

Immune Epitope Database Overview

www.iedb.org

Presented by: Alessandro Sette, IEDB Principal Investigator

Our goals for this user workshop

We want your input to make the IEDB better:

- Learn about real-life applications for the IEDB
- Identify and prioritize problems with the user interface, documentation, functionality etc.

We want to enable you to get the most out of the IEDB:

- The primary IEDB success metric is usage
- Best compliment for our program is if IEDB data & tools help in your research (citations)

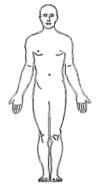
Immune Epitope Database

www.iedb.org

Database | Resource of experimentally-derived epitope information

- Allergens
- Infectious diseases
- Autoimmune diseases
- Transplantation / Alloantigens

... and more







Containing data on over **2.2 million unique structures** analyzed in over **6.6 million assays** from more than **24,000 curated references**.

Minimal Criteria for Epitope Inclusion

- Linear peptide <50 amino acids in length
- Tested as an immunogen or an antigen
- Discontinuous residues shown to be important in recognition
- Non-peptidic epitopes <5,000 Daltons
- Minimal information required (sequence, outcome, host, etc.)

Consistent data entry requires well defined data structure

Quantitation of CD8+ T Cell Responses to Newly Identified HLA-A*0201– restricted T Cell Epitopes Conserved Among Vaccinia and Variola (Smallpox) Viruses

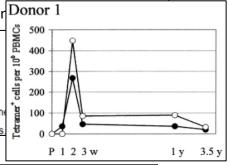
Masanori Terajima, John Cruz, Gregory Raines, Elizabeth D. Kilpatrick, Jeffrey S. Kenr

Francis A. Ennis

Materials And Methods

Donors.

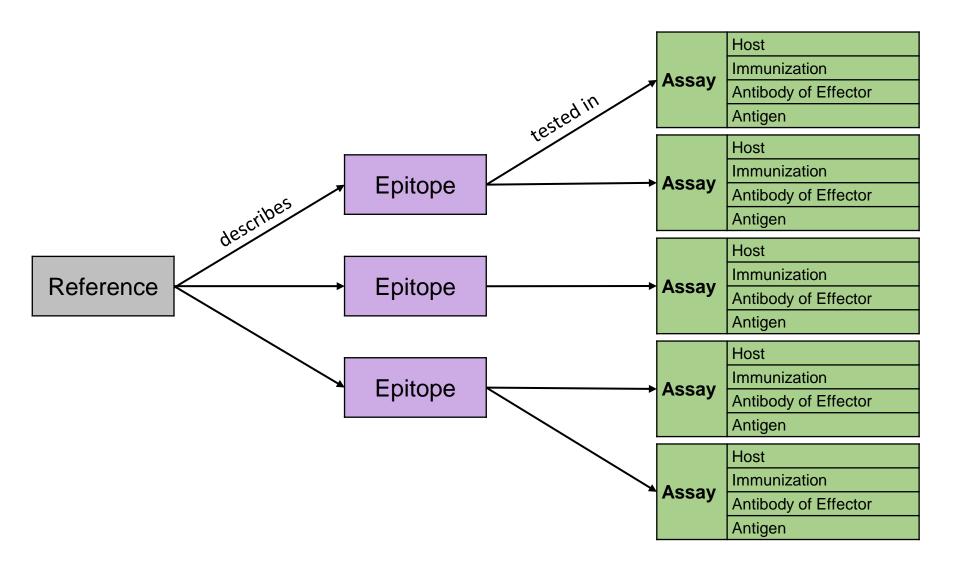
Donors in this study were three HLA-A*0201–positive laboratory workers received primary immunization by scarification with the licensed smallpox vaccine, Dryvax®, as recommended by the Centers for Disease Control and Prevention for laboratory pe



			•
		Name	74A
	Structure	Chemical Type	Peptide / Protein
		Sequence	CLTEYILWV
		Domain / Region	Defined Epitope
Epitope		Species	Vaccina Virus Ankara
		Strain	Ankara (MVA)
	Source	Antigen	Putative 21.7k protein
		Antigen Accession	2772819
		Antigen Positions	79-87
		Immunized Species	Homo sapiens
	Immunization	Immunogen Type	Source species
		Administration	Scarification
Context		Antigen Type	Epitope
	Assay	Assay Type	ELISPOT
		Response Measured	Cytokine Release-IFN-g
		MHC Allele	HLA-A*0201
		-	-

5

Data Structure: A Database of Experiments



Collaborations with Ontologies

- Provides standardized nomenclature, definitions, synonyms, and hierarchical relationships
- Makes curation easier → Finders
- Enhances user experience → Finders
- Ensures consistency and accuracy
- Finds errors
- Facilitates interoperability

