Finding Data in the IEDB Using the query interfaces

Nima Salimi. M.S. - Sr. Biocurator/Curation Mgr Kerrie Vaughan, PhD - Sr. Biocurator/Meta-Analysis





Overview

Introduction

- IEDB Home page search
- Exploring the query results
- Finders
- Specialized Searches (all fields)
 Break
- Immunome Browser
- Example queries & demos (live)
 Lunch
- Exercises (live)
- Participant-derived queries





How can the search interface be useful to me?



- 1. Finding epitopes (tested)
- 2. 'Big picture' analyses
- 3. Validation



Helps defines 'known universe' What has and has not been done

Part of prediction or informatics pipeline





Maximize the Available Information Pipeline







IEDB 'Jargon'

- Terms unique to the IEDB
 - A. Immunologic in nature
 - Immunogen, antigen, assay, in vivo admin, in vitro admin, occurrence of, IV1, IV2, process type
 - B. Ontological (database interoperability)
 - Relation, parent, child, sibling
 - C. Database necessity (explicit/constrained)
 - Source antigen, Source organism, Non-peptidic v. peptidic, B cell/Ab
- Link to Field Descriptions and Curation Manual: http://curationwiki.iedb.org/wiki/index.php/Main_Page





Assay-centric nature of the data





IEDB Home Page



Provide Feedback | Help Request | Solutions Center | Tool Licensing Information

IMMUNE EPITOPE DATABASE

AND ANALYSIS RESOURCE

Supported by a contract from the <u>National Institute of Allergy and Infectious Diseases</u>, a component of the Na Services. Data Last Updated: September 23, 2018



Home Page Search Interface



Search by:

- Epitope sequence
- Antigen
- Host
- MHC restriction
- Disease
- Assay type

Tailored to represent key elements of immune response - context and characteristics

Exploring the Results Page



Enables refinement of initial query
 Allows for more complex search using Finders (e.g. specific assay)
 Allows for Specialized Searches (e.g. mAb by name)

9 www.IEDB.org

Exploring the Results Page Current Filters: X Positive Assays Only Pending Filters Epitopes Search Reset References (522321) Epitope 🕐 (19554)Any Epitopes Export Results 522321 Records Found O Linear Epitope 25 Per Page Details v Epitope O Discontinuous Epitopes 123885 O Non-peptidic Epitopes cardiolipin 44920 NLVPMVATV 320 1028 7 275

- **Current Filters** track query parameters along the top
- Export of data per tab is available in Excel download





679



Epitope Sequence Search







Antigen Search



Pathogen (virus, bacteria) Mammals (self-Ag) Allergen source (grass)

m. m.

Protein of interest (source of epitope)	Antigen (?	MHC Restriction (?)		
	Organism HCV	Any MHC Restriction MHC Class I		
	A Hepatitis C virus (ID:111	103, HCV) sir		
	Hepatitis C virus (isolate I	olate H) (ID:11108, HCV-H)		
	Classical swine fever virus	Classical swine fever virus (ID:11096, hog cholera virus HCV)		
Click in field and start typing auto-	Hos Hepatitis C virus HCV-KF	(ID:31644)		
complete feature helps process	Hepatitis C virus genotype	e 1 (ID:41856, hepatitis C virus 1 HCV 1)		
	Hepatitis C virus subtype	1b (ID:31647, hepatitis C virus type 1b HC		
	O Rodents	O Allergic Disease		
MINNE EPITOPE DATABASE		12		
AND ANALYSIS RESOURCE		WWW.IEDB.ORG		

Receptor Data







Assay Search



Default setting (must de-select):

- 'All' assay types
- 'Positive only'

Click open **Finder** to access data tree if you cannot immediately find the one you want



14 www.IEDB.org

MHC Restriction Search







Host Search



Search for host of interest

- Human data
- Animal models





Search by Disease

Represents clinical status of a host:

- Patient history
- Known animal models of disease

Disease 🕐		
Any Disease		
O Infectious Disease		
O Allergic Disease		
O Autoimmune Disease		
🔿 Transplant Disease		
🔿 No Disease (Healthy)		
 Specific Disease 	Ex: asthma, diabetes	Finder

Unique feature

• Allergy

MMUNE EPITOPE DATABASE

AND ANALYSIS RESOURCE

- Autoimmunity
- Healthy controls

Can assess differential reactivity

• Animal model v. human



Reference Search

Reference (?) Any Reference Type Journal Article PubMed ID	Ex: 24196962	
Author	Ev: Galili, Grav	S NCBI Resources 🗹 How To 🖸
Title	Ex: Title Journal, Reyword	Publed gov PubMed
Date (Year)	Ex: 1952 SEx: Vresent	US National Library of Medicine Advanced
Search for interest By Author etc.	r paper of , , PMID,	Display Seturings, © Adsided Seturity: Prochronool. 2014 May;75(5):440-51. doi: 10.1016/j.humimm.2014.02.013. Epub 2014 Feb 12. Substantial gaps in knowledge of Bordetella pertussis antibody and T cell epitopes relevant for natural immunity and vaccine efficacy. Yauchan K ¹ , Seymour E ² , Peters B ² , Sette A ² . @ Author information Abstract The recent increase in whooping cough in vaccinated populations has been attributed to waning immunity associated with the acellular vaccine. The Immune Epitope Database (IEDB) is a repository of immune epitope data from the published literature and includes T cell and antibody epitopes for human pathogens. The IEDB conducted a review of the epitope literature, which revealed 300 Bordetella pertussis-related epitopes from 39 references. Epitope data are currently available for six virulence factors of B. pertussis: pertussis toxin, pertactin, fimbrial 2, fimbrial 3, adenylate cyclase and filamentous hemagglutinin. The majority of epitopes were defined for antibody reactivity; fewer T cell determinants were reported. Analysis of available protective correlates data revealed a number of studies defining epitopes; however few are defined in humans and few have been shown to be protective. Moreover, there are a limited number of studies defining epitopes from natural infection versus whole cell or acellular/subunit vaccines. The relationship between epitope location and structural features, as well as antigenic drift (SNP analysis) was also investigated. We conclude that the cumulative data is yet insufficient to address many fundamental questions related to vaccine failure and this underscores the need for further investigation of B. pertussis immunity at the molecular level.





Finders and Data Trees

Finders

- unique feature of IEDB
- provide standardization and hierarchical organization
- search enormous amount of data using collapsible 'data trees'
- enables big picture or granularity

Data tree

- ontological or taxonomical-based data organizer
- parent/child node relationships
- work 'behinds the scenes' and are accessible to the user
- used to organize:

Antigens (NCBI GenPept, SwissProt, PDB) Non-proteins (ChEBI) Organisms (NCBI taxonomy) Alleles (MRO) Assays (BAO) Disease (Disease Ontology, DO ids)









Accessing Finders

Results Page

۲



IMMUNE EPITOPE DATABASE

AND ANALYSIS RESOURCE

nding Filters Current Filters: 🗶 Positive Assays Only						
Reset	Search				Epitopes	
Epitope 🕐				_	(500301)	
					NH CH NH	
O Linear Epito	De Exact	Mat	Ex: SIIN	IFFKI		
O Discontinuo	us Epitopes	mar				
O Non-peptidic Epitopes Ex: penicillin						
3D structure	e available					
Amino Acid Modification Select Multiple Options					IGK	
		_	58560	SINFEKL		
Antigen			4602	ASNENME	тм	
Organism			112741	2,4-dinitro	ophenyl group	
Ex: influenza,	peanut		20788	GLCTLV	Mouro over	
Antigen Name			130694	1-O-(alpl		ningosine
Ex: core, cap	sid, myosin		24786	HSLGKW	pane to	
			130649	alpha-D-(expand	c-yl grouj
Receptor (?)			48237	PKYVKQ	слрана	
Has recepto	r sequence		112742	2,4,6-trinitrophenyl group		
Туре д			6435	CINGVCWTV		
			32208	KLVALGINAV		
Chain	Any Type 🔻		53112	RAHYNIV	TF	
Sequence	Exact Matches -		61086	SSIEFARL		
			61151	SSLENFR	AYV	
Assay 🕐	THE		16833	FLPSDFF	PSV	
Positive Ass	savs Only		30001	KAVYNFA	ATC	
			65748	TPRVTGG	GAM	
🖌 T Cell Assa	ys		6568	CLGGLLT	MV	
🕑 B Cell Assa	ys		7493	DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA		
MHC Ligand	d Assays		16878	FLRGRAY	′GL	
			17516	FQPQNGQFI		
MHC Restrictio	n 🔋 🚺		522321 Rec	ords Foun	d	
	astriction				-	



Non-Peptidic Epitope Finder

O Any Epitopes			NH CH
O Linear Epitope	Exact Matc 🗸	Ex: SIINFEKL	47.00 SEA
O Discontinuous Epitopes			
Non-peptidic Epitopes	Ex: penicillin	6	Finder

Click 'Finder' to open (below)

Non-protein chemical entities

- metal allergens
- drugs
- lipids, glycolipids, etc.

