

Ontology Analysis of Infectious Disease Genes at the Rat Genome Database



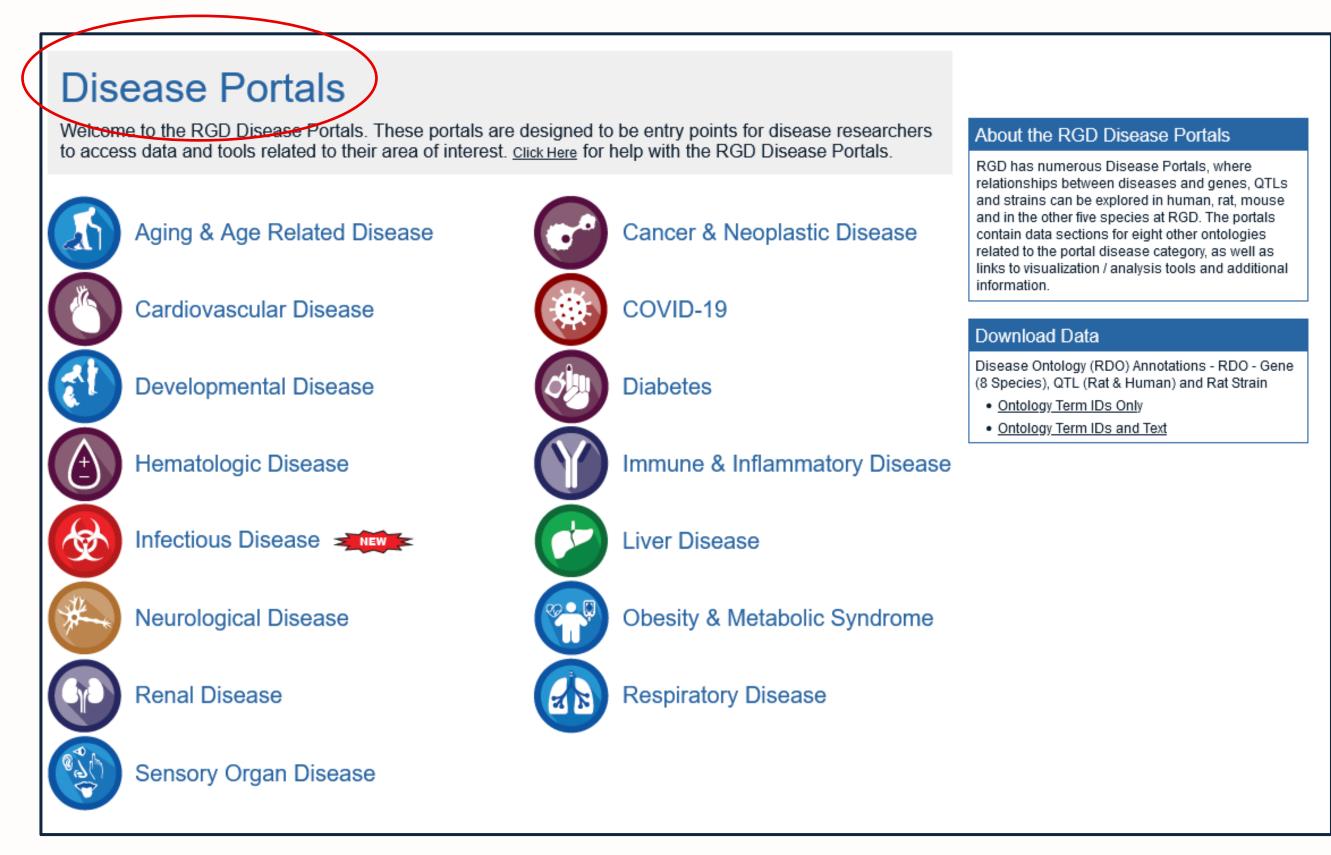


