

Assignment – II **(k-mer Analysis, Dotplots)**

Deadline: 30th March

1. Simulate observations having the binomial distribution with $p = 0.25$ and $n = 1000$. What is the probability of observing at least 240 A's in such a sequence? [Hint: Obtain 10,000 simulations and compute the number of A's in each run]. Compare your result with the normal approximation to binomial distribution.
2. Suppose X has a binomial distribution with $p = 0.3$ and $n = 10$. Compute $P(X=0)$, $P(X=2)$, $E(X)$ and $\text{Var}X$.
3. Briefly discuss the applications of k-mer analysis.
4. Show a dotplot of the following two sequences and give the conserved region:
(Make a $n \times m$ table and put '.' or 'x' for match)

GGCTGCAACTAGCTC
GGGTAAGCTTGC

5. Obtain the self-dotplot of the following sequence to identify repeat region:
TGGCACACTCACACCACACAGACAGTTA
6. Find self-complementary regions in the following RNA sequence:
AUGUGGCAUGCCAGG