Current Selection(s)				Reset Apply
Q Search By E Browse by Name: cardiolipin Molecule ID: Ex: 17334 Clear Search Type name molecule	Tree (Click dic material ical substanc (part of mole ule other tha Click	to select)	a Tree cal entities	
Search Results (Click to Select)				
13 Records Found		K < Page 1 of 3 >>		5 • Per Page
Molecule Name	~	Synonyms 🗸 🗸	Database ID 🗸 Organism Name	~
cardiolipin	+ 🎲	CARDIOLIPIN, 1',3'-bis(1,2-diacyl-sn-glycero-3-phospho)-sn-glycerol, Cardiolipin, Diphosphatidylglycerol, cardiolipins, DPG, 1',3'-Bis(1,2-diacyl-sn-glycero-3-phospho)-sn- glycerol	ChEBI:28494	
diphosphatidyl propylene glycol	+ 🎲	DPPG, deoxycardiolipin, 2'-deoxycardiolipin	ChEBI:60358	
reduced cardiolipin	🛨 🎲	reduced cardiolipins, hydrocardiolipin, CLred	ChEBI:60457	
acetyl cardiolipin	2'-O-acetyl-1',3'-bis(1,2-diacyl-sn-glycero-3-phospho)-sn-glycerol, acetylcardiolipin		ChEBI:60353	
2,2'-dilysocardiolipin	+ 🎲	2,2'-dilysocardiolipin, 2,2'-dilyso-1,3-diphosphatidylglycerol, 2,2'-dilyso diphosphatidyl glycerol, 2,2'-dilyso DPG, 2,2'-dilyso diphosphatidylglycerol, dilyso cardiolipin	ChEBI:60431	
13 Records Found		🔣 < Page 1 of 3 👂 🔰		5 Per Page
IMMUNE EPITOPE DATABASE			www.I	EDB.ORG

AND ANALYSIS RESOURCE

Organism Finder

Antigen 🕐			S
Organism			
Ex: influenza, peanut		Finder	
Antigen Name			
Ex: core, capsid, myosin		Finder	
	1946-		e 70 10 3

Click 'Finder' to open (below)

Links to NCBI taxonomy Used to search for:

- epitope source
- immunogen if organism
- assay antigen



Organism Finder

Current Selection(s) 🗶 Leishmania chagasi	<u> </u>	Res	set Apply
Q Search By	E Browse by Tree (Click to Select)		1
Name: leishmania Organism ID: Ex: 10002045	H H Leishmania H H Leishmania aethiopica H H Leishmania braziliensis	Click 'highlight in tree' to	*
Clear Search	 H Leishmania donovani species compl H Leishmania major H Leishmania mexicana species compl Leishmania mexicana species compl Leishmania panamensis Trypanosoma Viridiplantae (green plants) 	see breakdown of available genotypes	
Search Results (Click to Select) 20 Records Found			5 • Per Page
Organism Name		✓ Synchyms	✓ Organism ID ✓
Leishmania	H (Leishmania	5658 🖨
Leishmania major	H	Zeishmania major, Leishmania tropica major, Leishmania (Leishmania) major	5664 🕑
Leishmania pifanoi	+ (Leishmania pifanoi, Leishmania pifani, Leishmania (Leishmania) pifanoi	5682 🗳
Leishmania braziliensis	H	Z Leishmania braziliensis, Leishmania (Viannia) braziliensis, Leishmania brasiliensis, Leishmania viannia	5660 🕼
Leishmania chagasi		Leishmania chagasi, Leishmania chagasi subsp. chagasi, Leishmania chagasi chagasi, Leishmania (Leishmania) chagasi, Leishmania donovani chagasi, Leishmania infantum chagasi	44271 🗗
Can click 'Plus Apply to add cr Current Selecti	sign' and then iteria from on(s)	Page 1 of 4 22	5 • Per Page 23 B • B • B • C

Example of Organism Data Tree



Click to open successive nodes to find specific strain or use high node to look at all available





Antigen Finder

Antigen 🕐		5.3
Organism		
Ex: influenza, peanut	Finder	
Antigen Name		
Ex: core, capsid, myosin	Finder	

Click 'Finder' to open (below)

Links to NCBI GenPept Used to search for:

- epitope source
- immunogen if protein
- assay antigen source

MOLECULE FINDER 0	
Current Selection(s)	
Q Search By	E Browse by Tree (Click to Select)
Name:	E imaterial entity
	🕀 📄 non-peptidic material
Ex: p69710	🖂 🚞 protein
	Archeobacterium protein
Ex: influenza, pranut Finder	
Influenza virus (ID:10002045)	in
Influenza A virus (ID 11320)	
Influenza B virus (ID:11520)	tein
Influenza C virus (ID:1152)	
 Haemophilus Influenzae (ID: N27, Influenzae) Parainfluenza virus 5 (ID: 11207) 	nza-bacilide)
Mo recure manie	✓ Synonyms ✓ Database ID ✓ Organism Name
Ту	pe protein and organism





Antigen Finder

Q Search By	🔋 Browse by Tree (Click to Se	lect)		
Name: HA Molecule ID: Ex: P69710 Source Organism: Influenza A virus (ID: Finder Clear Search	Image: material entity Image: material enti	ein tein		
Use top	o row to sort			
Search Results (Click to Select)				
70 Records Found		K Page 1 of 14 🕨 🔰		
Molecule Name	✓ Synonyms	×	Database ID 👻	Organism Name
haemagglutinin [SRC293731]	🛨 🌠 hemagglutinin, HA		IEDB [293731]	Influenza A virus
Hemagglutinin ★★ (?)	HA2, HA1, Hemagglu Structure Of Broadly Arvetoria/361/2011	utinin precursor, hemagglutinin H5, Chain A, Crystal r Neutralizing Antibody F045-092 In Complex With (h3n2) Influenza Hemagglutinin, heamagglutmore	UniProt [P03452] 🕑	Influenza A virus
hemagglutinin gene [2271249]	🛨 🌠 на		GenPept [2271249] G	Influenza A virus
hemagglutinin gene [2271251]	Н А	Click Highlight in	GenPept [2271251]	Influenza A virus
hemagglutinin gene [2271273]	🛨 🌠 🗛 1, HA	Tree to see what's	GenPept [2271273]	Influenza A virus
70 Records Found		available		



Example of Antigen Tree

Click high node 'Hemagglutinin' to retrieve **all** HA from all flu







Receptors search (no finder)

otor sequence		
Any Type 🔻		18,292 total
Any Type Region CDR3 Katches Exact Matches Ex: CARNTGNQFYF]	
	Any Type Any Type Any Type Region CDR3	Any Type Any Type Region CDR3 Exact Matches Ex: CARNTGNQFYF







Receptors results summary

Epitopes					Receptors				
T Cell Rece (16949)	ptors B Cell F	Rece 1343)	eceptors 43)			(10292)			
			Go To						
16949 Record	ds Found				1< <				
Group ID 🔨	Species	~	Туре 🗸 🗸	Chain	1 CDR3	~	Chain 2 CDR3		
47 🌄	Homo sapiens (human)		αβ	IVVRS:	SNTGKLI		ASSQDRDTQY		
49 🌄	Mus musculus (mouse)		αβ	AASAN	ISGTYQR		ASGDAGGGYEQY		
50 🏹	Mus musculus C57BL/6		αβ	AAS			ASSL		
57 🌄	Homo sapiens (human)		αβ	AALIQ	GAQKLV		ASTYHGTGY		
94 🌄	Homo sapiens (human)		αβ	AVRPL	LDGTYIPT		ASSYLGNTGELF		
102 🌄	Mus musculus (mouse)		αβ	ALSEN	YGNEKIT		ASGDASGAETLY		
103 🌄	Mus musculus (mouse)		αβ	ALSEN	YGNEKIT		ASGDASGGNTLY		
104 🌄	Mus musculus (mouse)		αβ	AANSO	STYQR		ASGDFWGDTLY		
109 🌄	Homo sapiens (human)		αβ	IVWGG	YQKVT		ASRYRDDSYNEQF		
110 🏾 🌠	Homo sapiens (human)		αβ	AVTTD	SWGKLQ		ASRPGLAGGRPEQY		
111 🏾 🌠	Homo sapiens (human)		αβ	AVTTD	SWGKLQ		ASRPGLMSAQPEQY		
114 🏾 🌠	Mus musculus (mouse)		αβ	AVSDP	PPLLT		ASGGGGTLY		
115 🏾 🌠	Mus musculus (mouse)		αβ	AVSLE	RPYLT		ASGGGGTLY		
116 🏾 🌠	Homo sapiens (human)		αβ	ALSGF	YNTDKLI		ASPGLAGEYEQY		
117 🏾 🌠	Homo sapiens (human)		αβ	AVRPT	SGGSYIPT		ASSYVGNTGELF		
118 🏾 🌠	Mus musculus (mouse)		αβ	ALFLA	SSSFSKLV		ASSDWVSYEQY		
125 🏾 🏹	Homo sapiens (human)		αβ	ATDTT	SGTYKYI		SARDLTSGANNEQF		





Assay Finder







Allele Finder



MHC restriction defined in assay Alleles from 16 species Search at level of:

٤G

- Species
- Class
- Serotype
- Allele

ALLELE FINDER 0		×
Current Selection(s)		Reset Apply
Q Search By Name: Fx: HI A-A*02:01, H	E Browse by Tree (Click to Select)	
Organism:	MHC molecule	
Clear Search	bonobo (Pan paniscus) cattle (Bos taurus) chicken (Gallus gallus) chimpanzee (Pan troglodytes)	
	dog (Canis lupus familiaris) gorilla (Gorilla gorilla) horse (Equus caballus) human (Homo sapiens) mouse (Mus musculus)	
MHC class I MHC class II	organism (all species) pig (Sus scrofa) rat (Battus powericus)	
non-classical MHC	rhesus macaque (Macaca mulatta) sheep (Ovis aries)	31
AND ANALYSIS RESOURCE	ASE	www.IEDB.o