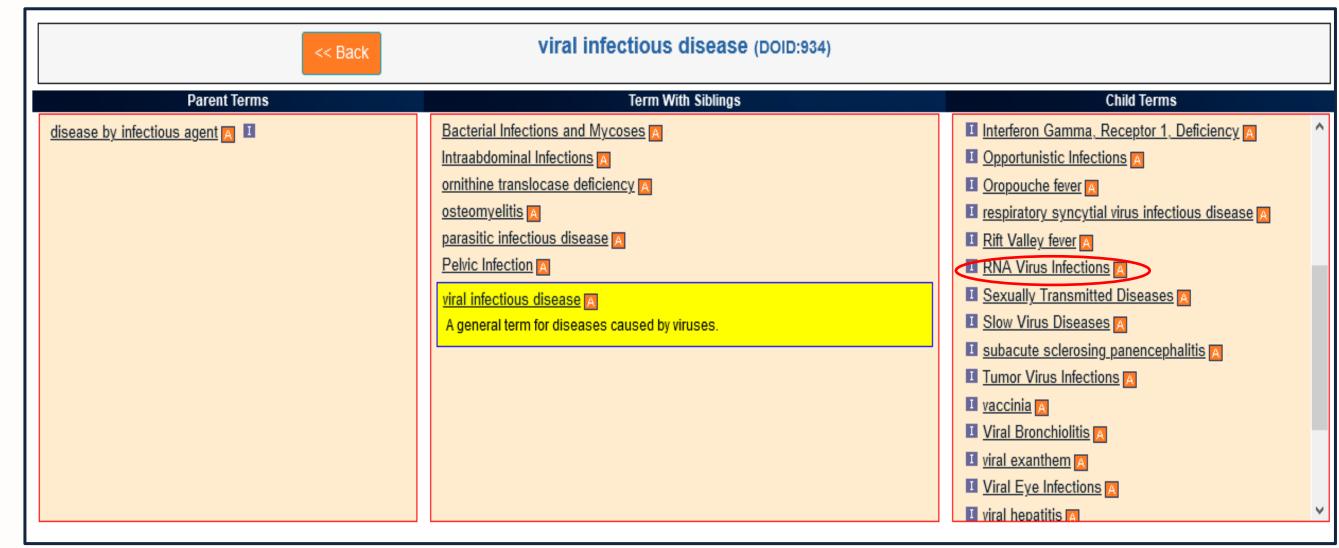
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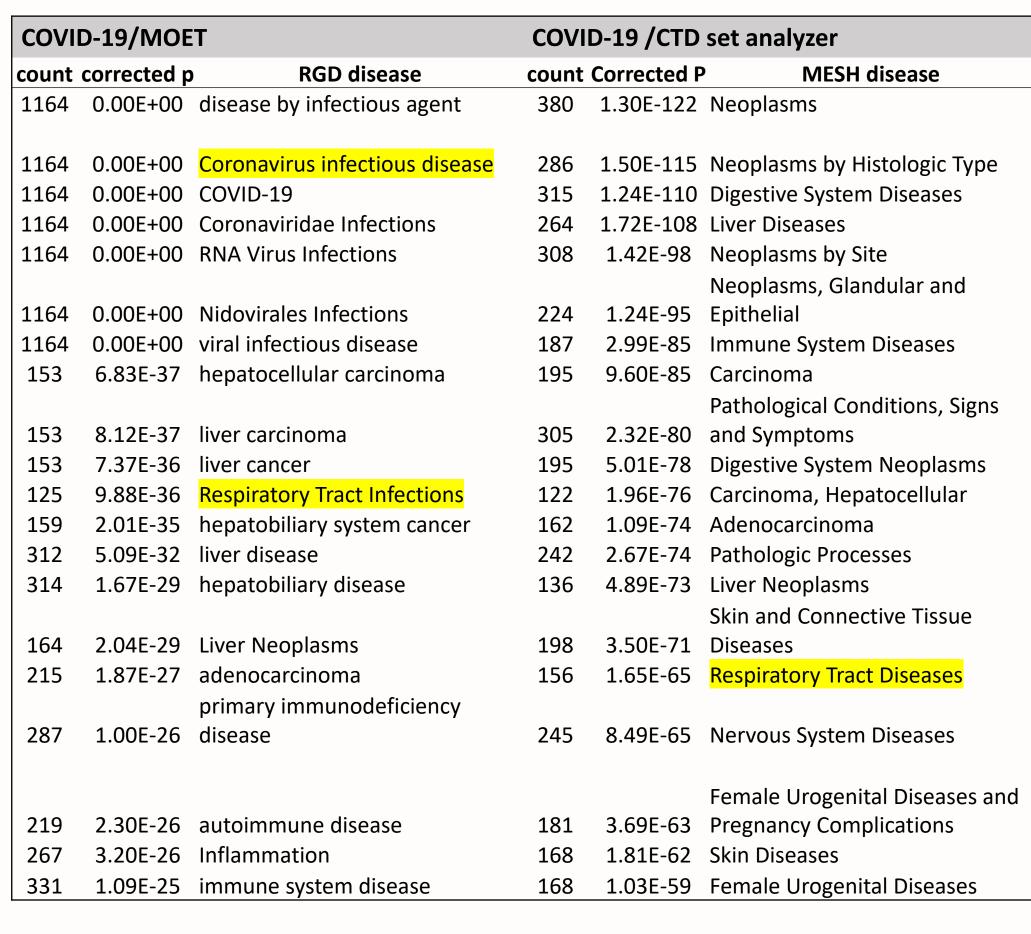
Abstract

