

Software & DBs

— Genome Informatics
— Molecular Function
— Biological Systems

GeneDecoder

Gene finding system for eukaryotes, based on HMMs

URL

<http://www.genecodecoder.org/>



ALN

Pairwise alignment of biological sequences supporting spliced alignment procedures

URL

http://www.genome.ist.i.kyoto-u.ac.jp/~aln_user/ALN/top.html



INOH

An event-oriented pathway database

URL

<http://www.inoh.org/>



ASTRA

The database of elementary patterns of alternative splicing and transcription

URL

<http://alterna.cbrc.jp/>



WoLF PSORT

Protein sub-cellular localization system

URL

<http://wolffpsort.seq.cbrc.jp/>



CellMontage

System for searching gene expression databases for similar cells or tissues to the query gene expression profile

URL

<http://cellmontage.cbrc.jp/>



SEVENS

Comprehensive collection of 7TM receptors in eukaryote genomes

URL

<http://sevens.cbrc.jp/>



TMBETA-NET

Prediction of membrane spanning β -strand segments in outer membrane proteins

URL

<http://psfs.cbrc.jp/tmbeta-net/>



FORTE

A profile-profile comparison tool for protein fold recognition

URL

<http://www.cbrc.jp/forte/>

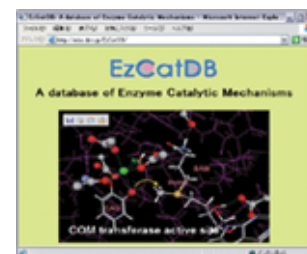


EzCatDB

A database of enzyme catalytic mechanisms

URL

<http://mbs.cbrc.jp/EzCatDB/>



CoCoozo

Parallel high-speed engine for peptide MS/MS ions search

URL

<http://www.cbrc.jp/cocoozo/>



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