

Announcements 04/02

- 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 4/7
 - Fine to email daifeng.wang@wisc.edu
 - Grading by preliminary results!