spa50_LiPMScov50_ent_genes | CG | GeneOverlap R1min R5min 196 (27.1%) 174 (24.0%) 72 (9.9%) 164 (22.7%) 22 (3.0%) 29 (4.0%) 67 (9.3%)

buff	OverlapClass	Count	Overlap	Bounds	pvalue
CG	R1min	174	0.240	(0.210, 0.272)	6.706e-01
CG	R5min	72	0.099	(0.079, 0.122)	1.000e+00
CG	R2hr	67	0.093	(0.073, 0.113)	3.908e-01
CG	R1min-R5min	196	0.271	(0.239, 0.302)	9.999e-05
CG	R1min-R2hr	29	0.040	(0.026, 0.055)	1.000e+00
CG	R5min-R2hr	22	0.030	(0.019, 0.043)	9.974e-01
CG	R1min-R5min-R2hr	164	0.227	(0.198, 0.258)	9.999e-05

R2hr