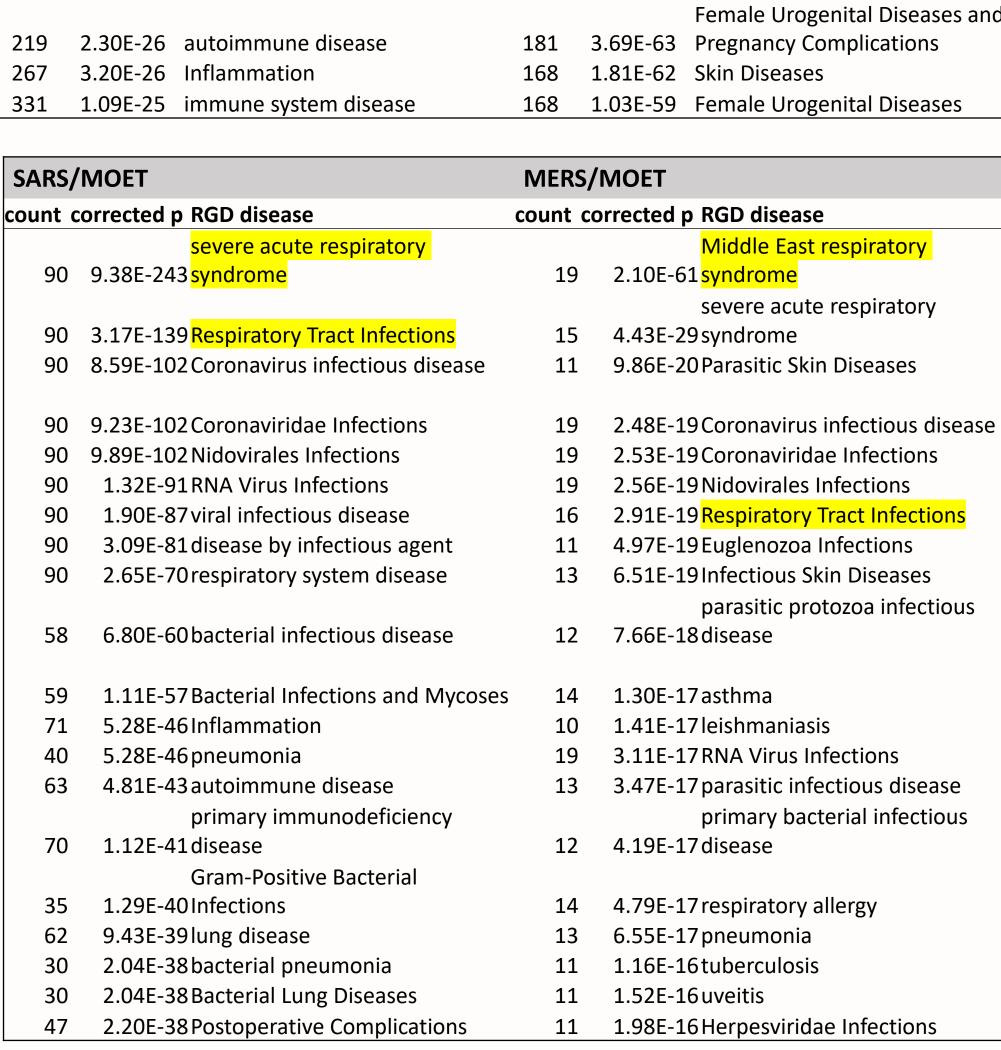
The Rat Genome Database (RGD) is the premier site for integrating rat genomic and genetic data with curated biological information. Multiple ontologies are used to annotate genes, strains, and QTLs. RGD has focused its manual curation efforts to annotating these genome objects with disease ontology terms. The RGD Disease Portals, with periodic addition of new portals and disease groups since they were launched, provide easy access to all genes, QTLs, strains, phenotypes, and biological functions associated with specific disease areas. The newly added "Infectious Diseases Portal" is the 15th portal to date. Here we present disease ontology enrichment analysis across selected disease gene sets within the Infectious Disease Portal using the Multi-Ontology Enrichment Tool (MOET). As expected, the COVID-19 disease gene set is highly enriched with coronavirus infectious disease and related diseases. Between coronavirus infectious disease and respiratory tract infections, several disease areas are also enriched among the COVID-19 genes. The enrichment profile of the COVID-19 gene set is compared to those of "Middle East respiratory syndrome" (MERS) and "severe acute respiratory syndrome" (SARS). Additional analysis of phenotypes and Gene Ontology were also performed using MOET and other tools at RGD. Using the Gene Annotator tool, we showed that close to one third of the COVID-19 genes were also annotated with "abnormality of nervous system" phenotypes. In this poster, we demonstrate the utility of RGD tools in analyzing coronavirus disease genes. We can look at not only the enrichment profile of a disease gene set, but also the gene distribution between two selected ontologies.

