

# Fig2-02 FLJ Human cDNA Database

**FLJ Human cDNA Database**

FLJ Human cDNA Database Ver1.0 : [Data set](#)

If you have questions or requests, please send the mail to [T\\_Isogai \(flj-cdna@nifty.com\)](mailto:T_Isogai@flj-cdna@nifty.com) Update 09/30/06

[Sequence ID search](#)  
You can search human cDNAs by sequence ID as a query.

[cDNA search by keyword](#)  
You can search human cDNAs by keywords as a query.

[cDNA cluster search by keyword](#)  
You can search human cDNA clusters mapped on human genome by keywords as a query.

[BLAST search](#)  
You can search human cDNAs using DNA sequence as a query by blastn.

[cDNA cluster search by genome locus](#)  
You can search cDNA clusters mapped on human genome by genome locus.

**Sequence ID search**

You can search human cDNAs in our DB by sequence ID as a query described below.

- DDBJ / EMBL / GenBank accession No. (ex. AK126746)
- FLJ ID (ex. FLJ44792)
- Sequence ID (ex. C-BRACE3039378; BRACE3039378 ...)

ID:

**cDNA search by keyword**

You can search human cDNAs by keywords\* as a query.

Search formula

Multiple words are combined with "AND" automatically.  
A character "\*" matches any symbols. For example, "abc\*" searches for words starting with "abc".  
Keyword search regards any sequence of true word characters (letters, digits, and underscores) as a word.  
Any word that is too short is ignored. The minimum length of words that are found by keyword searches is four characters.

Keyword:

**cDNA cluster search by keyword**

You can search human cDNA clusters mapped on human genome by keywords\* as a query.

Search formula

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Keyword:

**BLAST search**

You can search human cDNAs using DNA sequence as a query by blastn.

Sequence:

**cDNA cluster search by genome locus**

You can search cDNA clusters mapped on human genome by human genome locus.

Chromosome No:

Locus of chromosome:  (bp) (←→)  (bp)

\* Keyword search

All words described below in our DB will be effectively hit for keyword searches.

- cDNA definitions
- Blastp results against Swiss-Prot and RefSeq (1-E10 & up to three)
- Plan results (1-E10)
- PROSITE results
- GO (Gene Ontology) terms picked up from Blastp and Plan results

**FLJ Human cDNA Database - Netscape**

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**BLAST search**  
You can search human cDNAs using DNA sequence as a query by blastn.

Sequence:

**cDNA cluster search by genome locus**  
You can search cDNA clusters mapped on human genome by human genome locus.

Chromosome No:    
Locus of chromosome:   (-----)

\* Keyword search  
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- Pfam results (<1-E10)
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## \* Search formula

- Any word that is too short is ignored. The minimum length of words that are found by keyword searches is four characters.
- Multiple words are combined with "AND" automatically.
- A character "\*" matches any symbols. For example, "abc\*" searches for words starting with "abc".
- Keyword search regards any sequence of true word characters (letters, digits, and underscores) as a word.

# Fig2-02 FLJ Human cDNA Database

The image displays the FLJ Human cDNA Database website. On the left is a navigation menu with links for [Sequence ID search](#), [cDNA search by keyword](#), [cDNA cluster search by keyword](#), [BLAST search](#), and [cDNA cluster search by genome locus](#). A red dashed box highlights the **BLAST search** section in the menu. On the right is a screenshot of the BLAST search interface in a Netscape browser window. The interface includes a 'Keyword:' input field with 'Submit' and 'Reset' buttons. Below this is a blue header for 'BLAST search' with the text 'You can search human cDNAs using DNA sequence as a query by blastn.' A large text area labeled 'Sequence:' is provided for input, with 'Submit' and 'Reset' buttons below it. Further down is a blue header for 'cDNA cluster search by genome locus' with the text 'You can search cDNA clusters mapped on human genome by human genome locus.' This section includes a 'Chromosome No:' dropdown menu (set to 'chr1') and a 'Locus of chromosome:' input field (set to '1 [bp] <-----> 100000 [bp]'), with 'Submit' and 'Reset' buttons. At the bottom, a section titled '\* Keyword search' explains that all words in the database will be effectively hit by keyword searches, listing the types of results returned: cDNA definitions, blastp results against Swiss-Prot and RefSeq (<1-E10 & up to three), Pfam results (<1-E10), PROSITE results, and GO (Gene Ontology) terms picked up from blastp and Pfam results.