

Chapter 3. DNA Databases

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PPT slides and Message @ <http://jxpt.fafu.edu.cn/meol/homepage/common/>

Email: 1156743645@qq.com



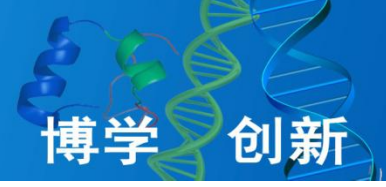
OUTLINES

Lecture 3.1 DNA Sequence Databases

1. DNA databases
2. GenBank overview

Lecture 3.2 Introduction to NCBI's query strategy

1. Database Query Tools
2. Query in NCBI



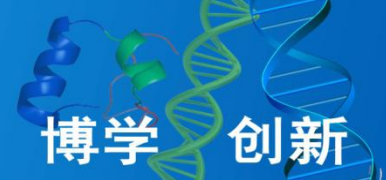
Lecture 3.1 DNA Sequence Databases



1. DNA databases

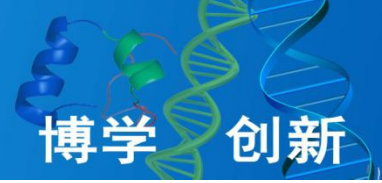
1.1 Primary sequence repositories:

- ① International Nucleotide Sequence Database Collaboration
GenBank (NCBI, USA), EMBL (European Molecular Biology Laboratory), DDBJ (DNA Data Bank of Japan).
- ② Daily data exchange and mirroring



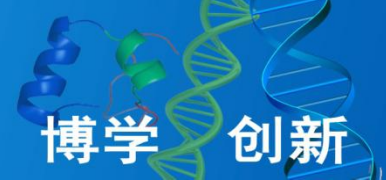
- ③ Sequences submitted to one become available in all.
- ④ Current holdings: 217,971,437,647 bases, 196,120,831 sequences in GenBank.

➔ Very high level of redundancy



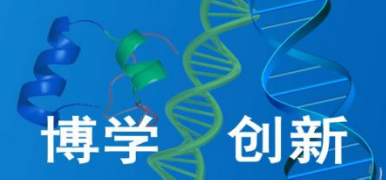
1.2 Other sequence repositories

- BIOSINO- (www.biosino.org, Chinese)
- [dbEST](#) - Database of Expressed Sequence Tags (NCBI)
- [dbSTS](#) - Database of Sequence Tagged Sites (NCBI)



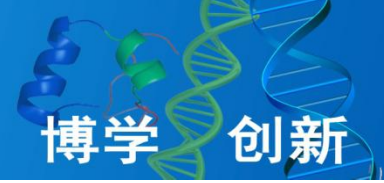
1.3 Genomic Databases

- **GenomeWeb (HGMP)** - A catalogue of useful Web sites for Genomics with a set of collections of genome databases
- **NCGR** – National Center for Genome Research (www.ncgr.org)
- **GOLD** - Genomes On-line db (<https://gold.jgi.doe.gov/>)



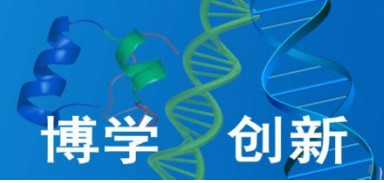
1.4 DNA 3D Databases

- **NDB** - Nucleic Acid Databank (3D structures)
- **BNASDB** - Nucleic acid structure db from University of Pune

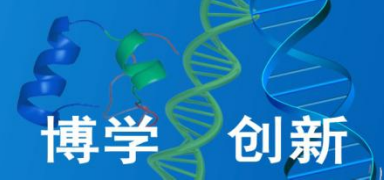


1.5 Others

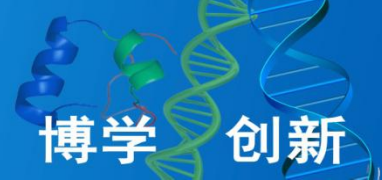
- ✓ [AsDb](#) - Aberrant Splicing db
- ✓ [ACUTS](#) - Ancient conserved untranslated DNA sequences db
- ✓ [Codon Usage Db](#) - The frequency of codon use in each organism is made searchable through this World Wide Web site.



- ✓ [EPD](#) - Eukaryotic Promoter db
- ✓ [PLACE](#) - Plant cis-acting regulatory DNA elements db
- ✓ [HOVERGEN](#) - Homologous Vertebrate Genes db
- ✓ [IMGT](#) - ImMunoGeneTics db [Mirror at [EBI](#)]
- ✓ [ISIS](#) - Intron Sequence and Information System
- ✓ [RDP](#) - Ribosomal db Project
- ✓ [gRNAs db](#) - Guide RNA db



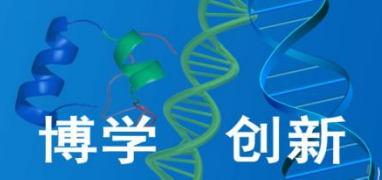
- ✓ [RNA editing](#) - RNA editing site
- ✓ [RNAmdb](#) - RNA modification db
- ✓ [TelDB](#) - Multimedia Telomere Resource
- ✓ [TRADAT](#) - TRAnscription Databases and Analysis Tools
- ✓ [Subviral RNA db](#) - Small circular RNAs db (viroid and viroid-like)



2. GenBank Overview

(www.ncbi.nlm.nih.gov)

- ① Corenucleotide: From whole genome to single gene sequences.
- ② Genomic database
 - *Saccharomyces cerevisiae* 酵母
 - *Caenorhabditis elegans* 线虫



- *Drosophila melanogaster* 果蝇
- *Arabidopsis thaliana* 拟南芥
- *Helicobacter pylori* 幽门螺旋杆菌

③ dbEST (Expressed Sequence Tags)

④ dbGSS (Genome Survey Sequences)



<http://www.ncbi.nlm.nih.gov/genbank/>

NCBI Resources How To Sign in to NCBI

GenBank Nucleotide Search

GenBank Submit Genomes WGS HTGs EST/GSS Metagenomes TPA TSA INSDC

NCBI has completed the https test for today that ran from 8:00-9:00 AM EDT (13:00-14:00 UTC). If you experienced problems with NCBI web sites during that time, please [read more](#) about our https testing.

GenBank Overview

What is GenBank?

GenBank[®] is the NIH genetic sequence database, an annotated collection of all publicly available DNA sequences ([Nucleic Acids Research, 2013 Jan;41\(D1\):D36-42](#)). GenBank is part of the [International Nucleotide Sequence Database Collaboration](#), which comprises the DNA DataBank of Japan (DDBJ), the European Molecular Biology Laboratory (EMBL), and GenBank at NCBI. These three organizations exchange data on a daily basis.

