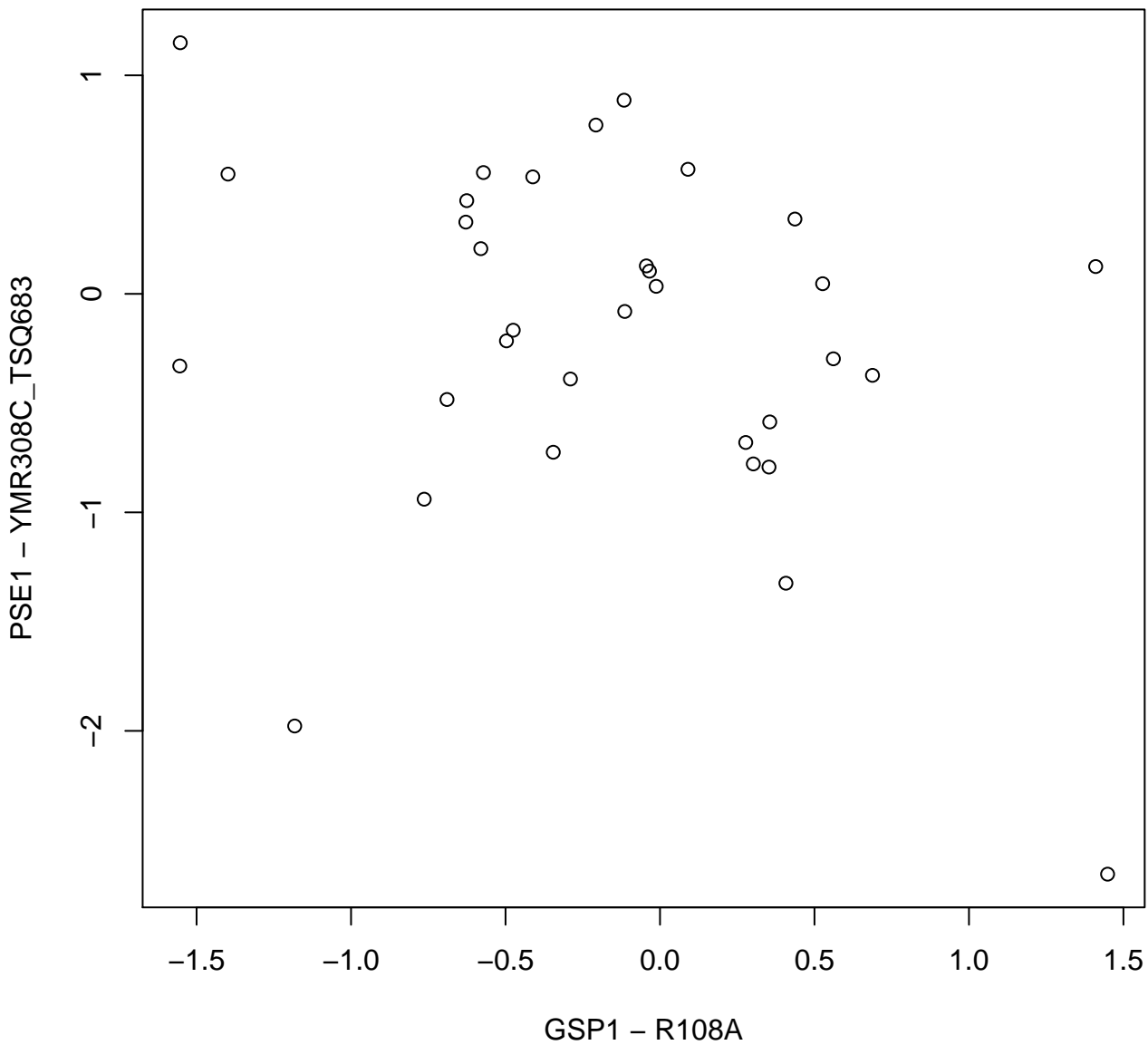
mitochondrion_GO_2



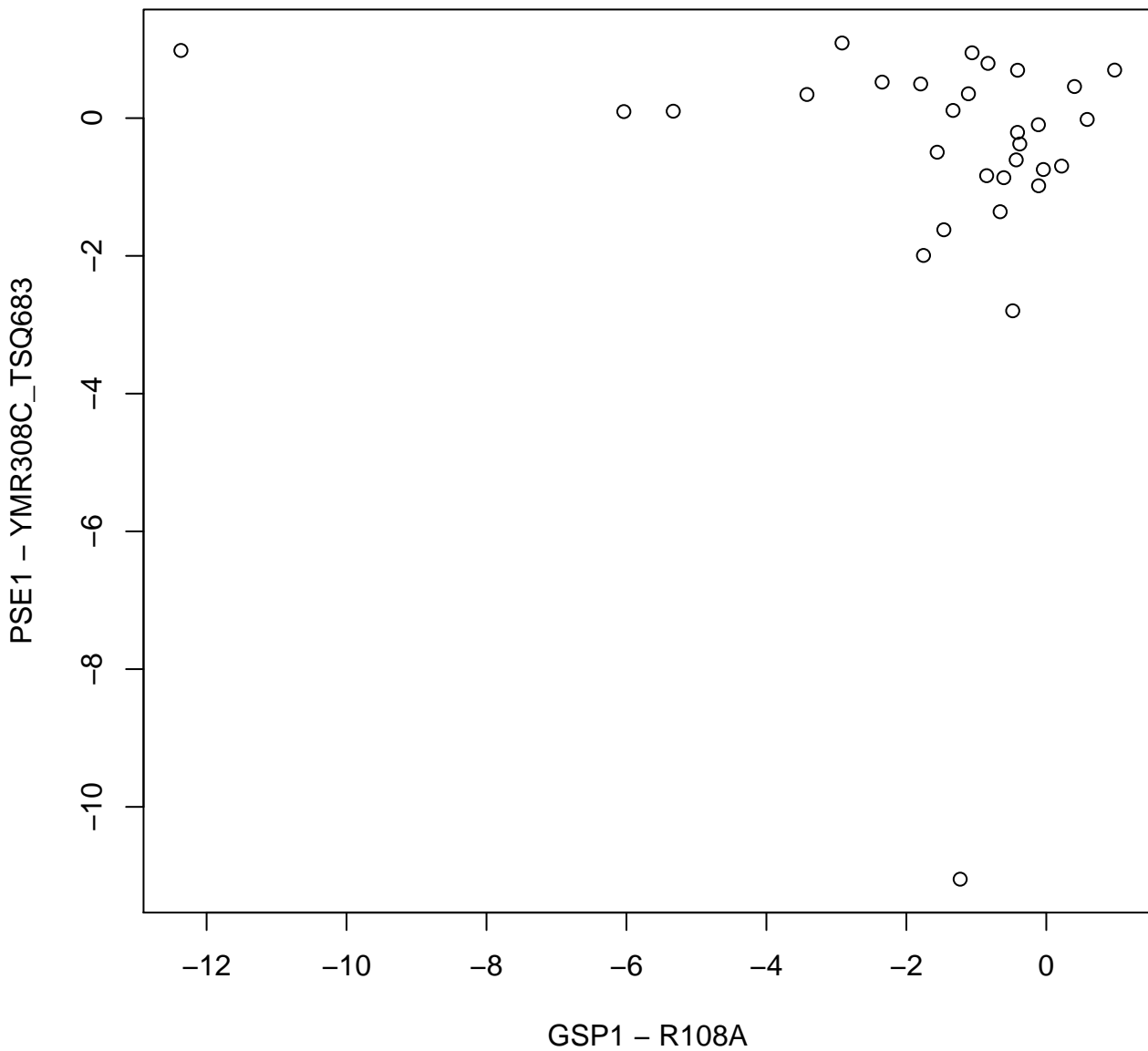
mitochondrion_GO_1



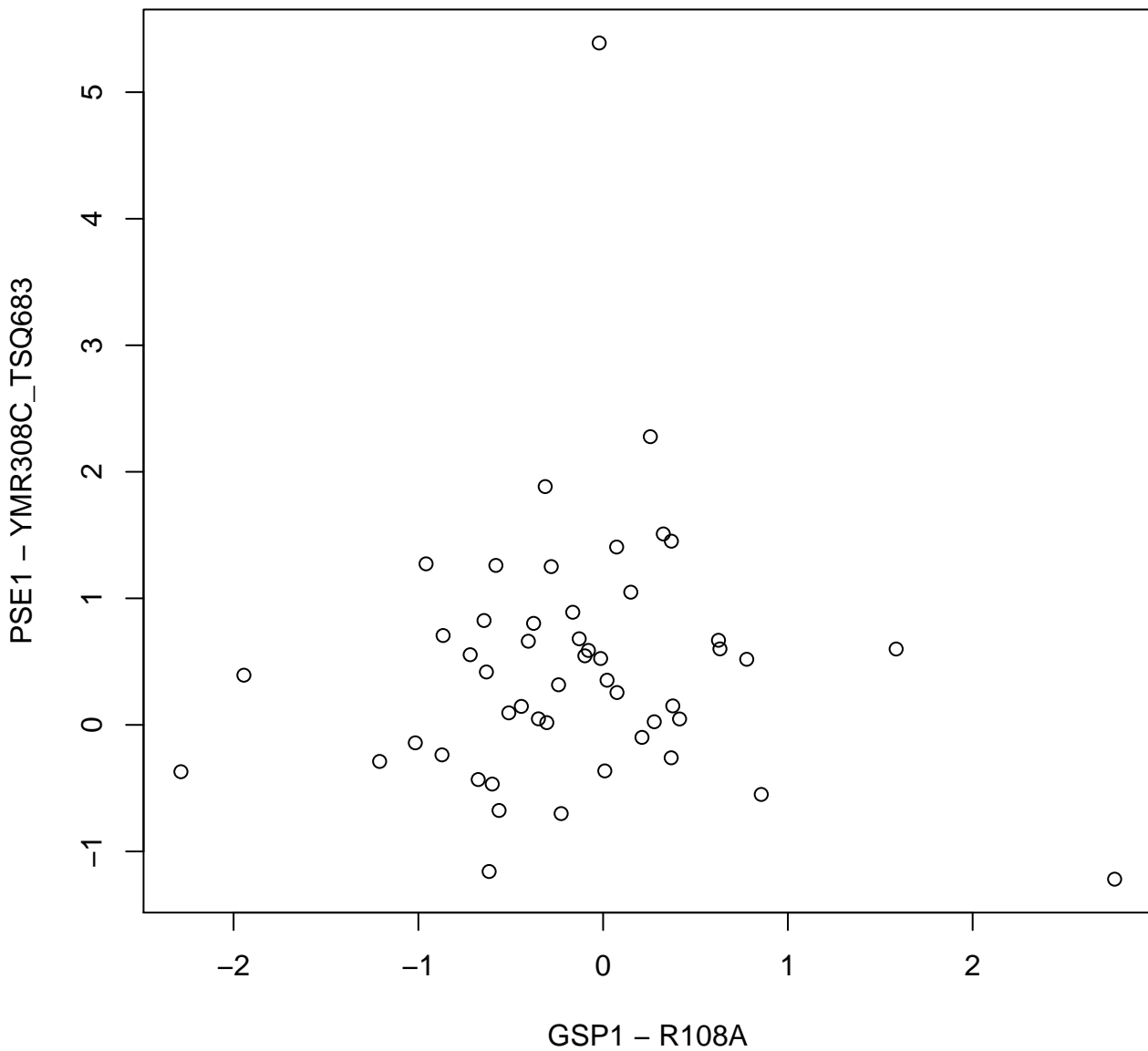
chromatin_GO_4



chromatin_GO_3

