Genes gained in OVER1 BP TreeMap

skeletal system morphogenesis	face development	face morphogenesis		negative regulation of osteoclast differentiation		cartilage development	hyaluronan catabolic process	U2–type prespliceosom assembly	ie RNA	\ capping	water transpo	regulation resport of reception clustering	
	Va	asculature developm mammalian	i <u>ent</u> ral	connective tissue development		vasculature development						of	establishment of protein localization
smooth muscle tissue development	in utero embryonic development	oogenesis stage	ovarian follicle growth	regulation of osteoclast differentiation		ovarian follicle development	nucleic acionspliceosomal snRNP assembly	l phosphodiester b	trans and hydrolysis RNA p	RNA polymerase		n targeting to pero	
		multicellular organism aging		negative regulation of myeloid leukocyte differentiation		osteoclast differentiation		hyaluronan biosynthetic	glycosaminoglyca	nucleic acid			argeting to eroxisome
cellular response to UV-B	response to antibiotic cellular response to tumor necrosis factor	cellular response to interleukin–1			response to pH	cellular response to platelet–derived growth factor stimulus	3'-UTR-mediated mRNA stabilization	hyaluronan metabolic process	nyaluronan		positive regu	isomal transport protein import into nucleus	
		response to antibiot fibroblast growth factor stimulus	platelet-	onse to derived n factor	response to molecule of bacterial origin	regulation of inflammatory response	fibroblast migration	regulation of neutrophil chemotaxis	regulatio	cell motility tr		negative regulation of G1/S transition ation of G1/S of mitotic cell cycle	
		cellular response to pH	response to lipopolysaccharide		negative regulation o BMP signalin pathway		negative regulation of cell migration	neutrophil re	negative reg	ulocyte of	of mitotic	1/S transition positive regulation of cell cycle G1/S	