A GenBank release occurs every two months and is available from the [ftp](#) site. The [release notes](#) for the current version of GenBank provide detailed information about the release and notifications of upcoming changes to GenBank. Release notes for [previous GenBank releases](#) are also available. GenBank growth statistics for both the traditional GenBank divisions and the WGS division are available from each release. GenBank growth [statistics](#) for both the traditional GenBank divisions and the WGS division are available from each release.

An [annotated sample GenBank record](#) for a *Saccharomyces cerevisiae* gene demonstrates many of the features of the GenBank flat file format.

Access to GenBank

There are several ways to search and retrieve data from GenBank.

- Search GenBank for sequence identifiers and annotations with [Entrez Nucleotide](#), which is divided into three divisions: [CoreNucleotide](#) (the main collection), [dbEST](#) (Expressed Sequence Tags), and [dbGSS](#) (Genome Survey Sequences).
- Search and align GenBank sequences to a query sequence using [BLAST](#) (Basic Local Alignment Search Tool). BLAST searches CoreNucleotide, dbEST, and dbGSS independently; see [BLAST info](#) for more information about the numerous BLAST databases.
- Search, link, and download sequences programmatically using [NCBI e-utils](#).
- The ASN.1 and flatfile formats are available at NCBI's anonymous FTP server: <ftp://ftp.ncbi.nlm.nih.gov/ncbi-asn1> and <ftp://ftp.ncbi.nlm.nih.gov/genbank>.

GenBank Resources

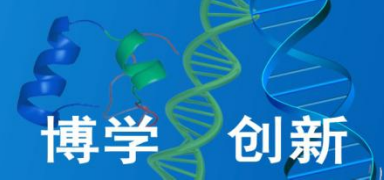
[GenBank Home](#)

[Submission Types](#)

[Submission Tools](#)

[Search GenBank](#)

[Update GenBank Records](#)



About NCBI



Our Mission

NCBI's contribution to the NIH mission of 'uncovering new knowledge'

Organizational Structure

The role of the branches within NCBI and the Board of Scientific Counselors.

Programs & Activities

NCBI's resources for genomic, genetic, and biomedical data

Researchers at NCBI

The basic research program conducted by our intramural investigators

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NCBI News

Leiden Open Variation Database to be retired September 30, 2016

Thursday, September 8, 2016

New on NCBI Insights: Find, Browse and Follow Biomedical Literature with PubMed Journals

Wednesday, September 7, 2016

September 7th webinar: The E-Utilities in an Age without GI Numbers

Thursday, September 1, 2016

October 24-26: Hackathon at Cold Spring Harbor Laboratory



NCBI Home

Resource List (A-Z)

All Resources

Chemicals & Bioassays

Data & Software

DNA & RNA

Domains & Structures

Genes & Expression

Genetics & Medicine

Genomes & Maps

Homology

Literature

Proteins

Sequence Analysis

Taxonomy

Training & Tutorials

Variation


Welcome to NCBI

The National Center for Biotechnology Information advances science and health by providing access to biomedical and genomic information.

[About the NCBI](#) | [Mission](#) | [Organization](#) | [NCBI News](#) | [Blog](#)


Submit

Deposit data or manuscripts into NCBI databases




Download

Transfer NCBI data to your computer




Learn

Find help documents, attend a class or watch a tutorial




Develop

Use NCBI APIs and code libraries to build applications




Analyze

Identify an NCBI tool for your data analysis task



Research

Explore NCBI research and collaborative projects



Popular Resources

PubMed

Bookshelf

PubMed Central

PubMed Health

BLAST

Nucleotide

Genome

SNP

Gene

Protein

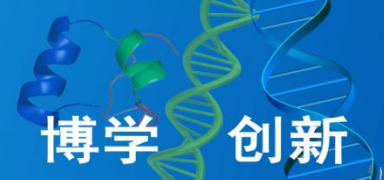
PubChem

NCBI Announcements

Genomes FTP site update (version 1.3) adds new data formats and more
14 Sep 2016

NCBI has released a comprehensive update of all current genome assemblies

RefSeq release 78 is now available
12 Sep 2016



Nucleotide

Nucleotide

U12345



Search

[Advanced](#)

[Help](#)

NCBI is phasing out sequence GI numbers in September 2016. Please use accession.version! [Read more...](#)



Nucleotide

The Nucleotide database is a collection of sequences from several sources, including GenBank, RefSeq, TPA and PDB. Genome, gene and transcript sequence data provide the foundation for biomedical research and discovery.

Using Nucleotide

[Quick Start Guide](#)

[FAQ](#)

[Help](#)

[GenBank FTP](#)

[RefSeq FTP](#)

Nucleotide Tools

[Submit to GenBank](#)

[LinkOut](#)

[E-Utilities](#)

[BLAST](#)

[Batch Entrez](#)

Other Resources

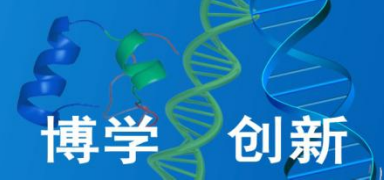
[GenBank Home](#)

[RefSeq Home](#)

[Gene Home](#)

[SRA Home](#)

[INSDC](#)



NCBI Resources How To

Nucleotide

Nucleotide

Advanced

NCBI is phasing out sequence GI numbers in September 2016. Please use accession.version! [Read more...](#)

GenBank

Send to:

Aepyceros melampus isolate am5 D-loop, partial sequence; mitochondrial

GenBank: U12345.1

[FASTA](#) [Graphics](#)

LOCUS AMU12345 426 bp DNA linear MAM 21-SEP-2007

DEFINITION Aepyceros melampus isolate am5 D-loop, partial sequence;
mitochondrial.

ACCESSION U12345

VERSION U12345.1 GI:540023

KEYWORDS

SOURCE mitochondrion Aepyceros melampus (impala)

ORGANISM [Aepyceros melampus](#)

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;
Pecora; Bovidae; Aepycerotinae; Aepyceros.

REFERENCE 1 (bases 1 to 426)

AUTHORS Arctander, P., Kat, P., Simonsen, B. and Siegismund, H.

