

Deterministic approach to Gibbs Sampling to find DNA Sequence Motifs

- **Goal:** Find regulatory motifs in DNA by looking at the most commonly repeated kmer (with mismatches) in DNA sequences
- **Method:** Gibbs Sampling with incorporation of a deterministic selection once the sample begins to converge
 - Deterministic = $\min(\text{HammingDistance}(\text{Consensus}, \text{k-mer}))$
 - Selection of motif = $\text{random}(\text{Deterministic}, \text{Gibbs})$
 - Probability of Gibbs = $(\tan^{-1}(\text{Score_Slope}))/90$
 - Probability of Deterministic = $1 - \text{Gibbs}$
 - Score_Slope = Is the average slope of the graph of the score at each iteration. Can be calculated via exponentially weighted moving average
- **Expected results:** The deterministic approach should give a faster and more accurate convergence
- **Testing:** Test both my method and plain Gibbs sampling method on DNA strands with motif to quantify both performance and accuracy
- **Related Work:** Gert Thijs, Kathleen Marchal, Magali Lescot, Stephane Rombauts, Bart De Moor, Pierre Rouzé, and Yves Moreau. Journal of Computational Biology. July 2004, 9(2): 447-464. doi:10.1089/10665270252935566.