

- Identified important genes which were found to be differentially expressed can be taken for further wet-lab experimentation and later can be developed as a therapeutic drug target for e.g. *MT-TI*.
- In near future with more availability of data the similar analysis can be carried out on bigger dataset.
- Similar analysis can be designed to understand other complex disease such as cancer, diabetes etc.
- GnRH pathway can be studied on priority with respect to Alzheimer as it was found to be most associated pathway with AD.