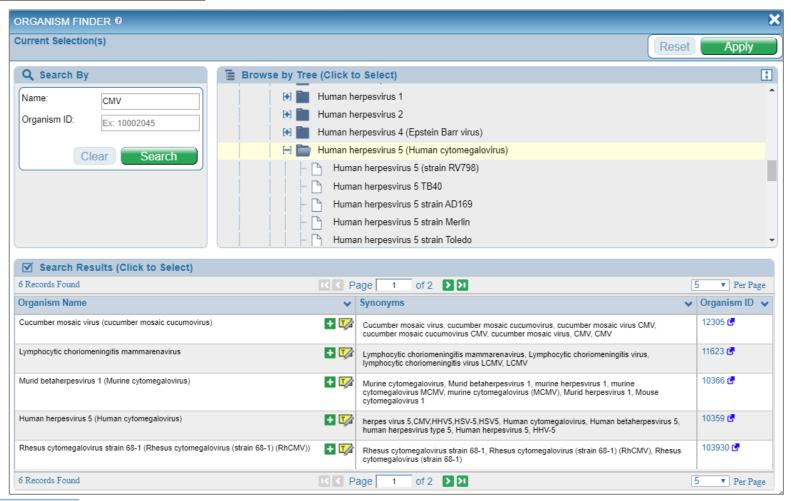
External Resources and Ontologies

Peptidic Epitope Amino acid sequence Protein source Organism source

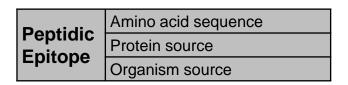


Human herpesvirus 5 (HHV-5) NCBI

taxon:10359

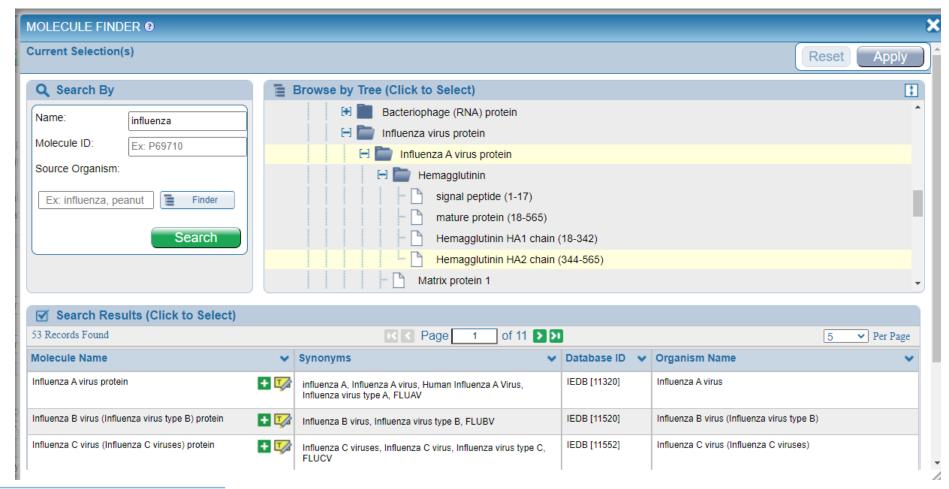


External Resources and Ontologies





PKYVKQNTLKLAT hemagglutinin HA1 GenBank/UniProt GI:AAL62329.1—UniProt:Q8V285



Literature Curation

PubMed / PDB

- Complex query
- Bi-weekly

240K retrieved

Classifier

- Content based categories
- Retrained annually

151K epitope related

Abstract Review

- Manual scan
- Confirmation of classification

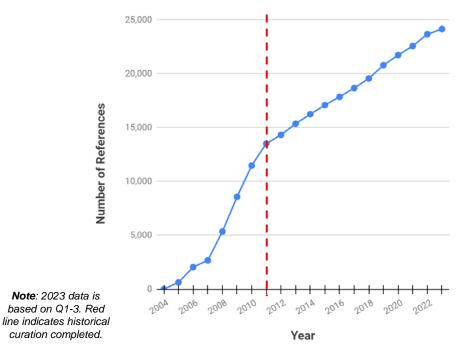
44K likely curatable

Manual Curation

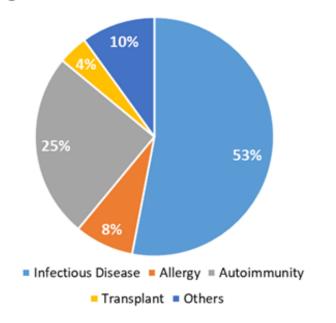
- Assigned to curators
- Peer review

24K curated

Growth of IEDB Curated References



Categorical Breakdown of Curated References



Consistency and Quality Control Measures

- Manually curated by a team of PhD-level scientists with specific expertise
- Formal curation guidelines and peer review
 Curation Manual: http://curationwiki.iedb.org/wiki/index.php/Main_Page
- External immunological experts
- Built in validation in the curation application

Data Submissions

- Primarily sourced through NIAID Epitope Discovery contracts
- Data deposition is open to the general research community on a case-by-case basis

As of October, 2022:

- Data from 340 submissions are publicly available & 90 submissions are in process or on-hold
- Submitted data comprises ~20% of epitopes available in the IEDB

Inquire by contacting: nblazeska@lji.org or submissionsupport@iedb.org!

Direct Submission Process

Submission

 Investigators submit files via IEDB data submission tool



Review

 Automated validation and manual curator review check data consistency

Submitter confirms final approval



Release

Public release date set by submitter



IEDB.org: Homepage & Cumulative Data



Home Specialized Searches Analysis Resource

Check out our new IEDB updates! (1) Learn how to customize your database exports and (2) test out the new Next-generation Tools site for all your analysis and prediction needs.

Welcome

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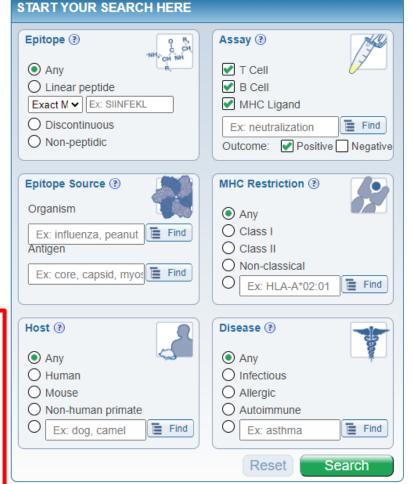
Learn More

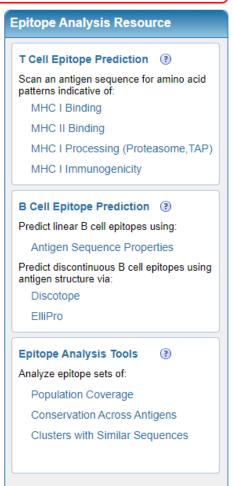
Upcoming Events & News

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AACR 2024 Apr 5-10, 2024 Festival of Biologics Apr 15-17, 2024 AAI 2024 May 3-7, 2024

