

# Announcements 04/03

- Required Reading
  - Jumper, J., Evans, R., Pritzel, A. *et al.* Highly accurate protein structure prediction with AlphaFold. *Nature* **596**, 583–589 (2021).  
<https://doi.org/10.1038/s41586-021-03819-2>
- Project progress report due 4/8
  - Email to [daifeng.wang@wisc.edu](mailto:daifeng.wang@wisc.edu) is fine
  - Show preliminary results!
  - To be graded