Genes gained in CYCO1 BP TreeMap

tricarboxylic a	cid cycle	/cle respons		mor cell	nucleoside diphosphate			AMP metabolic process		UDP metabolic process		riboi bio	rimidine nucleotide synthetic rocess	positive regulation of programmed cell death		of receptor
					biosynthetic process pyrimidine			line ribonucleotide metabolic p pyrimidine nucleobase biosynthetic process					metabolic process	positive regulation of glial cell migration	positive regulation of tumor necrosis factor–mediated signaling pathway	regulation of antigen receptor–mediated signaling pathway
positive regulation membrane permea	of cyste bility endop	vation of eine-type peptidase involved in tic process		vater eostasis								pyrimic trij meta	line nucleoside phosphate bolic process	regulation regulation of cell death	regulation of of glial cell pro factor-mediated signaling pathway	· .
					nucleoside monophosphate phosphorylation			ADP metabolic process		pyrimidine ribonucleotide metabolic process pyrimidine ribonucleosid triphosphate metabolic process		eoside meta phate olic ess de	yribonucleotide bolic process oxyribose hosphate	positive regulation of glial cell proliferation	regulation of glial cell proliferation	epidermal growth
dicarboxylic acid metabolic process	linid hor	ic process neostasis		netabolic ocess	water transport		pigment granule organization		organic ca		protein sec	retion	etrograde transport,	regulation of glial cell migration	glial cell proliferation	T cell receptor signaling pathway
renal system process	regulation o	regulate of rece	ptor	aintenance of lood-brain					transport			•	endosome to Golgi cilium	interleukin–2 production		regulation of interleukin–2 production
		proce		barrier			G	recept end mintransmembrane trans		s	protein targeting to lysosome	cell-cell adhesion	movement involved in cell motility		regulation of sterferon–gamma	cytokine production involved
aerobic electron transport chain	methylation	ovipositio	n	development	lysosome	transmembrane transport		transport	Wnt protei	า	protein localization	cilium-dependent	via plasma membrane	regulation of interleukin-12		2 production response
mitochondrial ATP synthesis coupled electron transport	germ cell proliferation	metencephalon development	receptor catabolic process	to virus	organization			etory granule ganization	cell-cell adhes via plasma-memb adhesion molec	sion orane	cilium or flagellum-dependent cell motility	proteir	rotein refolding	positive regulation cy of cytokine pro production involved invo	production	regulation of interleukin–12 action production