Summary Metrics Peptidic Epitopes 1,597,734 Non-Peptidic Epitopes 3,187 T Cell Assays 507.401 B Cell Assays 1,391,378 MHC Ligand Assays 4,777,560 Epitope Source Organisms 4,403 Restricting MHC Alleles 991 References 24,153





Help

More IEDB

IEDB.org: Homepage & Search Interface



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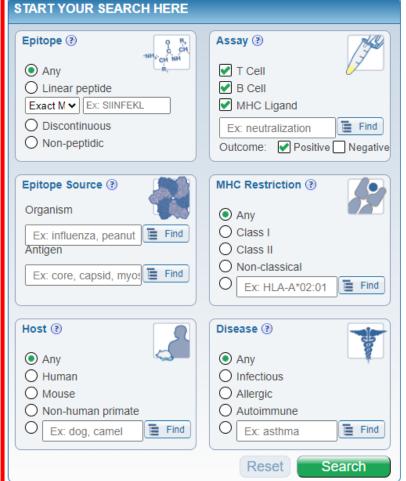
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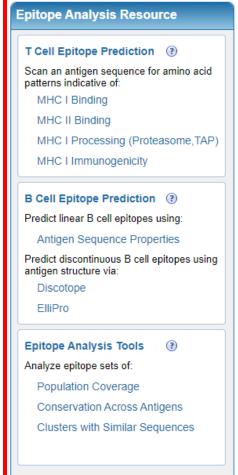
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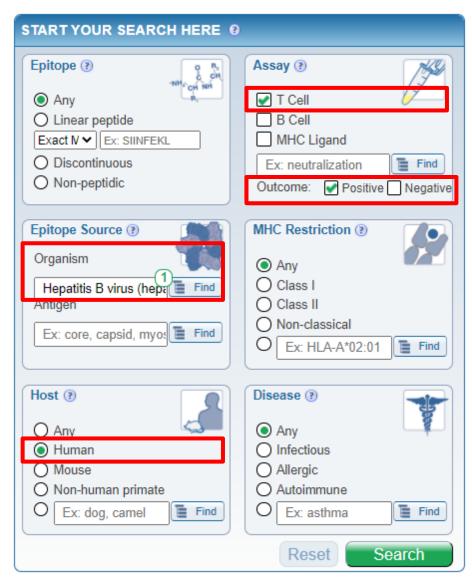
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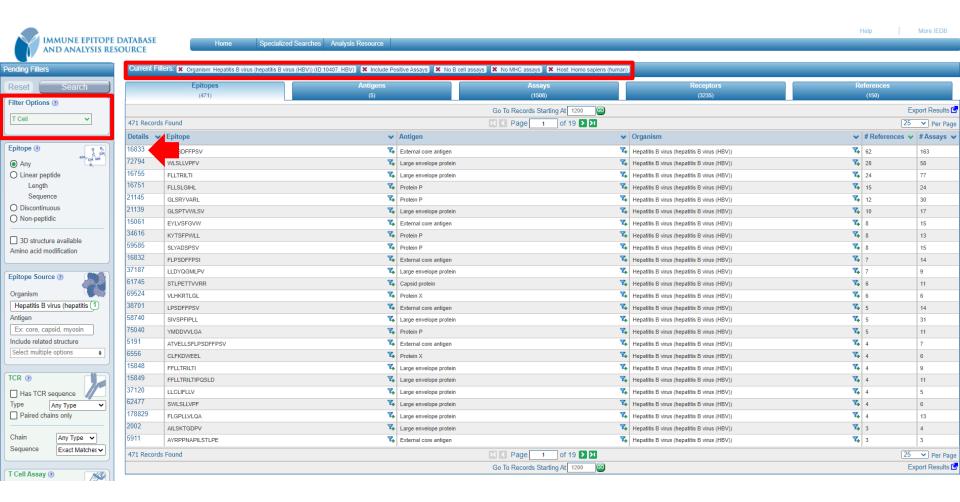


More IEDB

Example Query: HBV T Cell Epitopes in Humans



Results Summarized in Tables



Detail Pages Summarize Relevant Information

Details Pages:

- Epitope Summary
- Compiled Data
 - MHC Ligand Assay(s)
 - B Cell Assay(s)
 - T Cell Assay(s)
- External Resources

Also available for:

