Genes lost in HRPE1 MF TreeMap

phosphatidylinositol 3–kinase binding		co-receptor binding		lipopolysaccharide binding	structural constituent of virion	endopeptidas inhibitor activity	se peptidoglyca binding	acetylcholir an receptor regulator activity	piRNA binding	lipopolysaccharide immune receptor activity		une	transmembrane signaling receptor activity	
					bubble DNA binding	activity activity element binding								
receptor	pr	oteoglycan	TIR domain binding	DNA-binding transcription factor activity, RNA polymerase II-specific IgG binding	nutrient res RNA polymerase II transcription regulatory region sequence–specific DNA binding	DNA-binding transcription repressor activity, RNA polymerase II-specific	y biotin binding	nutrient reservoir activity	cyclic–GMP–AMP binding	transmembrar neurotransmitter receptor activity	orane sig er nı	tein-coupled naling rece unitergic ucleotide ptor activity	d ptor activity ^{ie} receptor activity	
antagonist activity		binding					metalloendopeptidase inhibitor activity	ligand-activated transcription factor activity	signaling receptor complex adaptor activity		G prote	ein-coupled	rotransmitter eptor activity neuropeptide	
sigr	naling receptors	otor binding	_{nma} antigen		scavenger receptor activity		translation repressor activity	nuclear receptor activity	regulatory RNA binding	G protein–couple receptor	a d	ctivity	involved in regulation receptor of postsynaptic activity	
peptide hormone receptor binding	growth fact	interferon-gamma receptor binding						RNA-DNA hybrid	endodeoxyribonuclease	activity		ctivity	GABA-A receptor activity	
laminin–1 binding	CARD domain	protease	signaling receptor					ribonuclease activity	1000	extracellula	ar lig	excitatory extracellular gand-gated ic hannel activit	activity	
	binding	binding epidermal	binding		endonuclea(endonuc l	ase activity		endodeoxyribonuclea activity, producing 5'-phosphomonoeste	activity, producing	ligand-gated channel acti	/ity ac	acetylcholine-gated cation-selective channel activity involved in regulation of postsynaptic ion channel activity by normal potential		
interleukin-7 caspase receptor binding binding			phosphatidylinositol 3-kinase catalytic subunit binding						tRNA-intron endonuclease		trans	ansmitter-gate	nucleobase d transmembrane transporter	
cytokine binding	growth fac	interleukin– binding	death receptor binding					endonuclease activity, active with either ribo- or deoxyribonucleic acids and producing 3'-phosphomonoesters	chalcium activated channel activity activity	annel	ansmitter-gate ion channel activity	activity d purine nucleobase transmembrane transporter activity		