Import of Human GWAS Data and Mapping of EFO to multiple ontologies at RGD

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Rat Genome Database
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Biocuration 2025

Ontologies used to curate phenotype and disease at the Rat Genome database (RGD)

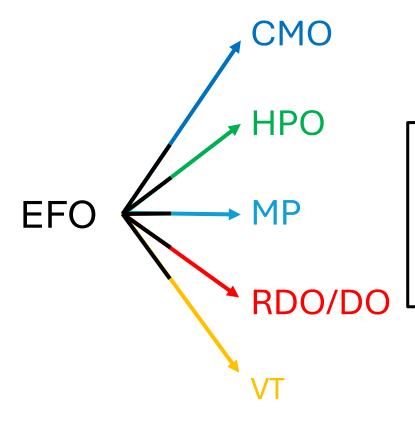
Data type at RGD	Ontology used at RGD		
rat phenotype (qualitative)	Mammalian Phenotype ontology (MP)		
rat phenotype (quantitative)	Vertebrate Trait ontology (VT)		
	Clinical Measurement Ontology (CMO)		
	Measurement Method Ontology (MMO)		
	Experimental Condition Ontology (XCO)		
	Uberon ontology (UBERON)		
	Cell Ontology (CL)		
human phenotype (qualitative)	Human Phenotype Ontology (HPO)		
	RGD disease ontology (RDO) (axiomatized		
	and extended version of the human		
disease (qualitative)	disease ontology (DO)		





Automated term mapping

- Text of term and synonyms
- 2. ID
- 3. Associated XREFs



- Text of term and synonyms
- 2. ID
- 3. Associated XREFs

Text of EFO/target term

Definition of EFO/target term

Synonyms of EFO/target term

Manual term mapping

Text of EFO term added to target term as synonym (exact, broad, narrow or related)

ID of EFO term added to target term as XREF

Number of EFO terms from GWAS records mapped to ontologies at RGD

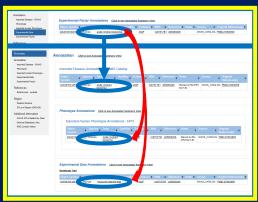
EFO term subsets	Subset Numbers as of March 2025
total # of unique EFO terms associated with imported GWAS	
records	7601
total # of unique EFO/HP/MONDO hybrid terms associated	
with imported GWAS records	2009
# of EFO terms mapped to RDO/DO	1930
# of EFO terms mapped to HPO	514
# of EFO terms mapped to CMO	469
# of EFO terms mapped to MP	209
# of EFO terms mapped to VT	900+

Term mapping output for users

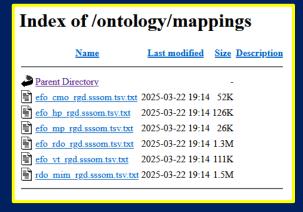
1. Entries in term browser (HPO, CMO, MP, RDO/DO, and VT)



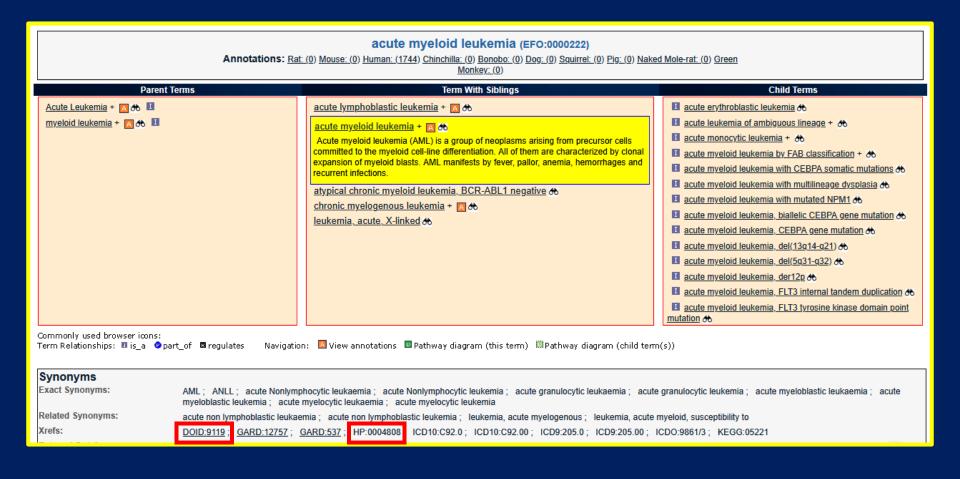
2. Transitive annotations on Human QTL pages (SNP variant pages)