TITLE Population genetics of Kenyan impalas - consequences for
conservation

JOURNAL (in) Wayne, R.K. and Smith, T.B. (Eds.);

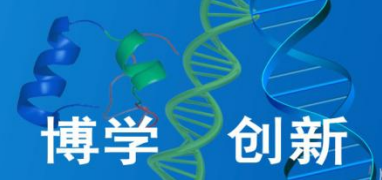
MOLECULAR GENETIC APPROACHES IN CONSERVATION: 399-412;
Oxford University Press, Oxford, U.K. (1996)

REFERENCE 2 (bases 1 to 426)

AUTHORS Arctander, P.

TITLE Direct Submission

JOURNAL Submitted (14-JUL-1994) Peter Arctander, Population Biology,
Zoological Institute, University of Copenhagen, Universitetsparken
15, Copenhagen, DK-2100 O, Denmark



FEATURES

source

Location/Qualifiers

1..426

/organism="Aepyceros melampus"

/organelle="mitochondrion"

/mol_type="genomic DNA"

/isolate="am5"

/db_xref="taxon:[9897](#)"

D-loop

1..>426

/note="control region"

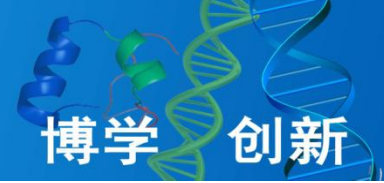
ORIGIN

```

1  actaccgcta tcaatatact cccacaaata tcaagagcct tcccagtatt aaatttgcta
61  aaatttttaa aattcaatac gaacttcaca ctccacagcc tcacgcgaaa ttaataatac
121  gtattttaat tctagagtac ataccatgaa ctatcgttta gtacatgaat ttacacacgt
181  cagcccgatc aaatgtttat gtacataaca cattatatat gtacatttca gtttgtgtat
241  atagacataa cattaatgta ataaagacat aatatgtata tagtacatta attgattgtc
301  ctcaagcata taagcaagta ctagacattc actagcggtg catagtacat ttcattgttc
361  atcgtaata ggcgatgtca gncaaatacg ttcttgtcaa catgcatatc ccgtccacta
421  gatcac

```

//



Sequence Formats

□ FASTA

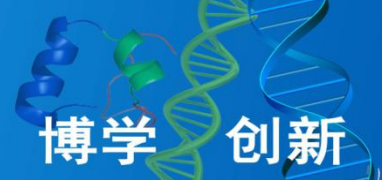
--Very simple “sequence only” format

--Specification:

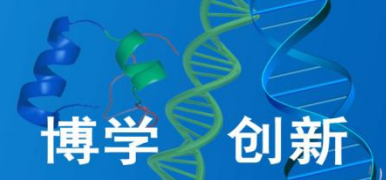
1. Start sequence entry with “>” on a new line
2. Provide sequence on next lines in single character code

□ Example:

```
>gi|14286176|sp|P00528|SRC1_DROME Tyrosine-protein kinase Src64B
```



MGNKCCSKRQDQELALAYPTGGYKKSDYTFGQTHINSSGGGNMGGVLG
QKHNNGGSLDSRYTPDPNHRGPLKIGGKGGVDIIRPRTTPTGVPGVVLKR
VVVALYDYKSRDESDLSFMKGDRMEVIDDTESDWWRVNLTTRQEGLIP
LNFVAEERSVNSEDWFFENVLRKEADKLLLAEEENPRGTFLVRPSEHNPN
GYSLSVKDWEDGRGYHVKHYRIKPLDNGGYYIATNQTFPSLQALVMAYS
KNALGLCHILSRPCPKPQPQMWDLGPELRDKYEIPRSEIQLLRKLGRGNF
GEVFYGKWRNSIDVAVKTLREGTMSTAAFLQEAAIMKKFRHNRLVALYA
VCSQEEPIYIVQEYMSKGSLLDFLREGDGRYLHFEDLIYIATQVASGMEYLE
SKQLIHRDLAARNVLIGENNVAKICDFGLARVIADDEYCPKQGSRFPVKW
TAPEAIYGKFSIKSDVWSYGILLMELFTYGGVPYPGHSREVIENIERGFRM
PKPTNHYPDPDNIYQLLLQCWDVPEKRPTFEFLNHYFESFSVTSEVPYRE
VQD



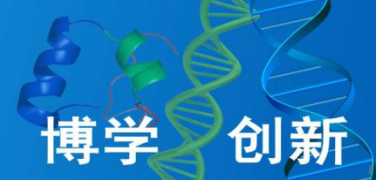
□ Full entry format

- Complete information for a particular entry
- Very annotation rich
- Many cross-links on WWW
- Varies between databases
- Most commonly used: GenBank & Swiss-Prot

□ Example: P00519

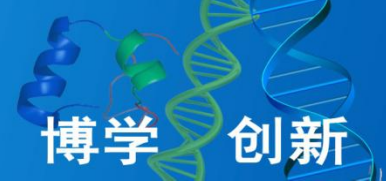


Lecture 3.2 Introduction to NCBI's query strategy



OUTLINES

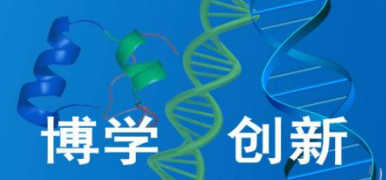
1. Database Query Tools
2. Query in NCBI



1. Database Query Tools

1.1 SRS (Sequence Retrieval System)

- EMBL, LION Bioscience
- Open query system, more than 300 databases in SRS servers.



- LION Bioscience Website

<http://www.lionbio.co.uk/publicsrs.html>

- <http://srs6.ebi.ac.uk/srs6/>
- <http://iron.hgmp.mrc.ac.uk/srs6/>
- <http://www.sanger.ac.uk/srs6/>
- <http://www.infobiogen.fr/srs6/>
- <http://www.cmbi.kun.nl/srs6>
- <http://srs.wehi.edu.au/srs6/>
- <http://genius.embnet.dkfz-heidelberg.de/menu/srs/>
- <http://www.cbr.nrc.ca/srs6/>
- <http://srs.pku.edu.cn/srs6bin/cgi-bin/wgetz>



SRS - 夏新笔记本电脑

文件(F) 编辑(E) 查看(V) 收藏(A) 工具(T) 帮助(H)

后退 搜索 收藏夹 媒体

地址(D) http://srs.ebi.ac.uk/srsbin/cgi-bin/wgetz?page+srsq2+-noSession 转到 链接

EMBL-EBI
European Bioinformatics Institute

Quick Search Library Page Query Form Tools Results Projects Views Databanks **HELP**

SRS **LION**

Start a Permanent Project

Tips

- ★ Want to know more about using SRS?
- go to the [Help Center](#) where you'll find all the searchable online help you need.
- ★ Where is the old library page?
- Click on the 'Library Page' tab on the menu bar.
- ★ Linking to SRS?
- Please read this [document](#) for important information regarding linking to our SRS server.
- ★ [Public SRS servers worldwide](#)

Quick Text Search [Search Tips](#)

Get **Nucleotide Sequences** matching :

Searches Databanks: EMBL Nucleotides

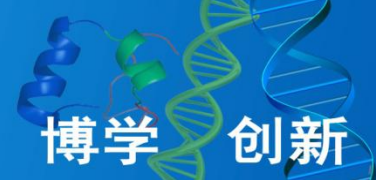
News and Announcements [Search Tips](#)

Important notes to all users.

