

(tertiary structure)

CASP

1.

(primary structure; DNA
)

(tertiary structure,
가

(three-dimensional)
(protein tertiary structure prediction)

[1,2,3,4].

(homology)
(high sequence

similarity)

(low sequence similarity)

(similarity)

가

(1) (Comparative Modeling)

(2) (Fold Recognition)

(3) (New Fold)

가

Comparative

Modeling (Homology Modeling)

가

Fold Recognition

. Fold Recognition

가

New

Fold (ab initio de novo ,

New Fold 가)

Comparative Modeling Fold Recognition

(, PDB, SCOP, CATH) ,

(homology)가 Comparative Modeling, Fold Recognition, Comparative Modeling, (homology)가 New Fold, (potential energy function), (global optimization), Modeling, (grand challenge), Fold Recognition, Comparative New Fold, New Fold, 가

2. (CASP)

Post-Genome 가 (CASP, <http://PredictionCenter.llnl.gov/>)가 1994 [2,3,4]. CASP community wide experiment on the Critical Assessment of techniques for protein Structure Prediction 6 CASP6 2004 가 2004 12 가 (Gaeta) CASP (Lawrence Livermore National Laboratory) 가

CASP6 가 266 가 2004 가(evaluation)

(Comparative Modeling, Fold Recognition, New Fold) 가 가 가

CASP6 가 가 Fold Recognition
 - Fold Recognition/Homology(FR/H) Fold
 Recognition/Analogy(FR/A) - FR/H
 (homology)
 , FR/A 가 FR/H
 . CASP6
 Comparative Modeling(CM)
 FR/H
 FR/A New Fold(NF)

CASP6 CM Ginalski (University of Texas, Dallas)
 1 , Venclovas (Institute of Biotechnology, Lithuania),
 Kolinski Bujnicki (Warsaw University), Skolnick (State University
 of New York, Buffalo) 2, 3, 4 . FR/H
 Ginalski GeneSilico (International Institute of Molecular and
 Cell Biology, Poland) 1, 2 . FR/A Baker
 (University of Washington, Seattle) Karplus (University of
 California, Santa Cruz) 1, 2 .

CASP6 NF 181 가 . 1 7
 (University of Washington, Seattle) Baker ,
 2 (Warsaw University)
 (International Institute of Molecular and Cell Biology)
 Kolinski Bujnicki . 3, 4, 5 Ginalski
 (University of Texas, Dallas), Karplus (University of California,
 Santa Cruz), Skolnick (State University of New York, Buffalo)
 . 6 (University
 College London) Jones .

[]

[1] M. J. E. Sternberg (editor), Protein Structure Prediction: A Practical Approach (1997) Oxford University Press.

[2] Proteins: Structure, Function, and Bioinformatics (1999) Volume 37, Number S3, p.1 – p.237.

[3] Proteins: Structure, Function, and Bioinformatics (2001) Volume 45, Number S5, p.1 – p.199.

[4] Proteins: Structure, Function, and Bioinformatics (2003) Volume 53, Number S6, p.333 – p.595.