

Žebříček CASP - [http://predictioncenter.org/casp10/groups\\_analysis.cgi?type=server&tbm=on&tbm\\_hard=on&tbfm=on&fm=on&submit=Filter](http://predictioncenter.org/casp10/groups_analysis.cgi?type=server&tbm=on&tbm_hard=on&tbfm=on&fm=on&submit=Filter)

Obecně o predikování struktury - <http://www.biostat.wisc.edu/bmi776/lectures/protein-structure.pdf>

Podobnostní modelování - <http://www.biochem.vt.edu/modeling/homology.html>

Navlékání - <http://www.biostat.wisc.edu/bmi776/lectures/threading.pdf>

Články o objevech Foldit - <http://www.nature.com/nsmb/journal/v18/n10/full/nsmb.2119.html>

[http://www.huffingtonpost.com/2011/09/19/aids-protein-decoded-gamers\\_n\\_970113.html](http://www.huffingtonpost.com/2011/09/19/aids-protein-decoded-gamers_n_970113.html)

<http://www.nature.com/nbt/journal/v30/n2/full/nbt.2109.html>

<http://www.scientificamerican.com/article.cfm?id=victory-for-crowdsourced-biomolecule2&page=2>

Doplňující informace k modelování - <http://biochem118.stanford.edu/Projects/2002/zhang.pdf>

RMSD - [http://en.wikipedia.org/wiki/Root-mean-square\\_deviation\\_\(bioinformatics\)](http://en.wikipedia.org/wiki/Root-mean-square_deviation_(bioinformatics))

Informace o predikovaném proteinu - <http://www.ebi.ac.uk/gxa/gene/FBgn0031459>

<http://www.yeastrc.org/pdr/viewProtein.do?id=661718>