- 12.04.05 - UniProt databases: EC numbers are now indexed and available from both the description as well as ECNumber indices.
- 01.04.05 - Please note that Description fields in the UniProt databases are now indexed as single words.
- 11.03.05 - EMBL release 82 is now on-line.
- 09.03.05 - ADVANCED NOTICE: As of EMBL release 82 Secondary Accession number ranges will be introduced. See: <http://www.ebi.ac.uk/embl/Documentation> for further details.
- 03.03.05 - ADVANCED NOTICE: As of the next release of EMBL (82) ...

LION
SRS is a product of Lion Bioscience AG

Internet 20:55



Entry Page - 夏新笔记本电脑

文件(F) 编辑(E) 查看(V) 收藏(A) 工具(T) 帮助(H)

后退 搜索 收藏夹 媒体

地址 http://srs.ebi.ac.uk/srsbin/cgi-bin/wgetz?id+29RVi1PuDt+e+[UNIPROT:'AATM_RABIT']+-qnum+1+-enum+1 转到 链接

EMBL-EBI

European Bioinformatics Institute

Quick Search Library Page Query Form Tools Results Projects Views Databanks HELP

[Text Entry](#) | [SwissEntry](#) | [NiceProt](#) | [iProClass](#) | [UniProtXML](#)

Reset Entry 1 of 2 from Query 1 Next Entry

[General](#) [Description](#) [References](#) [Comments](#) [Links](#) [Keywords](#) [Features](#) [Sequence](#)

Entry Information

Entry from: [UniProt](#)

Entry Options

Launch analysis tool:
BlastP Launch

Link to related information: Link

Save entry: Save

View: Printer Friendly

General information

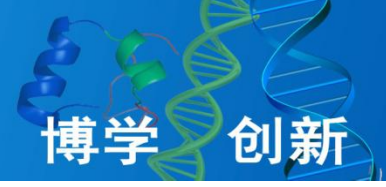
Entry name	AATM_RABIT
Accession number	P12345
Created	Rel. 12, 1-OCT-1989
Sequence update	Rel. 12, 1-OCT-1989
Annotation update	Rel. 44, 5-JUL-2004

Description and origin of the Protein

Description	Aspartate aminotransferase, mitochondrial (EC 2.6.1.1) (Transaminase A) (Glutamate oxaloacetate transaminase)
Gene name(s)	GOT2
Organism source	Oryctolagus cuniculus (Rabbit).
Taxonomy	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Leporidae; Oryctolagus.
NCBI TaxID	9986

References

[1] Kuramitsu,S., Inoue,K., Kondo,K., Aki,K., Kagamiyama,H.,
Aspartate aminotransferase isozymes from rabbit liver. Purification and properties.
(1985) *J. Biochem.* **97**:1337-1345



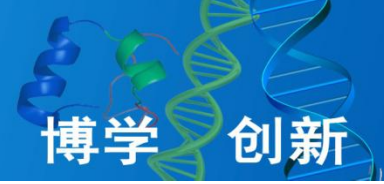
1.2 Query in NCBI

<http://www.ncbi.nlm.nih.gov/Entrez>



□ Overview:

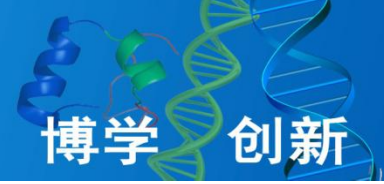
- Database structure
- Basic Searching
- Using related links/neighbors
- Details, Limits, Preview
- Entrez field qualifiers
- Advanced Boolean queries
- **Practical examples and hands-on work!**



NCBI databases

NCBI integrated databases include:

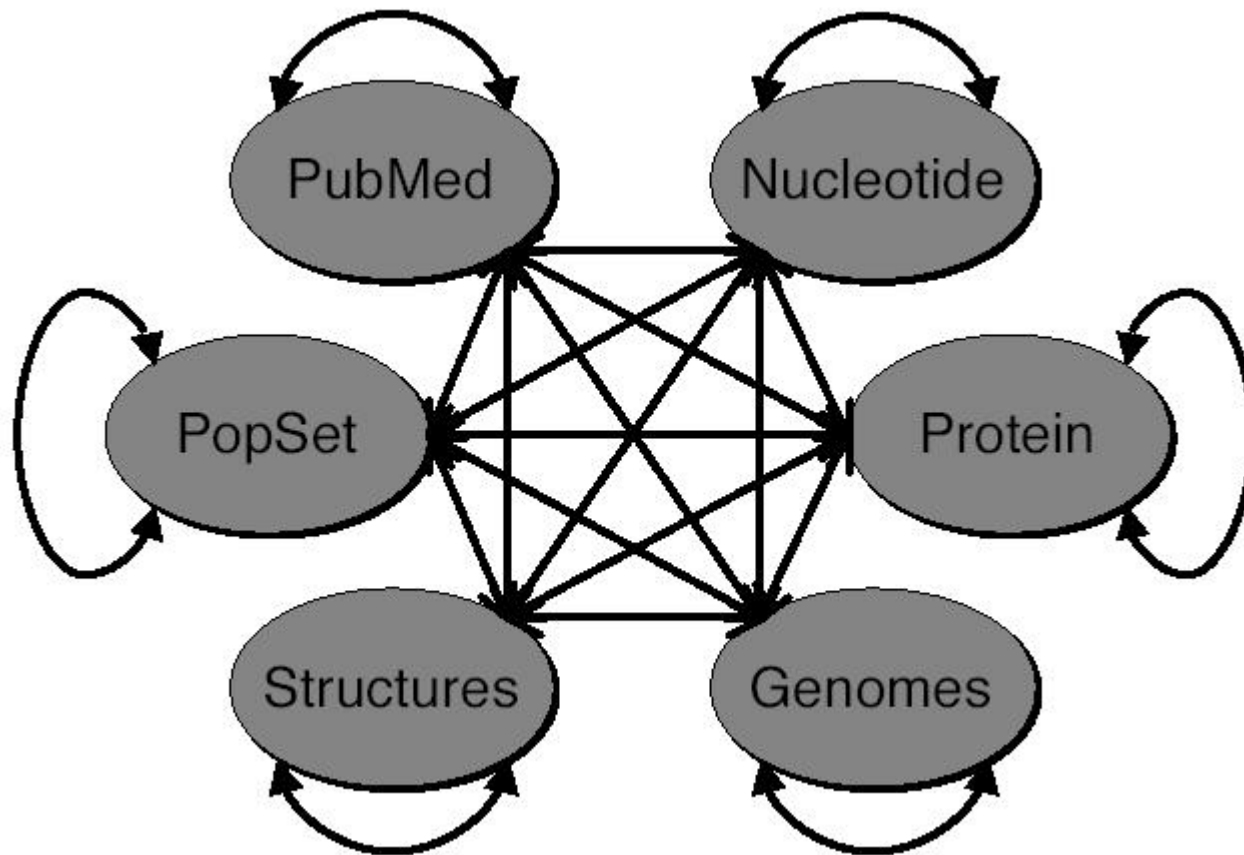
- ❑ PubMed: The biomedical literature (PubMed)
- ❑ Nucleotide sequence database (Genbank)
- ❑ Protein sequence database
- ❑ Structure: three-dimensional macromolecular structures
- ❑ Genome: complete genome assemblies
- ❑ PopSet: population study data sets
- ❑ OMIM: Online Mendelian Inheritance in Man

