

## Scientists at IRB Barcelona and BSC publish the world's largest vid proteins

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By EurekAlert

After four years of conducting intensive calculations in the supercomputer MareNostrum at the Barcelona Su scientists headed by Modesto Orozco at the Institute for Research in Biomedicine (IRB Barcelona) have pres data base on protein motions. Called MoDEL, this new database holds more than 1,700 proteins and is partial Internet to researchers worldwide. MoDEL has been developed to study the basic biology of proteins and to a the design of new pharmaceutical agents.

"Nowadays we design drugs as if the proteins against which they are to act were static and this goes a long was the development of new drug therapies because this is not a true scenario. With MoDEL this problem is solve user from 10,000 to 100,000 photos per protein, and these confer movement to these structures and allow a masays Orozco, head of the "Molecular modelling and bioinformatics" group at IRB Barcelona, director of the Lof the Barcelona Supercomputing Center and full professor at the University of Barcelona. According to this pharmaceutical companies are already using the MoDEL strategy to develop the first drugs against cancer and which could become available this year.

## A project in expansion

The scientists that develop MoDEL work from an international catalogue of static protein structures (approxi Data Bank (PDB). "1,700 videos of proteins from the 40,000 that make up the PDB may appear to be a small structures in the PDB are very similar. Therefore, following internationally established similarity criteria, we a proteins with a known structure".

But for Modesto Orozco the most relevant point is that MoDEL is now covering more than 30% of human propharmacological interest, that is to say, those that are potential targets of a new drug. "We obtained this datur consider that we are in fact covering more. However, MoDEL will continue to grow and this can be achieved well established". According to the researchers, the main objective is to focus on relevant proteins in human c and 3 years cover 80% of pharmaceutical targets.

In order to undertake the MoDEL project, Orozco and his group are supported by resources provided by IRB SuperComputing Center, the Marcelino Botín Foundation, the Fundación Genoma España, the National Bioir European projects.

## **SOURCE**

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