Instructions: -

1. The following are general purpose functions which are later called by R markdown codes. These functions are used to produce a structured output of the R markdown files. They are meant to be in the folder where the R markdown codes are and are called inside the R markdown codes:
2. Table\_Fun\_ComparsionCandiatevsBiomarkers 220114.R
3. Table\_Fun\_ComparsionCandiatevsBiomarkers.R
4. Table\_Fun\_ComparsionCandiatevsBiomarkers\_modified.R
5. Table\_Fun2\_Space 220114.R
6. Table\_Fun2\_Space.R
7. Table\_Fun2\_Space\_modified.R
8. Manuscript Table-1: - Selected 0utputs from the following R codes:
9. FNIH table1 and supp table-3.R
10. table-1 summary output MS.R
11. Manuscript Table-2: - Selected outputs from the R Markdown file:
12. FNIH-Report-20210820.Rmd
13. Manuscript Table-3: - Selected outputs from the R Markdown file:
14. FNIH-Report-TaskId5-20220420.Rmd
15. Supplemental Table 3: - Selected outputs from the following R code:
16. FNIH table1 and supp table-3.R
17. Supplemental Table 4 & 5: - Selected outputs from the R Markdown file
18. FNIH-Report-TaskID7&8-20220421-modified.Rmd