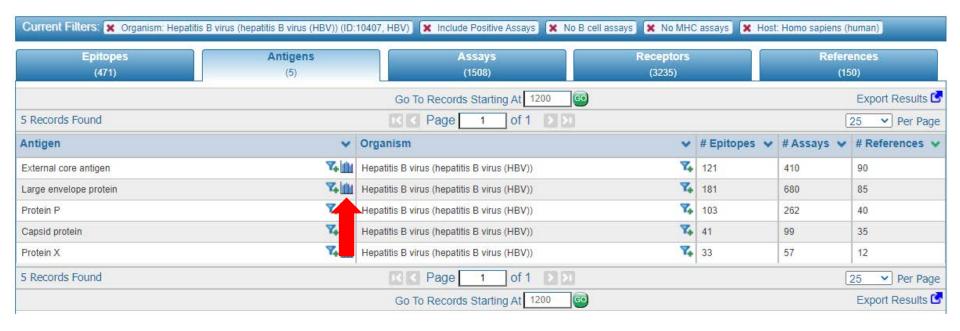
- Assays
- Receptors
- References

EPITOPE SUMMARY

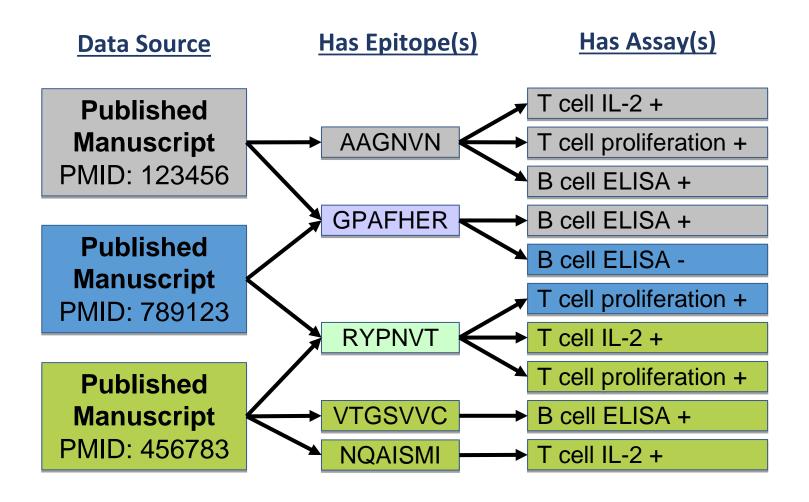
FLPSDFFPSV is a linear peptidic epitope (epitope ID 16833) studied as part of External core antigen from Hepatitis B virus (hepatitis B virus (HBV)) and Capsid protein from Hepatitis B virus (hepatitis B virus (HBV)). This epitope has been studied for immune reactivity in 100 publication(s), tested in 215 T cell assays, 8 B cell assays, 8 B MHC ligand assays and has 3D structure(s) 3OXR, 3OXS, 1HHH and 3OXR

COMPILED DATA		
MHC Ligand Assay(s) 88		
MHC molecul	e	Positive / All
HLA-A*02:01		35/35
HLA-A*02:03		8/8
HLA-A*02:06		8/8
HLA-A*02:02		6/6
HLA-A*02:07		4/4
HLA-A*68:02		4/4
HLA-A*03:01		0/3
HLA-A*11:01		0/3
HLA-A*02:05		2/2
HLA-A*01:01		0/2
HLA-B*07:02		0/2
HLA-B*15:02		0/2
HLA-A*02:04		1/1
HLA-A*02:11		1/1
HLA-A2		1/1
HLA-B44		1/1
H2-Kd		0/1
HLA-A*24:02		0/1
HLA-A*31:01		0/1
HLA-A*68:01		0/1
Mamu-A1*001:	01	0/1
B Cell Assay(s) 8		
Assay Type		Positive / All
qualitative binding		7/7
dissociation constant KD		1/1
T Cell Assay(s) 215		2 77 4 21
Assay Type		Positive / All
cytotoxicity		63/74
IFNg release		65/66
qualitative bindi	ng	45/50
TNFa release		7/8
IL-2 release		4/4
proliferation		4/4
IL-4 release		2/2
TNF release		2/2
activation		1/1
EXTERNAL RESOURCES		
Resource	Link	
ANALYSIS TOOLS IEDB.ORG IEDB-AR: MHC-I Processing	Predict MHC class I process	ing ௴
ANALYSIS TOOLS IEDB-AR: MHC-I	Predict MHC class I binding	affinity 🗳
ANALYSIS TOOLS IEDBORG IEDB-AR: B cell scales Predict B cell epitopes Predict B cell epitopes Predict B cell epitopes Predict B cell epitopes Predict B cell epitopes Predict		

