Genes gained in OVER1 BP TreeMap

collagen catabolic process		arginine metabolic proce	am	nine family ino acid olic process	chondrocyte proliferation	mitochondrial outer membrane permeabilization	lamellipodium I	small GTPas nediated sigr transductior	al matrix	protein	protein phosphorylation
					mitochondrion transport along microtubule	cellular response to environ regulation of mitochondrion organization	structure organization	cellular response environmer	ital or		_
glycine metabolic process	regulation of translation		acy	/I–CoA		cellular response to radiation	cellular	stimulus	depolymeriz	peptidyl-serin phosphorylatic	
				lic process			response to abiotic stimulus	external encapsulating structure organization			egulation of
	prote	protein maturation			endochondral ossification	embryonic cranial skeleton morphogenesis		bra	branching involved in labyrinthine layer morphogenesis		phosphorylation
nitrogen cycle metabolic process	regulation of alternative mRNA splicing, via spliceosome		fatty acid	d oxidation			liver developm	ent i			n
						animal organ rege animal organ regeneration				positive regulation	of phosphata
	translational initiation		peptidyl-serine	fatty acid			eneration development	layer	placent	1	
bile acid biosynthetic process			modification	metabolic process	face morphogenesis			morphogen	esis morphoger	positive	positive regulation of protein
	prote	ein processing				in utero embryonic	bone cell			regulation of phosphorus	
	protein processing		protein maturation			development	development	megakaryocyte differentiation		metabolic process	tyrosine kinase activity