Host Finder

Host (?) (Any Host Humans Rodents Non-human Primates Specific Host	Ex: dog, camel	Subject or animal model in which the immune response was defined
HOST ORGANISM FINDER 0		×
Current Selection(s)		Reset
Q Search By Name: human Organism ID: 9606 Clear Search Examples	 Browse by Tree (Click to Select) Vertebrate Ave (bird) Fish Fish Canual Glire (rodent or rabbit) Camelid Carnivore Marsupial 	
Search Results (Click to Select)		
1 Records Found	R Rage 1 of 1 D D	5 👻 Per Page
Organism Name	✓ Synonyms	✓ Organism ID ✓
Homo sapiens (human)	🛨 🌠 Homo sapiens, man, human	9606 🕑
1 Records Found	B Bage 1 of 1 DD	5





Disease Finder

Disease (?) Any Disease Infectious Disease Allergic Disease Autoimmune Disease Transplant Disease No Disease (Healthy) Specific Disease Ex: astignment 	hma, diabetes	 4 Broad d Specific d Includes a Easy way disease (A 	ise ise ani to AI)	ease categories ease types (RA, a mal models (e.g. find data for nor	asthm EAE n-infec	a)) ctious
DISEASE FINDER 0						×
Current Selection(s)		N			Reset	Apply
Q Search By	Browse by Tree (Click to	Select)				
Disease Name: asthma	-) 🚞 host health status [DTREE	E_00000185]				
Disease ID:	E Collocation disease [DOID:4]					
ID Source:	transplant-related o	disease and allo-reactivity [DTREE_00000016]				
	allergy [DOID: 1205]					
Clear Search	🕀 📄 autoimmune disea	se [DOID:417]				
	🕀 📄 additional diseases	s by category [DTREE_00000013]				
Example	Leithy [DTREE_0000	0014]				
Search Results (Click to Select)						
2 Records Found		Reference Page 1 of 1			5	✓ Per Page
Disease Name	Synony	ms	*	Disease ID		*
allergic asthma [DOID:9415]	+ 🔯 atopic as status as asthma, .	thma, extrinsic asthma with acute exacerbation, extrinsic asthma with sthmaticus, Extrinsic asthma with status asthmaticus (disorder), atopic 145.0		DO [DOID:9415] 🗗		
non-allergic/occupational asthma [DOID:2841]	toronchial bronchial chronic o with state	hyperreactivity, Bronchial hypersensitivity, chronic obstructive asthma, obstructive asthma with acute exacerbation, chronic obstructive asthma us asthmaticus, Exercise indumore		DO [DOID:2841] 🗗		
2 Records Found		I< < Page 1 of 1 >>>			5	- 3.3age
AND ANALYSIS RESOURCE				V	vww.L	EDB.OR

Specialized Searches

1. "Details" searches

- -Search all database fields
- -Formulate more specific queries

2. Identifier search

-Query using unique IDs

Browse by 3D Structure
 Structural data organized in a tree





Examples of Homepage vs. Details Search Capabilities

Query	Homepage	Details
Sequence	 ✓ 	~
Source antigen	 ✓ 	~
Source organism	~	~
MHC class	~	~
MHC allele	~	~
Host strain	 ✓ 	~
Specific assay type	 ✓ 	~
Immunized vs. naturally infected		~
Immunogen/Antigen		v
Effector cell type/Antibody isotype		V



www.IEDB.org

Specialized Search – Epitope Details


Epitope Detailed Search	Pending F	Filters				
Reset Search		Epitopes (543637)		Antigens (35440)		Assa (11867
Epitope 🕐		O R.		Go To Records	Starting At 1200	©
Epitope ID	Ex: 44920			K < Page	1 of 21746 >	21
Structure Type - Any Epitopes			~	Antigen	~	Organis
Question			7.			
Organism	Ex: influenza, Pean	iut 📔 Finder	7.	65 kDa phosphoprotein	7.	Human h
Antigen Name	Ex: core, capsid, m	yosin 📔 Finder	7.	Matrix protein 1	۳.	Influenza
Epitope Reference Details	\$		Y.	Myelin-oligodendrocyte glycoprotein	74	Mus mus
Epitope Structure Defines	Select Multiple Options	\$	7.	Gal d 2	7 .	Gallus ga
Evidence Code for Source	Select Multiple Options	\$	7.			L
Epitope Name	Ex: Fab-12 epitope		Search	hy enitone n	ame	nfluenza
Reference Start Position	Ex: 124 to Ex: 1	30			lame,	lus mus
Reference End Position			start/st	top positions	, etc.	łuman h
	EX: 130 EX: 1	30	anoj		•	
Reference Region	Ex: Reference Region		->4)-D-GICpNAc-yl group			
Comments	Ex: comments		'+	Canama polypratein		Llanatitia
Data Location in Reference	Ex: location_of_data_in	_reference	·+	Hemagglutinin	**	Influenza
Epitope Related Object			T.	Genome polyprotein		Henatitis
Related Object	Select Multiple Options	÷	7,	Other Human papillomavirus protein	7.	Human p
Type - Any Type	Select AllUnse					luman h
	The epitope is an an	alog of:	Search	tor analogs	or	lepatitis
Organism	The epitope is a neo	-epitope of:	mimot	0000		.ymphocy
Antigen Name				opes		luman h
		HRITINEV	74	Nucleoprotein	7 4	Influenza
Reference 🛞	7493	DAEFRHDSGYEVHHQKLVFFAE	DVGSNKGAIIGLMVGGVVIA 🏾 🏹	Amyloid beta A4 protein	7.	Homo sa
Author	61151	SSLENFRAYV	Y.	Polymerase acidic protein	7.	Influenza
Title	17516	FQPQNGQFI	7.	Nucleoprotein	7.	Lymphoc
1						

An Example Analog Epitope

Epitope		
Epitope ID	107004	_
Chemical Type	Linear peptide	-
Linear Sequence	YAKQATLALA S	equence studied

Epitope Reference Details						
Epitope Structure Defines	Exact Epitope					
Epitope Name	aHAP (308–317)					
Location of Data in Reference	Materials and Methods					

Epitope Related Object		
Related Object Type	The epitope is an analog of:	
Chemical Type	Linear peptide	
Linear Sequence	YVKQNTLKLA	Natural analog of the
Starting Position	308	
Ending Position	317	sequence studied
Source Molecule Name	hemagglutinin HA1	
Source Accession	223282 🗳	
Source Organism ID	11320	
Source Organism	Influenza A virus	





Antigen Name							
Epitope Reference		113645	MEVGWYRSPFSRVVHLYRNGK	. 🏹	Myelin-oligodendrocyte glycoprotein	74	Mus mu:
		20354	GILGFVFTL	7.	Matrix protein 1	74	Influenza
Epitope Structure Defines	5	58560	SIINFEKL	7.	Gal d 2	74	Gallus g
Evidence Code for Sourc	e	112741	2,4-dinitrophenyl gro	up 🏹			
Epitope Name		4602	ASNENMETM	7.	Nucleoprotein	7.	Influenza
Reference Start Position		24786	HSLGKWLGHPDKF	7.	Myelin proteolipid protein	74	Mus mu:
Reference End Position		130649	alpha-Gal epitope	7.			
Reference Region		20788	GLCTLVAML	7.	mRNA export factor ICP27 homolog	74	Human ł
Comments		112742	2,4,6-trinitrophenyl	group 🏹			
Data Location in Referent	ce	130694	1-O-(alpha-D-galactos N-hexacosanoylphytosp	yl)- 🏹			
		6435	CINGVCWIV	7.	Genome polyprotein	7.	Hepatitis
Reference 🕐				74	Hemagglutinin	7.	Influenza
Author	Eur Callin C			7.	Genome polyprotein	7.	Hepatitis
Aution The	EX: Gaill, G	rey		74	Envelope glycoprotein B	74	Human h
litte	Ex: Title, Jo	ournal, Keyword		74	Protein E7	74	Alphapa
Reference Details				7.	Pre-glycoprotein polyprotein GP complex	74	Lymphoo
Reference ID	Ex: 31512	.0		7.	Latent membrane protein 2	74	Human ł
Abstract	Ex: escap	e mutant			Annulaid bake Ad anatain	,	Lines) Sa
Affiliations	Ex: Nation	nal Jewish			arch by author(s)	
Date (Year)	E 4052	to				~ <i>)</i> ,	titis
Dute (reary	EX: 1952	Ev:	Procent	jou	rnal, etc.		172
		<u> </u> L.	Freschi	7.	Polymerase acidic protein	7.	Influenza
Type - Any			~	7 .	Nucleoprotein	7.	Lymphor
Type - Any Type - Journal Article				.	Enstein-Barr nuclear antigen 3	7	Human k
Type - Submission) `		••	
Poset Coard		429919 Re	ecoras Founa		K K Page 1	of 1/19/	
Search					Go To Records Starti	ng At 1200	60
IMMUNE EPITOPE DA AND ANALYSIS RESOU	TABASE JRCE				www.II	EDB.or	RG

Specialized Search – Assay Details (T, B, MHC)