Antigens: Identifying Protein Source of Epitopes



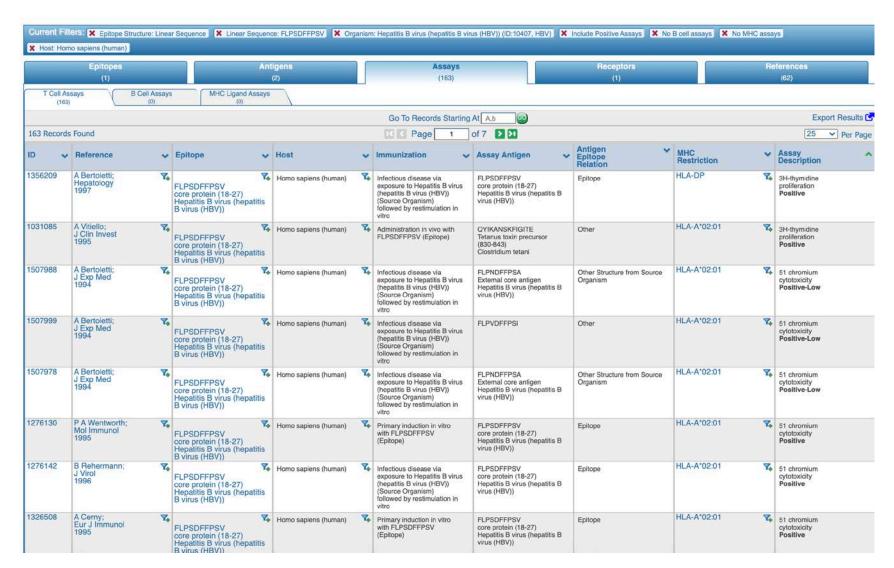
Data Aggregation



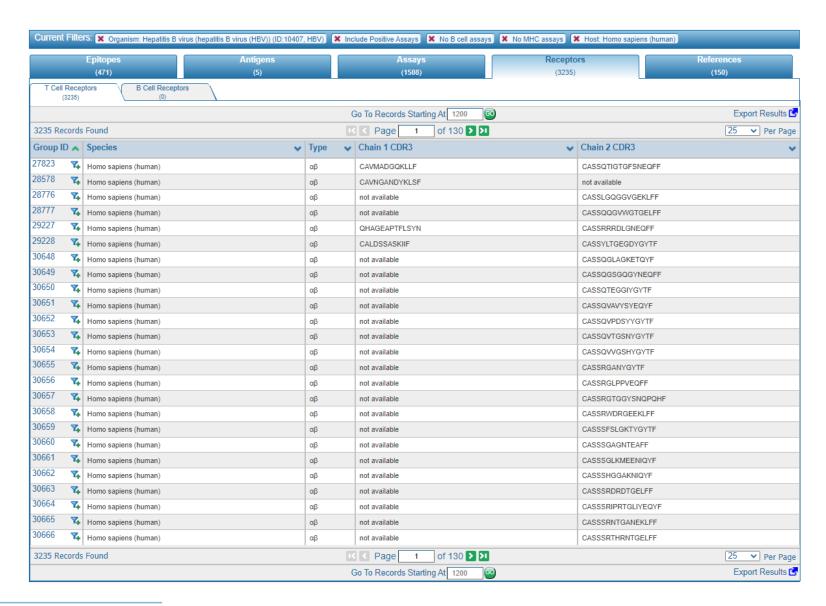
ImmunomeBrowser: Visualization on Reference Proteins



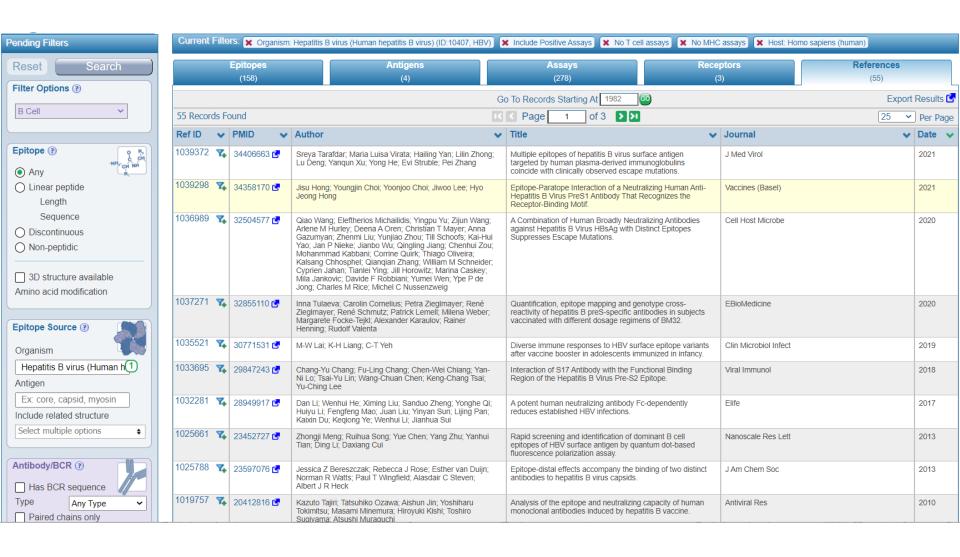
Assays: Experiments in which epitopes were tested



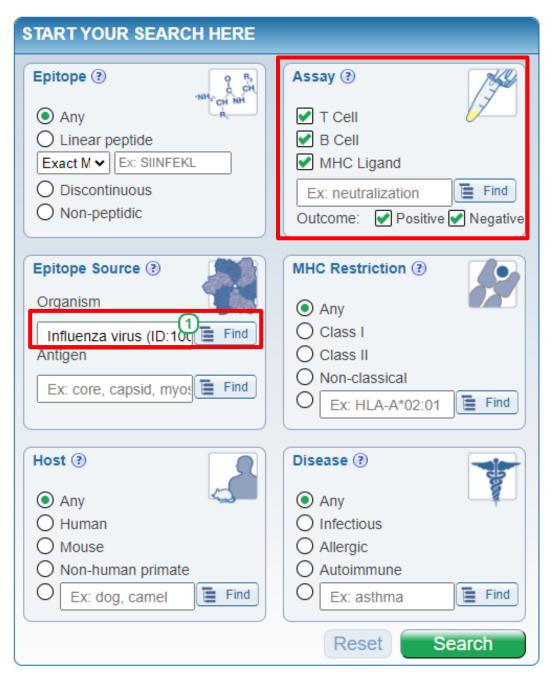
Epitope-specific B Cell and T Cell Receptors



