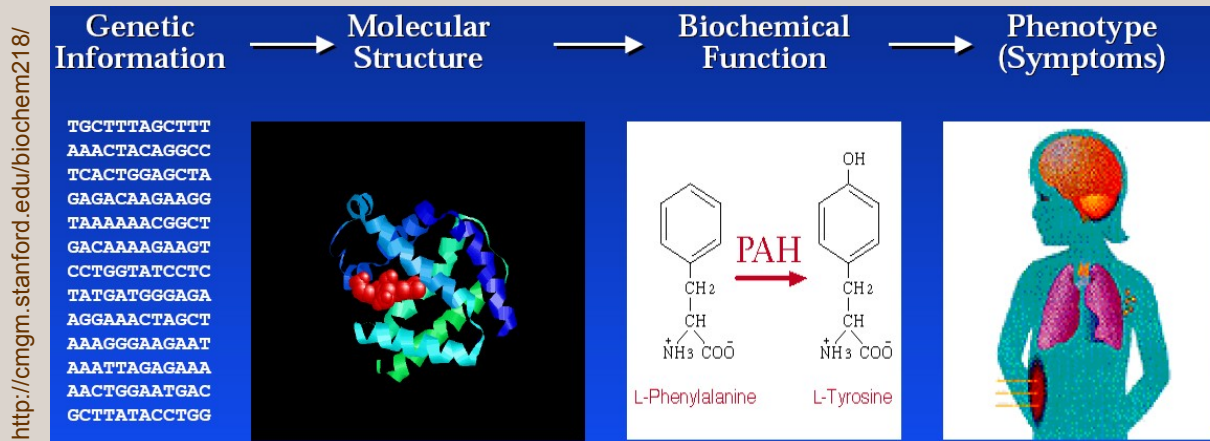# CSE 308-408 • Bioinformatics: Issues and Algorithms
Professor Daniel Lopresti • Fall 2007 • TuTh 2:35 - 3:50

"Biology easily has 500 years of exciting problems to work on."
– *Donald Knuth (Stanford Professor & famous Computer Scientist)*

By applying methods from Computer Science to the analysis of sequence data and related structures, we can help in understanding genetic nature of disease.

Topics we study in CSE 308-408:

- sequence comparison & alignment
- sequence assembly
- physical mapping of DNA
- phylogenetic (evolutionary) trees

- genome rearrangements
- RNA & protein structure prediction
- DNA microarray analysis
- Perl programming for bioinformatics

No prior knowledge of biology is required. For more details: dal9@lehigh.edu.

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