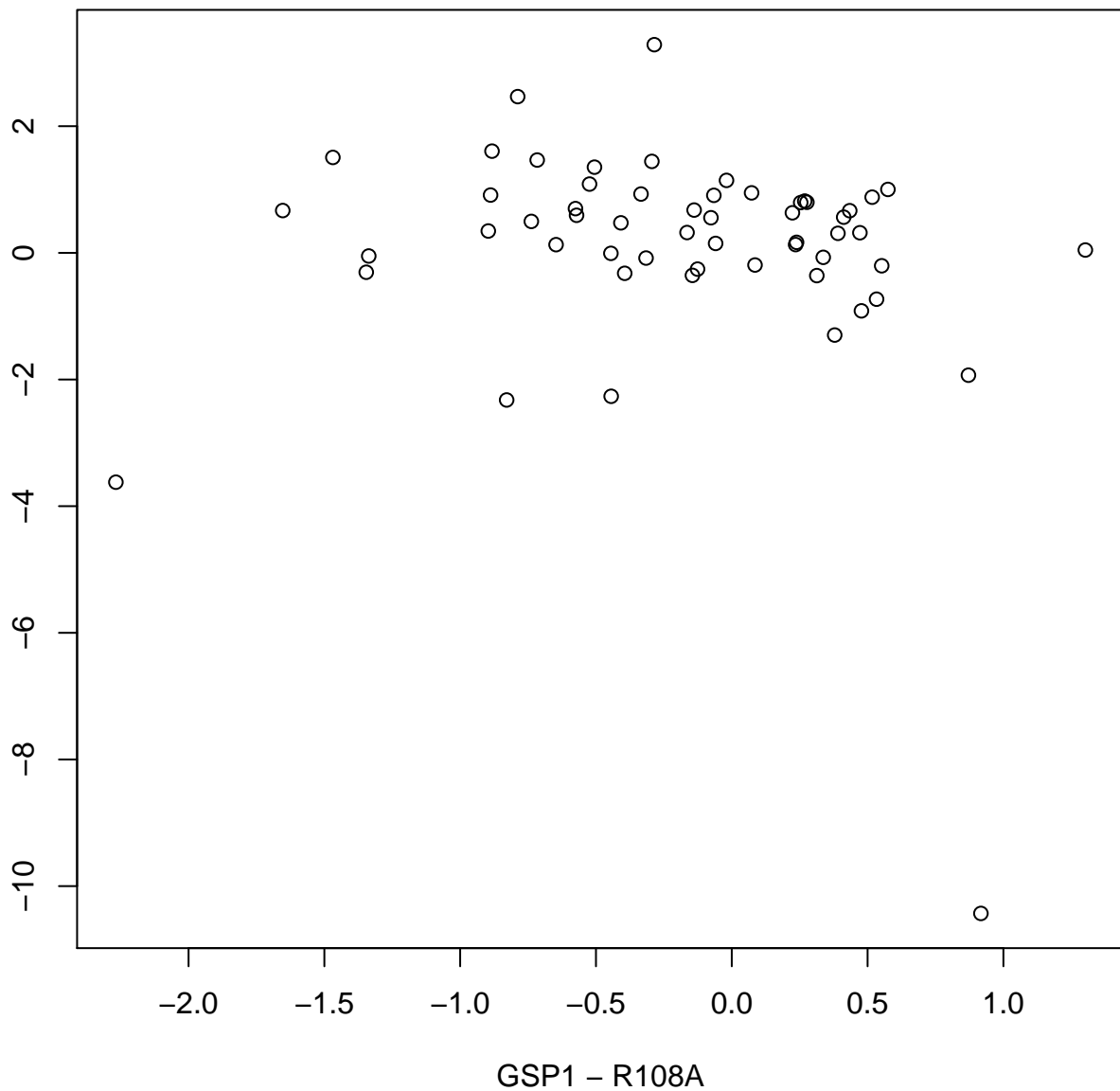


chromatin_GO_2

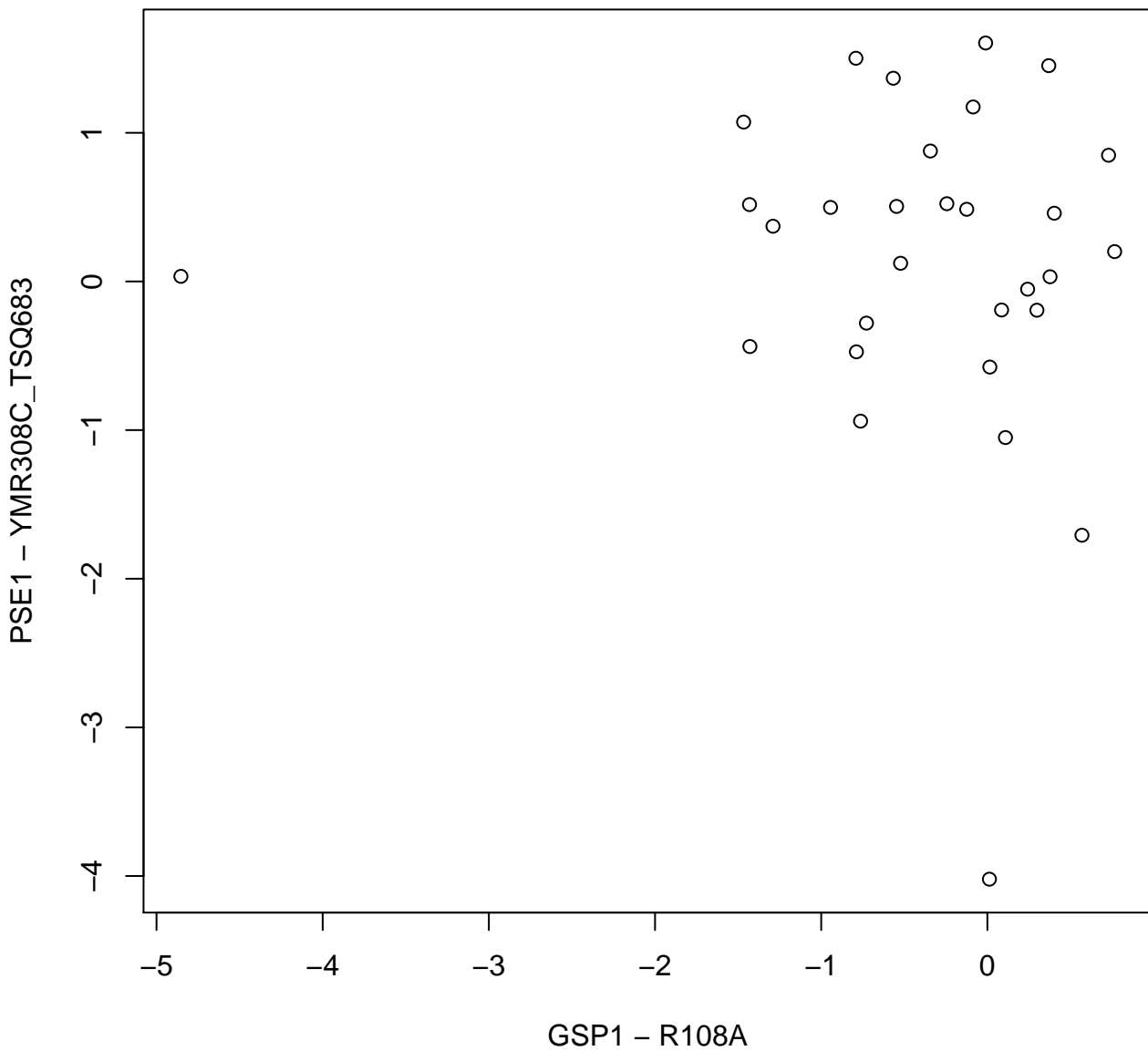


chromatin_GO_1

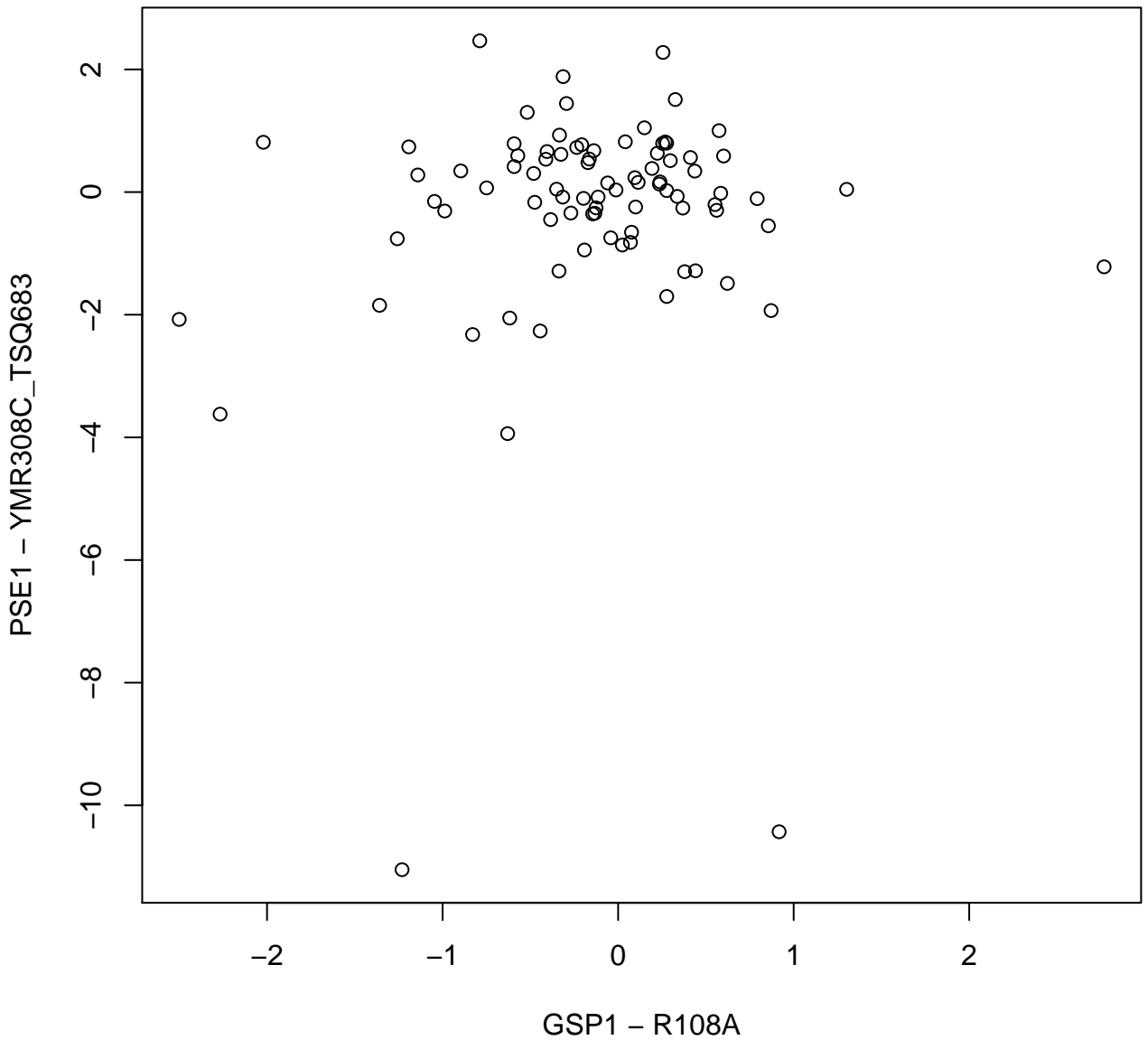
PSE1 - YMR308C_TSQ683



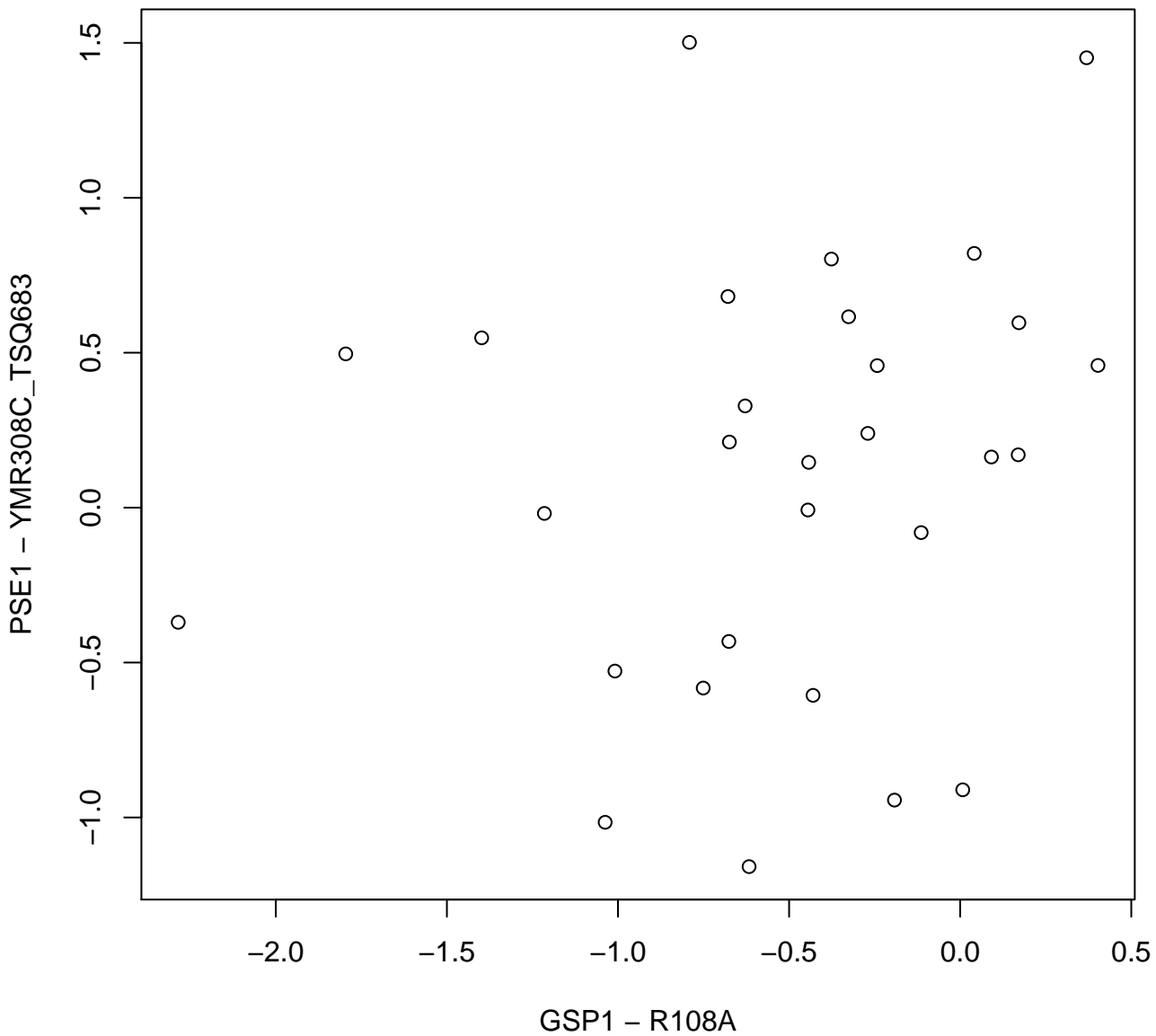
cytoskeleton_GO_2



