

1. Na PubMedu si najprej odpreš advanced search. iščemo s pomočjo leta izdaje in nahajanja besed v naslovu:

- a. *((("2000"[Date - Publication] : "2000"[Date - Publication])) AND (catalytic antibody[Title])).*

- b. Dobimo dva zadetka. Naš članek je prosto dostopen.

Povezava do članka: DOI: [10.1073/pnas.97.18.9892](https://doi.org/10.1073/pnas.97.18.9892)

## Structural evidence for a programmed general base in the active site of a catalytic antibody

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Affiliations + expand

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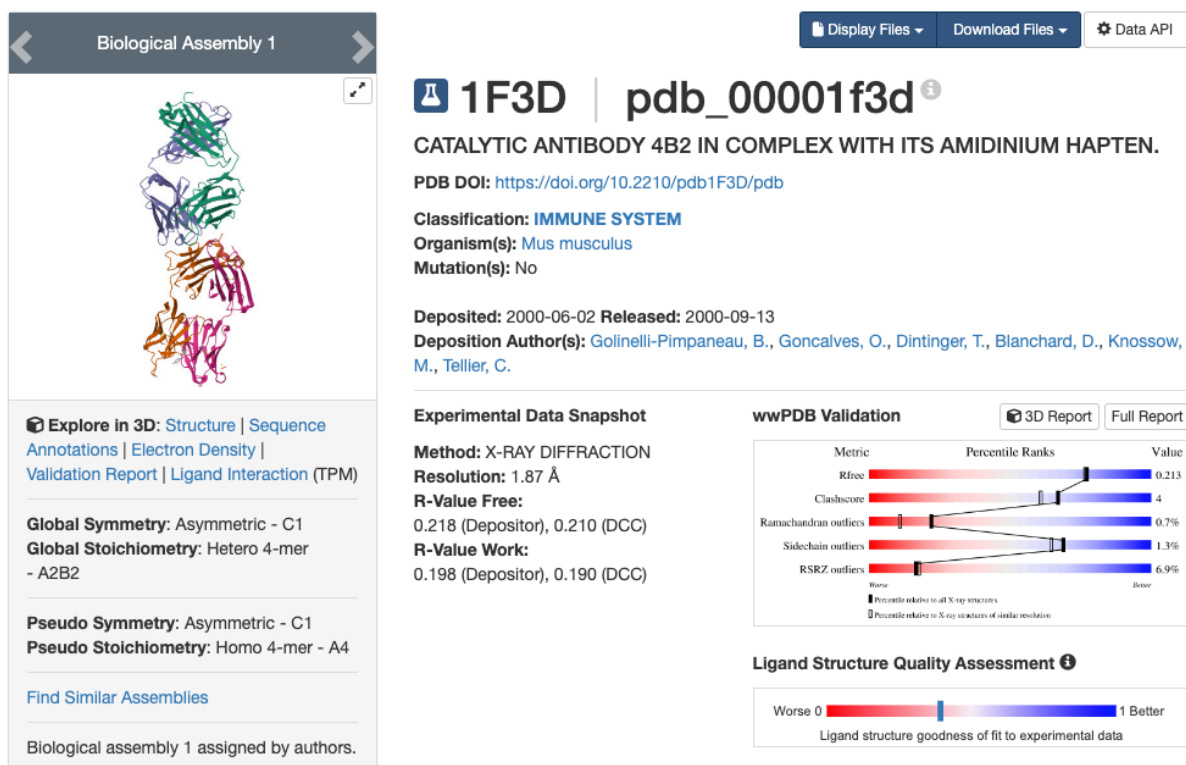
 Cite

### Abstract

The crystal structure of the complex of a catalytic antibody with its cationic hapten at 1.9-Å resolution demonstrates that the hapten amidinium group is stabilized through an ionic pair interaction with the carboxylate of a combining-site residue. The location of this carboxylate allows it to act as a general base in an allylic rearrangement. When compared with structures of other antibody complexes in which the positive moiety of the hapten is stabilized mostly by cation- $\pi$  interactions, this structure shows that the amidinium moiety is a useful candidate to elicit a carboxylate in an antibody combining site at a predetermined location with respect to the hapten. More generally, this structure highlights the advantage of a bidentate hapten for the programmed positioning of a chemically reactive residue in an antibody through charge complementarity to the hapten.

2. Na PDB lahko poiščemo 1F3D in dobimo zadetek našega proteina. Lahko pa vnesemo 4B2 in dobimo dva zadetka. Pravi protein je prvi.

Povezava do proteina na PDB: PDB DOI: <https://doi.org/10.2210/pdb1F3D/pdb>



- Organizem v katerem se katalitično protitelo nahaja je miška (*Mus muscuukus*). Struktura je bila določena z difrakcijo rentgenskih žarkov pri resoluciji 1,87Å. Naš protein ima 2 polipeptidni verigi torej gre za dimer. Ena polipeptidna veriga ima 3 alfa helikse in 19 beta ploskev.



