Genes lost in HRPE1 MF TreeMap

			NAD(P)+ nucleosidase activity			phosphatidylinositol 3–kinase binding	receptor antagonist activity		Tat protein binding	virion binding	lipopolysaccharide immune receptor activity		neuropeptide Y receptor activity
RNA-directed DNA polymerase activity			NAD+ nucleotidase, cyclic ADP-ribose generating			phosp TIR domain binding	proteoglycan hatidylinositol 3	translation initiation -kinase binding	death	interleukin–1 binding	neuropeptide recentor cytokine recep activity		interleukin–1 I receptor Ictivity activity
			glycine N-choloyltransferase activity				caspase binding	receptor	antigen	laminin-1	angiotensii	G protein-coupled	cytokine
		į,					binding	binding			receptor recep	purinergic nucleotide receptor activity	receptor activity
peptidoglycan muralytic act			:ivity			co-receptor binding	CARD domain binding		hinding	glutamate receptor binding	nucleotide receptor	receptor	transmembrane receptor protein nosphatase activity
very long chain acyl–CoA hydrolase activity choloyl–CoA hydrolase activity		myrid	myristoyl–CoA hydrolase activity		nd CoA						receptor activity	activity	protein tyrosine phosphatase activity
	medium–chain acyl–CoA hydrolase activity				oyl-CoA se activity					NA hybrid ease activity	DNA single-stranded replication DNA binding origin binding		sequence-specific double-stranded DNA binding
			rtic–type	DNA helicase activity	RNA–directed 5'–3' RNA polymerase activity	endonucl	ea <mark>endonuclease</mark>	endonuclease activity			sequence-specific -	DNA binding	siRNA binding
	galactosylceramidase activity	endopep	endopeptidase activity		lysozyme activity				endodeoxyribonucl activity, producir 5'-phosphomonoe:	g or deoxyribonucleic acids	mkina binding	NA secondary structure	transcription cis-regulatory
		protein tyrosine phosphata activity	DNA-directed DNA ase polymerase	peptidoglycan muralytic activity	N-acyltransferase activity				activity	yribonuclease , producing nomonoesters	binding	binding RNA polymerase II transcription regulatory region sequence–specific DNA binding	transcription regulatory region nucleic acid binding