**GUIDELINE TO USING EMBEDDING TECHNIQUES FOR OMICS DATA**

Users can refer to the schematic diagram to find the best suited embedding model for their purpose in analysing biological sequence data.

How to use the Guideline effectively:

1. Understand the domain of your research area in the sequential data analysis.
2. Trace the application area of your research area to find the data needed to accomplish the task (BLUE boxes).
3. Click on the desired dataset to find the database for them.
4. Find the options available for embedding your data for further analysis or classification.
5. Click on the desired embedding technique to find the package available for the embedding. (GREEN boxes)