

Introduction to Bioinformatics

1. **Course Code** : BLM3810
2. **Course Name** : Introduction Bioinformatics
3. **Instructor** : Nizamettin AYDIN
4. **Credit** : 3
5. **Nature of the course** : Lecture
6. **Assesments** :
 - Homeworks : 20%
 - Midterm 1 : 25%
 - Midterm 2 : 25%
 - Final : 30%
7. **Course Outline** :
 - Introduction
 - Definition of Bioinformatics, Importance of Bioinformatics
 - Overview of Molecular Biology
 - Cells, Chromosomes, DNA, RNA, Amino Acids, Proteins
 - Genome, Transcriptome, Proteome
 - Setting the Technological Scene
 - Perl as a Software Tool, Installing, Debugging, Programming
 - Pairwise Sequence Alignment
 - Relation of Sequences
 - Alignment methods (Visual, Brute Force, Dynamic Programming, Word-Based)
 - Dot plots, Global Alignment, Local Alignment
 - Scoring Matrices, Significance of Alignments
 - Advance Perl programming
 - Perl one liners, Database systems, Web Technologies
 - Multiple Sequence Alignment
 - Global Multiple Alignment, progressive global alignment, Iterative methods,
 - Alignments based on locally conserved patterns
 - Local Multiple Alignment, Profile Analysis, Block Analysis,
 - Pattern-searching or statistical methods
 - Sequence File Formats
8. **Recommended Texts** :
 - Introduction to Bioinformatics*. Arthur M. Lesk
 - Bioinformatics, Biocomputing and Perl*. Michael Moorhouse, Paul Barry
 - Fundamental Concepts of Bioinformatics*. Dan E. Krane, Michael L. Raymer
 - Beginning Perl for Bioinformatics*. James Tisdall
 - Bioinformatics: A Practical Guide to the Analysis of Genes and Proteins*. Andreas D. Baxevanis
 - Bioinformatics: Sequence and Genome Analysis*. David W. Mount
 - Biological Sequence Analysis: Probabilistic Models of Proteins and Nucleic Acids*. Richard Durbin G. Mitchison S. Eddy A. Krogh
 - Statistical Methods in Bioinformatics: An Introduction*. W. Ewens, G. Grant.
9. **Course Materials** : <http://www.yildiz.edu.tr/~naydin>