Assay Details – In Vivo Processes



Assay Details – In Vivo Process Types

Epitope Referenc		20354	GILGFVFTL		7.	Matrix protein 1
Epitope Related		58560	SIINFEKL			
		4602	ASNENMETM		"Adn	ninistration group"
Host @		24786	LIGE CRAWLOUDDING		Aun	initistration group
HUSI ()						
Host Organism	Ex: dog, can	nel	들 Finder	2		
Host Details				··· ·		
E 1st In Vivo Process				1.2 T		
Present in Search Results	May or may i	not be preser	nt 💌			Occurrence group"
In Vivo Process Type	Select Multip	le Options	\$			
Disease State	Select All	Unselec				
Disease Stage	Administra	ation in vivo ation in vivo to	o cause disease			
Administration Details	Administra	ation in vivo to	o prevent or	i <i>N</i> rih		"Exposure group"
🕒 1st Immunogen		e of infectiou	s disease 👻			
1 2nd In Vivo Process						
🕒 2nd Immunogen						
In Vitro Administration				Others:		
🕒 In Vitro Immunogen					tranaf	usion
H Immunization Comments					u ansil zation	121011
Adoptive Transfer				Unknown	Lauvii	
	-		VEH 1 151NEV 11 1511			
Assay 🕑 🛛 🦯	49)	23376	GYKDGNEYI	All process types	are defi	ned in the Curation Manual

Assay Details – Immunogen-Epitope Relation

Epitope Reference	20354	GILGFVFTL	74	Matrix protein 1
	58560	SIINFEKL	7.	Gal d 2
	4602	ASNENMETM	7.	Nucleoprotein
(m. 1.0)	24786		74	Myelin proteolipid protein
Host ③		2	osanoylphytosphingosine 🍒	
Host Organism Ex: d	og, camel	들 Finder	7,	mRNA export factor ICP27 homolog
Host Details			7,	
3 1st In Vivo Process			7.	Genome polyprotein
🗎 1st Immunogen			7,	Genome polyprotein
Present in Search Results	or may not be present		7,	Protein E7
May	or may not be present			Envelope alvcoprotein B
Epitope Relation - Any Epitope Relation - Epitope Epitope Relation - Source Antigen Epitope Relation - Source Organism Epitope Relation - Fragment of Source Epitope Relation - Derivative of Sour Epitope Relation - Other Structure fr Epitope Relation - Taxonomic Parent Epitope Relation - Taxonomic Sibling Epitope Relation - Taxonomic Child Epitope Relation - Structurally Relate Epitope Relation - Structurally Relate Epitope Relation - Other	ce Antigen ce Organism om Source Organism : ! ed		 What is the between the immunog Equivaler assay and wherever 	the epitope and the en in the assay? In fields exist in the tigen fields, and there are processes.
In Vitro Immunogen			14	Myelin basic protein
			7.	Listeriolysin O
Adoptive Transfer				43 C Page 1 of 4
AND ANALYSIS RESOURCE				WWW.IEDD.UKG

Identifier Search

A quick way to retrieve data by using an "inventory number"

Home	Specialized Searche	es Analysis Resource	_				
Tionic	operialized ocarent		-				
ositive Assays Only		_	_				
	IDENTIFIER SEA	RCH	×				
9852)			_				
	IEDB Identifiers	5					
	Epitope ID	Ex: 44920					
1	Reference ID	Ex: 315120		ļ	Unique identifiers c	reated by IED	В
pin	Submission ID	Ex: 1000548					
V	Assay ID	Ex: 1710106					
I.				K			
-	External Identifi	iers					
	PubMed ID	Ex: 24196962					
trophenyl group	PDB ID	Ex: 4NM8		ļ	 Unique identifiers fr 	om other resc	urces
M							
HPDKF	ChEBIID	Ex: CHEBI:28494					
герторе							
initrophenyl group			Search				
L							
ha-D-galactosyl)- sanoylphytosphingos:	ine 🍾						
v	🏹 Ge	enome polyprotein			44		
IMMUNE EPITO	OPE DATABASE S RESOURCE					www.IEDB.	ORG

Identifier Search

IEDB Identifiers	5						
Epitope ID	Ex: 44920						
Reference ID	Ex: 315120						
Submission ID	Ex: 1000548						
Assay ID	Ex: 1710106						
External Identifiers							
External Identifi	ers						
External Identifi PubMed ID	ers 22311355						
External Identifi PubMed ID PDB ID	ers 22311355 Ex: 4NM8						
External Identifi PubMed ID PDB ID ChEBI ID	22311355 Ex: 4NM8 Ex: CHEBI:28494						

AND ANALYSIS RESOURCE

Pending Filters

IDENTIFIER SEARCH

Current Filters: 🗶 Reference Type: Journal Article 🗶 PubMed Id: 22311355

Reset Search		Epitopes (9)	Antigens (1)		Assays (17)	Receptors
Epitope ?		· · · [Go To Records Starting At 1200	
Any Epitopes	9 Records Fo	ound			Page 1 of 1 D	
O Discontinuous Epitopes	Details 🗸	Epitope	*	Antigen	~	Organism
O Non-peptidic Epitopes	3078	AMDSNTLEL	7.	Nucleoprotein	۳.	Influenza A virus
	6615	CLPACVYGL	7.	Nucleoprotein	74	Influenza A virus
3D structure available	21255	GMDPRMCSL	7.	Nucleoprotein	7.	Influenza A virus
Amino Acid Modification	32157	KLSDYEGRL	7.	Nucleoprotein	74	Influenza A virus
	36516	LIFLARSAL	7.	Nucleoprotein	۲.	Influenza A virus
Antigen 🔋 🧷 🦾	42974	MVMELIRMI	7.	Nucleoprotein	Y.	Influenza A virus
Organism	54592	RLIQNSITI	7.	Nucleoprotein	۳.	Influenza A virus
Ex: influenza, peanut	144292	FQGRGVFEL	7.	Nucleoprotein	7.	Influenza A virus
Antigen Name	164335	QLSTRGVQI	7.	Nucleoprotein	¥.	Influenza A virus
Ex: core, capsid, myosin	9 Records Fo	bund			Page 1 of 1 D	
					Go To Records Starting At 1200 60	
Receptor (?)						
IMMUNE EPITOPE I	DATABASE					ITDD

www.IEDB.org

Browse by 3D Structure

Branches of tree organized by organism that is source of antibody, T Cell, and MHC molecule, respectively.





www.IEDB.org

Browse by 3D Structure

BROWSE BY STRUCTURE @

AND ANALYSIS RESOURCE





Browse by 3D Structure

Cur	rent Filte	ers: 🗙 Positive Assays	s Only 🗙	No B cell assays) 🗙 No MHC	ligand assays) 🔀 MH	IC Restriction	Type: HLA-A*02:01 🔀 3D struct	ure available		
		Epitopes (35)		A	ntigens (15)		Assays (62)		Receptors (32)	
	T Cell As (62)	says B Ce	ell Assays (0)	MHC Ligand Assays (0)						
							Go To Records Starting	At A,b 💿		
62 F	Records I	Found					K K Page 1	of 3 🔰 🎦		
ID	~	Reference	~	Epitope	Host	*	Immunization 🗸	Assay Antigen 🗸 🗸	Antigen Epitope Relation	* MHC Res
2119	9223 🕡	yych J himunol 2011	(🍾	AAGIGILTV Melanoma antigen recognized by T-cells 1 (27-35) Homo sapiens	Homo sapiens	Ϋ.	Occurrence of cancer (skin melanoma)	AAGIGILTV Melanoma antigen recognized by T-cells 1 (27-35) Homo sapiens	Epitope	HLA-
161	7229 🗊	Jennifer Buslepp; Immunity 2003	7.	ALWGFFPVL chromosome 15 open reading frame 24 (4-12) Homo sapiens	Mus musculus HLA- Tg	A*0201 🏹	Administration in vivo with chromosome 15 open reading frame 24 (Source Antigen) followed by restimulation in vitro	ALWGFFPVL chromosome 15 open reading frame 24 (4-12) Homo sapiens	Epitope	HLA-
188	3845 🇊	David K Cole; J Biol Chem 2009	7.	ELAGIGILTV	Homo sapiens	7.	Primary induction in vitro with ELAGIGILTV (Epitope)	ELAGIGILTV	Epitope	HLA-
198 [.]	1555 🗊	Florian Madura; J Biol Chem 2013	7.	ELAGIGILTV	Homo sapiens	7.	Primary induction in vitro with ELAGIGILTV (Epitope)	ELAGIGILTV	Epitope	HLA-
2118	3926 🇊	Brian G Pierce; PLoS Comput Biol 2014	7.	ELAGIGILTV	Homo sapiens	7.	Occurrence of cancer (skin melanoma)	ELAGIGILTV	Epitope	HLA-
2119	9220 🇊	Oleg Y Borbulevych J Immunol 2011	(; 7 4	ELAGIGILTV	Homo sapiens	7.	Occurrence of cancer (skin melanoma)	ELAGIGILTV	Epitope	HLA-
3134	4129 🗊	InYoung Song; Nat Struct Mol Biol 2017	7.	GILGFVFTL Saturna Satu	Homo sapiens	7.	Exposure to endemic/ubiqitious agent Influenza A virus (Source Organism) without evidence for disease followed by restimulation in vitro	GILGFVFTL Matrix protein 1 (58-66) Influenza A virus	Epitope	HLA-
3134	4130 🗊	InYoung Song;	7.	GILGFVFTL	Homo sapiens	7.	Exposure to	GIL GEVETI	Epitope	HLA-



Chains: MHC-alpha MHC-b2m TCR-Chain 1 TCR-Chain 2 Epitope Chain Calculated contacts: Epitope to MHC Epitope to TCR MHC to Epitope TCR to Epitope



Calculated contacts: Epitope to MHC Epitope to TCR I MHC to Epitope TCR to Epitope

Break 10:30-10:45am







The IB is a unique analysis tool that is integrated <u>into</u> query interface

• What is it?

