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| MSc supervisory meeting summary | | |
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| Location: | MB211L |  |
| Date: | 09/08/2023 |  |
| Time: | **13:00-13:30** |  |

# Summary of discussion

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|  | Item | **Brief Description** | **Action required** |
|  | BepiPred 3.0 | Using the holdout set, get the ordered protein sequences and input them into BepiPred 3.0 in FASTA format (manually type or code to do so) | Get BepiPred 3.0 results |
|  | Comparison | Compare the performance of BepiPred 3.0 to the genetic programming model | Finish comparison. |
|  | Documentation | Continue documentation | Finish documentation. |
|  |  |  |  |

## Additional information

## Sign off

Navroop Singh [ X ] Felipe Campelo [ X ]