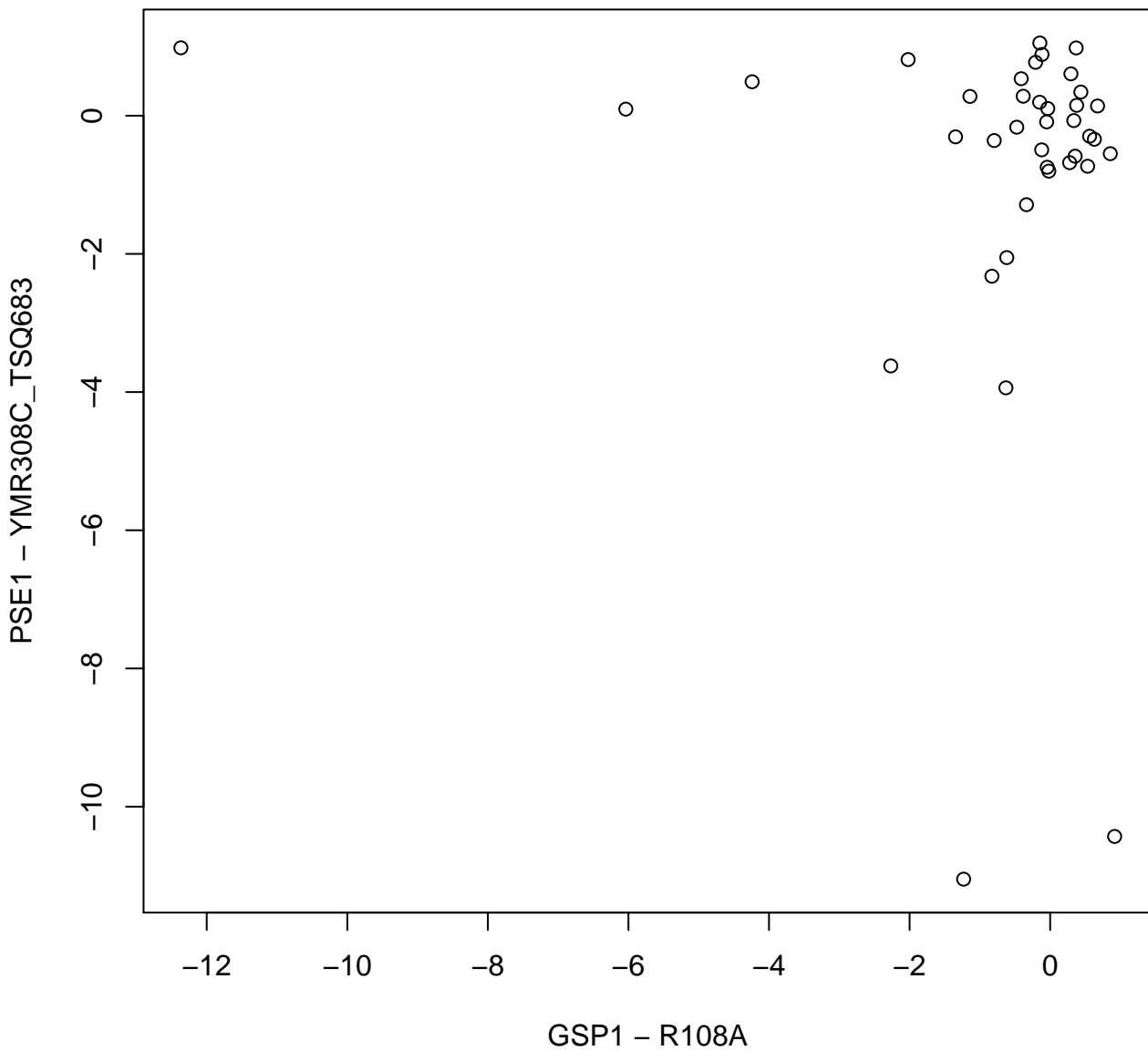
cytoskeleton_GO_1



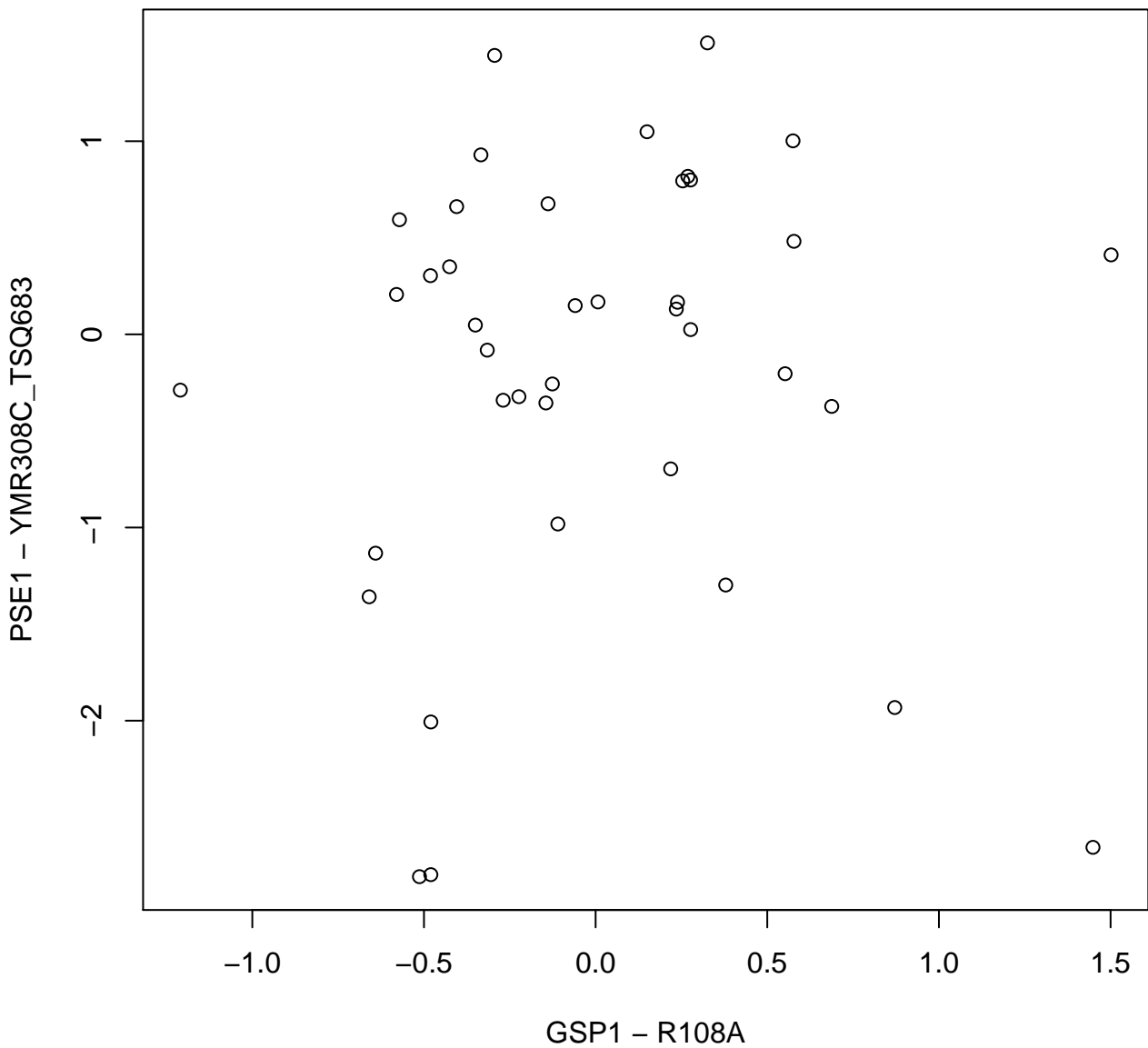
cell cycle_GO_4



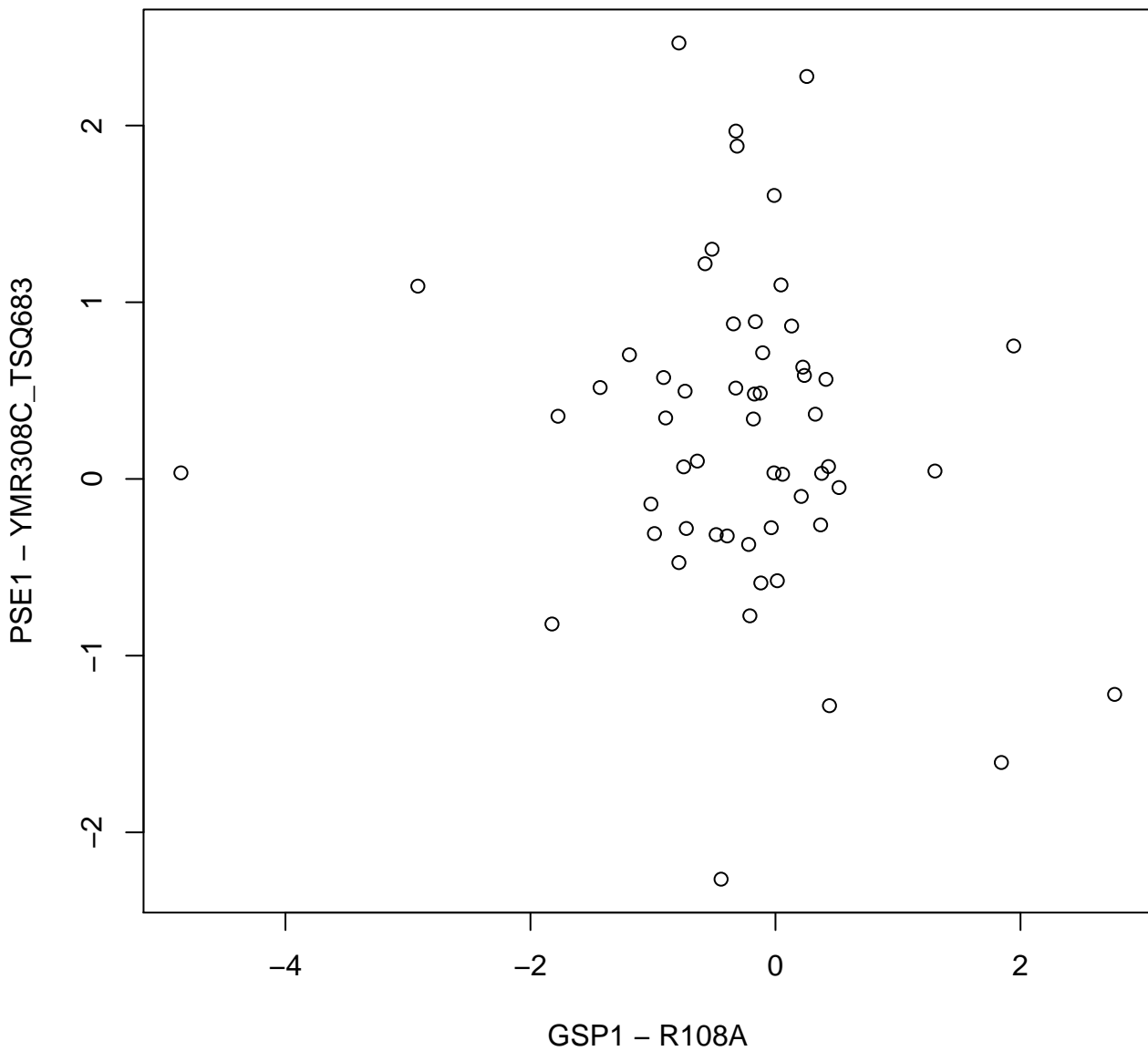
cell cycle_GO_3



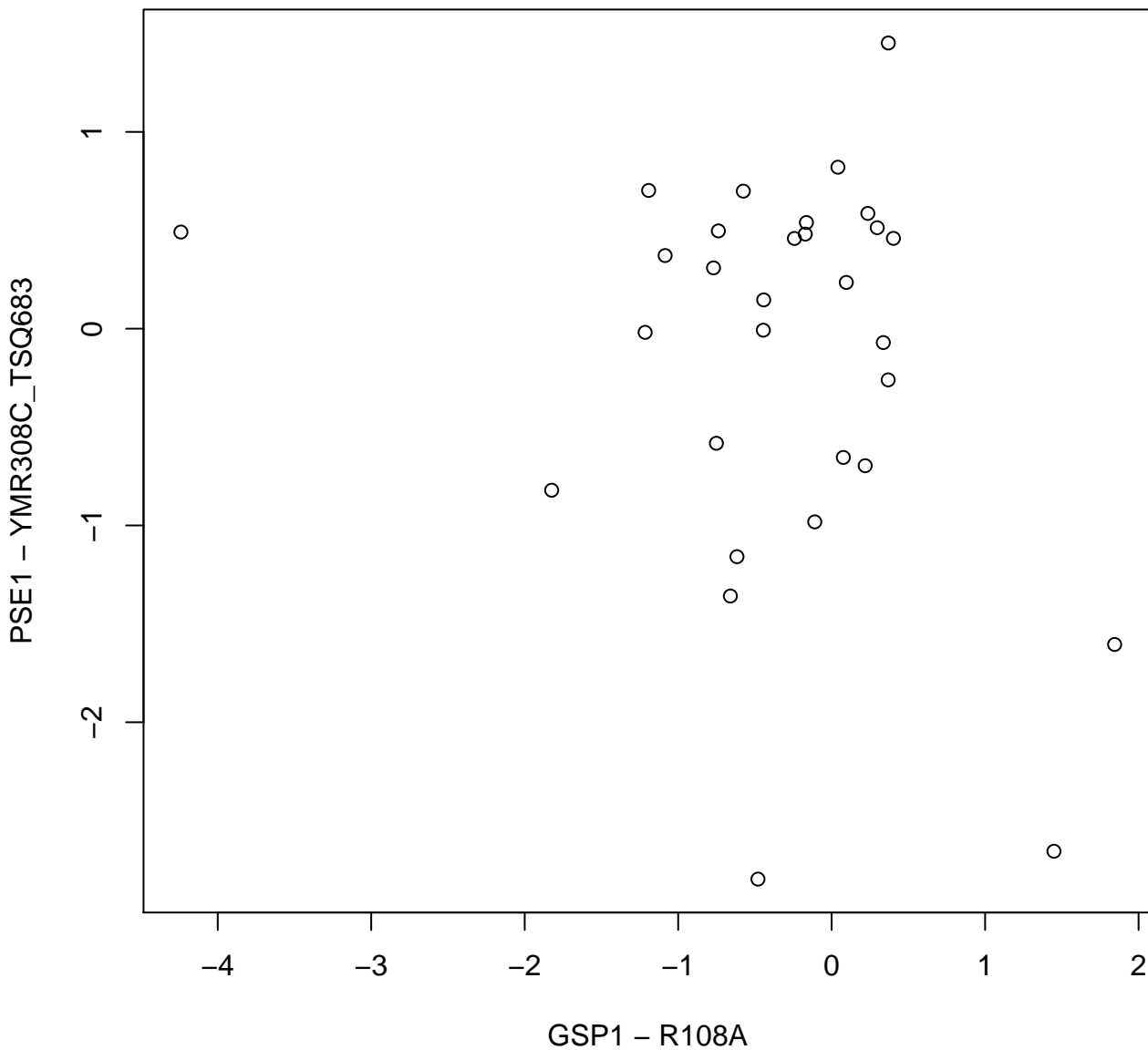
