Table S2. Parameters used in RedundancyMiner analyses

study	mode	.CIM file	genes in categories	genes in categories file
Retinal development	Custom	Exp1_BestClusterMap_LEIGS_KM_24.csv.join.22.txt.change.gce.CIM	.tvt file	total.txt.total.txt.total.tvt
Kinetochore genes	Custom	kinetochore.txt.change.gce.CIM	.gce file	kinetochore.txt.change.gce