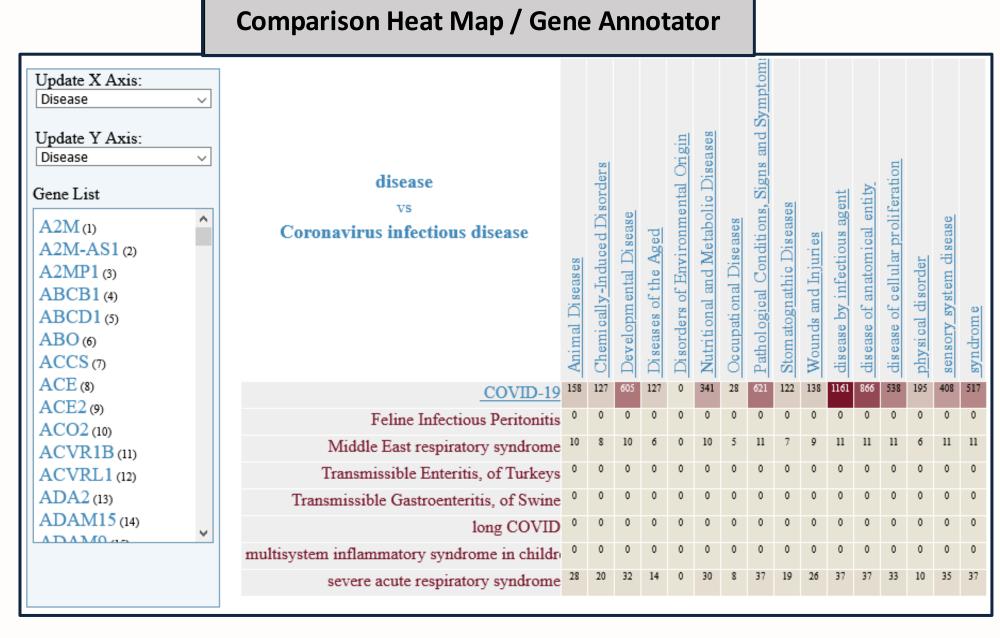




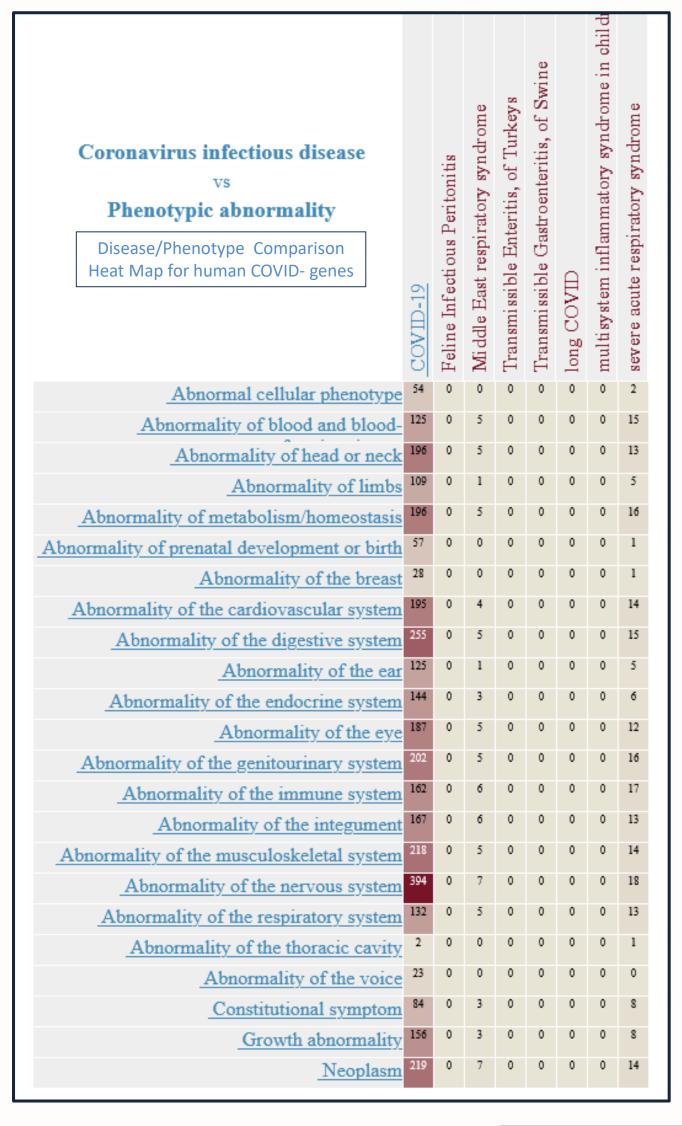


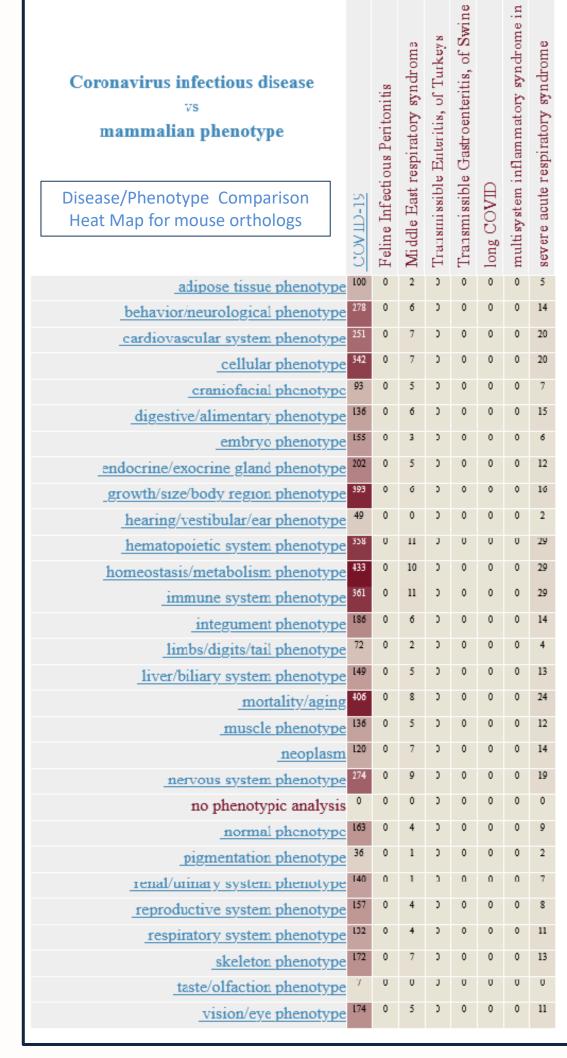


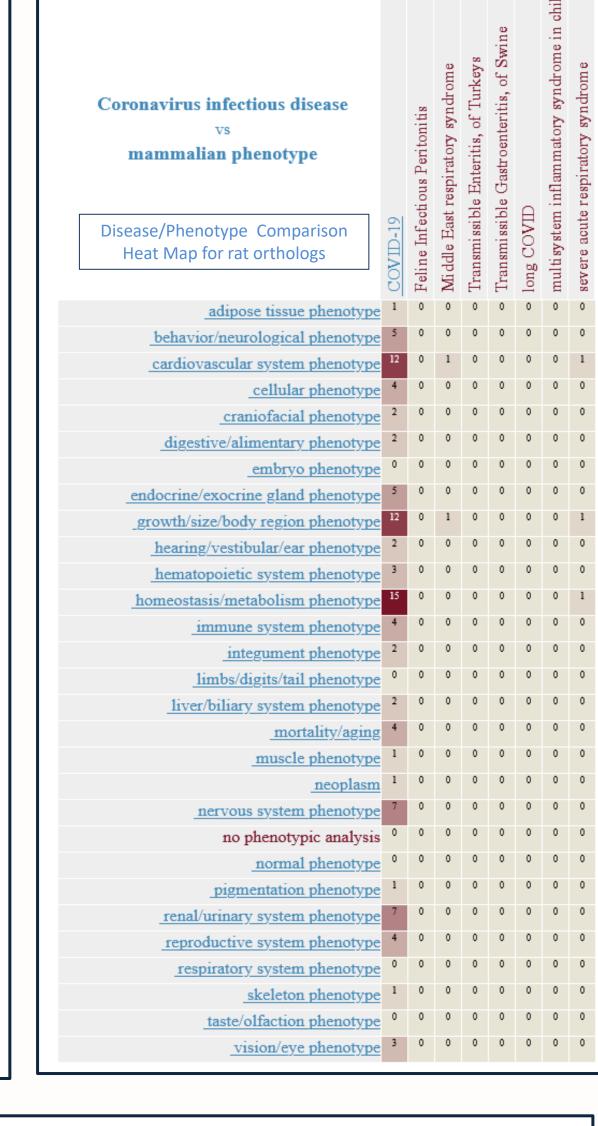




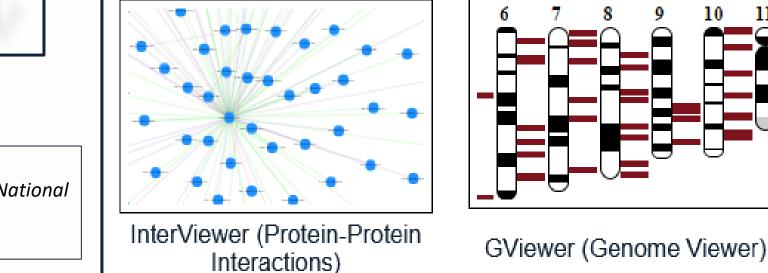
COVID-19/human disease genes			COVID-19 /mouse orthologs		
	corrected			Corrected	
count	р	Human phenotype	count	р	Mammalian phenotype
562	0.00E+00	Phenotypic abnormality	209	1.12E-12	abnormal adaptive immunity
		Abnormality of the nervous			abnormal cell-mediated
395	4.79E-193	system	205	3.72E-12	immunity
		Abnormal nervous system			abnormal immune cell
345	1.34E-149	physiology	203	6.11E-12	physiology
