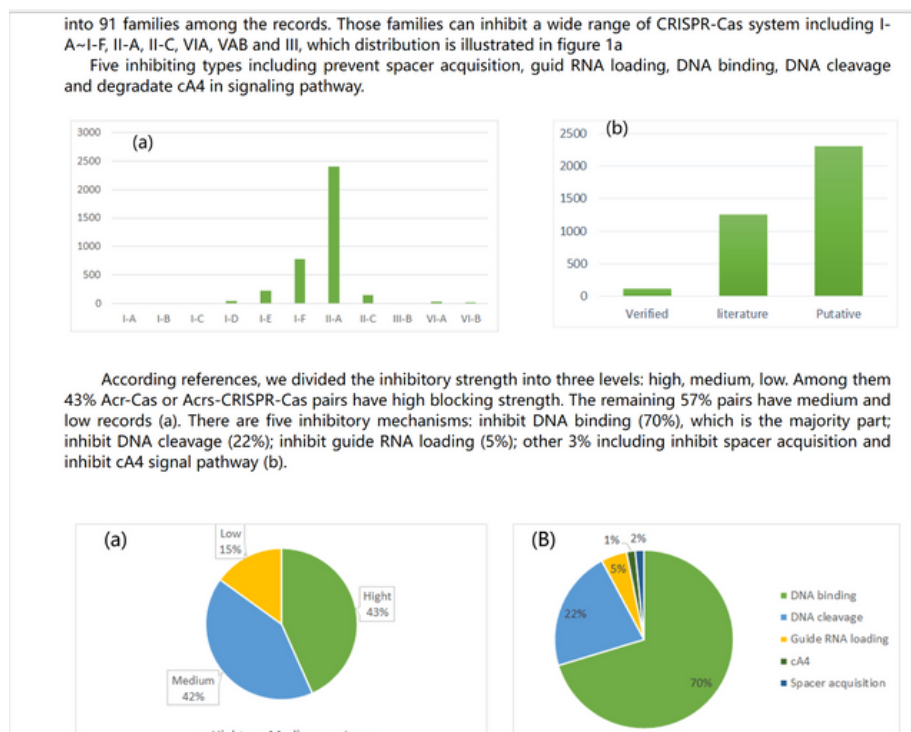


# Phage anti-defence database

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[Verified  
anti-defense proteins](#)
[Predicted  
anti-defense proteins](#)
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## Welcome to Phage anti-defence database


[Anti-CRISPR](#)
[Anti-RM](#)
[Anti-Abi](#)
[Anti-Adsorption  
inhibition](#)
[Other](#)

设置链接，点击可以进入各自的防御系统，展示实验和预测的各种信息

Home	Verified anti-defense proteins	Predicted anti-defense proteins	Search	Prediction	Help
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System family	ID	protein	Inhibited Type	Phage	Host	Length	reference(s)
Anti-Preventing adsorption	APA0001						文章链接
Anti-Restriction–modification	ARM0001		RM I				
Anti-CRISPR/Phage-encoded CRISPR							
Anti-Abi							
Phage nucleus-like compartment							

做一个链接，点进去可以看到这个蛋白有关的信息，例如表格里列举的基础信息、蛋白序列、在基因组上位置的可视化以及二级结构等等

Q Detailed Information

- > Basic Information
- > Protein Structure
- > Secondary Structure
- > Disorder Area
- > Similarity Analysis
- > Multiple Sequence Alignments
- > Phylogenetic Analysis
- > Homology Network Analysis

Basic Information (Experimental Results) [↑ Top](#)

AcrHub ID	Acr00001
Name	-
Gene Name	-
Family	AcrIB
Inhibited CRISPR Type(s)	I-B
Inhibited Stage	-
Molecular Mechanism	-
Targeted Protein	-
Cellular Context	Leptotrichia buccalis DSM 1135 ?
Source Of Species	Leptotrichia buccalis DSM 1135
Function	The protein inhibits type I-B CRISPR-Cas in Leptotrichia buccalis.
Sequence	LESKNLRKLLNEYEEIDINEMLKNFRSIKNSGTKNDIEIFLHEKAIFEKSSISSTYVVFSEDNEILGYFTIANRSLVIPKENFGILSKTQQK KLGNSAAILKNGDLMTSSFLLGQLGKNYSDDIENLITGRELLTFAYDLFLKIKELINVKYIWLECNQNEPKLISFYQNFGFKMLESLTSEEG KVMIMELK
Length	193 amino acids
UniProt ID	-
PubMed ID	<a href="#">32348779</a>

Protein Structure (Experimental Results) ? [↑ Top](#)

Home	Verified anti-defense proteins	Predicted anti-defense proteins	Search	Prediction	Help
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System family	ID	protein	Inhibited Type	Phage	Host	Length	reference(s)
Anti-Preventing adsorption	APA0005						文章链接
Anti-Restriction–modification	ARM0006		RM I				
Anti-CRISPR/Phage-encoded CRISPR							
Anti-Abi							
Phage nucleus-like compartment							
Anti-superinfection exclusion(SE)							
Anti-Thoeris							
Anti-quorum sensing(QS)							
Anti-CBASS							
Anti-Pyscar							

Home	Verified anti-defense proteins	Predicted anti-defense proteins	Search	Run	Help
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## Anti-CRISPRdb

Database of anti-CRISPR proteins



[Home](#) [Browse](#) [Search](#) [Download](#) [Statistics](#) [Help](#) [Version](#) [Reference](#)

### Search by keywords:

Keywords:  Search field:

#### For example:

Search by family using family name such as **AcrIF**.