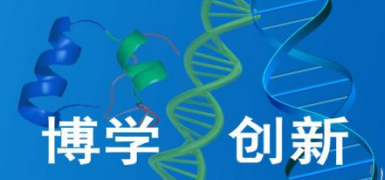



NCBI databases

- ❑ Taxonomy: organisms in GenBank
- ❑ Books: online books
- ❑ ProbeSet: gene expression and microarray datasets
- ❑ 3D Domains: domains from Entrez Structure
- ❑ UniSTS: markers and mapping data
- ❑ SNP: single nucleotide polymorphisms
- ❑ CDD: conserved domains

Entrez database connection






地址 (U)  <http://www.ncbi.nlm.nih.gov/entrez/query.fcgi>

All Databases	PubMed	Nucleotide	Protein	Genome	Structure
Search	PubMed	for <input type="text"/> <input type="button" value="Go"/> <input type="button" value="Clear"/>			
	PubMed	<input type="button" value="Preview/Index"/> <input type="button" value="History"/> <input type="button" value="Clipboard"/> <input type="button" value="Details"/>			
	Protein	get started, enter one or more search terms.			
	Nucleotide	Search terms may be topics , authors or journals .			
	CoreNucleotide	Enhancement for Preferred Method of			
	GSS				
	EST				
	Structure				
	Genome				
	Books				
	CancerChromosome				
	Conserved Domains				
	3D Domains				
	Gene				
	Genome Project				
	GENSAT				
	GEO Profiles				
	GEO DataSets				
	HomoloGene				
	Journals				
	MeSH Da				
	Single Ci				
	MeSH				
	NCBI Web Site				
	NLM Catalog				
	OMIA				
	OMIM				
	PMC				
	PopSet				
	Probe				
	PubChem BioAssay				
	PubChem Compound				
	PubChem Substance				
	NLM Gateway				

Choose a database



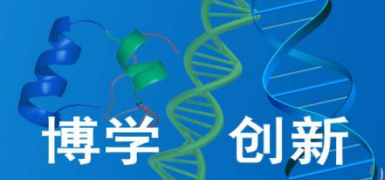
Need to find a specific citation or journal issue?


Click [Single Citation Matcher](#) on the PubMed sidebar.


Enter what you know and click Go.

Journal and author boxes include an autocomplete feature that suggests titles and as you type. Read the [PubMed Help](#) for additional information.

is a service of the [National Library of Medicine](#) that includes over 16
 citations from MEDLINE and other life science journals for biomedical
 back to the 1950s. PubMed includes links to full text articles and other
 sources.







A service of the National Library of Medicine
and the National Institutes of Health

All Databases PubMed Nucleotide Protein Genome Structure

Search Protein for camp activated protein kinase

Limits Preview/Index History Clipboard Details

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- To get started, enter one or more search terms
- Search terms may be [topics](#), [authors](#) or [pubmed IDs](#)

Enter Query Term

Clinical Advisory: [NCI Issues Clinical Announcement for Preferred Method of Treatment for Advanced Ovarian Cancer](#)

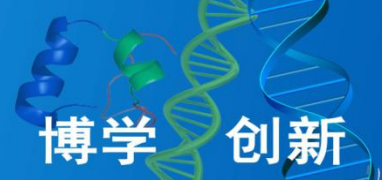


Need to find a specific citation or journal issue?

- (1) Click [Single Citation Matcher](#) on the PubMed sidebar.
- (2) Enter what you know and click Go.

The journal and author boxes include an autocomplete feature that suggests titles and names as you type. Read the [PubMed Help](#) for additional information.

PubMed is a service of the [National Library of Medicine](#) that includes over 16 million citations from MEDLINE and other life science journals for biomedical articles back to the 1950s. PubMed includes links to full text articles and other related resources.



地址: <http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?CMD=search&DB=protein>

NCBI

Entrez Protein

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All Databases PubMed Nucleotide Protein Genome Structure PMC Taxonomy

Search Protein for camp activated protein kinase Go Clear Save Search

Limits Preview Clipboard Details

Display Summary Show 20 Sort by Relevance Send to

All: 514 bacteria: 1 RefSeq: 421

Items 1 - 20 of 514

1: [NP_059043](#) Reports BLink, Conserved Domains
protein kinase, mitogen activated 3 (extracellular-signal-regulated kinase 1, ERK1) [Rattus norvegicus]
gi|8393331|ref|NP_059043.1|[8393331]

2: [NP_619726](#) Reports BLink, Conserved Domains
peroxisome proliferative activated receptor gamma isoform 1 [Homo sapiens]
gi|20336235|ref|NP_619726.1|[20336235]

3: [NP_619725](#) Reports BLink, Conserved Domains
peroxisome proliferative activated receptor gamma isoform 1 [Homo sapiens]
gi|20336233|ref|NP_619725.1|[20336233]