180	1.42E-73	Behavioral abnormality	198	2.25E-11	abnormal leukocyte physiology
					abnormal hematopoietic
93	1.28E-55	Schizophrenia	219	1.04E-10	system physiology
		Neurodevelopmental			abnormal immune serum
204	2.33E-37	abnormality	165	2.20E-10	protein physiology
		Abnormality of higher mental			abnormal immune system
218	6.96E-29	function	293	3.64E-10	physiology
					abnormal professional antigen
168	1.29E-25	Intellectual disability	148	7.96E-10	presenting cell physiology
		Abnormality of the digestive			
255	7.10E-24	system	203	2.21E-09	abnormal blood cell physiology
		Abnormal nervous system			abnormal lymphocyte
220	1.64E-22	morphology	151	4.21E-09	physiology
		Morphological central nervous			
205	1.03E-21	system abnormality	79	5.18E-09	abnormal IgG level
216	3.74E-21	Neoplasm by anatomical site	104	1.00E-08	abnormal response to infection
219	4.73E-21	Neoplasm	105	1.03E-08	abnormal response to antigen
		Abnormality of the			
218	1.28E-20	musculoskeletal system	79	1.41E-08	autoimmune response
					abnormal immunoglobulin
276	2.23E-20	Mode of inheritance	93	1.57E-08	level
		Abnormality of the			
190	9.57E-18	gastrointestinal tract	79	1.75E-08	abnormal self tolerance
					abnormal susceptibility to
