ReadME

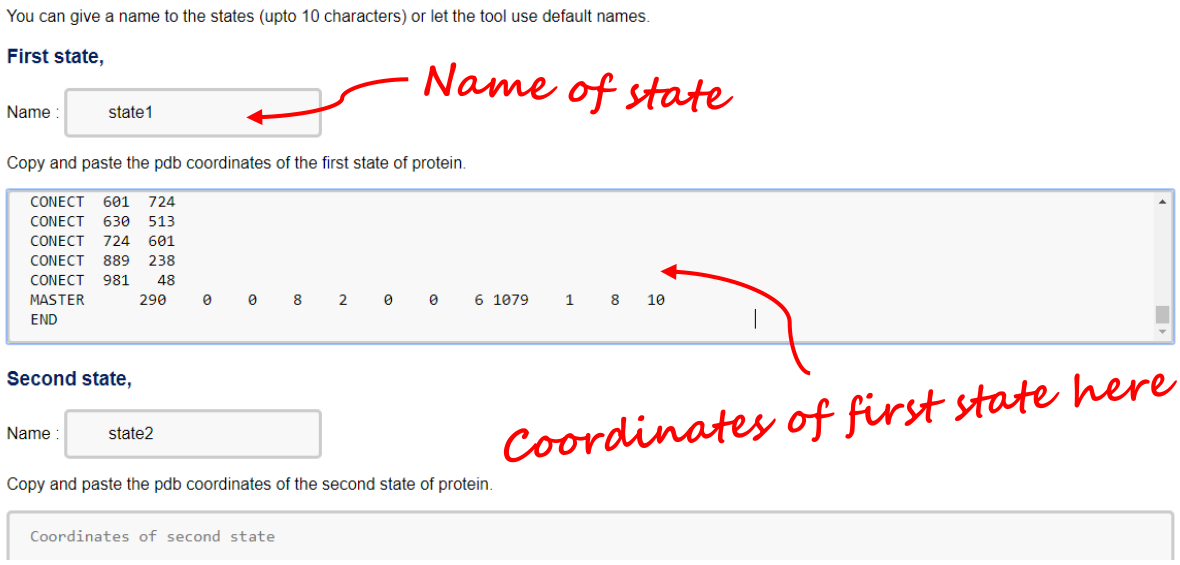
This ReadME documentation provides a guideline on how to perform secondary structure comparison between two states of a protein using CompASS. Follow the step by step instructions that are made easy to comprehend by pictorial representation.

CompASS requires the atomic coordinates of the protein in both states to compare the secondary structure elements in them.

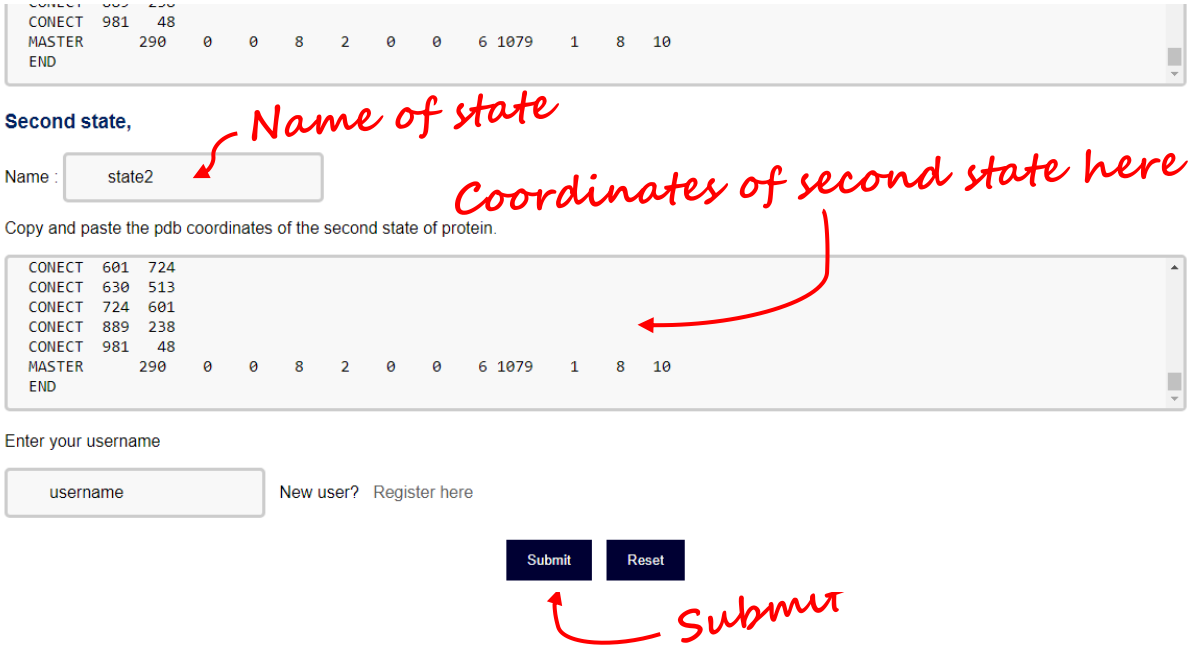
CONTENT

How to give inputs

STEP 1 Give a name to the first state of your protein. This will help you to identify this state in the alignment. Copy and paste the atomic coordinates of this state of protein in the first text area.



STEP 2 Now give name to the second state of protein. Copy and paste the atomic coordinates of protein in this state at the second text area.



STEP 3 Enter your username and submit the job to get results.