Maps the immunome - visualize the totality of immune reactivities

- individual proteins and polyproteins

• What is it for?

What are potentially 'immunodominant' epitopes? What regions on an antigen are well-characterized?

Where is it located?

IB icon found on the Antigen Tab of Results page

How does it work?

For a given query, the IB plots the response frequency scores (RFscores) for epitopes along the length of antigen using a reference proteome (per residue).

Designed to **visualize** the entirety or an optimized subset of data





Immunome Browser Example



Hepatitis C virus (HCV)

- Genomic polyprotein ~3,000 aa
- >4,000 epitopes captured in the IEDB
- Abundance reflects iterative coverage
 - multiple labs, assays, hosts
- Can view overall coverage
- Can identify 'choice trees' within the 'forest'
- Selection criteria can be broad or very stringent







Immunome Browser – How it works

A '**HCV**' search from the Home page

- 4,645 epitopes
- >13,700 assays from 598 papers
- Click Immunome Browser icon

to map

Epitopes (4645)	Antigens (3)	Assays (13713)	Receptors (162)	Receptors (162)		
		Go To Records Starting At 1200	60			Export Re
3 Records Found		K K Page 1 of 1 D D			[25 r Pe
Antigen	v	Organism	v	# Epitopes 🗸	# Assays 🗸	# Referen
Genome polyprotein	~	Hepatitis C virus	۳,	4610	13582	596
Other Hepatitis C virus protein	7,	Heparos C virus	7.	20	82	13
Other Hepatitis C virus protein	%	Hepa us C virus	۳,	20	82	13



Response Frequency Graph

Response Frequency 📀



Core	E1	E2	Ns2	Ns3	Ns4A	Ns4B	Ns5A	Ns5B

Top graph: Y-axis = RFscore X-axis = epitope residue position





Assay Counts Graph



Bottom graph: Positive and Negative assay count





IB Data Table

Results Returned	d: 7349 Displaying: 7349 Display Graphed Residue Positions							Export Results
Epitope ID	Epitope Sequence	Mapped Position	Identity	Subjects Tested	Subjects Responded	Assays Positive	Assays Negative	Response Freq. (95% CI)
42674	MSTIPLPZRLTKRNTNRRPZ	1-20	75%	10	6	1	0	0.60 (0.30:0.84)
42704	MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRL	1-44	100%	1	1	1	0	1.00 (0.04:1.00)
42688	MSTNPKPQKKNKRNTNRRPQDVKFPGGGQI	1-30	93%	5	4	1	0	0.80 (0.34:0.99)
42691	MSTNPKPQR	1-9	100%	3	0	0	1	0.00 (0.00:0.61)
42701	MSTNPKPQRKTKRNTNRRPQDVKFPG	1-26	100%	10	10	1	0	1.00 (0.74:1.00)
42693	MSTNPKPQRKIKRNTNRRPQDVKFPGGG	1-28	96%	1	1	1	0	1.00 (0.04:1.00)
42703	MSTNPKPQRKTKRNTNRRPQDVKFPGGGQI	1-30	100%	5	4	1	0	0.80 (0.34:0.99)
42685	MSTNPKPQKKNKRNTNRRPQ	1-20	90%	23	1	1	1	0.04 (0.00:0.18)
42702	MSTNPKPQRKTKRNTNRRPQDVKFPGGG	1-28	100%	60	52	1	0	0.87 (0.76:0.93)
42705	MSTNPKPQRKTKRNTSRRPQDVKFPGGGQI	1-30	96%	30	4	1	0	0.13 (0.05:0.27)
42684	MSTNPKPQKKNKRNTNRR	1-18	88%	14	14	1	0	1.00 (0.81:1.00)
42673	MSTIPKPQRKTKRN	1-14	92%	60	3	1	0	0.05 (0.02:0.14)
42700	MSTNPKPQRKTKRNTNRRPQDV	1-22	100%	1	1	1	0	1.00 (0.04:1.00)
150453	MSTLPKPQRKTKRNTIRRPQ	1-20	90%	1	0	0	1	0.00 (0.00:0.94)
462400	MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVY	1-35	100%	2	2	1	0	1.00 (0.22:1.00)
42681	MSTNPKPQIKTKRNTNRR	1-19	94%	1	0	0	1	0.00 (0.00:0.94)
	Sequence	tion			+/- assay	r counts	5	RFscore





K

IB - stand alone version

A tool to aggregate and visualize <u>user supplied</u> epitopes and antigen [PMID: 29878047]

Home Help Example Reference Contact

ImmunomeBrowser

step 1/3 Specify input data	
	Specify Epitope Data
Enter epitope data in specified format	MALWMRLLPLLALLAL332MALWMRLLPLLALLALWGPDPAAA11ALWMRLLPL5419ALWMRLLPLL5012WMRLLPLLAL73RLLPLLALL7219LLPLLALLALWGPDPAA332PLLALLALWGPDPAA64LALLALWGPDPAAAFVV11LALWGPDPAAAFVNQHLCGS11
Or upload epitope data from a file	Choose File No file chosen
Please select a format for epitope data	White spaced delimited Epitope Data
Please check to autofill missing epitope response data	
	Specify source antigen sequnce/s
Paste source protein sequence in Fasta or plain format	<pre>>seq_P01308.1 MALWMRLLPLLALLALWGPDPAAAFVNQHLCGSHLVEALYLVCGERGFFYTPKTRREAEDLQVGQVEL GGGPGAGSLQPLALEGSLQKRGIVEQCCTSICSLYQLENYCN >seq_NP_000198.1 MALWMRLLPLLALLALWGPDPAAAFVNQHLCGSHLVEALYLVCGERGFFYTPKTRREAEDLQVGQVEL</pre>
Or upload source antigen sequence(s) from a file	Choose File No file chosen
Spe	ecity identity threshold for epitope mapping
Please select identity threshold for epitope mapping	70% 🔻
	Next Step Reset

Located under 'Epitope Analysis Tools' or at 'tools.iedb.org/immunomebrowser'

58

www.IEDB.org



Cancer

Programmatically, cancer is not currently within the scope of the IEDB.

IEDB does contain data as part of in-scope papers: cancer from pathogens and ligand elution MS data

How would I find cancer epitopes?

On Main Page

- Search by antigen e.g. Pmel
- Search by disease ovarian cancer, breast cancer, melanoma, etc.

In Specialize Searches – Epitope Details

Under 'Epitope Related Object'

• 'Neo-epitope of'





Queries with vaccine development in mind

• Select for human data

•	 Natural infection Occurr of infect disease, immunogen = organism Vaccination <i>in vivo</i> administration, immunogen = protein or organism 	 Specialized Search - Process type
•	<i>In vitro assays that define 'correlates of protection'</i> Neutralization, CTL, IFNg, etc. 	
•	 In vivo challenge assays demonstrating protection Survival, decreased pathogen burden Treatment Assays (prophylactic/therapeutic) 	 Assay specific Query
٠	'Coverage' and/or Immunoprominence	ulte Dago Antigon

- Immunome Browser

DATABASE

MMUNE FPIT

AND ANALYSIS RESOURCE

Results Page Antigen Tab

60

WWW.IEDB.ORG



Example Queries





Example query for the epitope sequence ASNENMETM (NP Flu A)



Elements highlighted

- Sequence search
- Homology search
- Filter and drill down
- Results table Summary Data details





Results Page - summarized on tabs

	Epitopes	Antigens		Assays	Receptors	Re	ferences	
2	(5)	(1)		(401)	(190)		(145)	
				Go To Records Starting At 1200			Exp	ort Results 🛃
5 Records F	ound			KK Page 1 of 1 D DI			25	✓ Per Page
Details 🗸	Epitope	۷	Antigen	*	Organism	*	# References 🗸	# Assays 👻
4602	ASNENMETM	¥,	Nucleoprotein	Υ.	Influenza A virus	7.	144	397
4600	ASNENMETM + MCM(E7)	7.					1	1
460 <mark>1</mark>	ASNENMETM + MCM(M6)	7,					1	1
161084	ASNENMETM + GLYC(E4)	T.	Nucleoprotein	7.	Influenza A virus	Y,	1	1
161085	ASNENMETM + GLYC(M6)	7.	Nucleoprotein	7 ,	Influenza A virus	7.	া	1
5 Records F	ound			R Rage 1 of 1 DD			25	✓ Per Page
				Go To Records Starting At 1200 🚳			Exp	ort Results 🕑

- 5 structures, includes structures with modifications
- Source antigen = Nucleoprotein
- 401 assays
- 190 receptors (TCR)
- 145 references
 Can then click through the tabs to review data





Assay 1	ab
---------	-----------



- 351 T cell assays
- 50 MHC Ligand assays
- Use columns to review high level
- Click on assay ID to see all details





