## **Assignment-4**

- Q.1 Compute the complete graph G=(V, E) of the project dataset.
- Q.2. Construct MST neighborhood graph of 3-round i.e. G'=(V, E').
- Q.3 Let A' be the adjacency matrix of G'. Compute the adjacency matrix A'.
- Q.4 Find the degree matrix D of G', where each D(i,j)=0,  $i\neq j$  and  $D(i,i) = \sum_{j=1}^{j=n} A'(i,j)$
- Q.5 Compute the Laplacian Matrix L= D-A'.
- Q.6 Compute first k eigenvectors  $v_1, v_2, v_3, \dots, v_k$  of L, where k is the number of clusters in your dataset.
- Q7. Form a n\*k matrix U, where each eigenvector  $v_i$ ;  $1 \le i \le k$  is stacked as a column in U. Treat each row  $u_i$   $1 \le i \le n$  of U as a data point.
- Q.8 Apply k-means and MST based clustering algorithm to partition the points  $u_{i=1,2,3,...,n}$  into k clusters.
- Q.9 Compute the accuracy and execution time of each algorithm.

## Reference:

[1] Jothi, R., Mohanty, S. K., & Ojha, A. (2016). Functional grouping of similar genes using eigenanalysis on minimum spanning tree based neighborhood graphs. *Computers in biology and medicine*, 71, 135-148.