#### FAIR IMPLEMENTATION FOR NI4OS-EUROPE SERVICE PROVIDERS

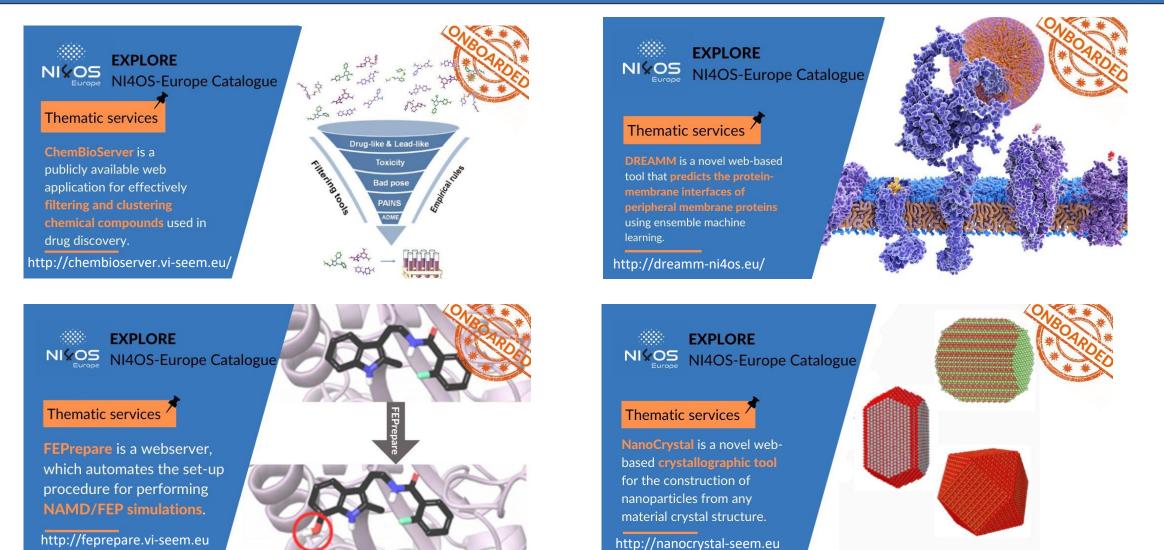
29 April 2022

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# Drug discovery tools onboarded on NI4OS-Europe





FAIR Implementation for NI4OS-Europe Service Providers, 29 April 2022

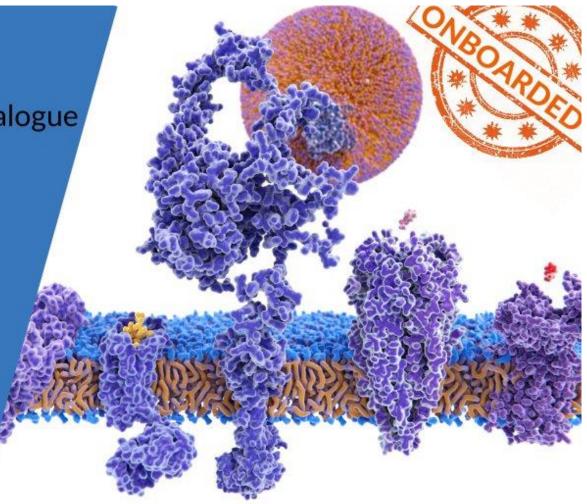
#### DREAMM: Predicting cavities at protein-membrane interfaces



#### Thematic services >

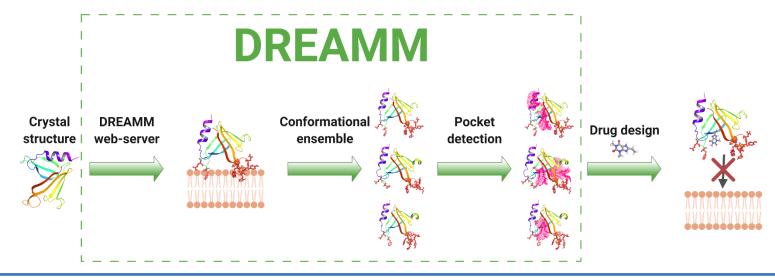
DREAMM is a novel web-based tool that predicts the proteinmembrane interfaces of peripheral membrane proteins using ensemble machine learning.

http://dreamm-ni4os.eu/



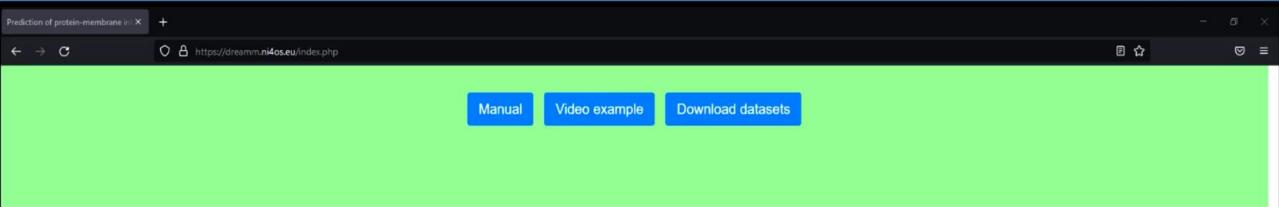
#### **DREAMM Main Features**

- Use an ensemble machine learning model that predicts the protein-membrane interfaces of peripheral membrane proteins
- Predict binding sites in the predicted protein-membrane regions in conformational ensembles using P2Rank
- Cluster the predicted binding sites for all protein conformations based on the binding sites center coordinates
- □ Create a novel web-based tool for predicting and drugging protein-membrane interfaces



## DREAMM Video Example





Check this box to search for binding sites (using P2Rank) near the predicted membrane-penetrating residues in protein ensembles:



#### DREAMM: Predicting cavities at protein-membrane interfaces

Information in NI4OS-Europe Agora: <u>https://catalogue.ni4os.eu/? =/resources/bc445004-2869-43c9-9ef3-e5aa6b44d2e3</u>

Access: <u>https://dreamm.ni4os.eu/</u>

Training Material: <u>https://training.ni4os.eu/mod/scorm/view.php?id=1187</u>

## DREAMM FAIR demands



#### □ Findability: Onboarded to NI4OS-Europe

#### **Accessibility**:

- Persistent identifier (PID) assigned to each job and URL of the results
- Results are only findable and accessible by the user who ran the prediction and deleted after 2 days
  - Possible Solution: Create a web-database storing the results, enabling search for relevant information through their metadata, by linking them
- □ Interoperability: Only PDB structures are interoperable.
- □ PyMOL visualizations and csv, txt, xlsx containing clustering results are not.
  - Onboarded
  - Solution: Add the binding site spheres as dummy atoms in the PDB structure and export clustering results in JSON or XML format
- **Reusable**: A detailed video example and a well-documented manual is available

containing:

- □ A short description of what data it includes
- Definitions of column headings and row labels for tabular data
- Whom to contact with questions
- Data are released with a clear and accessible data usage license:
  - https://wiki.ni4os.eu/images/7/77/DREAMM-Terms\_of\_use.pdf



# Thanks for your attention!

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