References tab

	Epitop	es	Antigens		Assays		Receptors		References
	(5)		(1)		(401)		(190)		(145)
					Go To Records Starting At 1982	2 😡			
145 Records	s Found				🔣 🗹 Page 🔢 🚺 of 6 👂	>1			
Ref ID 🗸 🗸	PMID	Author	×	Title	×	Abstract	•	Date	
1032692	28872670	Akihiro Yoshizawa; Kevin Bi; D Reinhold; Ellis L Reinherz	lerin B Keskin; Guanglan Zhang; Bruce	TCR-pMHC encounter resident CD8 T cells.	differentially regulates transcriptomes of tissue-	To investigate the tissue-resident T compared agains	role of TCR-pMHC interaction in regulating lung CD8 cell (TR) differentiation, polyclonal responses were t NP366-374 /Db and PA224-233 /Db , two immore	2018	
1(29 🏹	28974367 C	M Fehlings; S Chakarov; Y Sim Newell	oni; B Sivasankar; F Ginhoux; E W	Multiplex peptide-MHC analysis of the influen	tetramer staining using mass cytometry for deep za-specific T-cell response in mice.	Antigen-specific immunity against cytometry togeth more	T cells play a crucial role for the host protective viruses and other diseases. The use of mass er with a combinatorial multiplex tetramer staining ha	2018	
1033108 🍸	29274250 C	Simona Anticoli; Francesco Ma Arenaccio; Eleonora Olivetta; F Emiliana Falcone; Anna Ruggie	nfredi; Chiara Chiozzini; Claudia iavia Ferrantelli; Antonio Capocefalo; ri; Maurizio Federico	An Exosome-Based V Immunity Against Vira	′accine Platform Imparts Cytotoxic T Lymphocyte Antigens.	Exosomes are 50 cells. The authors exosomes in vivo more	I-150 nm sized nanovesicles released by all eukaryotic s very recently described a method to engineer with the E7 protein of Human Papilloma Virus (HPV)	2018	
1031880 🍸	28636592	Pradyot Dash; Andrew J Fiore- Shalini Sharma; Aisha Souquet Clemens; Thi H O Nguyen; Kath Philip Bradley; Paul G Thomas	-Gartland; Tomer Hertz; George C Wang; te; Jeremy Chase Crawford; E Bridie terine Kedzierska; Nicole L La Gruta;	Quantifiable predictive repertoires.	e features define epitope-specific T cell receptor	T cells are define receptor (TCR), ti epitopes through more	d by a heterodimeric surface receptor, the T cell hat mediates recognition of pathogen-associated interactions with peptide and major histocompatibi	2017	
1031759 🍸	28615708 C	Lana Vandersarren; Cedric Bo James J Moon; Andrew J East Bart N Lambrecht; Mary J van I	steels; Manon Vanheerswynghels; on; Gert Van Isterdael; Sophie Janssens; Helden	Epitope mapping and of mice in the C57BL/6	inetics of CD4 T cell immunity to pneumonia virus strain.	Pneumonia virus rodent model to s virus (hRSV). Wh more	of mice (PVIM) infection has been widely used as a tudy the closely related human respiratory syncytial nile T cells are indispensable for viral clearanc	2017	
1032055 🍸	26873987 C	Alison J Carey; Donald T Graci Ogan K Kumova; Yvonne M Mu David B H van Zessen; Peter D	as; Jillian L Thayer; Alina C Boesteanu; ielier; Jennifer L Hope; Joseph A Fraietta; Katsikis	Rapid Evolution of the	CD8+ TCR Repertoire in Neonatal Mice.	Currently, there is animal model to s cell (CTL) respon more	s little consensus regarding the most appropriate tudy acute infection and the virus-specific CD8(+) T ses in neonates. TCR high-throughput sequenci	2016	

- Listed by most recent pub year
- Link to PubMed (blue PMID)
- Use columns to review high level
- Click on Ref ID to summary table





Filtering feature

5 Records F	bund	
Details 👻	Epitope	~
4602	ASNENMETM	
4600	ASNENMETM + MCM(E7)	74
4601	ASNENMETM + MCM(M6)	7.
161084	ASNENMETM + GLYC(E4)	Y 4
161085	ASNENMETM + GLYC(M6)	7.

Click on 'funnel' to select an epitope of interest



Tabs will re-set data for only that epitope





Looking for homologous peptides







Refined query at 70% homology reveals variants as well as analogs the flu epitope

			Epitopes (176)		Antigens (2)			Assays (780)
					Go To Records Sta	rting At	1200 💿	
	176 Records	Found			R R Page 1	of 8	> >1	
	Details 🗸	Epitop	e	~	Antigen		~	Organism
	4602	ASNENN	METM .	7.	Nucleoprotein		7.	Influenza A virus
	4578	ASNENN	IDAM	74	Nucleoprotein		74	Influenza A virus
	4630	ASNEN	/ETM	7.	Nucleoprotein		7,	Influenza A virus
	25439	IASNENI	METMESSTLE	74	Nucleoprotein		74	Influenza A virus
	4629	ASNENT	ETM	74	Nucleoprotein		7,	Influenza A virus
	4580	ASNENN	IDTM	74	Nucleoprotein		74	Influenza A virus
	4581	ASNENN	IEAM	74	Nucleoprotein		7,	Influenza A virus
	25434	IASNENI	MDAMESSTL	74	Nucleoprotein		74	Influenza A virus
	4564	ASNENA	AETM	74	Nucleoprotein		7.	Influenza A virus
	4573	ASNENI	ETM	74	Nucleoprotein		7,	Influenza A virus
	4605	ASNEN	MEVM	74	Nucleoprotein		74	Influenza A virus
	25435	IASNEN	MDAMESSTLE	74	Nucleoprotein		74	Influenza A virus
	318	AANENI	NETM	74				
	733	ADNEN	IETM	74				
	1092	AENENN	IETM	74				
	1619	AGNENI	NETM	74				
			ETM	7.	Filter on this			
Identify	analo	bg	тм	74				
of inter	oct	-	ETM	74	epitope			
or mer	031		ETM.	74				
	3356	ANNENN	IETM .	74				
	3662	APNENN	IETM	74	Nucleoprotein		74	Influenza A virus
	4012	AQNENI	METM .	74				
	4200	ARNENN	1ETM	7.				
	4304	ASAENI	NETM	7.				





4 assays reported for this analog – 2 T cell and 2 MHC ligand Explore T cell first....

		Epitopes (1)			Anti (gens 0)		Assa (4)	ys		Ref	eren (2)	ices	
T Cell /	As: 2)	says B Cell As:	says	MHC Ligand	Assays									
							Go To Records Starting	g At Ex: Ab 😡			Ð	port	T Cell Assays Res	uits 🗙
2 Records	Fo	ound					IS S Page 1	of 1 2 2					25 • Pe	er Page
ID V	,	Reference 🗸	Epitop	e v	Host	*	Immunization 🗸	Assay Antigen	*	Antigen ¥ Epitope Relation	MHC Restriction	*	Assay Description	^
1004030		R A Uger; J Immunol 1999	ADNEN	Y. Metm	Mus musculus C57BL/6	₹,	Administration in vivo with Influenza A virus (Structurally Related) followed by restimulation in vitro	ADNENMETM		Epitope	H-2-Db	7,	51 chromium cytotoxicity Positive	
1004029		R A Uger; J Immunol 1999	ADNEN	METM	Mus musculus C57BL/6	7.	Administration in vivo with Influenza A virus (Structurally Related) followed by restimulation in vitro	ADNENMETM		Epitope	H-2-Db	7.	51 chromium cytotoxicity Positive	I
2 Records	Fo	ound			1		IK K Page 1	of 1 2 2			1		25 • Pe	er Page

- Click on the **assay ID** to drill down into details of assay
- Click other assay tabs (MHC Ligand Assays) to review all relevant assays

69

www.IEDB.org



	Deference Information	
	Reference	
	Article Authors:	R A Uger; S M Chan; B H Barber
Evill dataile far	Article Title:	Covalent linkage to beta2-microglobulin enhances the MHC stability and antigenicity of suboptimal CTL epitopes.
Full details for	Reference Detail	
this accove	Reference ID:	1000159
 Reference Epitope Host Immunization 	Abstract:	Many CTL epitopes of clinical importance, particularly those derived from tumor Ags, display relatively poor MHC binding affinity and stability. Because in vivo immunogenicity, and thus the efficacy of peptide-based vaccines, is thought to be determined by MHC/peptide complex stability, there is a need to develop a simple strategy for enhancing the binding of suboptimal epitopes. Toward this goal, the ability to enhance suboptimal peptides through covalent linkage to beta2-microglobulin (beta2m) was explored. Two suboptimal variants of a high-affinity Db-restricted influenza nucleoprotein peptide were covalently linked, via a polypeptide spacer, to the amino terminus of human beta2m and the recombinant fusion proteins expressed in Escherichia coli. When compared with their uncoupled counterparts, the beta2m-linked epitopes display enhanced MHC stabilization and antigenicity. Thus, tethering epitopes to beta2m provides a simple method for augmenting the biological activity of suboptimal peptides and could be useful in the design of peptide-based vaccines or immunotherapeutics.
 Assay type 	Affiliations:	Department of Immunology, Medical Sciences Building, University of Toronto, Toronto, Canada.
	Date:	1999
Antigen	Reference Type:	Literature
	PubMed ID:	10229842
	Journal:	J Immunol
	Journal Volume:	162
	Article Pages:	6024-8
	Journal ISSN:	0022-1767
	Article Chemical List:	Antigens, Viral; Epitopes; Nucleoproteins; Recombinant Fusion Proteins; Vaccines, Synthetic; Viral Core Proteins; beta 2-Microglobulin
	Article MeSH List;	Animals; Antigens, Viral(immunology); Cytotoxicity, Immunologic; Drug Design; Epitopes(genetics; immunology); Humans; Major Histocompatibility Complex(immunology); Mice; Mice, Inbred BALB C; Mice, Inbred C57BL; Nucleoproteins(genetics; immunology); Orthomyxoviridae(genetics; immunology); Protein Binding; Recombinant Fusion Proteins(immunology); T-Lymphocytes, Cytotoxic(immunology); Vaccines, Synthetic; Viral Core Proteins(genetics; immunology); beta 2-Microglobulin(genetics; immunology)