4. Na UniProt pridemo tako, da vpišemo kodo proteina 1F3D. V vezavnem mestu je potreben  $\text{Zn}^{2+}$  ion. Naš monomerni protein tehta 24165Da, dimer pa 48330Da. Imamo pa dva disulfidna mostička. Naša izoelektrična točka je 7,06.

## A2NHM3 · A2NHM3\_MOUSE

Protein <sup>i</sup>	Ig kappa light chain	Amino acids	219 ( <a href="#">go to sequence</a> )
Gene <sup>i</sup>	Igkc	Protein existence <sup>i</sup>	Evidence at protein level
Status <sup>i</sup>	UniProtKB unreviewed (TrEMBL)	Annotation score <sup>i</sup>	1/5
Organism <sup>i</sup>	Mus musculus (Mouse)		

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5. S pomočjo blastp smo poiskali homologa. Prvi vrnjen zadetek, dobimo identičen protein zato izberemo drugega, ki ima najmanjše E vrednost in največji % ujemanja. Naš homologen protein je dolg 219 aminokislinskih ostankov.

Descriptions

Graphic Summary

Alignments

Taxonomy

Sequences producing significant alignments

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100 sequences selected

GenPept

Graphics

Distance tree of results

Multiple alignment

MSA Viewer

	Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
<input checked="" type="checkbox"/>	Ig kappa chain (monoclonal antibody MabA34) - mouse (fragment) [...]	Mus musc...	455	455	100%	3e-161	100.00%	219	PC4203
<input checked="" type="checkbox"/>	Chain B, Light chain of S2H5 Fab [Mus musculus]	Mus musc...	454	454	100%	7e-161	99.54%	219	8K9B_B
<input checked="" type="checkbox"/>	anti-panflavirus glycoprotein specific immunoglobulin light chain [Mu...	Mus musc...	453	453	100%	4e-160	99.09%	238	WIW78149.1
<input checked="" type="checkbox"/>	Chain L, S9.6 Fab LC [Mus musculus]	Mus musc...	451	451	100%	1e-159	99.09%	219	7XLT_L
<input checked="" type="checkbox"/>	Chain F, IgG1 Kappa Light Chain [Mus musculus]	Mus musc...	451	451	100%	3e-159	99.09%	238	6CNJ_F
<input checked="" type="checkbox"/>	Chain B, Fab Light Chain [Mus musculus]	Mus musc...	450	450	100%	3e-159	99.09%	219	3U9U_B
<input checked="" type="checkbox"/>	Chain L, IgG2a Fab fragment Heavy Chain [Mus musculus]	Mus musc...	449	449	100%	6e-159	98.63%	219	2IPT_L
<input checked="" type="checkbox"/>	immunoglobulin kappa light chain [Mus musculus]	Mus musc...	449	449	100%	7e-159	98.63%	238	BCQ06380.1
<input checked="" type="checkbox"/>	Chain K, IgG2a Fab fragment Heavy Chain [Mus musculus]	Mus musc...	448	448	100%	2e-158	98.17%	219	2IPU_K

6. Proteina sta 100% podobna, do edine razlike prihaja v identičnosti in sicer na 30 in 32 mestu.

20-238	1	DVLMQTPLSLPVSLGDQASISCRSSQSILHSNGNTYLEWYLQKPGQSPK	50
		:	
A2NHM3_MOUSE	1	DVLMQTPLSLPVSLGDQASISCRSSQSIVHTNGNTYLEWYLQKPGQSPK	50
20-238	51	LLIYKVSNRFSGVPDRFSGSGSGTDFTLKISRVEAEDLGVYYCFQGSHVP	100
A2NHM3_MOUSE	51	LLIYKVSNRFSGVPDRFSGSGSGTDFTLKISRVEAEDLGVYYCFQGSHVP	100
20-238	101	RTFGGGTKLEIKRADAAPTIVSIFPPSSEQLTSGGASVVCFLNNFYPKDIN	150
A2NHM3_MOUSE	101	RTFGGGTKLEIKRADAAPTIVSIFPPSSEQLTSGGASVVCFLNNFYPKDIN	150
20-238	151	VKWKIDGSRQNGVLNSWTDQDSKSTYSMSSTLTCLKDEYERHNSYTCE	200
A2NHM3_MOUSE	151	VKWKIDGSRQNGVLNSWTDQDSKSTYSMSSTLTCLKDEYERHNSYTCE	200
20-238	201	ATHKTSTSPIVKSFNRECE	219
A2NHM3_MOUSE	201	ATHKTSTSPIVKSFNRECE	219