cell cycle_GO_2



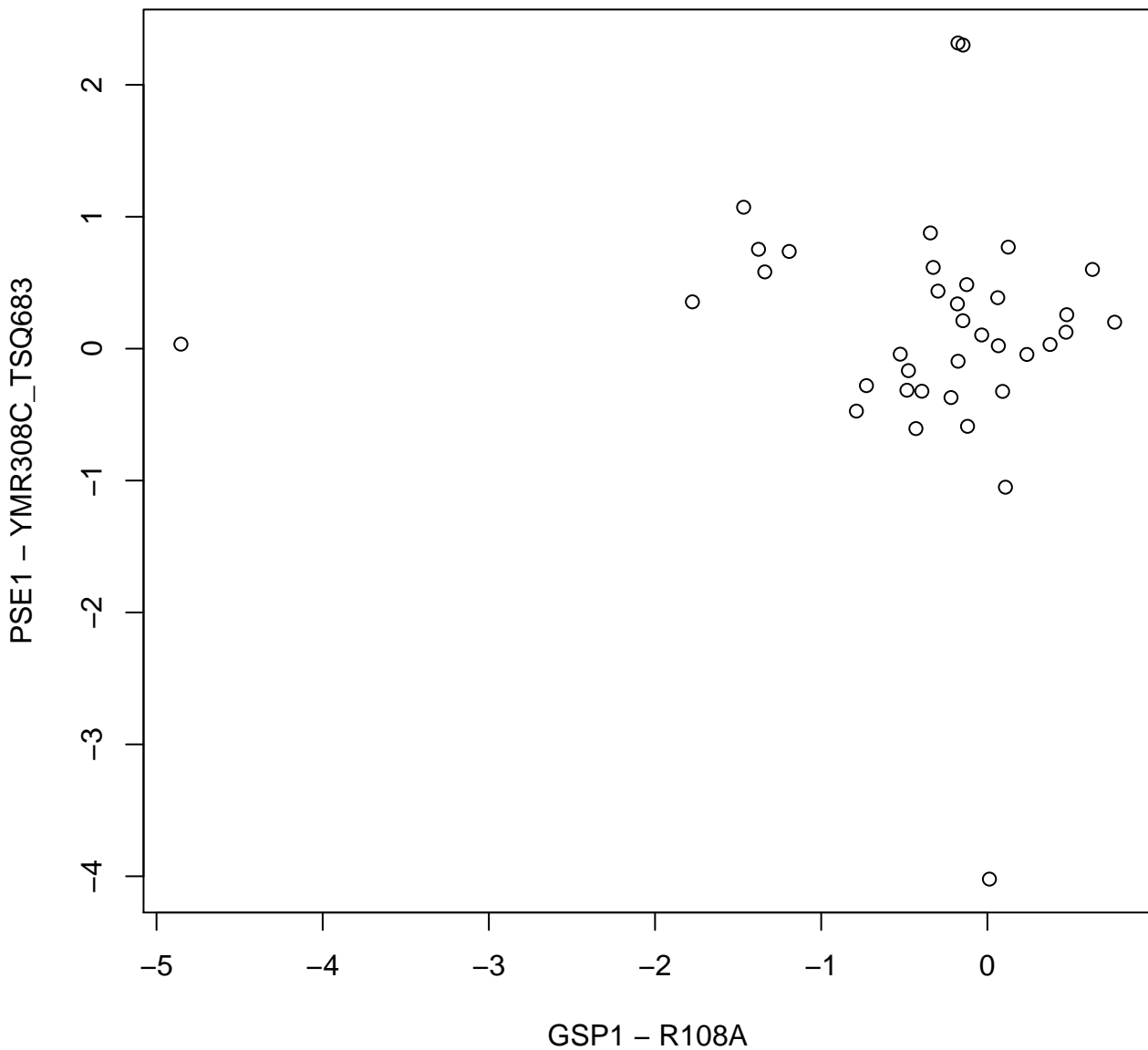
cell cycle_GO_1



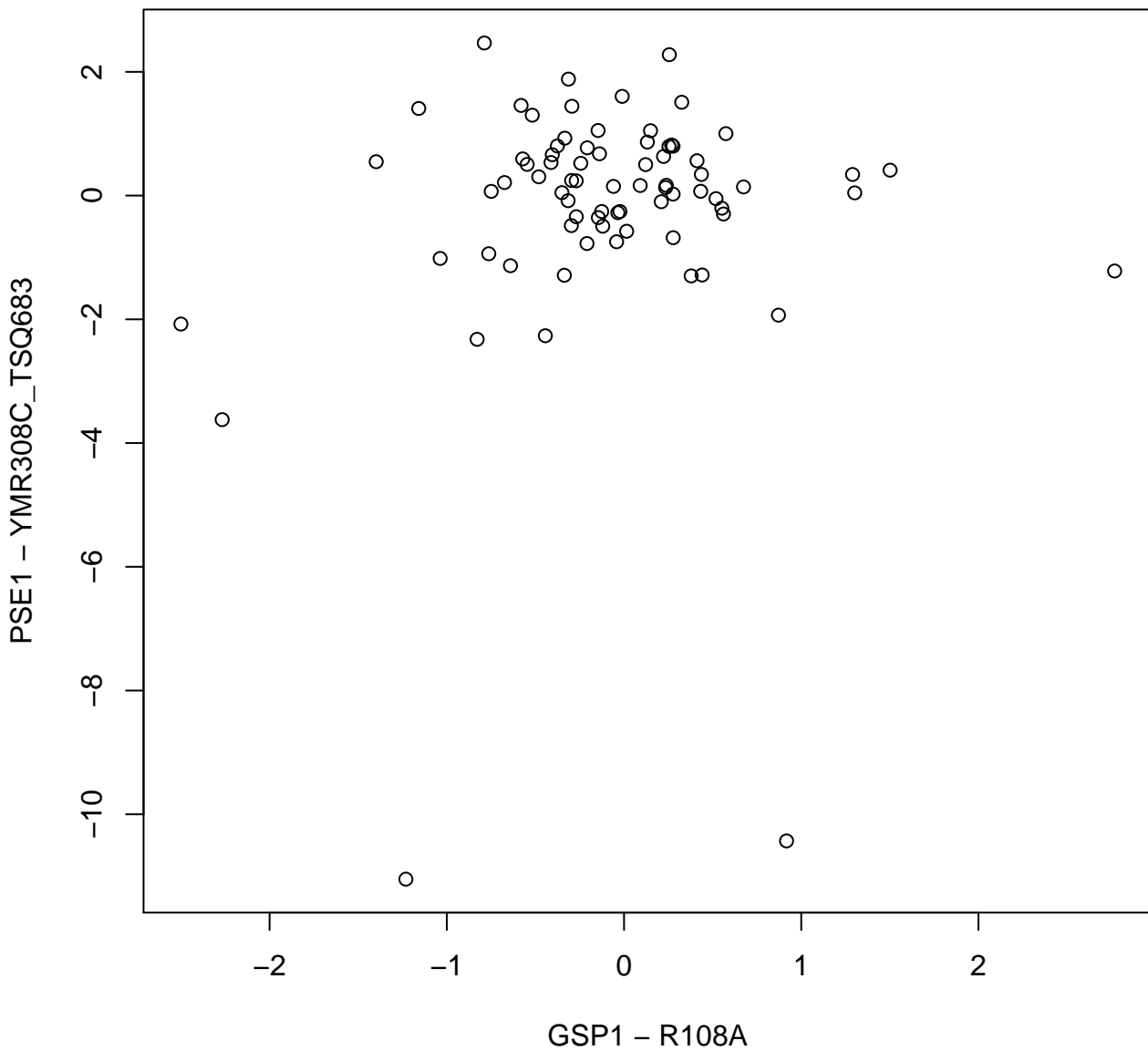
budding_GO_3



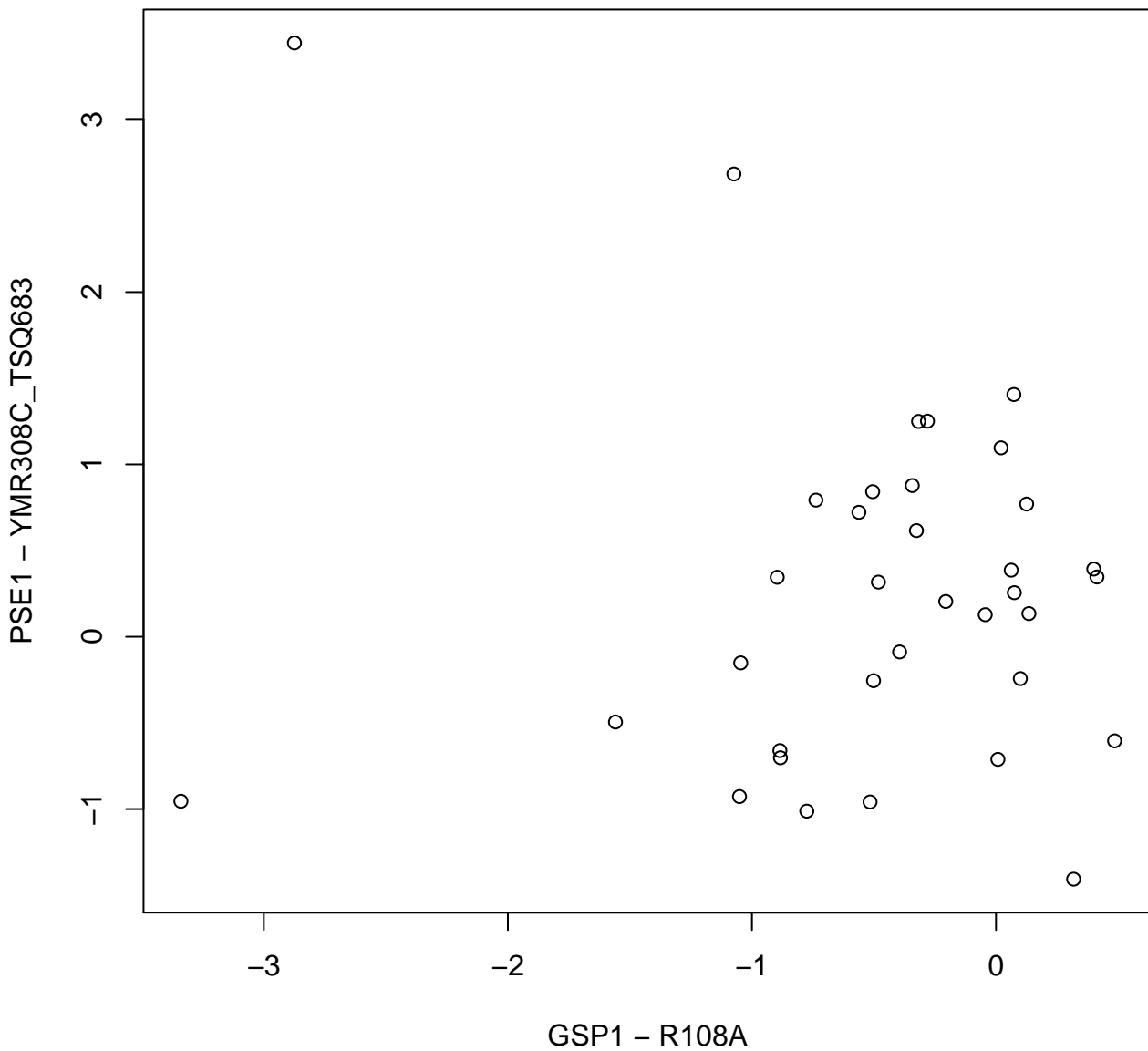
budding_GO_2