References: Source of Information

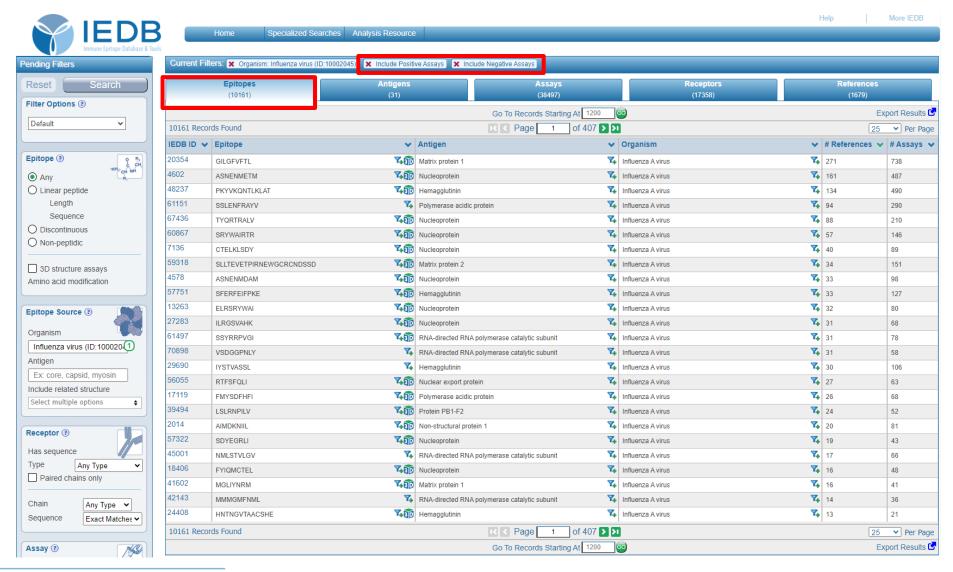


Example Query II: Influenza Virus Epitopes



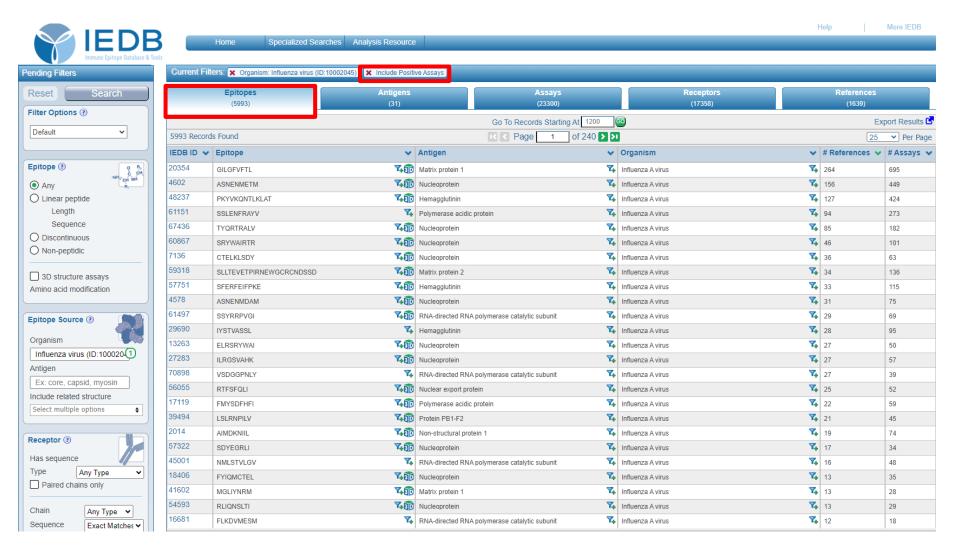
Results: All influenza virus epitopes

Both positive and negative assay outcomes



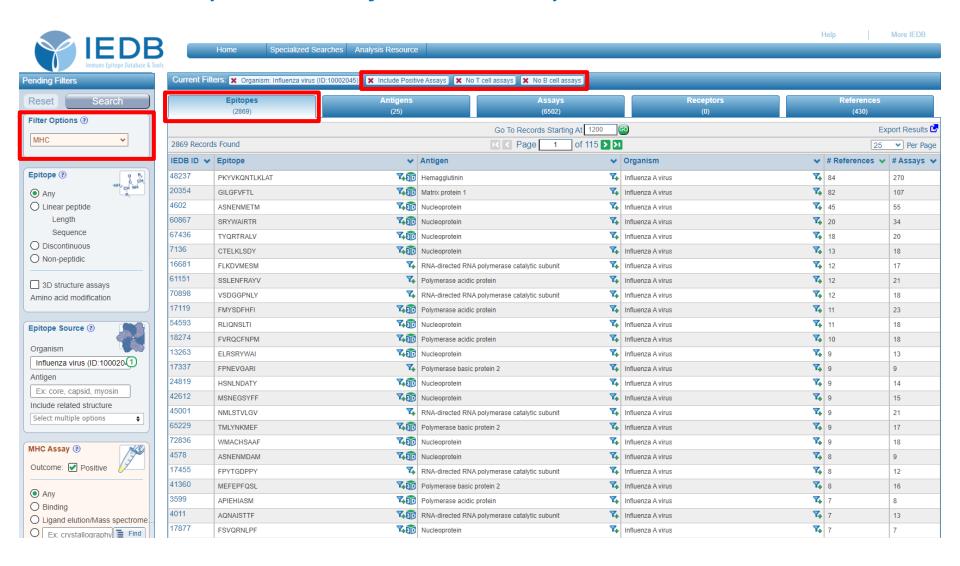
Results: Positive influenza virus epitopes

Positive assay outcomes



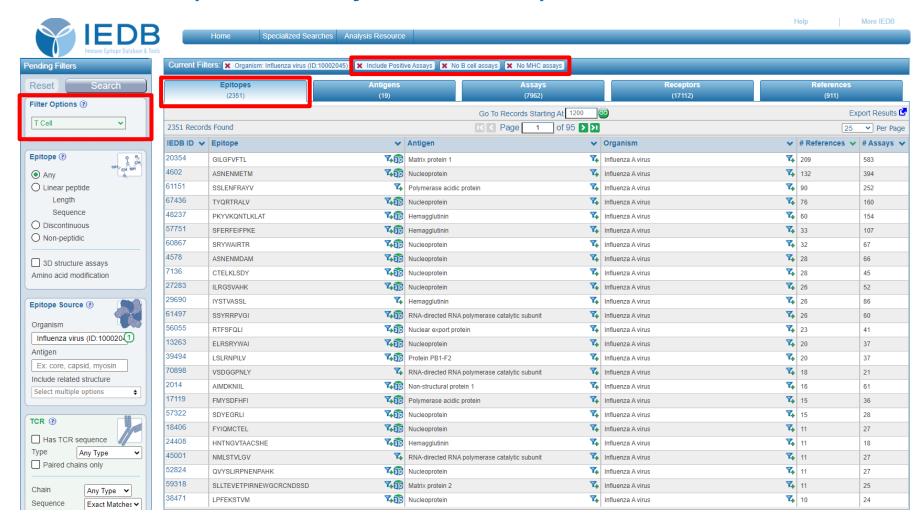
Results: Influenza virus epitopes that bind