156	3.23E-15	Growth abnormality	99	1.88E-08	infection
		Abnormality of the			
156	4.93E-15	musculature	79	4.11E-08	abnormal immune tolerance
					abnormal humoral immune
109		Abnormality of limbs	96	8.34E-08	•
121	8.02E-15	Seizure	107	9.23E-08	abnormal cytokine secretion





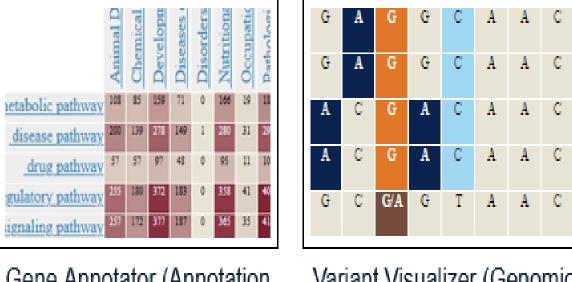




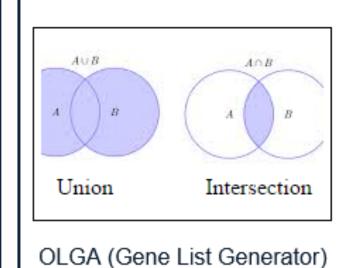


MOET (Multi-Ontology

Enrichement)



Variant Visualizer (Genomic Gene Annotator (Annotation Distribution) Variants)

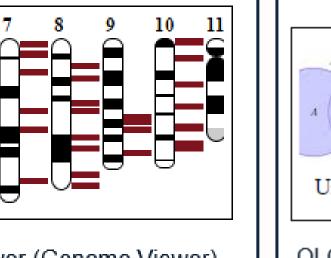




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GOLF (Gene-Ortholog Location

Finder)