Epitope Information		
Epitope		
Epitope ID:	733	
Chemical Type:	Linear peptide	
Linear Sequence:	ADNENMETM	
Epitope Reference Details		
Epitope Structure Defines:	Exact Epitope	
Epitope Name:	D2	
Epitope Related Object		
Related Object Type:	The epitope is an analog of:	
Chemical Type:	Linear peptide	
Linear Sequence:	ASNENMETM	
Starting Position:	366	
Ending Position:	374	
Source Molecule Name:	Nucleoprotein]
Source Accession:	SRC124]
Source Organism ID:	11320	70
Source Organism:	Influenza A virus	ITDD
	¥¥ ¥¥ ¥	IEDB



T Cell Assay I	Information				
Immunization					
	Host Organism ID:	10000067			
	Host Organism:	Mus musculus C57BL/6			
1st In Vivo Pro	ocess				
	In Vivo Process Type:	Administration in vivo			
Administration	n Details				
	Dose Schedule:	1			
1st Immunogen					
Epitope Relation: Structurally Related					
Object Type: Organism					
Organism ID: 11320					
Organism: Influenza A virus					
Immunogen Details					
Immunogen Evidence Code: Representative selection					
In	munogen Reference Name:	Influenza A virus			
In Vitro Administration					
	In Vitro Process Type:	Restimulation in vitro			
	Responder Cell Type:	Splenocyte			
	Stimulator Cell Type	Splenocyte	Effector Cells		
In Vitro Immu	nogen	opicitocyte	Effector Cell Tissue Type:	Spleen	
III HUO IIIIIU	Enitone Relation:	Enitone	Effector Cell Type:	Splenocyte	
	Chemical Type	Linear pentide	Effector Cell Culture Conditions:	Short Term Restimulated	
	Linear Sequences		Antigen Presenting Cells		
Immunoren Details		Cell Tissue Type:	Lymphoid		
Immunogen Details		Cell Type:	: EL-4 cells-Lymphoblast		
Immuniogen Evidence Code: Exact match to reference		Cell Culture Conditions:	Iture Conditions: Cell Line / Clone		
Innunization	Comments	The yearts of immunization	Autologous or Syngeneic:	Y	
	Immunization Comments:	immunization and CTL der	MHC Allele		
	initialization connector	with 1 µM NP 366-374 per	MHC Allele Name:	H-2-Db	
T Cell Assay			MHC Evidence Code:	Cited reference	
	Oualitative Measurement:	Positive	Antigen		
	Method/Technique:	51 chromium release	Epitope Relation:	Epitope	
	Measurement of:	cytotoxicity	Chemical Type:	Linear peptide	
		-//	Linear Sequence:	ADNENMETM	
			Antigen Details		
			Antigen Evidence Code:	Exact match to reference information	
			Antigen Reference Name:	D2-hβ2m	
			Antigen Containing Object		
			Complex Type:	Protein conjugate	
			Molecule Name:	Beta-2-microglobulin precursor	
			Molecule Accession:	48428791	
			Molecule Source Organism ID:	9606	
			Molecule Source Organism:	Homo sapiens	
			Assay Reference Details		
				CTL specific for NP 366-374 were co-c soluble human β 2-microglobulin or 2)	ultured with EL4 target cells loaded with either 1) D2 peptide plus a fusion protein comprised of the D2 peptide covalently linked to human
			Assay Comments:	β 2-microglobulin (D2-h β 2m). Lysis of the epitope expressed within the fusio The D2-h β 2m fusion protein induced a peptide/protein concentrations.	the two different targets was assessed in a ⁵¹ Cr-release assay to see if on protein could enhance target cell lysis as compared to free peptide. a higher level of specific lysis than the free peptide at lower
IMMUNE EPITOPE DATABASE			Location of Assay Data in Reference:	Figure 4A	
AND ANALYSIS RESOURCE					

Example B Cell Details Search

Find Ebola virus epitopes recognized by neutralizing human monoclonal antibodies.






IMMUNE EPITOPE DATABASE AND ANALYSIS RESOURCE



Home

IMMUNE EPITOPE DATABASE AND ANALYSIS RESOURCE



Home

Immunization Com		112142	2,4,6-trinitrophenyi group		••		
		115009	phosphocholine		74		
Adoptive Transfer		135393	phosphatidylcholine(1+)		74		
		138830	hote D Col (1 x2) hote D ColN	Ac-(1->4)-bata-D-Cal-(1->4)-bata-D-Cic-	7.		
Assay (?)			144				
Qualitative Measurement	Positive		<u></u>		Υ.		7
Assay	Select All	Unselect A		4. Filter for p)OS	itive	
🕀 Measurement Details	Positive Positive-Lov	v		responses	20	nlv	
Assayed Antibody	Positive-Inte	ermediate		Тезропзек			
🕂 Antigen	Positive-Hig Negative	h	v	lpha-N-acetylneuraminosyl-(2->3)-beta osyl-(1<->1')-N-acylsphingosine	i-		
3D Structure of Complex	[DVGSNKGAIIGLMVGGVV	74	Amyloid beta A4 pro	otein
Assay Reference Details	;				74	Myelin-oligodendro	cyte gly
Reference ?		139630	alpha-Neu5Ac-(2->3)-beta-D-G Neu5Ac-(2->8)-alpha-Neu5Ac-((1<->1')-Cer	al-(1->3)-beta-D-GalNAc-(1->4)-[alpha- (2->3)]-beta-D-Gal-(1->4)-beta-D-Glc-	7.		
Author		7491	DAEFRHDSGYEVHHQK		74	Amyloid beta A4 pro	otein
Title		135396	phosphatidic acid		7 4		
Reference Details		59318	SLLTEVETPIRNEWGCRCNDS	SD	74	Matrix protein 2	
Reference ID		319526 Red	cords Found				
Abstract							
Affiliations							
Date (Year)							
Type - Any							
Reset Search							

Immunization Com					
		115009	74		
Adoptive Transfer		135393	phosphatidylcholine(1+)	7.	
		138830	bate D. Col. (1 > 2) bate D. Col.NAc-(1->4)-beta-D-Gal-(1->4)-beta-D-Glc-	7.	
Assay 🕐					
Qualitative Measurement	Positivo			74	
Quantative measurement	rositive			261	av type
Assay	neutraliza	ation biological	act Finder	220	ay type
Measurement Details	0			7.	
Assayed Antibody			>3)-beta-D-galactosyl-(1->3)-N-acetyl-	7.	
+ Antigen			lpha-N-acetylneuraminosyl-(2->3)]-beta- osyl-(1<->1')-N-acylsphingosine		
3D Structure of Complex	(DVGSNKGAIIGLMVGGVV	74	Amyloid beta A4 protein
Assay Reference Details	5			74	Myelin-oligodendrocyte
	_	139650	alpha-Neu5Ac-(2->3)-beta-D-Gal-(1->3)-beta-D-GalNAc-(1->4)-[alpha-	7.	
Reference ③			Neu5Ac-(2->8)-alpha-Neu5Ac-(2->3)]-beta-D-Gal-(1->4)-beta-D-Gic- (1<->1')-Cer		
Author		7491	DAEFRHDSGYEVHHQK	74	Amyloid beta A4 protein
Title		135396	phosphatidic acid	74	
Reference Details		59318	SLLTEVETPIRNEWGCRCNDSSD	74	Matrix protein 2
Reference ID		319526 Re	cords Found		
Abstract					
Affiliations					
Date (Year)					
Type - Any	-				
Deast County					
Kesel Search					

6-2			phospharaylethanola	mine			
Immunization Com		112742	2,4,6-trinitrophenyl gro	oup		74	
Adoptive Transfer		115009	phosphocholine			74	
		135393	phosphatidvlcholine(1	1+)		7.	
Assay 🕐				49 I	c-(1->4)-beta-D-Gal-(1->4)-beta-D-Glc-	74	
Qualitative Measurement	Positive		• 🖉			▼.	
Assay	neutra	lization biological a	t 들 Finder			•• •	
Heasurement Details	1)				ha-Neu5Ac-(2->3)-beta-Gal-(1->3)-beta-GalNAc		
Assayed Antibody						7.	
Source Material	Select	Multiple Options	\$		>3)-beta-D-galactosyl-(1->3)-N-acetyl-	7.	
Immunoglobulin Domain	Select	Multiple Options	¢		osyl-(1<->1')-N-acylsphingosine		
Antibody Purification Status	Monoc	lonal	¢		DVGSNKGAIIGLMVGGVV	7.	Amyloid beta A4 protein
Assayed Antibody Name	Se	ect All] Unselect /	All)			~	••••••••••••••••••••••••••••••••••••••
Heavy Chain Type	Mor Roh	ioclonal			6. Select and	tib	ody
Light Chain Type		clonal-Monospecific			purificatio	on	status
🛃 Assayed Antibody Object	Disp Disp	olay Library Day Library (monoclon	al) 🔽		– "Monoc		nal"
🕒 Antigen							
3D Structure of Complex					3D	1+	Matrix protein 2
🕒 Assay Reference Details							
					1		
Reference 🔅							
Author							

