IntAct: The molecular interactions database at EMBL-EBI: webinar

Pablo Porras Millan [1]
Systems
Proteins
Beginner
1 hour

This two-part webinar from Pablo Porras Millan provides you with an introduction to protein-protein interactions and the molecular interaction database IntAct [2] (recorded on 30th September 2015). IntAct is a central, public repository where molecular interactions data can be stored and accessed.


NB: These videos works best using Google Chrome and when viewed in full screen.

Find out about upcoming webinars [4].

Learning objectives:

- Comprehend what protein-protein interactions are and how they are represented
- Search for molecular interaction on the IntAct website

IntAct webinars

An introduction to protein-protein interactions

An introduction to EMBL-EBI's molecular interaction database IntAct

Your feedback

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Contributors
Pablo Porras Millan [1]

EMBL-EBI
Scientific curator in IntAct

Pablo Porras got his PhD in 2006 in the University of Córdoba, Spain, having done research about trans-membrane protein translocation and redox homeostasis. After that, he moved to Berlin to work in the Neuroproteomics group of the Max Delbrueck Center, getting involved in projects dealing with interactomics, neurodegenerative diseases and the ubiquitin-proteasome system. During this postdoc, he faced the problem of how to represent and analyze molecular interactions data. This experience proved to be of great value once he joined the EBI to work as a scientific curator in the molecular interactions database IntAct [2] in 2011.

Birgit Meldal [5]

EMBL-EBI
Scientific Database Curator (Molecular Interaction & Complexes) - Orchard Team: Molecular Interactions

Birgit Meldal joined EMBL-EBI in 2012 as a Scientific Curator for the molecular interactions database IntAct. Her work focuses on curating macromolecular complexes into the Complex Portal, and she also contributes to the Gene Ontology. Birgit is a trainer for the IntAct and Reactome databases. She gained her PhD from the University of Southampton (in collaboration with the Natural History Museum, London) in 2004, having built the first comprehensive molecular phylogeny of the phylum Nematoda. She followed this with postdoctoral positions at the University of Cambridge working on the function of the breast cancer gene EMSY and on molecular epidemiology and host-virus interactions of transfusion-transmissible, blood-borne viruses (Hepatitis B and Hepatitis E virus).
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