321 Numerical methods in computational molecular biology

Lecture: TR Phillips 203 1:25-2:15PM, Section: Hollister B14 R 2:30-3:20

Web site: http://www.cs.cornell.edu/Courses/cs321/2005fa/

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Walk-in on Thursdays 10:00, or by appointment

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Statistics and Numerical methods motivated by problems from computational molecular biology.

Grades: 5% quiz; 15% homework; 50% projects; 20% final

## Textbooks:

M.H. DeGroot and M.J. Schervish, "Probability and Statistics", Addison Wesley Rudra Pratap, "Getting started with MATLAB 5", Oxford University Press Charles F. Van Loan, "Introduction to Scientific Computing", Prentice-Hall

- (\*) Probability: Continuous variables
  - 1. Random variables
  - 2. Probability density function
  - 3. Uniform, normal and exponential distribution
  - 4. Expectations and variance
  - 5. Law of large numbers
  - 6. Central limit theorem
  - 7. More than one random variables
  - 8. Rejection and transformation methods for sampling distributions
  - 9. Transformation methods for sampling distributions (section)
- (\*) Statistics:
  - 1. Estimators: Mean, standard deviation
  - 2. Maximum likelihood
  - 3. Confidence interval
  - 4. chi-2 statistics
  - 5. Regression
  - 6. Goodness of fit
- (\*) Matlab preliminaries:
  - 1. What is Matlab?
  - 2. Sample of simple programs
  - 3. Plotting in one and more dimensions
  - 4. Writing functions
- (\*) Analysis of protein shapes
  - 1. Plotting protein shapes (vectors, matrices, PDB files)
  - 2. Smoothing and interpolation (linear interpolations, linear problems, cubic splines)
  - 3. Rotations and overlapping structures (least square fit, Lagrange multipliers, Matrix diagonalization, singular value decomposition)
  - 4. Clustering protein shapes (building RMS matrices, eigenvalue analysis, identification of families)
- (\*) Sequence analysis
  - 1. Manipulation of characters in Matlab
  - 2. Alignment of sequences with gaps
  - 3. Dynamic programming
- (\*) Optimization
  - 1. Steepest Descent
  - 2. Conjugate gradient
  - 3. Discrete optimization by Monte Carlo: Design of an optimal sequence

Each student in this course is expected to abide by the Cornell University Code of Academic Integrity. Any work submitted by a student in this course for academic credit will be student's own work.