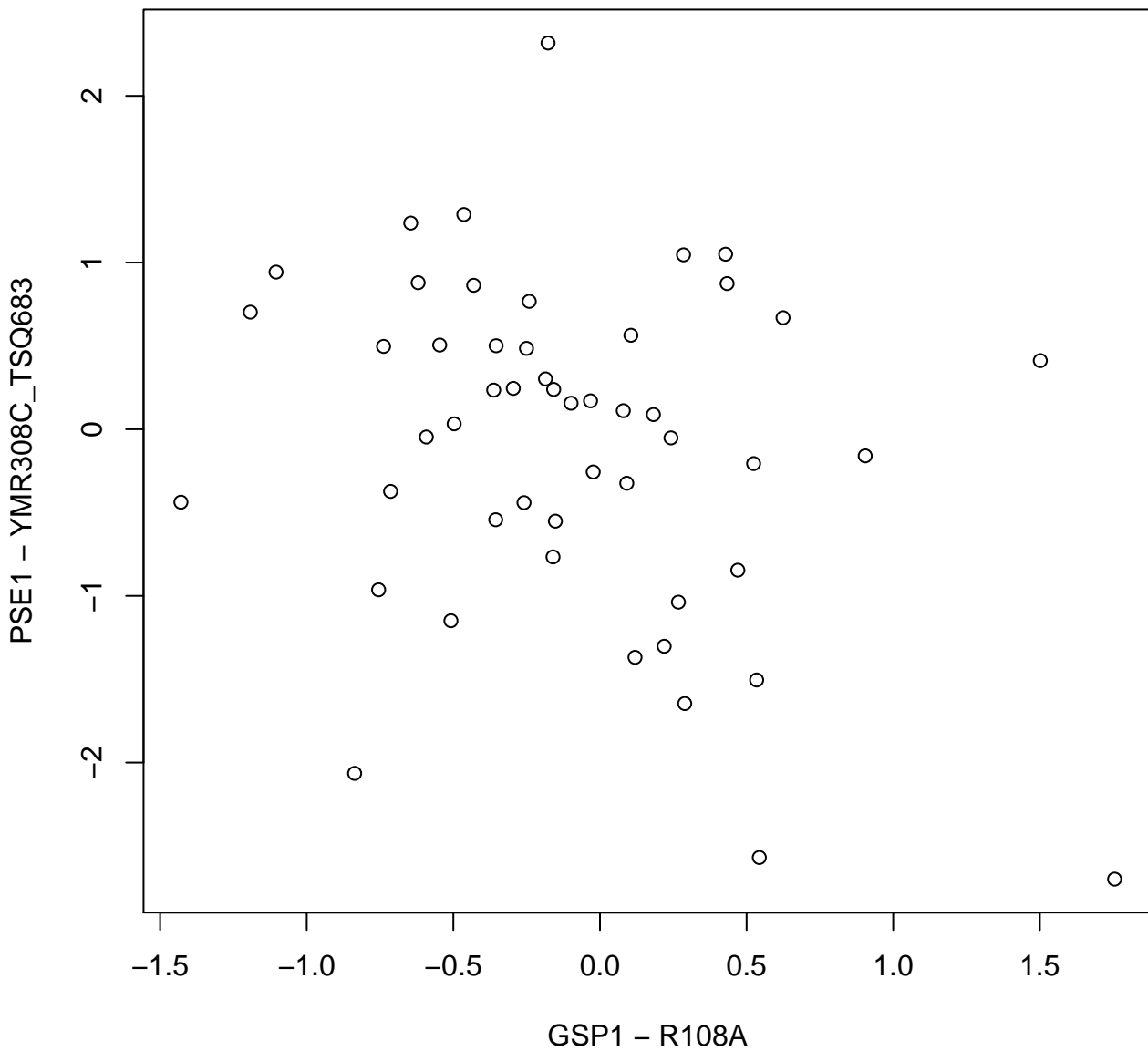
budding_GO_1



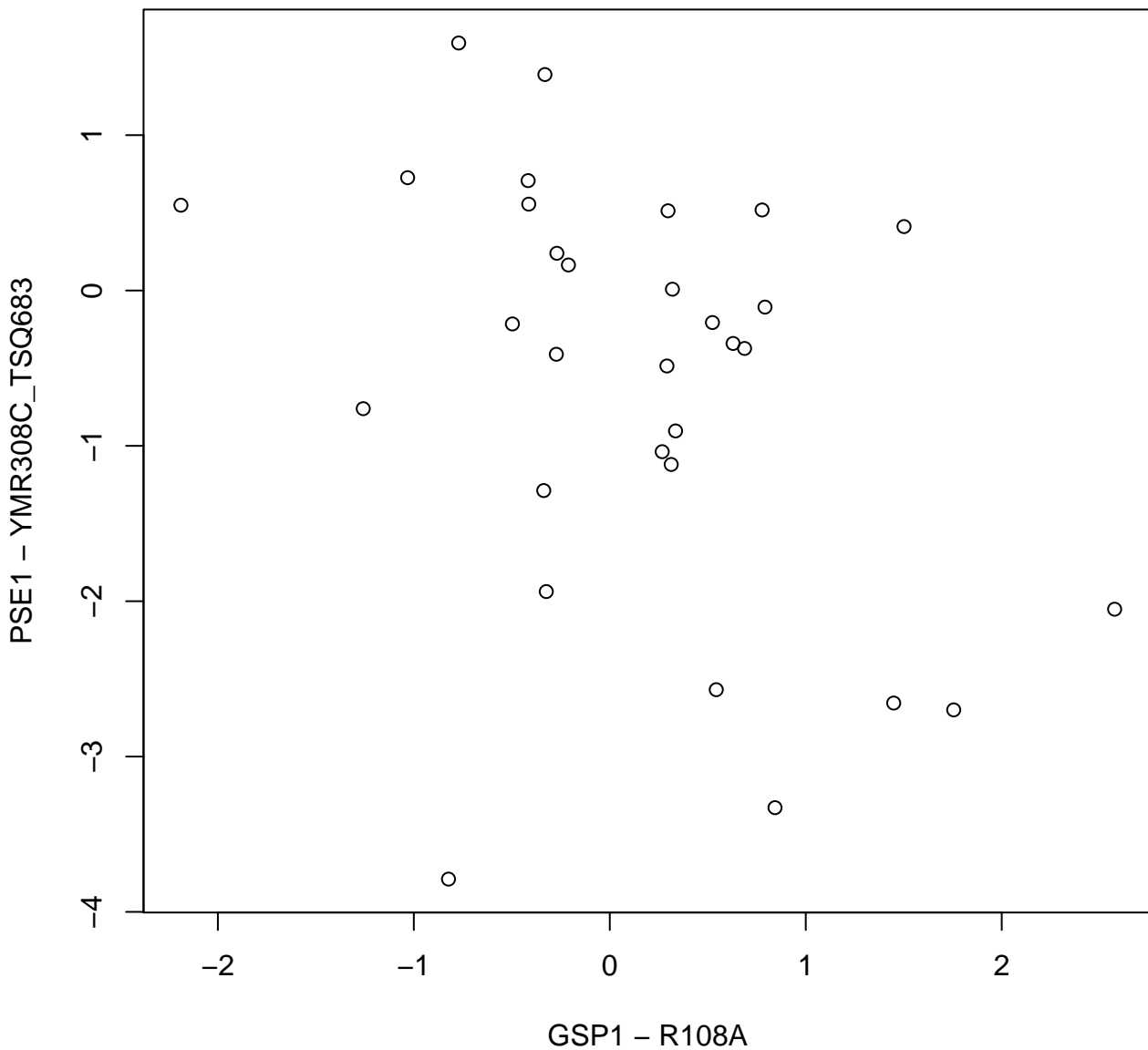
lipids_GO_2



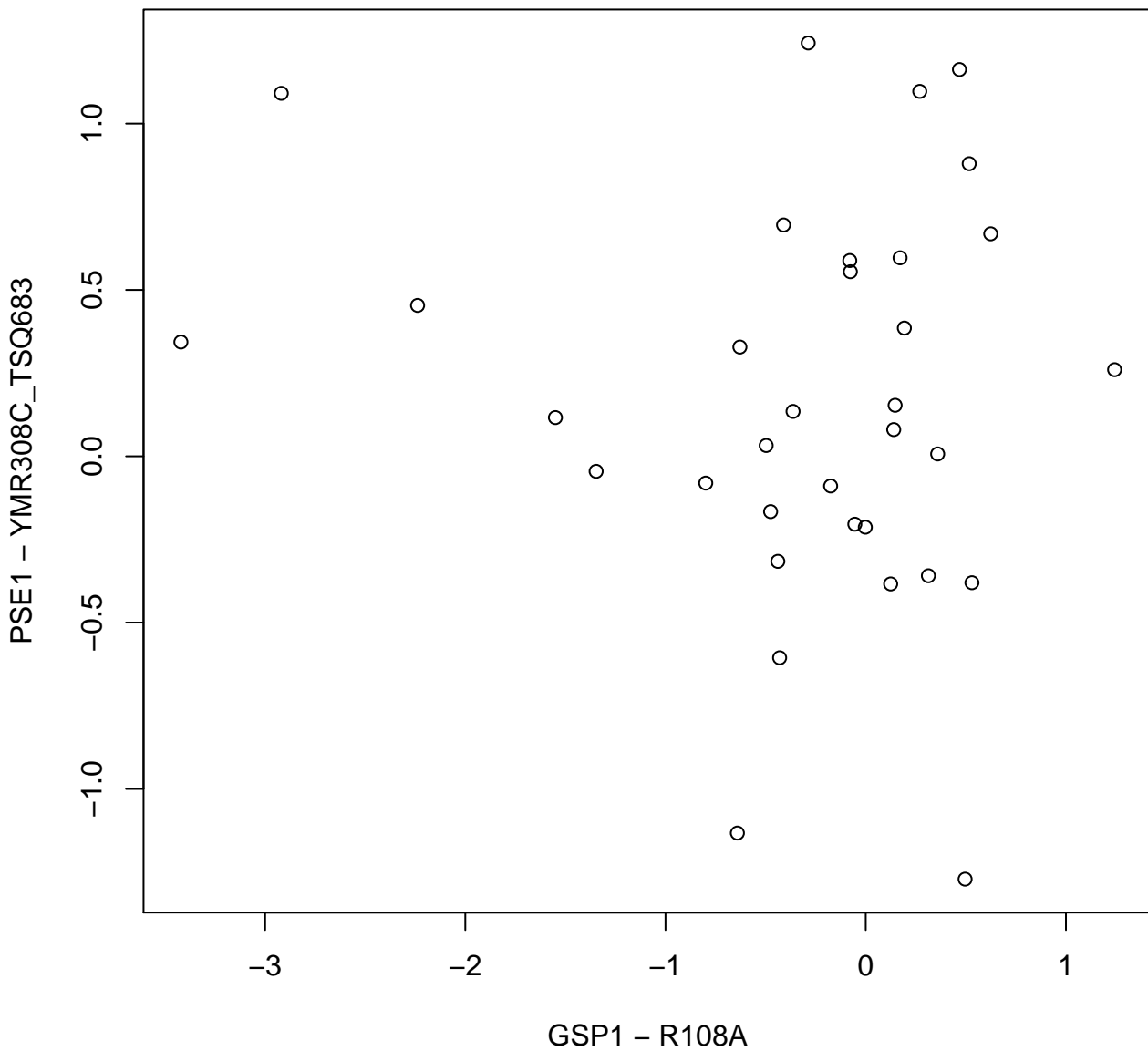
lipids_GO_1



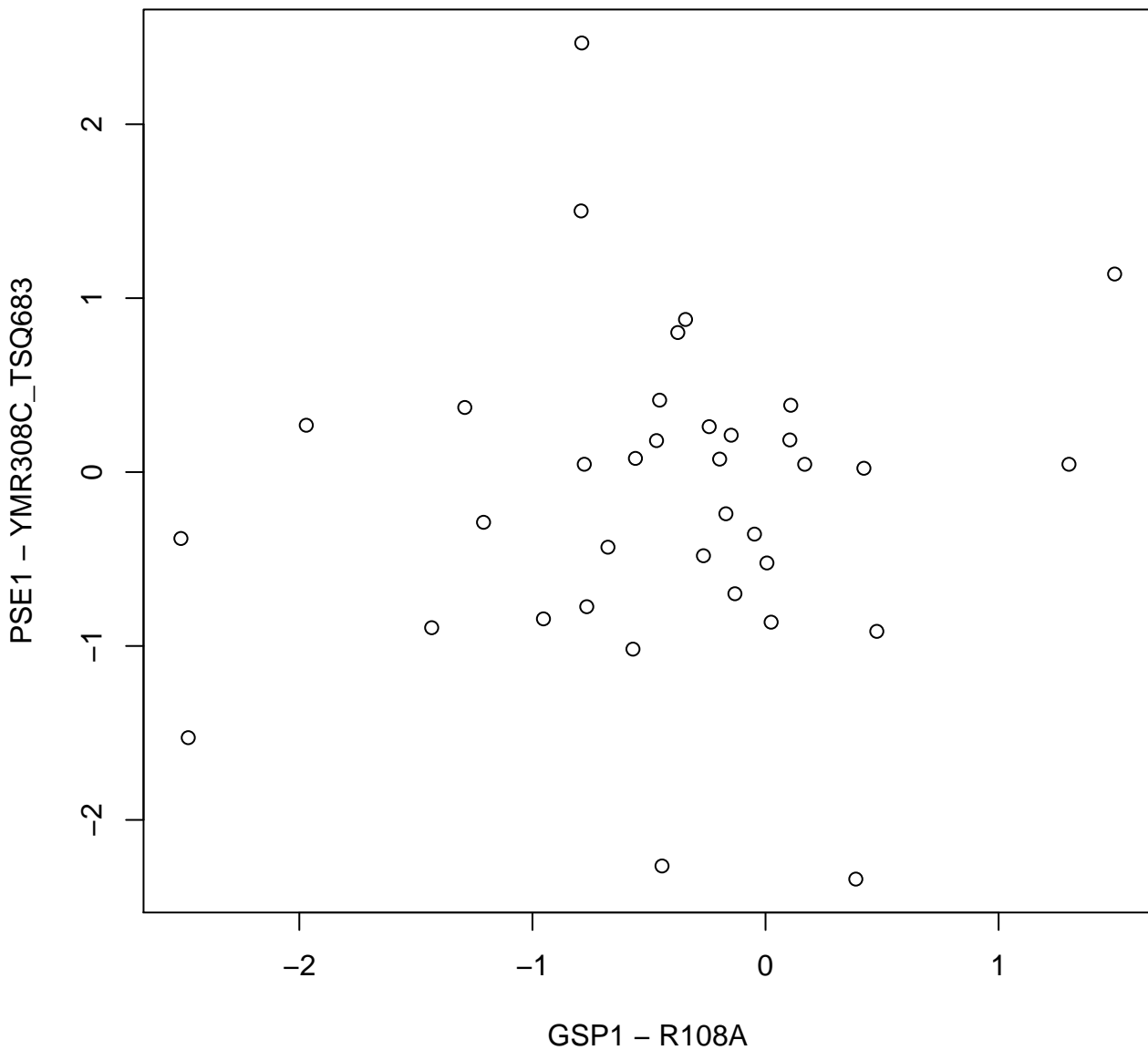
nuclear transport_GO_9



