Exploring models for human disease with UniProt

Sangya Pundir [1]

- Proteins
- Beginner
- 0.5 hour

UniProt is the Universal Protein Resource covering human variants and their disease association, in addition to general protein function and sequence data. This webinar will illustrate how to explore models for human disease variants to support experimental design.

You can learn more about UniProt in our UniProt: Quick tour [2].

This webinar was recorded on 5th April 2017. It is best viewed in full screen mode using Google Chrome. The slides from this webinar can be downloaded below.

See the EMBL-EBI training pages for a list of upcoming webinars [3].

Learning objectives:

- Search for diseases to discover related proteins
- Search for proteins and view related diseases
- View disease and related variants in a protein entry

Your feedback

Please tell us what you thought about this webinar. Your feedback is invaluable and helps us to improve our courses and thus enhance your learning experience.

Contributors

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Sangya Pundir is a User Experience (UX) Manager in EMBL-EBI’s UniProt team, where she established a user-centred process for the redesign of the world’s leading protein resource. To make UniProt easy for researchers to explore, Sangya conducts usability testing and information-gathering methods such as card sorting, contextual studies and workshops. Before she came to EMBL-EBI, Sangya worked at a healthcare consultancy, designing bespoke management systems. She holds an MSc in Biotechnology, Bioprocessing and Business Management from the University of Warwick.

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**Links**
[1] [https://www.ebi.ac.uk/training/online/trainers/sangya.pundir](https://www.ebi.ac.uk/training/online/trainers/sangya.pundir)
[2] [https://www.ebi.ac.uk/training/online/course/uniprot-quick-tourversion-0](https://www.ebi.ac.uk/training/online/course/uniprot-quick-tourversion-0)
[3] [http://www.ebi.ac.uk/training/webinars](http://www.ebi.ac.uk/training/webinars)