4: [NP_005028](#) Reports BLink, Conserved Domains
peroxisome proliferative activated receptor gamma isoform 1 [Homo sapiens]
gi|20336231|ref|NP_005028.3|[20336231]

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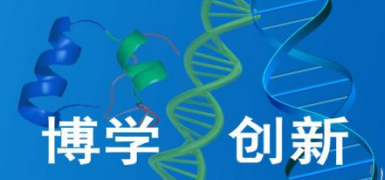
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Reference sequence project

Search for Genes

Clusters of



地址 <http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?CMD=Details&DB=protein>

All Databases PubMed Nucleotide Protein Structure

Search Protein for camp activated protein kinase [Save Search](#)

[Limits](#) [Preview/Index](#) [History](#) [Clipboard](#) **Detail**

Query Translation:

```
camp[All Fields] AND activated[All Fields] AND protein  
kinase[All Fields]
```

Result:

[514](#)

Database:

Protein

User query:

camp activated protein kinase

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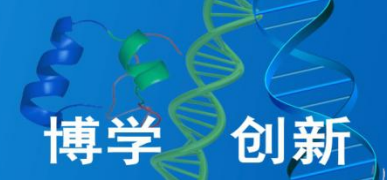
BLAST

Reference sequence
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Search for Genes

Detail Screen

Exact Query Syntax



地址 (D) <http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?CMD=search&DB=protein>

Entries added to Clipboard,
Text,File,Printer

Search Protein for camp activated protein kinase

Limits Preview/Index History Clipboard Data

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OMIA Links

OMIM Links

BioAssay Links

PubChem Compound Links

PubChem Substance Links

PMC Links

PopSet Links

PubMed Links

SNP Links

Gene Genotype Links

Number of Entries/Page

Format, Neighbors & Links

BLink, Conser

BLink, Conser

BLink, Conser

BLink, Conser

BLink, Conser

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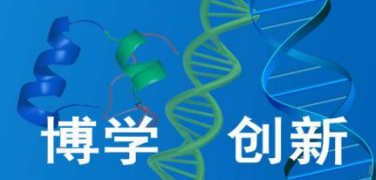
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project

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Clusters of
orthologous groups

Protein reviews on the
web

Search for full length



地址 (Q) <http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?CMD=search&DB=protein>

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Search Protein for "camp activated protein kinase" Go Clear Save Search

LimitsPreview/IndexHistoryClipboardDetails

Quoted phrase not found. See Details. !!!

Display Summary Show 20 Sort by Relevance Send to

All: 514 bacteria: 1 RefSeq: 421

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☐ 1: [NP_059043](#) Reports BLink, Conserved I

protein kinase, mitogen activated 3 (extracellular-signal-regulated kinase 1, ERK 1) [Rattus norvegicus]

gi|8393331|ref|NP_059043.1|[8393331]

☐ 2: [NP_619726](#) Reports BLink, Conserved I

peroxisome proliferative activated receptor gamma isoform 1 [Homo sapiens]

gi|20336235|ref|NP_619726.1|[20336235]

☐ 3: [NP_619725](#) Reports BLink, Conserved I

peroxisome proliferative activated receptor gamma isoform 1 [Homo sapiens]

gi|20336233|ref|NP_619725.1|[20336233]

☐ 4: [NP_005028](#) Reports BLink, Conserved I

peroxisome proliferative activated receptor gamma isoform 1 [Homo sapiens]

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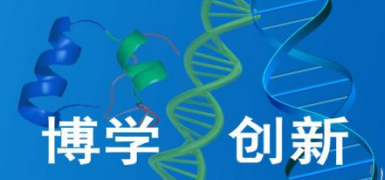
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Clusters of



地址 (D) <http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?CMD=Details&DB=protein>

All DatabasesPubMedNucleotideProteinGenomeStructure

Search Protein for "camp activated protein kinase" [Save Search](#)

LimitsPreview/IndexHistoryClipboardDetails

Query Translation:
camp[All Fields] AND activated[All Fields] AND protein kinase[All Fields]

Result:
[514](#)

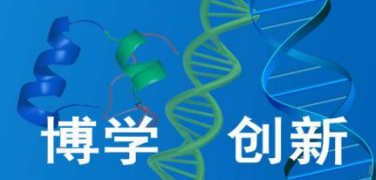
Quoted phrase not found:
"camp activated protein kinase"

Database:
Protein



User query:
"camp activated protein kinase"

Same as without quotes

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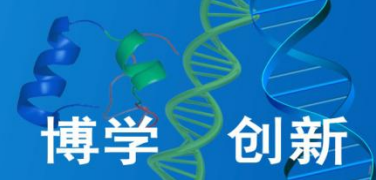
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Query Translation:



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Search PubMed for "camp dependent protein kinase" Go Clear Save Search

LimitsPreview/IndexHistoryClipboardDetails

Query Translation:
"camp dependent protein kinase"[All Fields]

Search

Result !!!

6303

Database:
PubMed

User query:
"camp dependent protein kinase"

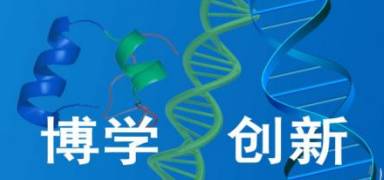
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All DatabasesPubMedNucleotideProteinGenomeStructureOMIMPMC

SearchPubMedfor immunoglob*PreviewGoClear

LimitsPreview/IndexHistoryClipboardDetails

- Enter terms and click Preview
- To combine searches use # b
- Click on query # to add to strategy

Search

Most Recent Queries

#2 Search immunoglob*

#1 Search "camp dependent protein kinase"

Add Term(s) to Query or View Index:

- Enter a term in the text box, use the pull-down menu to specify a search field.
- Click Preview to add terms to the query box and see the number of search results, or click Index to vie

All FieldsPreviewIndex

Click AND OR NOT to add a term to the query box.