Title

E Reference Details

Reference ID

Abotract

Current Filte	RTS: 🗶 Epitope Organisms: Ebolavirus (ID:186536)	🗙 Host Organism: Hon	no sapiens (human) (ID:9606	, Homo sapian) 🛛 🗶	assay-qualitative_measure	ment: Positive 🔀 assay-as	say_b_cell_data: neutralization biological ac	ctivity (n	eutralization)	
🗙 Assayed A	ntibody Purification Status: Monoclonal									
	Epitopes (15)		Antigens (2)			Assays (39)		Refe	rences (5)	
				Go To Records	Starting At 1200 💿				Ex	port Results 🖒
15 Records F	ound			Reference Page	1 of 1 >>				25	Per Page
Details 🗸	Epitope	~	Antigen		~	Organism		~	# References 🗸	# Assays 🗸
500035	IWKVNPTVD	7.	spike glycoprotein		7.	Bundibugyo ebolavirus		74	1	1
502007	RSNTTGTLIWKV	7.	spike glycoprotein		7.	Bundibugyo ebolavirus		74	1	1
502927	TIYTNGRR	7.	spike glycoprotein		7.	Bundibugyo ebolavirus		74	1	1
503951	L273, W275	7.	spike glycoprotein		7	Bundibugyo ebolavirus		74	1	3
503952	W275	7.	spike glycoprotein		7,	Bundibugyo ebolavirus		74	1	3
503953	Y241, W275	74	spike glycoprotein		7,	Bundibugyo ebolavirus		74	1	3
534854	K114, K115, P116, D117, G118, E120, S142, G143, T1		Envelope glycoprotein		7,	Zaire ebolavirus		74	1	1
539006	Q508, C511, N550, D552	7.	Envelope glycoprotein		7	Zaire ebolavirus		74	1	1
606555	E231, R247, L254, G271, K272, P279	7.	Envelope glycoprotein		7.	Zaire ebolavirus		74	1	4
606556	G528	7.	Envelope glycoprotein		7	Zaire ebolavirus		74	1	5
606557	H628, D632	74	Envelope glycoprotein		7.	Zaire ebolavirus		74	1	3
606558	K510	74	Envelope glycoprotein		7,	Zaire ebolavirus		74	1	3
769835	D624	7.	spike glycoprotein		7.	Bundibugyo ebolavirus		74	1	3
769836	D624, D632	7.	spike glycoprotein		7.	Bundibugyo ebolavirus		7 4	1	3
769837	K633	7.	spike glycoprotein		7.	Bundibugyo ebolavirus		74	1	4
15 Records F	Found			Reference Page	1 of 1 >>				25	Per Page
				Go To Records	Starting At 1200 💿				Ex	port Results 🖒





Current Filters: 🗶 Epitope Organisms: Ebolavirus (ID:186536)	🗙 Host Organism: Homo sapiens (human) (IE	:9606, Homo sapian)	🗙 assay-qualitative_measurement: Positive	🗙 assay-assay_b_cell_data: neutralization	biological activity (neutralization)	
X Assayed Antibody Purification Status: Monoclonal							
Epitopes (15)	AntigensAssaysReferences(2)(39)(5)						
		Go To Record	Is Starting At 1200 00				Export Results 🗳
2 Records Found		Reference Page	1 of 1 >>			[25 Per Page
Antigen	*	Organism		~	# Epitopes 🗸	# Assays 🗸	# References 🗸
Envelope glycoprotein	7. 🔟	Zaire ebolavirus		Υ.	6	17	3
spike glycoprotein	7. 🔟	Bundibugyo ebolavirus		7,	9	22	2
2 Records Found		K Page	1 of 1 >>			[25 Per Page
		Go To Record	Is Starting At 1200 💿				Export Results 🔂





Current Filters: 🗶 Epitope Organisms: Ebolavirus (ID:186536) 🗶 Host Organism: Homo sapiens (human) (ID:9606, Homo sapian) 🗶 assay-qualitative_measurement: Positive 🗶 assay-assay_b_cell_data: neutralization (biological activity (neutralization))										
X Assayed Antibody Purification Status: Monoclonal										
	Epitopes (15)		Antigens (2)		Assays (39)	Refe	rences (5)			
T Cell As (0)	B Cell Assays (39)	MHC Ligand Assays (0)								
	Go To Records Starting At A,b 💿 💿 Export Results 🕑									
39 Records Found I Page 1 of 2 2 2										
ID 🗸	Reference	✓ Epitope	Host 🗸	Immunization 🗸	Assay Antigen 🗸 🗸	Antigen Epitope 🗸 🗸	Assay Assay			
3218082	Anna Z Wec; Cell 2017	G528 GP Zaire ebolavirus	Homo sapiens 🏹	Infectious disease via exposure to Zaire ebolavirus (Source Organism)	Envelope glycoprotein precursor Envelope glycoprotein precursor Reston ebolavirus - Reston	Taxonomic Sibling	biological activity neutralization Positive-Intermediate			
3218080	Anna Z Wec; Cell 2017	Y Y E231, R247, L254, G271, K272, P279 GP Zaire ebolavirus Zaire ebolavirus	Homo sapiens 🏹	Infectious disease via exposure to Zaire ebolavirus (Source Organism)	Envelope glycoprotein precursor Envelope glycoprotein precursor Reston ebolavirus - Reston	Taxonomic Sibling	biological activity neutralization Positive-Intermediate			
2683559	Andrew I Flyak; Cell 2016	Y. L273, W275 Y. envelope glycoprotein Bundibugyo ebolavirus Y.	Homo sapiens 🏹	Infectious disease via exposure to Bundibugyo ebolavirus (Source Organism)	Sudan ebolavirus Sudan ebolavirus	Taxonomic Sibling	biological activity neutralization Positive			
3218077	Anna Z Wec; Cell 2017	K K E231, R247, L254, G271, K272, P279 GP Zaire ebolavirus Control	Homo sapiens 🏹	Infectious disease via exposure to Zaire ebolavirus (Source Organism)	virion spike glycoprotein virion spike glycoprotein Tai Forest ebolavirus	Taxonomic Sibling	biological activity neutralization Positive			
2683556	Andrew I Flyak; Cell 2016	RSNTTGTLIWKV envelope glycoprotein (266-277) Bundibugyo ebolavirus	Homo sapiens 🏹	Infectious disease via exposure to Bundibugyo ebolavirus (Source Organism)	Bundibugyo ebolavirus Bundibugyo ebolavirus	Source Organism	biological activity neutralization Positive			
3218071	Anna Z Wec; Cell 2017	K H628, D632 GP Zaire ebolavirus	Homo sapiens 🏻 🍾	Infectious disease via exposure to Zaire ebolavirus (Source Organism)	envelope glycoprotein envelope glycoprotein Bundibugyo ebolavirus	Taxonomic Sibling	biological activity neutralization Positive			
2683552	Andrew I Flyak; Cell 2016	W275 envelope glycoprotein Bundibugyo ebolavirus	Homo sapiens 🏹	Infectious disease via exposure to Bundibugyo ebolavirus (Source Organism)	Bundibugyo ebolavirus Bundibugyo ebolavirus	Source Organism	biological activity neutralization Positive			
3218072	Anna Z Wec; Cell 2017	H628, D632 To GP Zaire ebolavirus	Homo sapiens 🏻 🏹	Infectious disease via exposure to Zaire ebolavirus (Source Organism)	virion spike glycoprotein virion spike glycoprotein Tai Forest ebolavirus	Taxonomic Sibling	biological activity neutralization Positive			
2683560	Andrew I Flyak; Cell 2016	Y241, W275 envelope glycoprotein Bundibugyo ebolavirus	Homo sapiens 🏻 🏹	Infectious disease via exposure to Bundibugyo ebolavirus (Source Organism)	Ebola virus Ebola virus	Taxonomic Sibling	biological activity neutralization Positive			
3218068	Anna Z Wec; Cell 2017	X E231, R247, L254, G271, K272, P279 GP	Homo sapiens 🏹	Infectious disease via exposure to Zaire ebolavirus (Source Organism)	GP GP Zaire ebolavirus	Source Antigen	biological activity neutralization Positive			





Live Demo (KV) - Dengue virus

Once I run a query, how can I best use the data?

- a. Navigating the tabs
- b. Understanding the summary pages
- c. Downloading the results





Live Demo (NS) - Allergy

How would I search for epitopes defined for food allergens?

- a. Disease Finder data tree
- b. Antigen Finder

e.g. parvalbumin (Gad m 1; Sal s 1)





Live Demo (KV) – T1D

How would I find data related to type 1 diabetes?

- a. patients with diagnosed T1D
- b. auto-antigens related to
- c. compare mouse and human data





Live Demo (NS) – Linear Epitope Search

How to determine known epitopes from a protein sequence of interest.





Live Demo (NS) - T Cell Details Search

Find *Plasmodium falciparum* T cell epitopes tested specifically on CD8⁺ T cells and defined in humans.





Live Demo (NS) - T Cell Details Search

Find *Plasmodium falciparum* T cell epitopes tested specifically on CD8⁺ T cells and defined in humans.

Have tetramers been made with any of these epitopes and used to detect CD8+ T cells?





Lunch 12:30-1:30pm





Exercises continued...





Participant Exercises

1. What antibody epitopes have been defined for dengue virus in humans? Any mAbs? Protective epitopes?

2. What epitopes have been defined for peanut (*Arachis hypogaea*)?

3. What tetramers have been used in the context of influenza A virus?





Participant Exercise

Find HLA-A*02:01 epitopes tested in MHC binding assays, then find HLA-A*02:01 epitopes defined with MHC elution assays.





If time permits





Live Demo (KV)– Receptor data

How would I identify epitopes TCR-MHC interactions?

- a. how many? What diseases?
- b. kinetic measurements
- c. filtering data