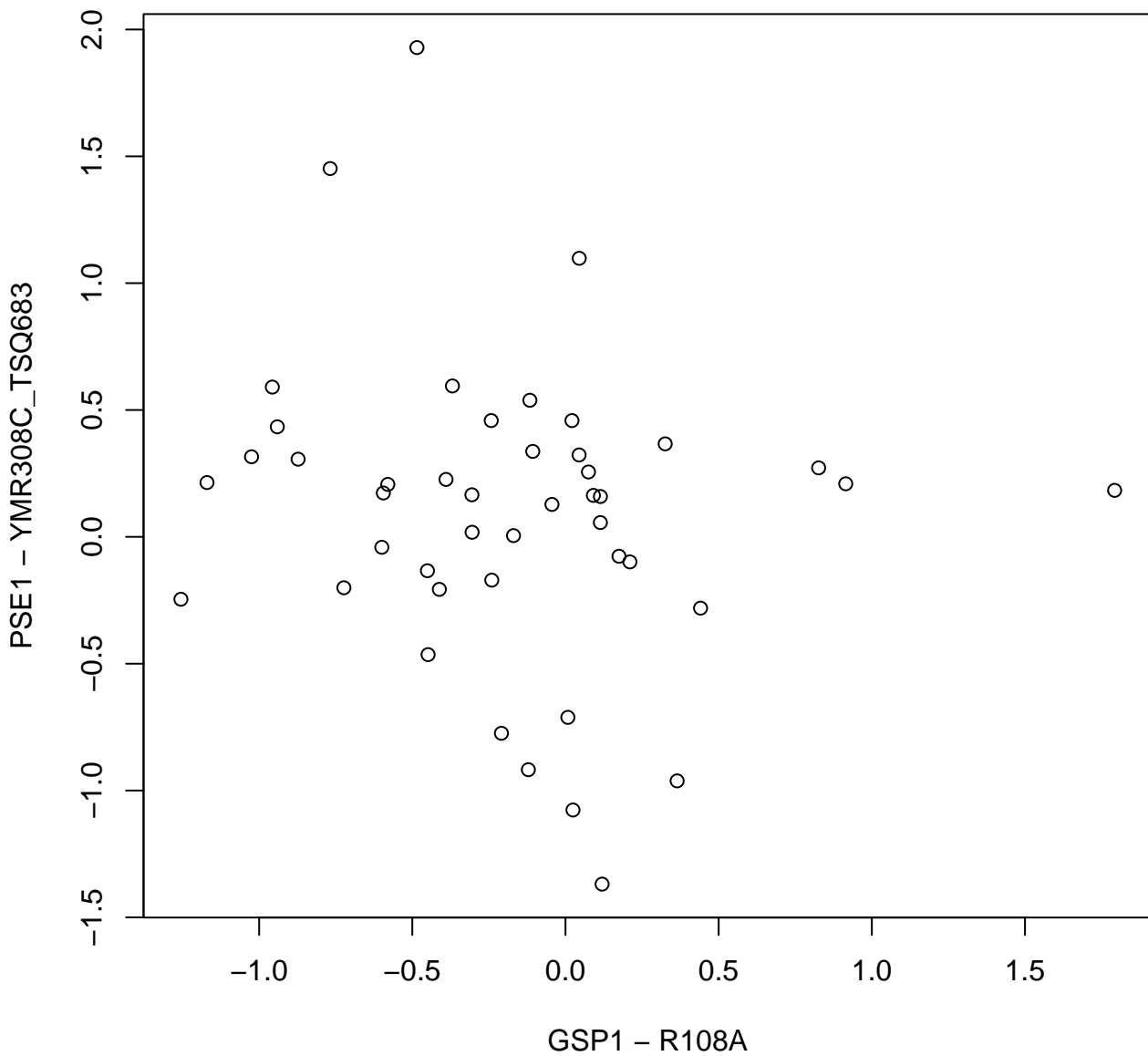
nuclear transport_GO_8



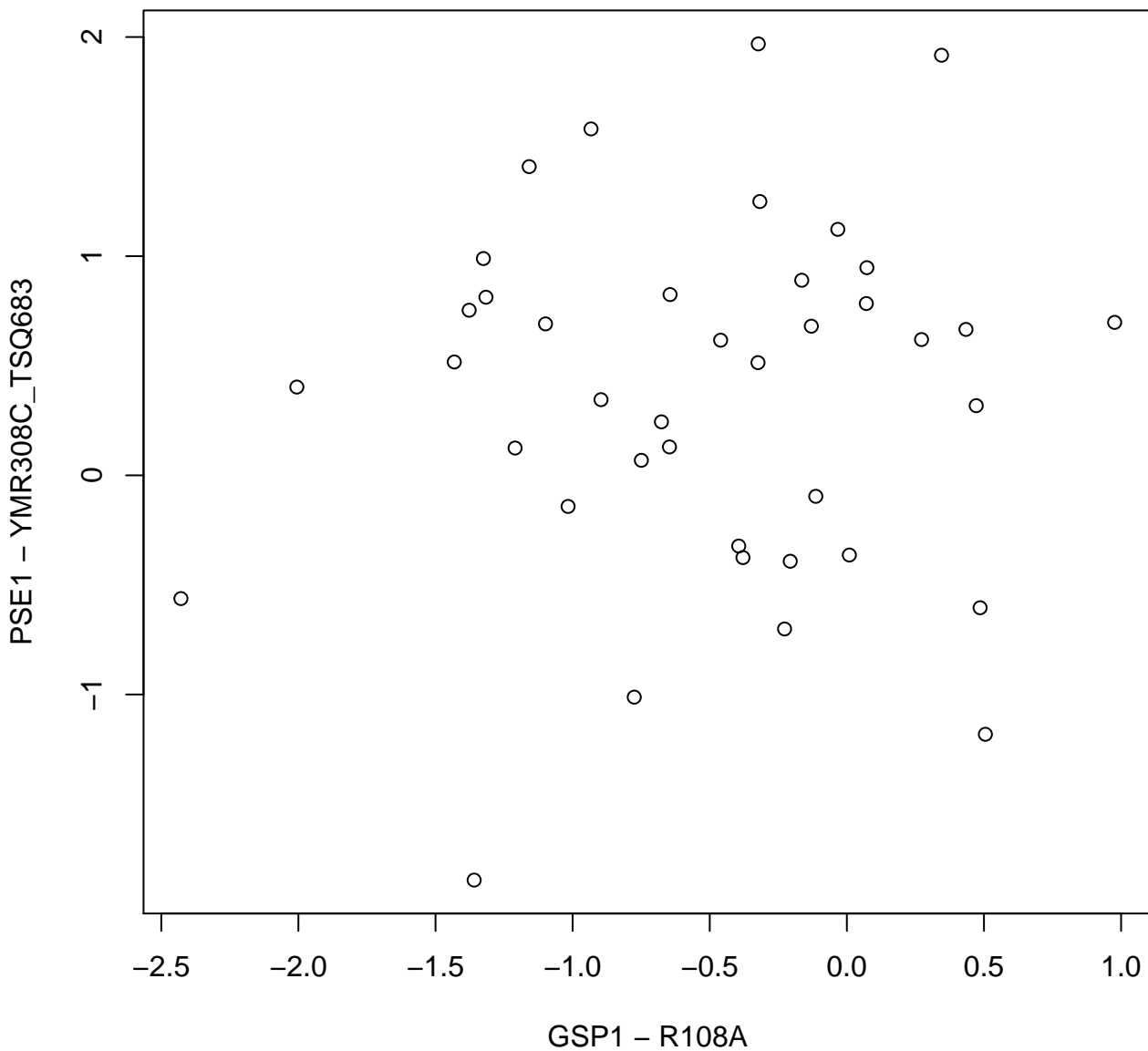
nuclear transport_GO_7



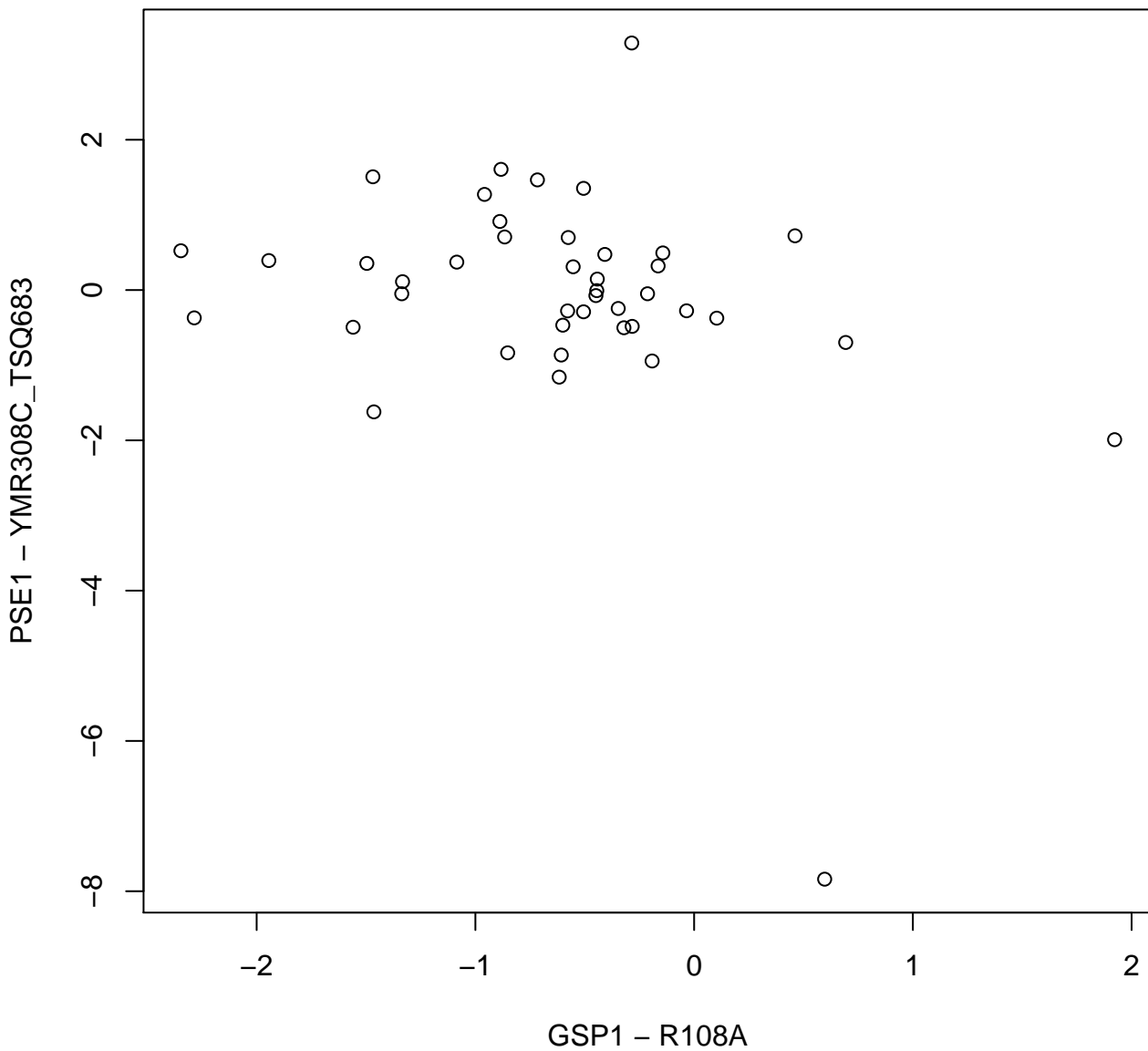
nuclear transport_GO_6