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NCBI Toolbar

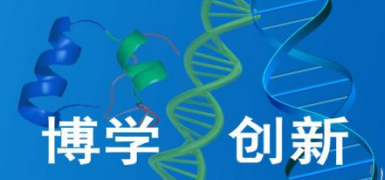
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Preview/Index Screen



Search PubMed for immunoglob* Preview Go Clear

Limits **Preview/Index** History Clipboard Details

- Enter terms and click Preview to see only the number of search results.
- To combine searches use # before search number, e.g., (#2 OR #3) AND asthma.
- Click on query # to add to strategy

Search Most Recent Queries

[#3](#) Search immunoglob* AND kinase
[#2](#) Search immunoglob*
[#1](#) Search "camp dependent protein kinase"

Add Term(s) to Query or View Index:

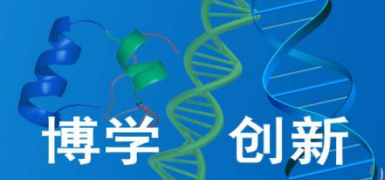
- Enter a term in the text box; use the pull-down menu to specify a search field.
- Click Preview to add terms to the query.
- Multiple terms selected from Index will

Index for "kinase"

All Fields kinase Preview Index

Click AND OR NOT to add terms selected from Index to the query box.

kinase(230354)	Up
kinase/5 fluoro 2 deoxyuridine(1)	
kinase/5' fluoro 2' deoxyuridine(1)	
kinase/90 kda(2)	
kinase/90 kilodalton(1)	
kinase/acetyltransferase(1)	
kinase/act(1)	
kinase/actin/hsc 70(1)	
kinase/actin/hsp(1)	
kinase/actin/hsp70(2)	Down



Search PubMed for immunoglob* NOT kinase Preview Go Clear

Limits Preview/Index History Clipboard

- Enter terms and click Preview to see only
- To combine searches use # before search number, e.g., (#2 OR #3) AND asthma.
- Click on query # to add to strategy

Boolean term inserted

Search Most Recent Queries

- #3 Search immunoglob* AND kinase
- #2 Search immunoglob*
- #1 Search "camp dependent protein kinase"

Add Term(s) to Query or View Index:

- Enter a term in the text box; use the pull-down menu to specify a search field.
- Click Preview to add terms to the query box and see the number of search results, or
- Multiple terms selected from Index will be ORed; click AND to add to search.

All Fields Preview Index

Click AND OR NOT to add terms selected from Index to the query box.

kinase(230354) Up

kinase/5 fluoro 2 deoxyuridine(1)

kinase/5' fluoro 2' deoxyuridine(1)

kinase/90 kda(2)

kinase/90 kilodalton(1)

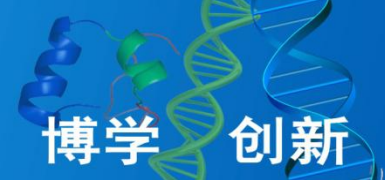
kinase/acetyltransferase(1)

kinase/act(1)

kinase/actin/hsc 70(1)

kinase/actin/hsp(1)

kinase/actin/hsp70(2) Down



The screenshot shows the NCBI PubMed homepage. The search bar contains the text 'immunoglob* NOT kinase'. Below the search bar, the 'Limits' tab is selected. A light blue box labeled 'Limits Screen' points to the search criteria. Another light blue box labeled 'Database dependent field' points to the 'All Fields' dropdown menu. The 'Limited to:' section includes various filters such as 'Publication Types', 'Ages', 'Entrez Date', 'Publication Date', 'Languages', 'Humans or Animals', 'Subsets', and 'Gender'. The 'only items with abstracts' checkbox is also visible.

NCBI

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All Databases PubMed Nucleotide Protein Genome Structure OMIM PMC Journals

Search PubMed for immunoglob* NOT kinase Go Clear

Limits Preview/Index History Clipboard Details

- specify a field.
- If search field T must be in upper case.
- In search field operator square brackets, e.g., rubella [ti].
- Search limits in 操作人员, 接线员 publisher supplied citations

Limited to:

All Fields

only items with abstracts

Publication Types

Languages

Subsets

Ages

Humans or Animals

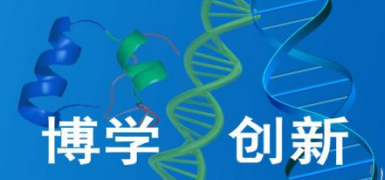
Gender

Entrez Date

Publication Date From To

Use the format YYYY/MM/DD; month and day are optional.

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地址 (U) <http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?CMD=Details&DB=pubmed>

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Search PubMed for taylor [Save Search](#)

☒ Limits Preview/Index History Clipboard **Details**

Field: **Author**

Query Translation:

taylor[Author]

Result:

[46911](#)

Database:

PubMed

User query:

taylor

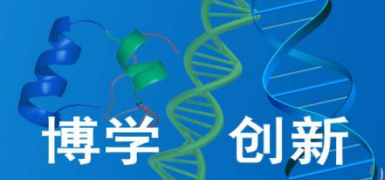
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Field Qualifier



NCBI

PubMed www.pubmed.gov

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Search PubMed for **taylor[Author] AND "protein kinase"** [Save Search](#)

Limits Preview/Index History Clipboard

Display Summary Show 20

All: 467 Review: 47

Items 1 - 20 of 467

1: [Day JR, Taylor KM, Lidington EA, Mason JC, Haskard DO, Randi AM, Landis RC.](#)
 Aprotinin inhibits proinflammatory activation of endothelial cells by thrombin through the protease-acti
 J Thorac Cardiovasc Surg. 2006 Jan;131(1):21-7.
 PMID: 16399290 [PubMed - in process]

2: [Taylor AJ, Ye JM, Schmitz-Peiffer C.](#)
 Inhibition of glycogen synthesis by increased lipid availability is associated with subcellular redistributi
 J Endocrinol. 2006 Jan;188(1):11-23.
 PMID: 16394171 [PubMed - in process]

3: [Vigil D, Lin JH, Sottriffer CA, Pennypacker JK, McCammon JA, Taylor SS.](#)
 A simple electrostatic switch important in the activation of type I protein kinase A by cyclic AMP.
 Protein Sci. 2006 Jan;15(1):113-21. Epub 2005 Dec 1.
 PMID: 16322564 [PubMed - in process]

4: [Taylor EB, Ellingson WJ, Lamb JD, Chesser DG, Compton CL, Winder WW.](#)
 Evidence Against Regulation of AMP-Activated Protein Kinase and LKB1-STRAD-MO25 Activity
 Am J Physiol Endocrinol Metab. 2005 Nov 8; [Epub ahead of print]
 PMID: 16278246 [PubMed - as supplied by publisher]

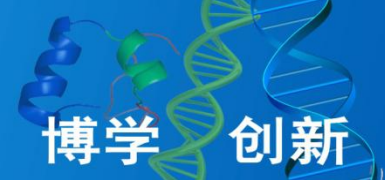
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“Advanced Boolean Query”



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History Screen

All Databases PubMed Nucleotide Structure OMIM

Search PubMed for taylor[Author] AND "protein kinase" Preview Go Clear

Limits Preview/Index **History** Clipboard Details

- Search History will be lost after eight hours of inactivity.
- To combine searches use # before search number, e.g., #2 AND #6.
- Search numbers may not be continuous; all searches are represented.
- Click on query # to add to strategy