Search by reference using PubMed ID such as **PMID: 23242138**

Search by strength using **low, medium or high**

高级搜索，通过限定Verified/Predicted anti-defense proteins页面中列举的不同类别的关键词，搜索噬菌体反防御系统的相关信息

Home	Verified anti-defense proteins	Predicted anti-defense proteins	Search	Prediction	Help
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- 1、输入蛋白序列或者fasta格式的文件；
- 2、填写邮箱，预测结果出来后发至邮箱；
- 3、在下方列出示例文件



Run PADLOC for a user-supplied genome.

Versioning: Currently running PADLOC v1.1.0 with defense systems included in padlocdb [v1.4.0](#)

Please enter a name for the job:\*

Required - maximum 80 characters

Please choose how your genome is formatted:\*

NCBI-formatted genome (.gb or .gbff) - preferred format.

☒ Run CRISPRDetect to predict CRISPR arrays? (only available for GenBank and nucleotide input options) **Warning: run times are substantially longer with CRISPRDetect.**

Select files with the file dialog or by dragging and dropping them onto the dashed region.



Select appropriate file(s) for the input format you have chosen above

Example NCBI genome

Example nucleotide fasta

Example protein fasta with GFF

Reset form

IN A NUTSHELL

1. ANTI-CRISPR ANALYSIS

- 1.1 Browse
- 1.2 Search
- 1.3 Statistics
- 1.4 Download
- 1.5 Detailed Information

2. PREDICTION

3. RELATIONSHIP ANALYSIS

- 3.1 Similarity Analysis
- 3.2 Phylogenetic Analysis
- 3.3 Homology Network Analysis

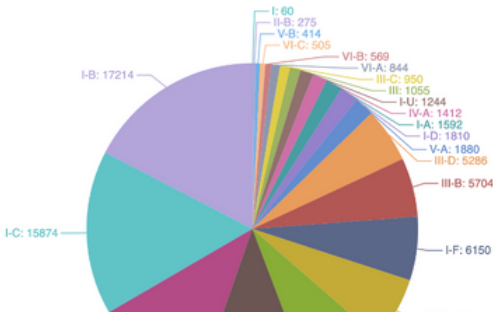
4. ANALYSIS PIPELINE

- 4.1 From Prediction To Relationship Analysis
- 4.2 From Results To Known Acrs

AcrHub ID	Family	Inhibited CRISPR Type	Inhibited Stage	Targeted Protein	Molecular Mechanism	Cellular Context	Source Of Species	Length	PDB	PubMed ID
<a href="#">Acr00308</a>	AcrVA1	V-A	DNA binding	Cas12a	Trigger cleavage of the target-recognition sequence of the Cas12a-bound guide RNA to inactivate the Cas12a complex (block DNA binding)	Pseudomonas aeruginosa PAO1; Human U2-OS-EGFP cells; Moraxella bovoculi strain 58069; Lachnospiraceae bacterium ND2006; Acidaminococcus sp. BV3L6	Moraxella bovoculi	170	-	<a href="#">30190308</a> <a href="#">30190307</a>
<a href="#">Acr00309</a>	AcrVA1	V-A	DNA binding	Cas12a	Trigger cleavage of the target-recognition sequence of the Cas12a-bound guide RNA to inactivate the Cas12a complex (block DNA binding)	-	Moraxella bovoculi	174	-	<a href="#">31155345</a>
<a href="#">Acr00310</a>	AcrVA1	V-A	DNA binding	Cas12a	Trigger cleavage of the target-recognition sequence of the Cas12a-bound guide RNA to inactivate the Cas12a complex (block DNA binding)	-	Moraxella bovoculi	165	-	<a href="#">31155345</a>
<a href="#">Acr00311</a>	AcrVA2	V-A	-	-	-	Pseudomonas aeruginosa PAO1	Moraxella bovoculi	322	-	<a href="#">30190308</a>
<a href="#">Acr00312</a>	AcrVA3	V-A, I-C	-	-	-	-	Moraxella bovoculi	168	-	<a href="#">30190308</a>
<a href="#">Acr00313</a>	AcrVA4	V-A	DNA binding	Cas12a	Recognize the Cas12a pre-crRNA processing	HEK293T human embryonic kidney cells; Moraxella bovoculi strain 58069; Lachnospiraceae bacterium ND2006	Moraxella bovoculi	234	-	<a href="#">30190307</a> <a href="#">31467167</a> <a href="#">31155345</a> <a href="#">31397669</a>
<a href="#">Acr00314</a>	AcrVA5	V-A	DNA binding	Cas12a	Function as an acetyltransferase to modify Cas12a	HEK293T human embryonic kidney cells; Moraxella bovoculi strain 58069; Lachnospiraceae bacterium ND2006	Moraxella bovoculi	92	-	<a href="#">30190307</a> <a href="#">30936526</a>

Showing 1 to 7 of 7 rows

1.3.2 Distribution of predicted Acrs according to their inhibited CRISPR types



数据库的用法介绍