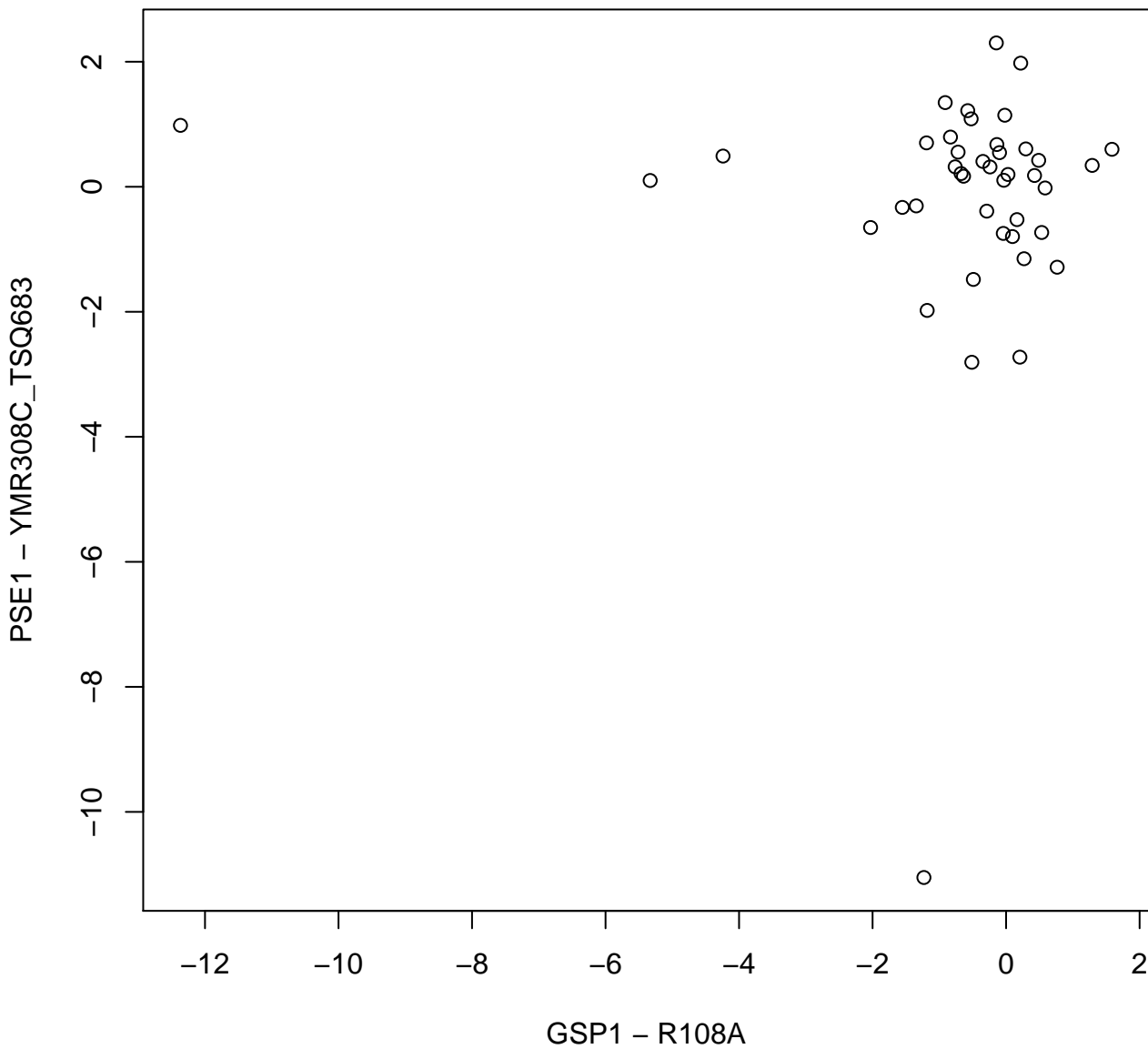
nuclear transport_GO_5



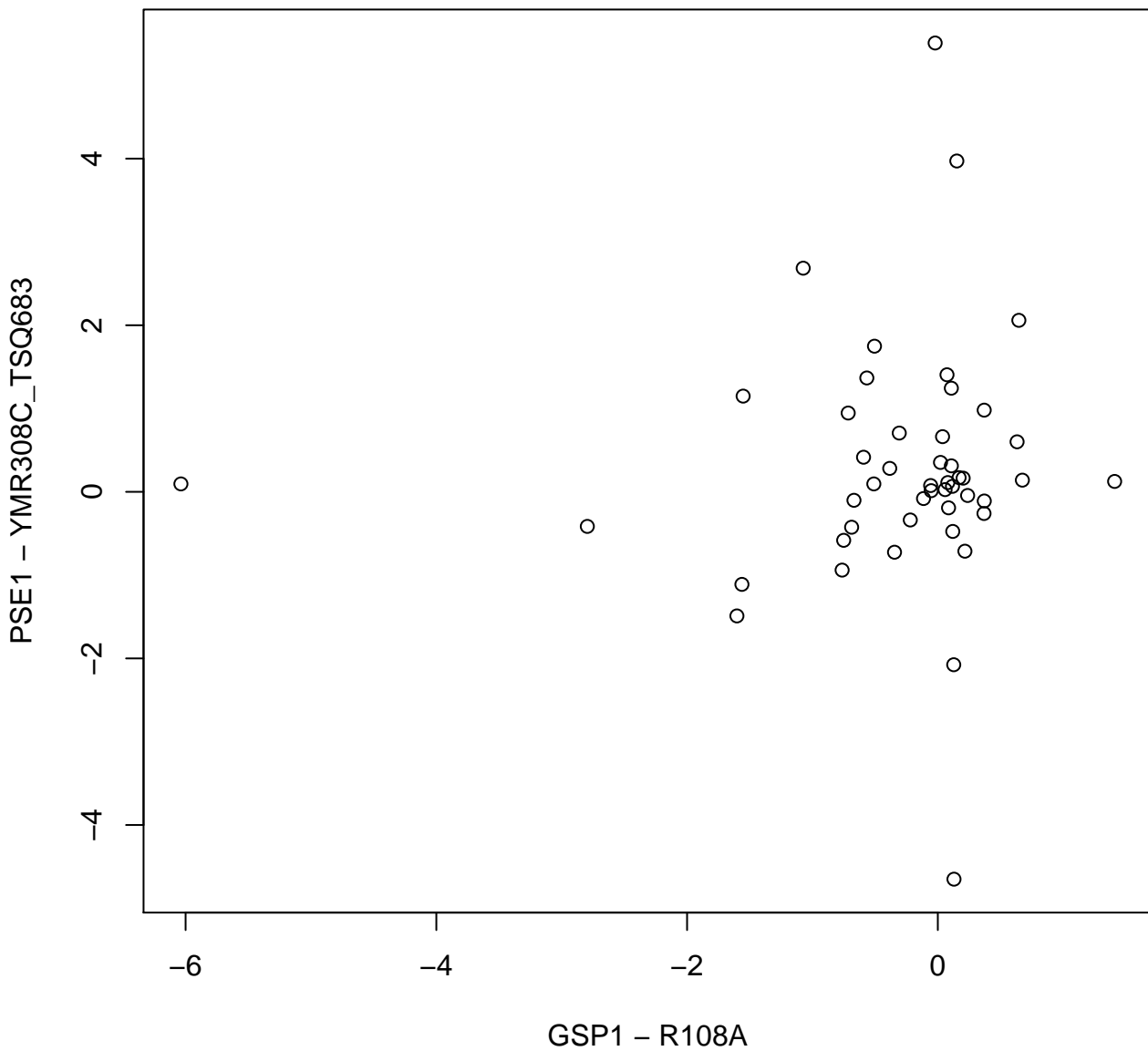
nuclear transport_GO_4



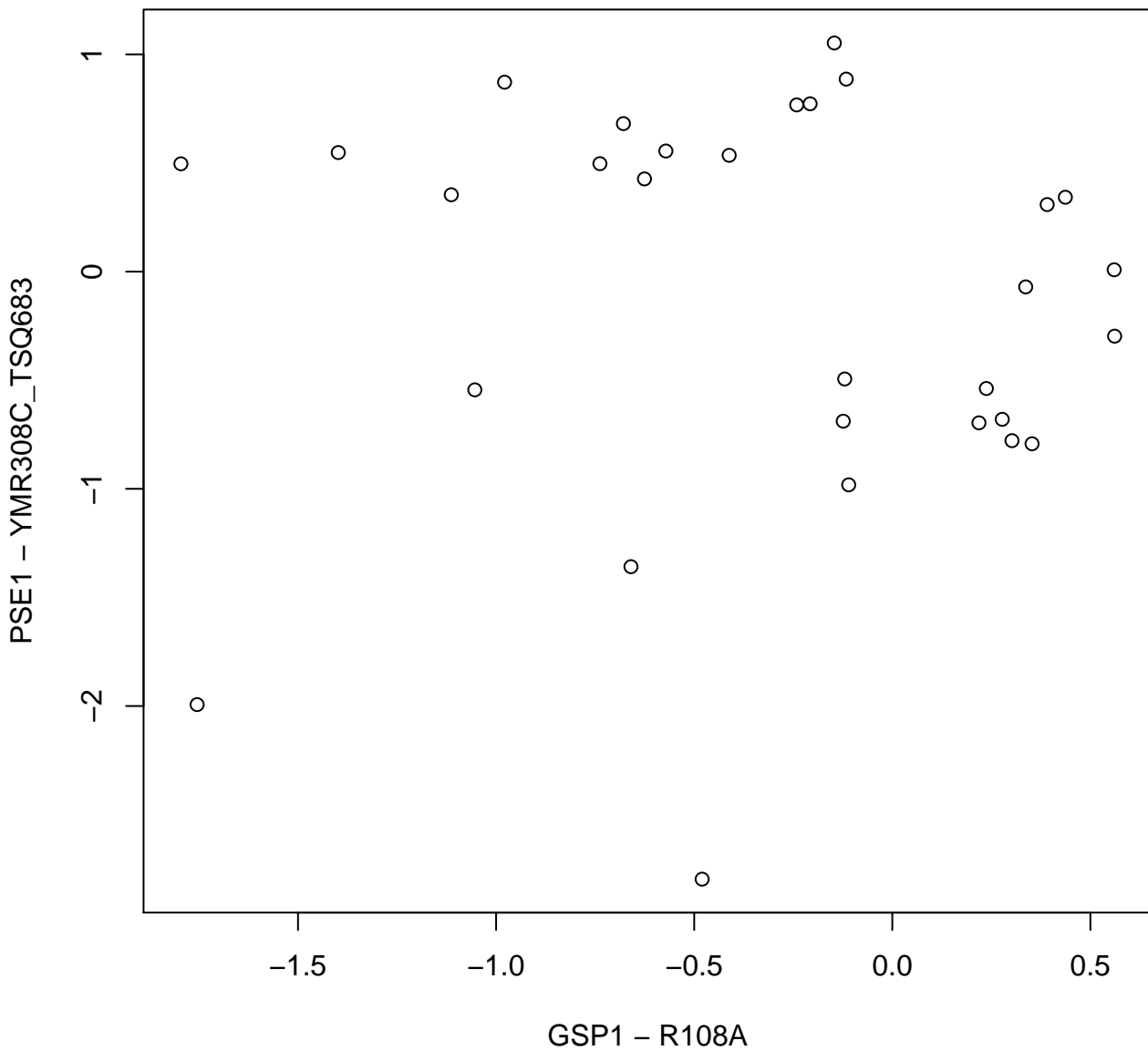
nuclear transport_GO_3



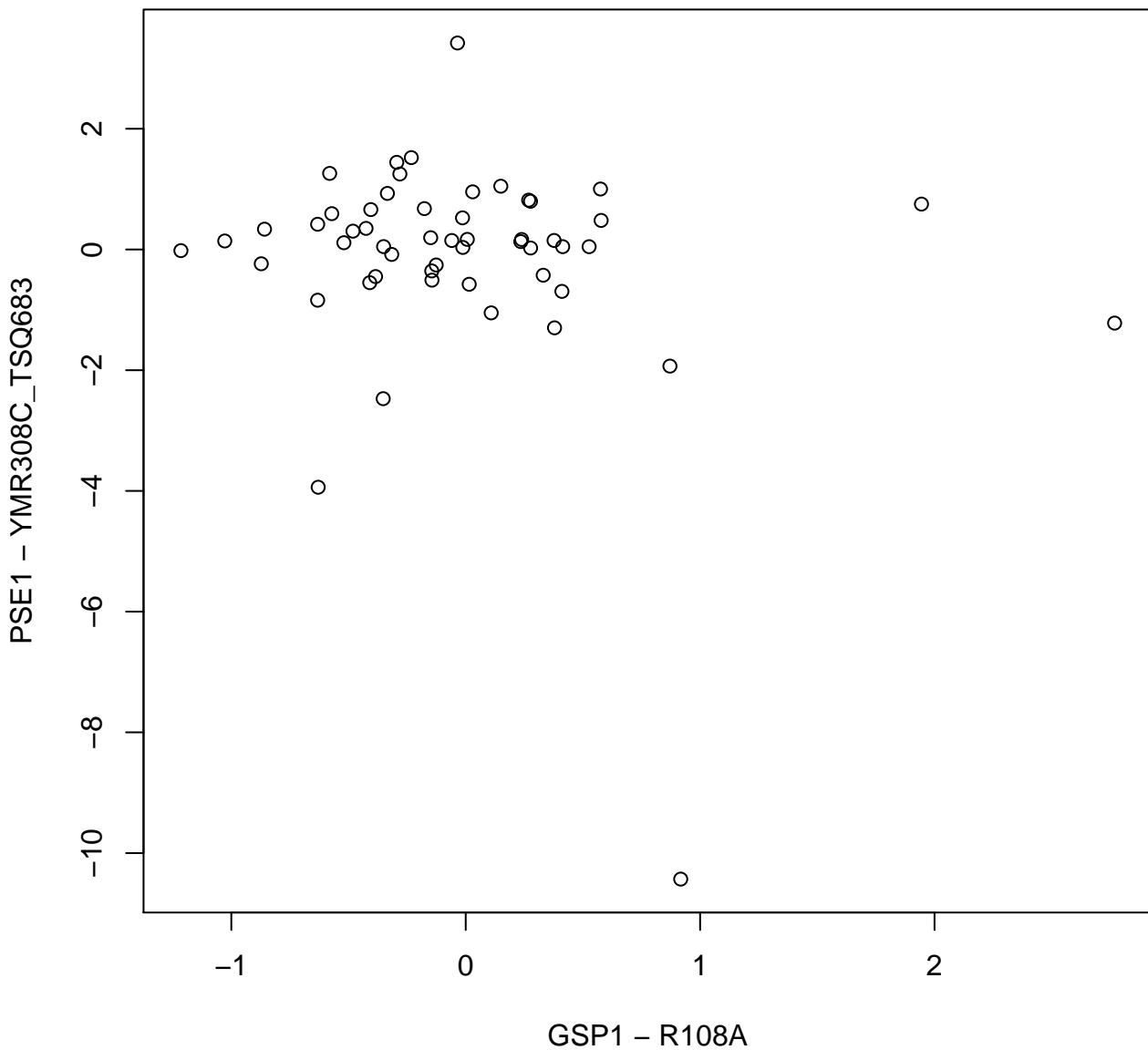