Search	Most Recent Queries
#7	Search taylor[Author] AND "protein kinase"
#6	Search taylor[Author] AND "protein kinase" Field: Author
#5	Search taylor Field: Author
#4	Search tarlor Field: Author
#3	Search immunoglob * AND kinase
#2	Search immunoglob *
#1	Search "camp dependent protein kinase"

Clear History

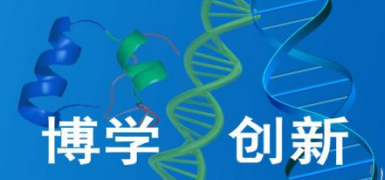
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Search PubMed for #7 AND #1 [Preview](#) [Go](#) [Clear](#)

[Limits](#) [Preview/Index](#) **[History](#)** [Clipboard](#) [Details](#)

- Search History will be lost after eight hours of inactivity.
- To combine searches use # before search number, e.g., #2 AND #6.
- Search numbers may not be continuous; all searches are represented.
- Click on query # to add to strategy

Search

Most Recent Queries

[#7](#) Search **taylor[Author] AND "protein kinase"**

[#6](#) Search **taylor[Author] AND "protein kinase" Field: Author**

[#5](#) Search **taylor** Field: **Author**

[#4](#) Search **tarlor** Field: **Author**

[#3](#) Search **immunoglob* AND kinase**

[#2](#) Search **immunoglob***

[#1](#) Search **"camp dependent protein kinase"**

[Clear History](#)

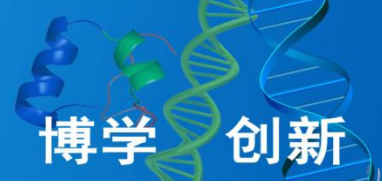
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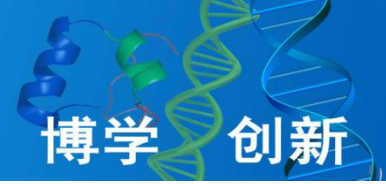
Basic Searching (Summary)

- ❑ Choose database
- ❑ Enter query term
- ❑ Hit “Go” button
- ❑ Examples:
 - camp activated protein kinase
 - “camp activated protein kinase”
 - “camp activated” “protein kinase”
 - “camp dependent protein kinase”
 - “immunoglob*”



Details, Limits, Preview ...

- ❑ Details: review Entrez's interpretation of the query term
- ❑ Limits: ways to restrict a search to a defined subset of the database.
- ❑ Indexes: alphabetical lists of terms from searchable database fields
- ❑ History: review, recall and combine past searches
- ❑ Clipboard: temporarily store results



Field Qualifiers

❑ Format: **term [qualifier]**

❑ Where:

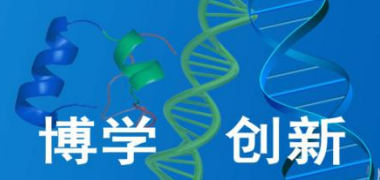
-- **Term** is the string to search for

-- **Qualifier** specify the field to search in

-- DB specific

-- one of WORD, TITL, MESH, MAJR, AUTH, JOUR, ECNO, GENE, DATE, PDAT, MDAT, PAGE, VOL, KYWD, ORGN, ACCN, PROT, SLEN, SQID, SUBS, PROP, FKEY, and PTYP

❑ Example: **Bourne PE [AUTH]**



ACCN accession number

FKEY feature key

MAJR MeSH major topic

PROP property

TITL title word

WORD text word

ECNO EC number

KYWD keyword

ORGN organism

PROT protein name

SUBS substance

DATE publication year

JOUR journal name

PAGE first page

PTYP publication type

SLEN sequence length

AUTH author name

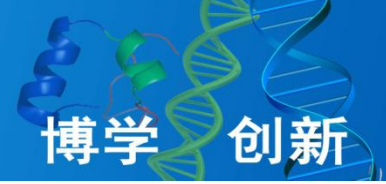
GENE gene name

MDAT modification date

PDAT publication/creation date

SQID sequence id

VOL volume



Advanced Boolean Queries

- Format:

 - **term [field] operator term [field] ...**

- Where **Operator** is any of :

 - **AND** (intersection)

 - **OR** (union)

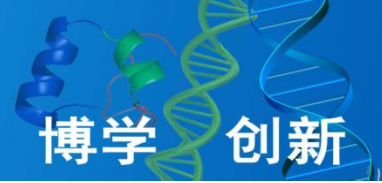
 - **BUTNOT** (difference)

- Example:

 - Taylor P [AUTH] AND “protein kinase”**

- Ranges:

 - **term1:term2 [field]**



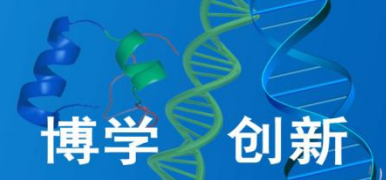
Using “Related” Links (aka “Neighbors”)

- ❑ PubMed Neighbors
 - Text & MeSH
- ❑ Protein and Nucleotide Neighbors
 - BLAST
- ❑ Macromolecular Structure Neighbors
 - VAST
- ❑ Non similarity based: references within
 - text/annotation
- ❑ VERY helpful in identifying related records



Exercises

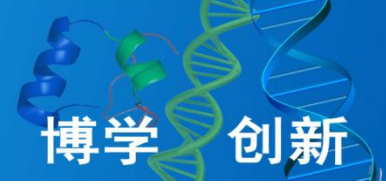
1. Work through the NCBI query tutorial at
-- <http://www.ncbi.nlm.nih.gov/Database/tut1.html>
2. Find references about shingles and facial paralysis.
-- Display the records in the format that shows the abstract and the MeSH headings. How does PubMed map the term, shingles?



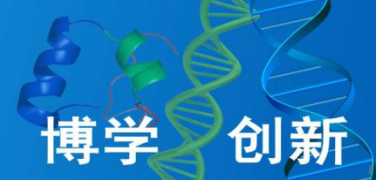
3. Find murine nucleic acid sequences involved in apoptosis and cancer.

-- Display all of the retrieved records on one Web page.

4. Find protein sequences related to AIDS with publications in Nature. Save this search strategy so the search can be run again at a later date.



5. Find all mouse or human nucleotide sequences added to Entrez in 1997.
 - How many structures correspond to your result set?



Thanks for your attentions !