Jalview

a multiple alignment editor

Jalview Documentation

Jalview (2004) is a fast Java multiple alignment editor and analysis tool. It features many of the t of **AMAS**, for the analysis of sub-families and the prediction of functional sites, but is fully inter (View the **Jalview homepage**).

If you use Jalview in your work, please cite the bioinformatics paper:

Clamp, M., Cuff, J., Searle, S. M. and Barton, G. J. (2004), "The Jalview Java Alignment Editor", Bioinformatics, 20, 426-7.