



上海交通大学  
SHANGHAI JIAO TONG UNIVERSITY



# Programming Language for Bioinformatics

## 生物计算编程语言

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- Programming languages for Bioinformatics
- Course information
  - Goal
  - Contents
  - Organization
  - Grading



How many programming languages you know?

C

C++

Pascal

Java

Basic

Perl

Python

Rubby

...

Matlab

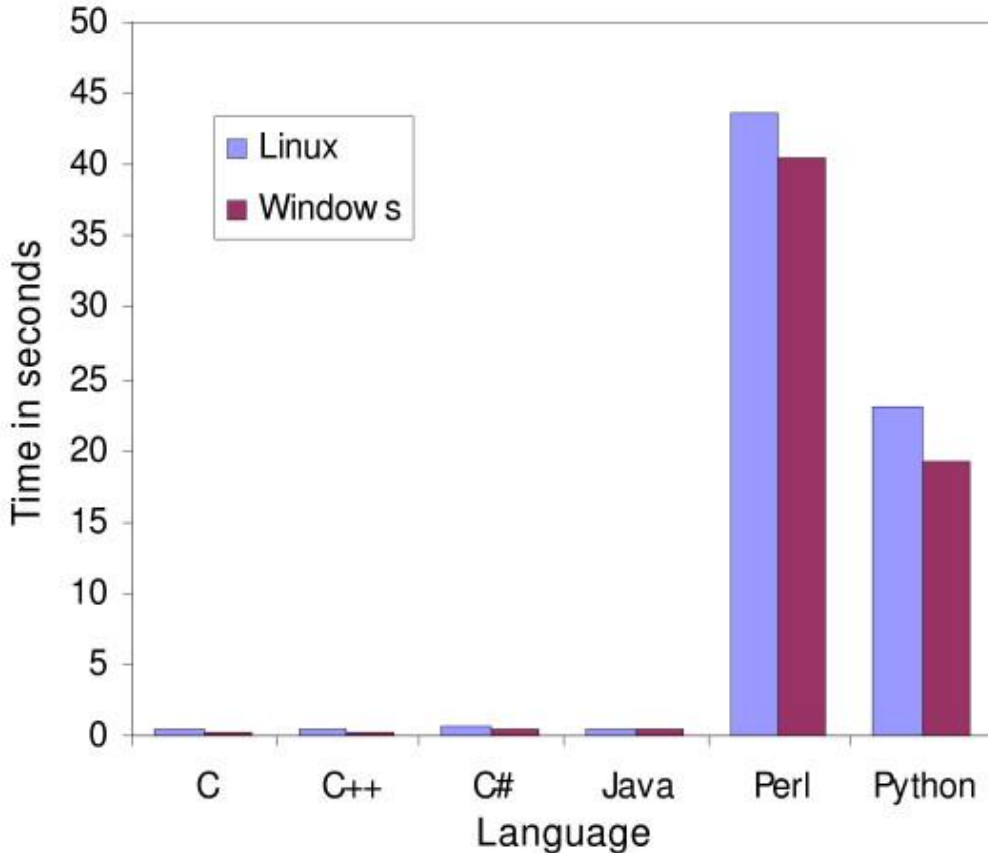
