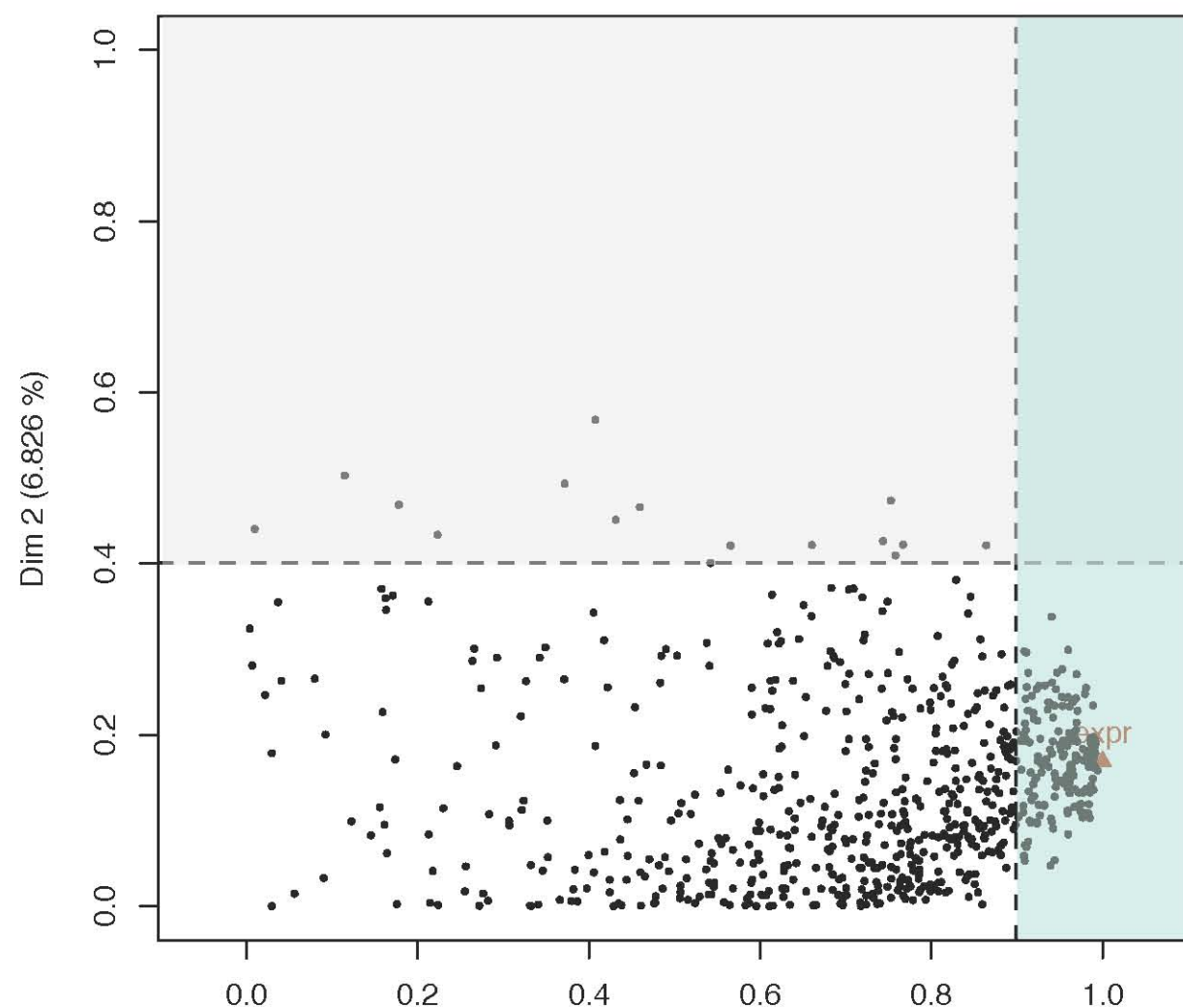


GO Biological Processes (coordinate > 0.4) correlated to axis 2

GO terms	Axis 1
cytoskeleton-dependent intracellular transport	0.578
tyrosine phosphorylation of STAT protein	0.493
bone remodeling	0.493
STAT protein nuclear translocation	0.473
peptidyl-tyrosine phosphorylation	0.473
Notch signaling pathway	0.471
meiosis I	0.465
pyrimidine deoxyribonucleoside metabolism	0.465
cortical actin cytoskeleton organization and biogenesis	0.465
regulation of smooth muscle contraction	0.465
response to extracellular stimulus	0.464
pathogenesis	0.440
positive regulation of nucleocytoplasmic transport	0.421
respiratory tube development	0.421
pyrimidine deoxyribonucleotide biosynthesis	0.421
histone acetylation	0.416
detection of stimulus	0.404



GO Biological Processes (coordinate > 0.95) correlated to axis 1

GO terms	Axis 1
* positive regulation of biological process	0.978
cell surface receptor linked signal transduction	0.970
biosynthesis	0.969
lipid metabolism	0.968
* protein transport	0.967
cell adhesion	0.966
* response to external stimulus	0.966
* apoptosis	0.965
* intracellular transport	0.965
phosphorus metabolism	0.963
* protein localization	0.963
* protein kinase cascade	0.962
immune response	0.961
positive regulation of physiological process	0.960
cell cycle	0.959
establishment of cellular localization	0.957
cellular biosynthesis	0.957
ubiquitin cycle	0.957
* cell death	0.956
response to pest, pathogen or parasite	0.956
* carboxylic acid metabolism	0.956
* small GTPase mediated signal transduction	0.956
morphogenesis	0.955
cell-cell signaling	0.954
* death	0.953
organic acid metabolism	0.953
locomotion	0.952
* programmed cell death	0.951
* cell differentiation	0.951
phosphate metabolism	0.950

* enriched term (p<0.05)