This course will provide an introduction to Bioinformatics and the primary methods for data and text mining for computational biology.

**Course Objectives:**
- Understand the fundamentals of the field of bioinformatics.
- Learn the main bioinformatics problems and algorithms proposed to solve them.
- Design and apply computational methods to biological data.
- Read/present review papers on state-of-the-art research in bioinformatics.
- Prepare for original research in bioinformatics.

**Prerequisites:** Medium level programming skills in any high level programming language (e.g. C/C++, Java, Perl, Python, etc.), background on data structures and algorithms.

**Textbooks (Reference):**
  Supplementary website: http://bioinformaticsalgorithms.com

**Course Web Site:**
We will use the Moodle Course Management System for lecture notes, announcements, grades, and project submissions: [http://moodle.boun.edu.tr/](http://moodle.boun.edu.tr/)

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**Tentative List of Topics:**
- Review of relevant background material from molecular biology
- Dynamic Programming and Sequence Alignment
- Motif finding and Randomized Algorithms
- Combinatorial Pattern Matching and Suffix Trees
- Heuristic Sequence Similarity Search Algorithms (FASTA and BLAST)
- Graph algorithms and genome assembly
- Gene expression analysis, Clustering and Classification Algorithms
- Phylogenetic Trees
- Hidden Markov Models
- Text mining for biology

**Grading:**
- Assignments: 20%
- Midterm Exam: 20%
- Paper summaries: 10%
- Paper presentation: 15%
- Term Project: 30%
- Class Participation: 5%