
DYNAMEOMICS.ORG ACCESS REQUEST

TO: Dr. Valerie Daggett

FROM:

Department of Bioengineering
University of Washington

DATE:
6/8/2010

FAX NUMBER:
(206) 685-3300

TOTAL NO. OF PAGES INCLUDING COVER:
3

INSTRUCTIONS:

Please complete and sign the following form and return both pages via fax to using this cover sheet or mail to:

Dr. Valerie Daggett
Department of Bioengineering
University of Washington
Foegen Building N310B, Box 355061
1705 NE Pacific Street
Seattle, WA 98195-5061

Dynameomics.org Database Access Agreement

Research Topic:	
User Name:	
Job Title:	
Name of PI:	
Department:	
Institution:	
Address:	
Phone Number:	
Email Address:	

I request access to the Dynameomics.org database and understand that if my request is granted, I will be bound by the following terms and conditions:

Purpose of database: The Dynameomics.org database is a research project managed by Dr. Valerie Daggett at the University of Washington. The purpose of the project is to characterize the native state dynamics and the folding / unfolding pathway of representatives from all known protein folds by molecular dynamics simulation.

Access: Access is granted for six (6) months from when we set up an account for you, and can be limited or terminated at anytime. Only the user named above is authorized to access the database and use the data. Should the user change jobs, PI, department, or institution, a new access agreement must be signed and submitted. Other individuals must request access; no sharing of logins is allowed. Access to the database is intended for research purposes only. All materials and services provided on the database are protected by copyright or other intellectual property rights owned and controlled by Dr. Daggett and University of Washington.

Data use: Access to our database is limited to performing queries and retrieving simulation and analysis data. Any other use may result in immediate cancellation of access. You agree to use the data from our database for internal research purposes only, and not distribute, publish, transfer, or otherwise make available the data to users other than those in your lab. If you are preparing a scientific publication or presentation and would like to publish or disclose the data from our database, you will need to obtain written permission from Dr. Daggett.

Feedback: You agree to provide us feedback on your use of our database and we are allowed to use any information you provide in improving our database.

Monitoring: We reserve the right to monitor all activities taking place on the server. We may publish aggregate or specific usage information that includes your queries.

Confidentiality: All information submitted to our database may become publicly accessible, and University of Washington does not guarantee confidentiality.

Citation: Any publications that utilize data from dynamomeomics.org must cite at least the first two of the following papers, and any or all of the last three as appropriate:

- Van der Kamp MW, Schaeffer RD, Jonsson AL, Scouras AD, Simms AM, Toofanny RD, Benson NC, Anderson PC, Merkley ED, Rysavy S, Bromley D, Beck DAC, and Daggett V., *Dynamomeomics: A comprehensive database of protein dynamics*. Structure, 18: 423-435, 2010
- Beck, D. A. C., Jonsson, A. L., Schaeffer, R. D., Scott, K. A., Day, R., Toofanny, R. D., Alonso, D. O. V., Daggett V., *Dynamomeomics: Mass Annotation of Protein Dynamics by All-Atom Molecular Dynamics Simulations*. Protein Engineering Design & Selection 21: 353-368, 2008.
- Simms, A. M., Toofanny, R. D., Kehl, C., Daggett, V., *The Molecular Dynamics Data Warehouse: Design of a Computational Lab Workflow and Scientific Data Repository*. Protein Engineering Design & Selection 21: 369-377, 2008.
- Day, R., Beck D. A. C., Armen R. S., Daggett V., *A consensus view of fold space: Combining SCOP, CATH, and the Dali Domain Dictionary*. Protein Science, 2003. 12(10): 2150-2160.
- Kehl, C., Simms, A. M., Toofanny, R. D., Daggett, V., *The Dynamomeomics OLAP Database: A Multi-Dimensional Analysis-Optimized Database for Molecular Dynamics Simulation Data and Metadata*. Protein Engineering Design & Selection 21: 379-386, 2008.

We would also like to promote your publication(s) based on these data on our website, please forward an appropriate reference.

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Limitation of Liability: The entire risk for use of this database lies with the user. The University of Washington reserves the right to modify the database or reduce or discontinue the database at any time. This database access is provided for educational and informational purposes only.

I AGREE TO THE TERMS AND CONDITIONS

User Signature:

Signature

Date

User: Printed Name

Principal Investigator Signature:

Signature

Date

PI: Printed Name