nuclear transport_GO_2



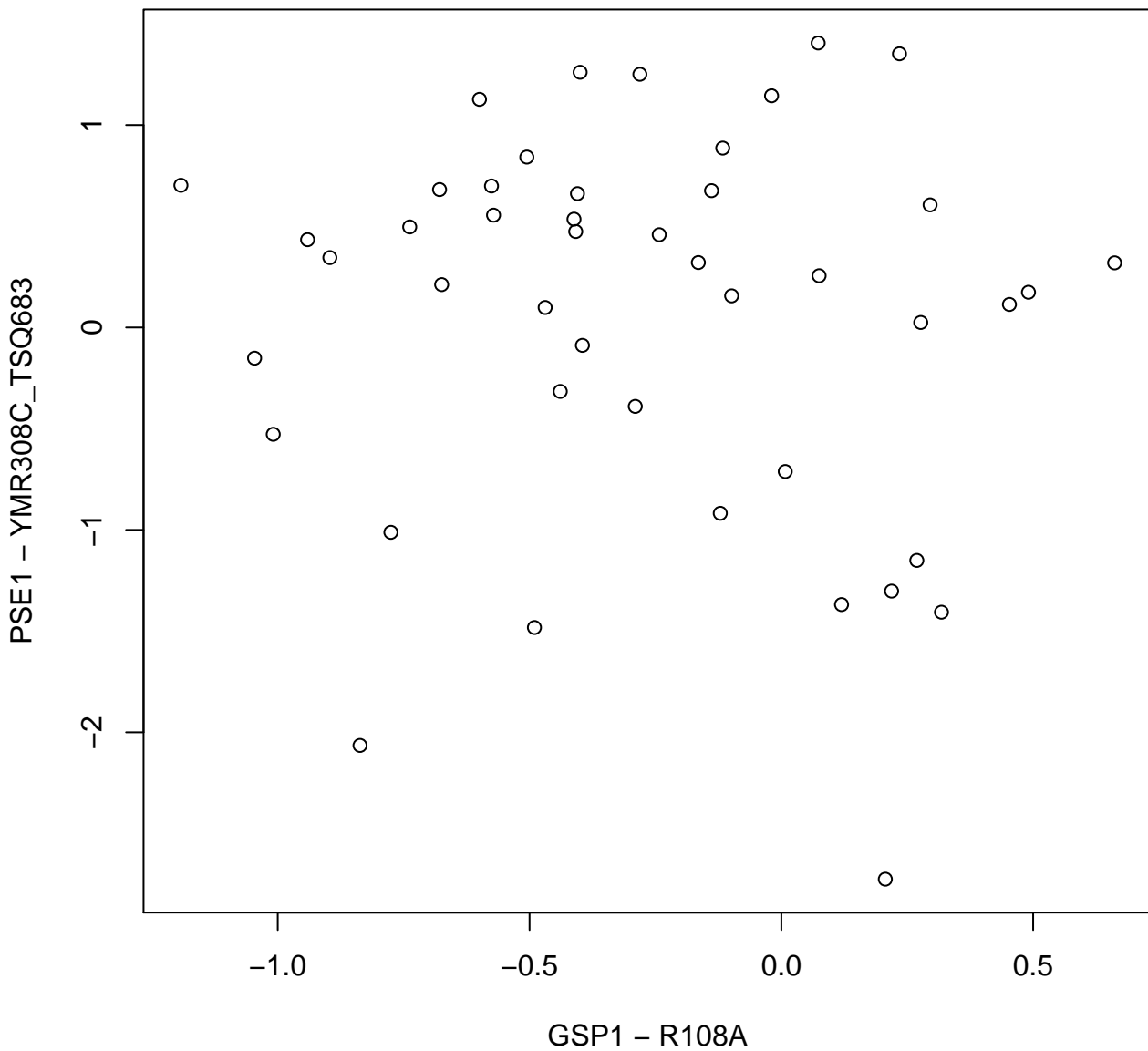
nuclear transport_GO_10



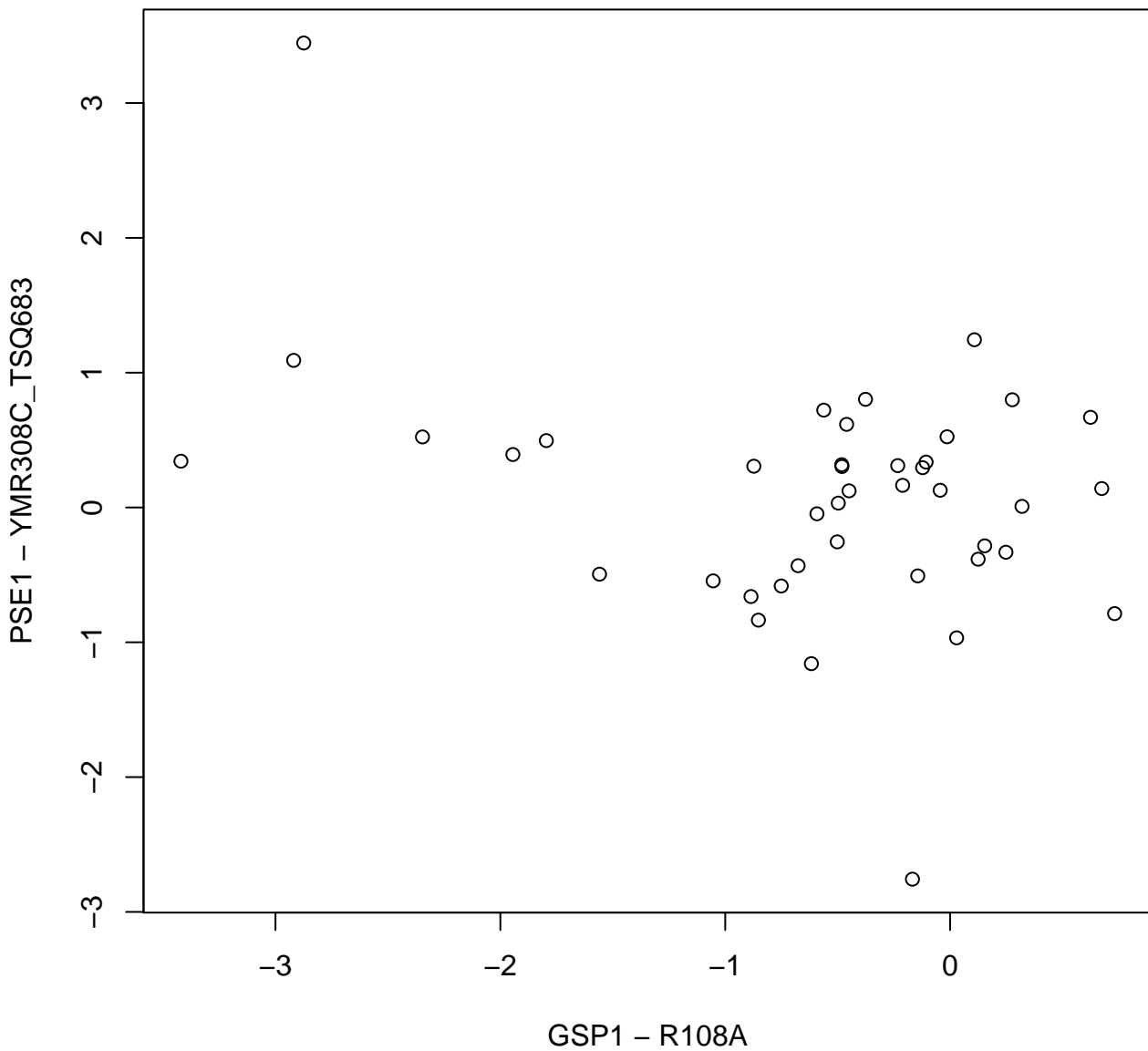
nuclear transport_GO_1



metabolic_GO_3



metabolic_GO_2



metabolic_GO_1

