Genes lost in PELE1 MF TreeMap

RNA-directed DNA polymerase activity DNA-directed I		DNA-binding transcription factor activity, RNA polymerase II-specific		aspartic-type	nvpria	neuropeptide hormone activity		growth factor		ein kinase binding		neuropeptide	e he	heparin binding	
				endopeptidase activity	ribonuclease activity			activity		erleukin–1 oinding	odorant binding	binding			
		DNA polymerase ac activity	DNA-binding transcriptic activator activity, RNA Ctivity II-specific	kinase kinase activity	olinesterase activity protein-membrane adaptor activity	SMAD binding	K63–linked	scaffold protein binding d protein binding phosphatase binding	transmembra receptor pro- serine/threor kinase bindi	ein receptor binding DNA-binding transcription	calcium ion binding	phosphatidylserine pe odorant bindin		zinc ion binding	
endonuclease activity		DNA-binding transcription repressor activity, RNA polymerase II-specific	DNA polymerase activity protein tyrosine	NAD+ nucle nucleosidase c activity ADP gen	AD+ cotidase, cyclic 2-ribose erating	receptor tyrosine kinase binding	polyubiquitin modification-depende protein binding	ion channel binding	acetylcholine receptor do	binding BH3 glucocorticoid receptor binding	DNA-binding transcription factor activity	immunoglobulin binding	structural constituent of cuticle	cargo receptor activity	
		RNA-directed 5'-3' RNA polymerase	phosphatase activity NAD(P)+ nucleosidase	acetylcholinesterase very activity 3-sy 17-beta-hydroxysteroid dehydrogenase cy	rry-long-chain -ketoacyl-CoA rnthase activity activity	calcium-dependent	Notch binding transforming	protein-coupled i receptor binding	immunoglobulin receptor binding	receptor nuclear antagonist receptor activity binding	dynein complex	toxin activity	galactoside binding	lgA binding	
		activity	activity (NAD		activity involved in poptotic signaling pathway abrane Signaling	protein binding	growth factor beta binding	II-specific DNA-binding	growth factor receptor binding	epidermal growth factor receptor binding	binding	calcium–dependent phospholipid binding	structural constituent of virion	IgG binding	
	neuropeptide Y receptor activity	receptor activity	ceptor positivity re	receptor receptor receptor receptor receptor rectivity erotonin	receptor activity	voltage–gated potassium channel activity cation channel a		neurotransmitte transmembrane transporter activity	cation seq	RNA polymeras	y RNA	lipoprotein lipase activator	signaling receptor inhibitor	nentidase	
G protein-coupled receptor activity		G protein-coupled purinergic chemo	noattractant otor activity	eceptor prot activity tyrosine acti	ein protein-coupled amine receptor activity			kainate selective glutamate		region sequence-spec DNA binding	stem-loop ic binding	activity	activity peptidase		
	neurotransmitter	signaling receptor a	eceptor neu	transme postsynaptic receptor urotransmitter tyros peptor activity	receptor receptor			calcium		RNA polymerase	ecific DNA bindin	g peptidase activity	activator		
tumor necrosis factor–activated receptor activity	interleukin–1 receptor activity	receptor cher activity rec	ceptor ctivity	eceptor E reci	glandin prostanoid	acetylcholine transmembrane transporter activity	extracellular	on	transmitter-gated channel activity potassium	transcription regulatory region sequence—specion DNA binding		serine-type endopeptidase inhibitor activity	ndopeptidase regulator activity	cysteine-type endopeptidase regulator activity involved in apoptotic process	
		olfactory receptor G prote se	ein-coupled trar erotonin rec	activity nsmembrane ceptor protein hosphatase activity activity prostag rece acti	glandin immune receptor		channel activition ligand-gated sodium channel activity	transmitter-gated ion channel activity hem	channel activity neme transmembrane transporter activity	sequence-spec	cis-regulatory region double-strand telomeric sequence-specific	nentidase	cysteine-type endopeptidase activator activity involved in apoptotic process	peptidase activator activity involved in apoptotic process	