## Case study I: How to publish a database easily for small/medium size sequencing projects

Input: A number of DNA/ Protein sequences

(+ Images)

blast2GO, KAAS etc.

to assign GO, KO etc. to the input

## **ANNOTATOR**

for a small number of sequences which are difficult to be predict their functions

## **BioMart**

to make annotated sequence tables of input data with related Ensembl sequences

## TogoDB

to publish annotated sequences as a database

