#### Biomedical Informatics: Lecture 1

Alberto Paoluzzi

Tue, Mar 3, 2015

## Outline: Syllabus, GitHub

- Syllabus
- Student projects
- Grading
- GitHub
- 6 Assignments
- 6 Introduction to bioinformatics

# Syllabus



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General information



- General information
- Course notes and student home



- General information
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- Programming tools



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- Programming tools
  - Pyton



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  - pyplasm (PLaSM for Python)



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  - Pyton
  - pyplasm (PLaSM for Python)
  - Javascript
  - plasm.js (PLaSM for JavaScript)

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#### Scheduling

1 Project design: due Fri, May 2, 2014



#### Scheduling

- Project design: due Fri, May 2, 2014
- Working prototype: due Fri, June 6, 2014



#### Scheduling

- Project design: due Fri, May 2, 2014
- Working prototype: due Fri, June 6, 2014
- Project deadline: Before final exam

# Grading



#### Exam requirements

• Programming project (informatics: prof. Paoluzzi) ( $\leq 15$ )



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## Exam requirements

- Programming project (informatics: prof. Paoluzzi) ( $\leq 15$ )
- 2 Term Paper (biomedical: prof. Ceccanti) ( $\leq 10$ )



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# Exam requirements

- **1** Programming project (informatics: prof. Paoluzzi) ( $\leq 15$ )
- **2** Term Paper (biomedical: prof. Ceccanti) ( $\leq 10$ )
- **3** Project design and prototype presentations ( $\leq 5$ )



## GitHub



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# SignOn GitHub

GitHub is a web-based hosting service for software development projects that use the Git revision control system.

GitHub offers both paid plans for private repositories, and free accounts for open source projects.

• configure git and github (guide)



# Assignments



#### Enroll to the course !!



To:	Alberto Paoluzzi <apaoluzzi@gmail.com></apaoluzzi@gmail.com>	
Cc:		
Bcc:		
Reply To:		
Subject:	[Informatica biomedica] iscrizione al corso 2014	
≣ ▼ From:	Alberto Paoluzzi <apaoluzzi@me.com> †   iCloud (iCloud)</apaoluzzi@me.com>	<b>‡</b>
ognome Nome rimo anno laurea magistrale (oppure: secondo) ngegneria informatica (oppure: altro) atricola: xxxxxx mail: accountr@provider rafica computazionale: SI (oppure: NO) nteressato a tesi di laurea: SI (oppure: NO)		

send an email to me ... NOW!

 Essay (Pre-thesis) on bioinformatics topics at the Mendel Institute with Dr. Valerio Mazza



- Essay (Pre-thesis) on bioinformatics topics at the Mendel Institute with Dr. Valerio Mazza
- Import/export of image/graphics/CAD formats to/from LAR format



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Your choice to be discussed with me (after April, 8)



#### References

Course syllabus

Principles of Biomedical Informatics, Second Edition

Handbook of Biomedical Informatics



#### Introduction to bioinformatics

#### **Bioinformatics**

Oxford English Dictionary, 2001

(Molecular) bio – informatics: bioinformatics is conceptualising biology in terms of molecules (in the sense of physical chemistry) and applying "informatics techniques" (derived from disciplines such as applied maths, computer science and statistics) to understand and organise the information associated with these molecules, on a large scale. In short, bioinformatics is a management information system for molecular biology and has many practical applications.



states that DNA is transcribed into RNA, which is then translated into proteins

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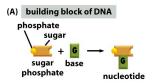
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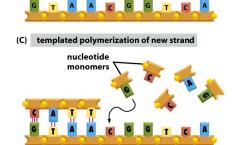
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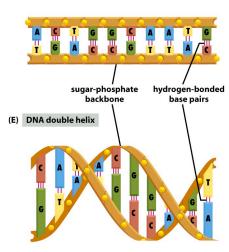
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- The genetic code is the mapping from codons to amino acids
- A gene is a sequence of codons and "non-coding" sequences that determine when the gene is active (expressed).
- There are stop and start codons which signal the beginning and end of the protein coding sequence
- Proteins are organic compounds made of amino acids arranged in a linear chain and folded into a globular form.



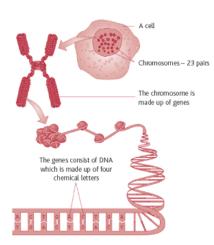
(B) DNA strand



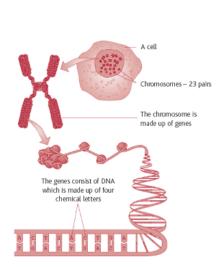
(D) double-stranded DNA

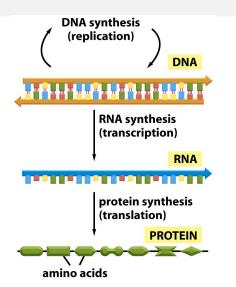


### **DNA** structure

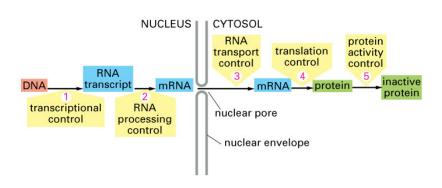


### **DNA** structure





### Gene expression process



MyPDB Login

A MEMBER OF THE PDB

### SPDB PROTEIN DATA BANK

## An Information Portal to Biological Macromolecular Structures

(WHAT'S NEW) | HELP | PRINT

PDB ID or keyword

Search 2

Advanced Search

### **‡** Home

News & Publications Policies FAQ Contact Feedback About Us

#### Deposition

All Deposit Services Electron Microscopy NMR Validation Server BioSync Beamline Related Tools

#### **≜** Search

Advanced Search Latest Release Latest Publications Sequence Search Ligand Search Unreleased Entries Browse Database Histograms

### ‡ Tools

File Downloads
File Formats
Services: RESTful | SOAP

### A Resource for Studying Biological Macromolecules

The PDB archive contains information about experimentally-determined structures of proteins, nucleic acids, and complex assemblies. As a member of the wwPDB, the RCSB PDB curates and annotates PDB data according to agreed upon standards.