Positive assay outcomes for MHC assays



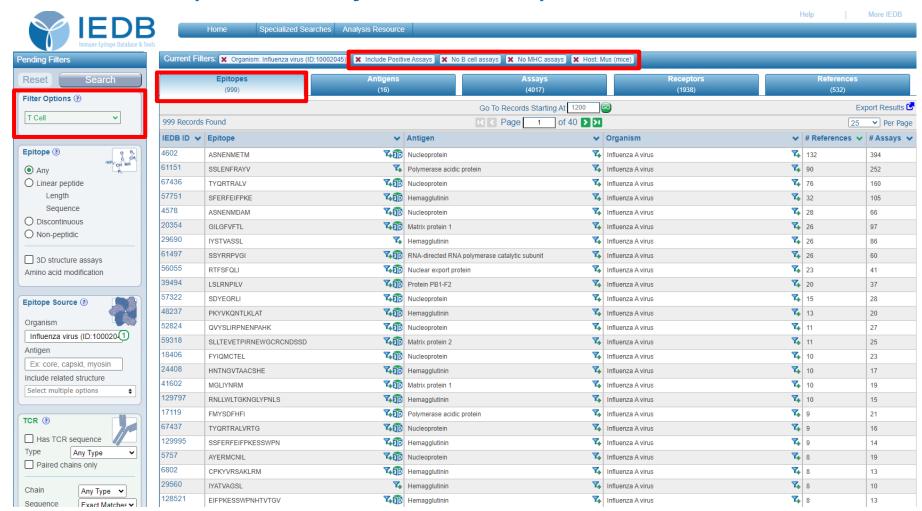
Results: Influenza virus epitopes that are immunogenic

Positive assay outcomes for T cell assays



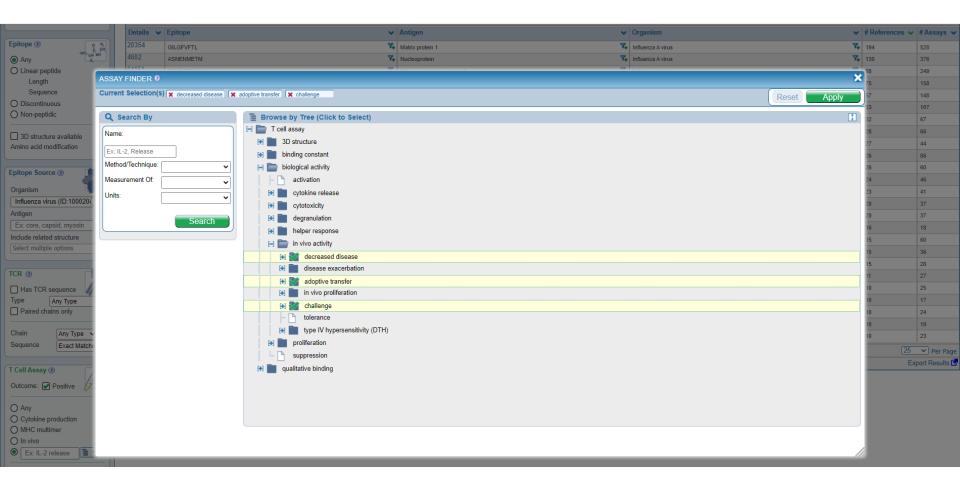
Results: Influenza virus mouse epitopes that are immunogenic

Positive assay outcomes for T cell assays in a mouse host



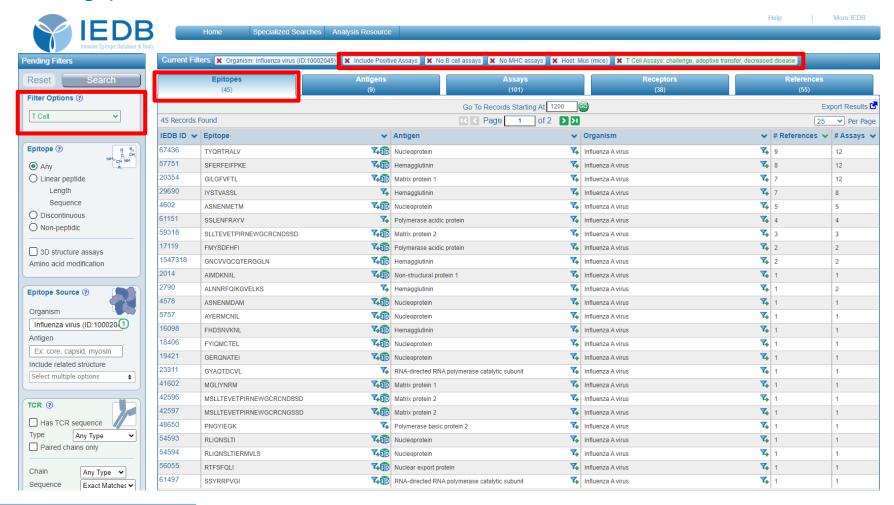
Results: Influenza virus epitopes that offer protection from disease

Positive assay outcomes for T cell assays (decreased disease, adoptive transfer, challenge) in a mouse host



Results: Influenza virus epitopes that offer protection from disease

Positive assay outcomes for T cell assays (decreased disease, adoptive transfer, challenge) in a mouse host



More Exports



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Meta-Analyses

Database Export

More IEDB

Citing the IEDB Check out our new IEDB updates! (1) Learn how to customize your database exports and (2) test out the new Next-generation Tools site for all your

Release Notes

Links

T Cell Epitope Prediction ②

Scan an antigen sequence for amino acid

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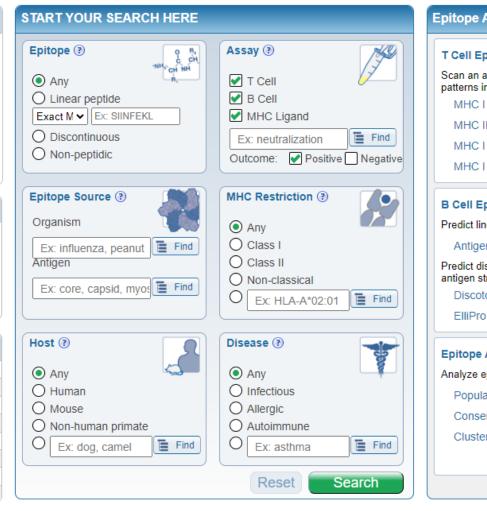
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T Cell Assays	507,656
B Cell Assays	1,391,572
MHC Ligand Assays	4,777,564
Epitope Source Organisms	4,403
Restricting MHC Alleles	991
References	24,173





B Cell Epitope Prediction ②

Predict linear B cell epitopes using:

Antigen Sequence Properties

Predict discontinuous B cell epitopes using antigen structure via:

Discotope

ElliPro

Epitope Analysis Tools



(?)