Mathematica

R





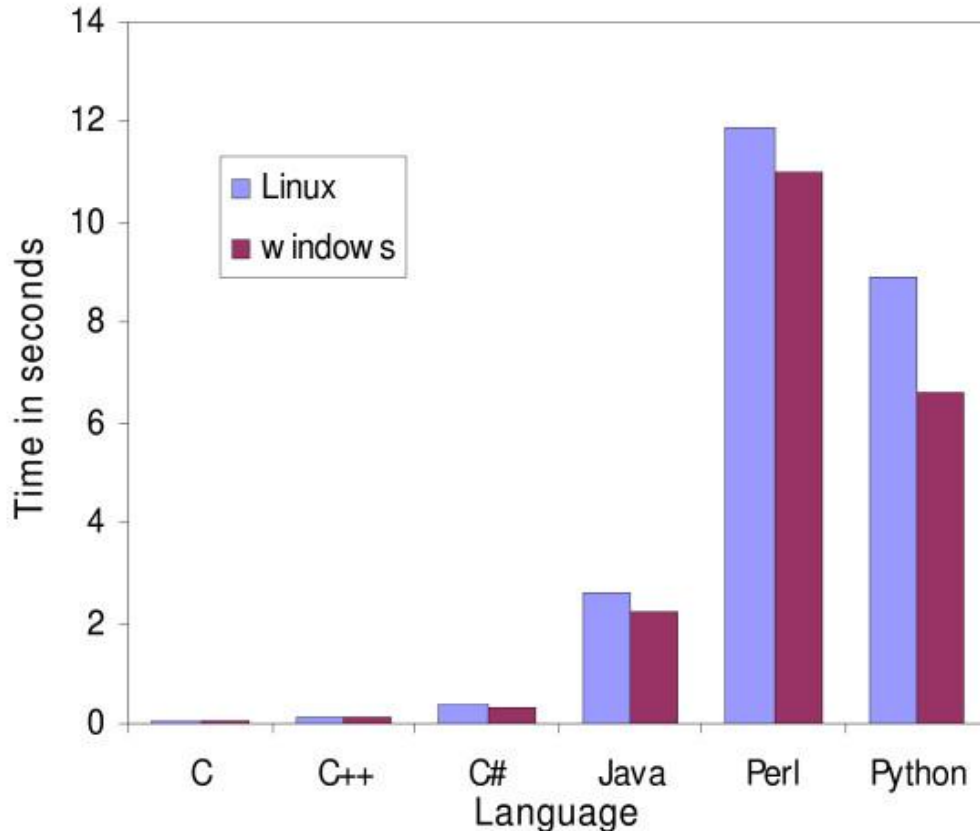
# Comparison of different programming languages for bioinformatics



**Speed comparison of the global alignment program.** Speed comparison of the global alignment algorithm using a gap penalty of 10 implemented in C, C++, C#, Java, Perl and Python. The programs were run on Linux and Windows platforms. Two DNA sequences of 3216 bp and 3217 bp were used.



