Questions for PEAK

1. Can we record meetings
2. When is good to meet for you
3. How long and frequent should meetings be
4. How often should we report back outside of meetings? Frequent communication will be mutually beneficial
5. Have you worked with students before, how did it go? Any recommendations?
6. How to communicate and who is allocated emailer
7. Add to GitLab
8. Minimum viable product
9. How should it looks in terms of UI and branding, examples would be good
10. What is the front end going to look like? How are we receiving the protein requests? How are people adding proteins?
11. Can you give us some user stories for the product, who is going to be using it
12. Approx. how large is the database going to be
13. A Kubernetes cluster appears to be a difficult thing to set up and maintain, can you explain how you expect us to use this with your system
14. What do you expect to see (prototype) by the next time we see you, what should we accomplish in the first sprint?
15. How do we filter out the ‘junk’ proteins from protein data bank without knowledge of that field
16. How do we decide which is the ‘best’ protein prediction to select from the Alpha Fold Database
17. Can you elaborate on what happens when a protein structure needs to be predicted. You mention this is resource intensive but would this be handled by us? Mentioned a request goes to another group
18. Do you want to search database for individual structures or search based on criteria to return multiple results
19. Any monetary incentive?
20. Licensing of the app