Analyze epitope sets of:

Population Coverage

Conservation Across Antigens

Clusters with Similar Sequences

More Exports

https://www.iedb.org/database_export_v3.php

XML Database Export	
Complete Database Export	510MB
<u>ledbAccessionList.zip</u>	80kB
MhcAlleleNameList.zip	533kB
OrganismList.zip	39MB
AssayTypeList.zip	5kB
GeoLocList.zip	3kB

CSV Metric Exports		
epitope full v3.zip	78MB	
antigen full v3.zip	2MB	
toell full v3.zig	37MB	
boell_full_v3 (single_file.zip) (multi_file.zip)	66MB	
mhc_ligand_full (single_file.zip) (multi_file.zip)	231MB	
reference full v3.zig	змв	
receptor full v3.zip	6MB	
tor full v3.zip	5MB	
ber full v3.zip	967kB	
jedb 3d full.zip	4MB	

IEDB Schema	
Curation.xsd (Primary IEDB schema)	49kB
<u>CurationSimpleTypes.xsd</u>	323kB
<u>ledbAccessionList.xsd</u>	909B
MhcAlleleNameList.xsd	1kB
<u>OrganismList.xsd</u>	751B
<u>AssayTypeList.xsd</u>	771B
<u>GeoLocList.xsd</u>	642B

TSV Metric Exports		
epitope full v3 tsv.zip	78MB	
antigen full v3 tsv.zig	2MB	
toell full v3 tsv.zig	37MB	
boell full v3 tsv.zip	66MB	
mho ligand full tsv.zig	231MB	
reference full v3 tsv.zip	змв	
tor full v3 tsv.zip	5MB	
bor full v3 tsv.zip	967kB	

MySQL Database Export		
SQL Statement Export 51	16MB	
MylSAM Binary Export 10	3B	

31kB

Excel Metric Exports		
epitope full v3.xlsx	172MB	
antigen full v3.xlsx	4MB	
toell full v3.xlsx	154MB	
boell full v3.xlsx	318MB	
mho ligand full.xlsx	805MB	
reference full v3.xlsx	4MB	
tor full v3.xlsx	17MB	
ber full v3.xlsx	2MB	

Physical Entity Relationship Diagram	
iedb_public_erd.pdf	

	Json Metric Exports		
	epitope full v3 ison.zip	114MB	
	antigen full v3 ison.zip	2MB	
	toell full v3 json.zip	73MB	
	hoell full v3 ison zin	130MB	

Help integrated throughout the website



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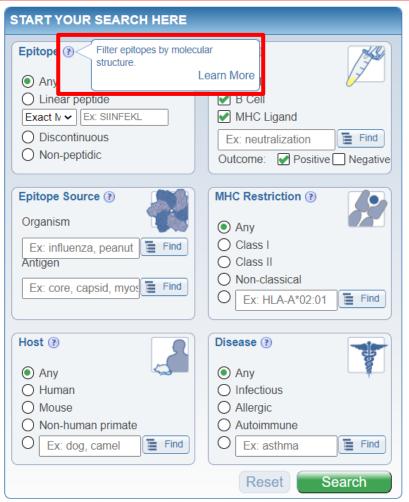
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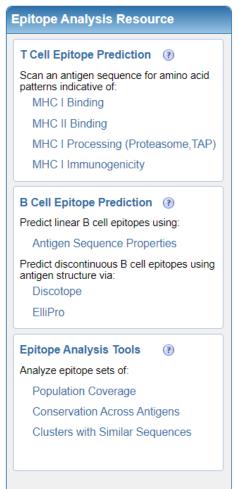
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Help More IEDB

Support

Help Request

Provide Feedback

Video Tutorials

and prediction needs.

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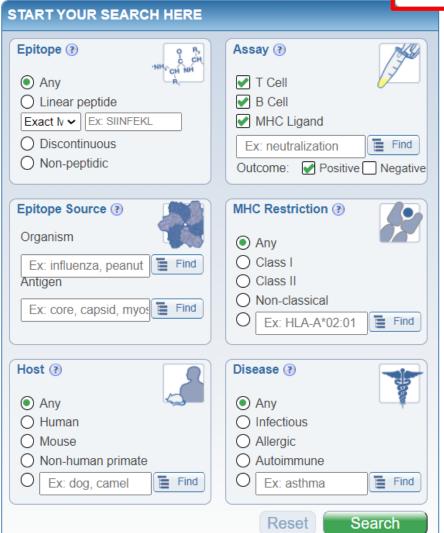
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Epitope Analysis Resource T Cell Epitope Prediction

Scan an antigen sequence for amino acid patterns indicative of:

MHC I Binding

MHC II Binding

MHC | Processing (Proteasome, TAP)

MHC I Immunogenicity

B Cell Epitope Prediction (?)



Predict linear B cell epitopes using:

Antigen Sequence Properties

Predict discontinuous B cell epitopes using antigen structure via:

Discotope

ElliPro

Epitope Analysis Tools



Analyze epitope sets of:

Population Coverage

Conservation Across Antigens

Clusters with Similar Sequences

Solution Center: help.iedb.org

Accessible through header and footer on every page or submit via email to help@iedb.org



