Some tips for the exam presentation

Technical:

- Make sure to number your slides!
- Start by briefly saying which paper you are talking about (title, where is it from, which year).
- Avoid excessive whitespace; can you enlarge that figure or increase the font of that bit of text?
- Avoid excessive text or figures/text that is too small to see well
- If you have too much on one slide, think about whether you can split up the content in two slides.
- Use colors/highlighting to show where you are on a slide, for example a red box around part of a complex workflow figure to highlight that region
- Use animations as well, if you have three parts of text, then only show the first part when the slide appears, then the second when you start talking about it, … . Can also be done with figure highlights (or by incrementally showing parts of the figure)
- If you 'pull' figures from the internet (or elsewhere), give the source.

Content:

- Make sure you introduce and situate your topic broadly, focusing on the essential concept that the method is addressing. For example, if talking about protein structure prediction, start very briefly with 'what do proteins do', 'how does sequence relate to structure', and 'how does this enable the function of proteins'. Core concepts only!
- The dataset information is important, say which kind of data, how many data, … so describe the data in some detail, and how reliable it is in your view.
- Try to give concrete examples, especially when talking about abstract concepts. For example, if you are talking about a feature important for the method, then show a real example of such a feature.
- If you think the method has pitfalls or shortcomings, or if there are aspects of it that are not well explained, please discuss those. Same for figures that might not have all the relevant information or might have been done differently. Think critically!