PCA of hyperLOPIT yeast data with pointer scaled to discovery probability

Legend:
- Cytosol
- ER
- Golgi
- Histone
- Mitochondrion
- Nucleus
- Phenotype 1
- Phenotype 10
- Phenotype 2
- Phenotype 3
- Phenotype 4
- Phenotype 5
- Phenotype 6
- Phenotype 7
- Phenotype 8
- Phenotype 9
- Plasma membrane
- Proteasome catalytic subunit
- Proteasome regulatory subunit
- Ribosome
- V ATPase
- Vacuole

PC1 (47.87%)

PC2 (23.32%)