

Window Size (SNPs)	Corner	GO biological process	REFLIST	#Found Genes	Expected Genes	Over/under	Fold enrichment	Raw p-value	False Discovery Rate
1000	1	endocytosis (GO:0006897)	527	13	2.53	+	5.14	2.00E-06	5.20E-03
	1	import into cell (GO:0098657)	571	14	2.74	+	5.11	8.47E-07	4.40E-03
	2	NO SIGNIFICANT ENRICHMENT	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	3	import into cell (GO:0098657)	571	16	2.48	+	6.45	5.22E-09	2.71E-05
	3	endocytosis (GO:0006897)	527	14	2.29	+	6.11	9.64E-08	2.51E-04
	3	vesicle-mediated transport (GO:0016192)	903	14	3.92	+	3.57	4.32E-05	3.74E-02
	3	transport (GO:0006810)	2895	32	12.58	+	2.54	5.18E-07	8.98E-04
	3	establishment of localization (GO:0051234)	2939	32	12.77	+	2.51	7.22E-07	9.39E-04
	3	localization (GO:0051179)	3075	32	13.37	+	2.39	2.53E-06	2.63E-03
1500	1	NO SIGNIFICANT ENRICHMENT	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	2	NO SIGNIFICANT ENRICHMENT	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	3	endocytosis (GO:0006897)	527	15	2.01	+	7.46	2.17E-09	5.64E-06
	3	import into cell (GO:0098657)	571	16	2.18	+	7.34	7.47E-10	3.88E-06
	3	vesicle-mediated transport (GO:0016192)	903	15	3.45	+	4.35	1.99E-06	3.45E-03
	3	transport (GO:0006810)	2895	27	11.05	+	2.44	8.67E-06	1.13E-02
	3	establishment of localization (GO:0051234)	2939	27	11.22	+	2.41	1.77E-05	1.84E-02
	3	localization (GO:0051179)	3075	27	11.74	+	2.3	2.92E-05	2.53E-02
2500	1	endocytosis (GO:0006897)	527	12	1.47	+	8.14	3.06E-08	7.95E-05
	1	import into cell (GO:0098657)	571	13	1.6	+	8.14	7.64E-09	3.97E-05
	1	vesicle-mediated transport (GO:0016192)	903	13	2.53	+	5.15	1.36E-06	2.35E-03
	1	transport (GO:0006810)	2895	21	8.1	+	2.59	3.14E-05	4.08E-02
	1	establishment of localization (GO:0051234)	2939	21	8.22	+	2.55	3.93E-05	4.08E-02
	2	NO SIGNIFICANT ENRICHMENT	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	3	cell adhesion (GO:0007155)	465	14	1.88	+	7.45	8.38E-09	4.35E-05