0

The RCSB PDB also provides a variety of tools and resources. Users can perform simple and advanced searches based on annotations relating to sequence, structure and function. These molecules are visualized, downloaded, and analyzed by users who range from students to specialized scientists.

#### Molecule of the Month: Xanthine Oxidoreductase



Our diet includes a wide variety of different molecules. Many of these molecules are broken down completely and used to generate the metabolic energy that powers our cells. Others are disassembled piece-by-piece and recycled to build our own proteins and nucleic acids. The ones that are left over are broken down and discarded. Xanthine oxidoreductase, shown here from PDB entry 16-4, is the last stop for extra purine nucleotides (ATP and GTP) in our cells. Purines are broken down in several steps.

ultimately yielding uric acid, which is excreted from the body.

Biomedical Informatics: Lecture 1

■ Read more ... ■ Previous Features

### News

#### ■ Complete News

- Newsletter
- Discussion Forum
- Job Listings

29-September-2009 SF-Tool: A Tool for Crystallographic Experimental Data Validation

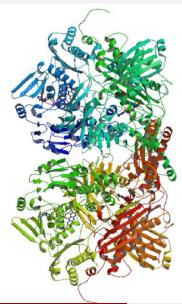
A streamlined, web-based tool is available for validating crystallographic experimental data. **SF-Tool** can be used to:

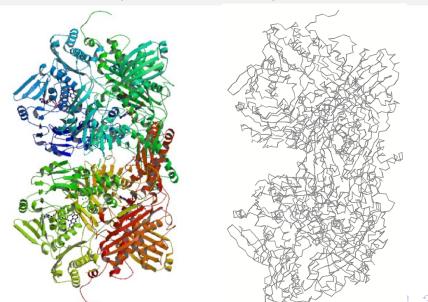
- Validate model coordinates against structure factor data (using SFCheck)
- Easily translate your structure factor file between different formats (mmCIF, MTZ, CNS/CNX, XPLOR, SHELX, TNT, HKL2000, SCALEPACK, D\*Trek, SAINT, or OTHER format)

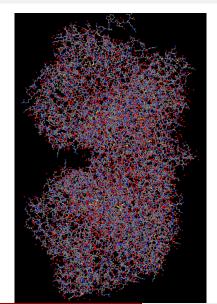
More >>

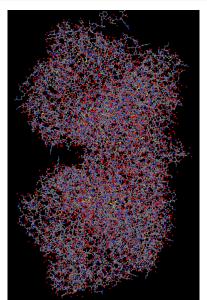
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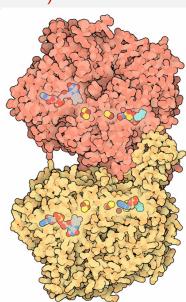
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         4 SYNONYM: XD;
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         5 EC: 1.1.1.204
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         3 ORGANISM COMMON: CATTLE:
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KEYWDS
          XANTHINE DEHYDROGENASE, FAD, MOLYBDOPTERIN, 2FE-2S IRON
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         2 SULFUR CENTERS, SALICYLATE, OXIDOREDUCTASE
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                                                                                                                                  20.572
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AUTHOR
          C.ENROTH.B.T.EGER.K.OKAMOTO.T.NISHINO.T.NISHINO.E.F.PAI
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REVDAT
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            TITL 2 DEHYDROGENASE AND XANTHINE OXIDASE:
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TRNI
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REMARK
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REMARK
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                           BRUNGER, ADAMS, CLORE, DELANO, GROS, GROSSE-
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REMARK
                           KUNSTLEVE.JIANG.KUSZEWSKI.NILGES. PANNU.
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REMARK
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            REFINEMENT TARGET : ENGH & HUBER
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REMARK
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REMARK
            DATA USED IN REFINEMENT.
                                                                                                 ATOM
             RESOLUTION RANGE HIGH (ANGSTROMS): 2.10
REMARK
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                                                                                                                                  30.712
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             RESOLUTION RANGE LOW (ANGSTROMS) : 25.00
REMARK
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REMARK
             DATA CUTOFF
                                     (SIGMA(F)): 0.000
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             DATA CUTOFF HIGH
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                                                                                                                   DHE A
                                                                                                                                          28.777
                                                                                                                                                  21,269 1,00 26,36
REMARK
         3
             DATA CUTOFF LOW
                                       (ABS(F)) : NULL
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REMARK
             COMPLETENESS (WORKING+TEST)
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REMARK
             NUMBER OF REFLECTIONS
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### Beyond central dogma: epigenetics

Genetic Science Learning Center – University of Utah WHAT IS EPIGENETICS?

Science 29 October 2010: Vol. 330 no. 6004 p. 611

DOI: 10.1126/science.330.6004.611

INTRODUCTION TO SPECIAL ISSUE

What Is Epigenetics?