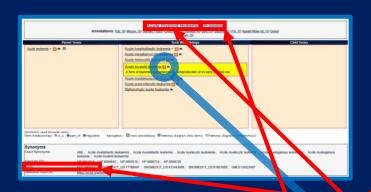
3. SSSOM file on the RGD download page (https://download.rgd.mcw.edu/ont ology/mappings/)

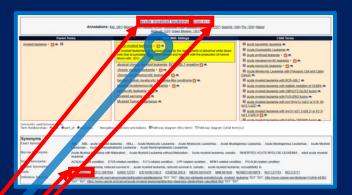


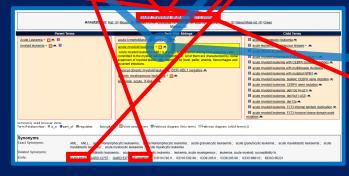
EFO entry in term browser: acute myeloid leukemia

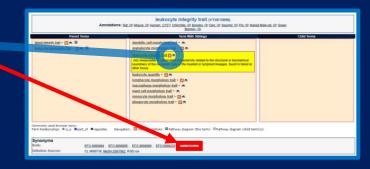


Interrelationships in entries in term browser (EFO, HPO, RDO/DO, and VT)

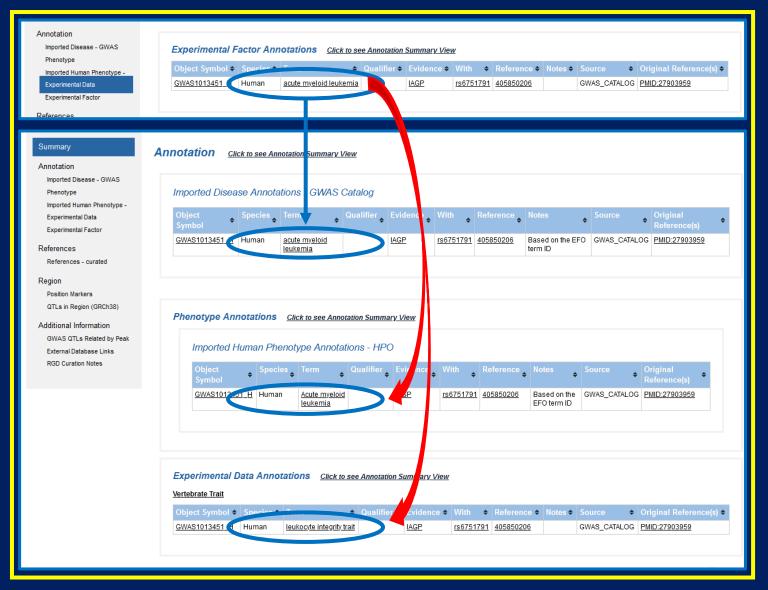








Transitive annotations on Human QTL pages (SNP variant pages)



SSSOM files on the RGD download page

Index of /ontology/mappings

	<u>Name</u>	Last modi	<u>fied</u>	Size D	<u>Description</u>
Pare	nt Directory			-	
efo	cmo rgd.sssom.tsv.txt	2025-03-22	19:14	52 K	
efo :	hp_rgd.sssom.tsv.txt	2025-03-22	19:14	126K	
efo:	mp_rgd.sssom.tsv.txt	2025-03-22	19:14	26K	
efo	rdo rgd.sssom.tsv.txt	2025-03-22	19:14	1.3M	
efo	vt_rgd.sssom.tsv.txt	2025-03-22	19:14	111K	
<u>rdo</u>	mim rgd.sssom.tsv.txt	2025-03-22	19:14	1.5M	

Acknowledgements





Rat Genome Database

Management:

Anne Kwitek Mindy Dwinell

Software Developers:

Jeff De Pons Marek Tutaj Jyothi Thota Logan Lamers Akhilanand Kundurthi Adam Gibson Varun Gollapally

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