# Comparison of different programming languages for bioinformatics

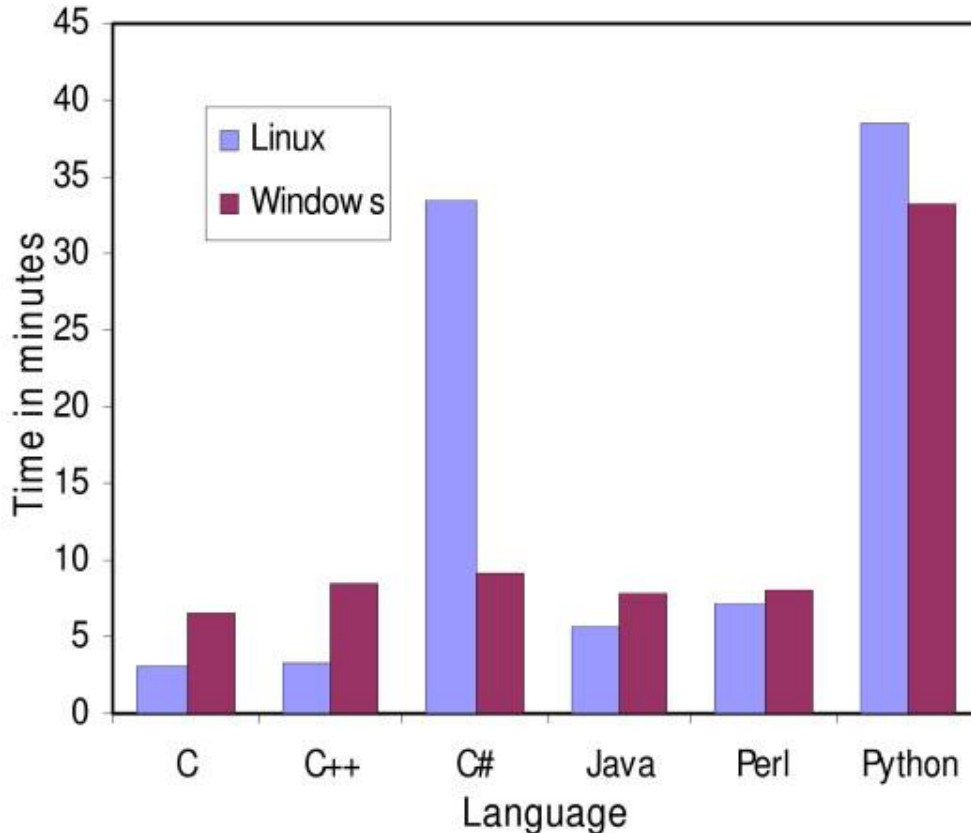


## Speed comparison of the Neighbor-Joining program.

Speed comparison of the Neighbor-Joining algorithm using the Jukes-Cantor evolutionary model implemented in C, C++, C#, Java, Perl and Python. The programs were run on Linux and Windows platforms. The input file was an alignment of 76 DNA sequences.



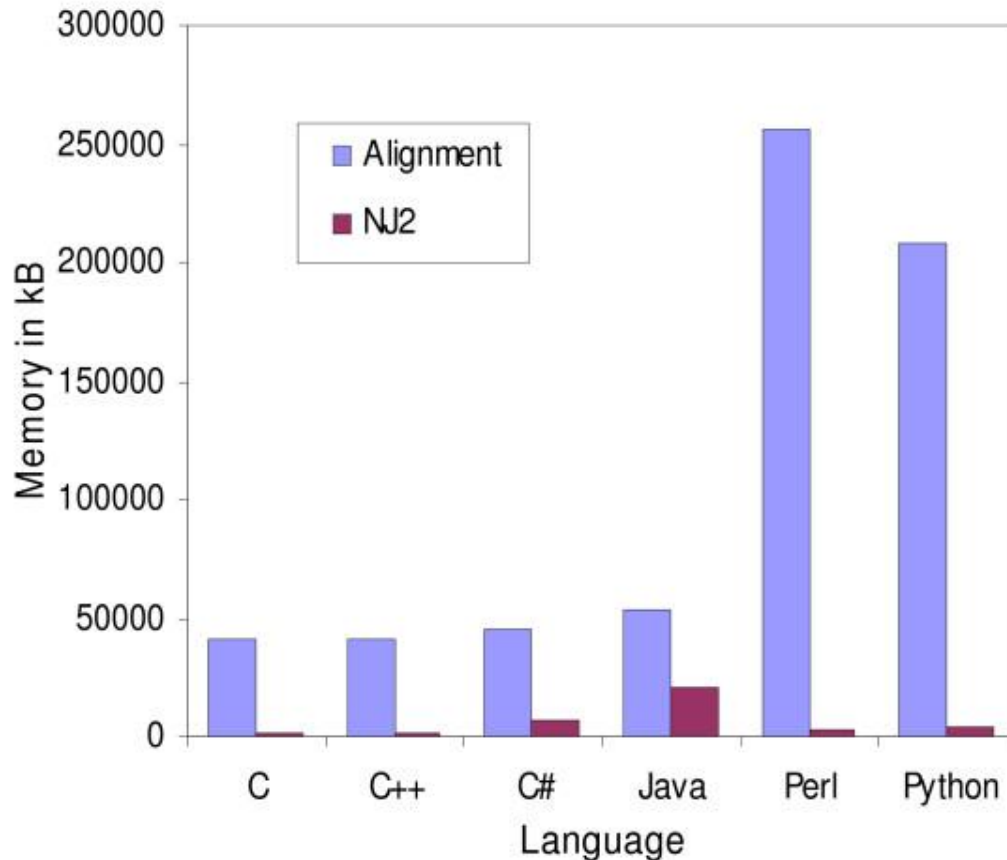
# Comparison of different programming languages for bioinformatics



**Speed comparison of the BLAST parsing program.** Speed comparison of the BLAST parsing program implemented in C, C++, C#, Java, Perl and Python. The programs were run on Linux and Windows platforms. The input file was a 9.8 Gb file from a BLASTP run.



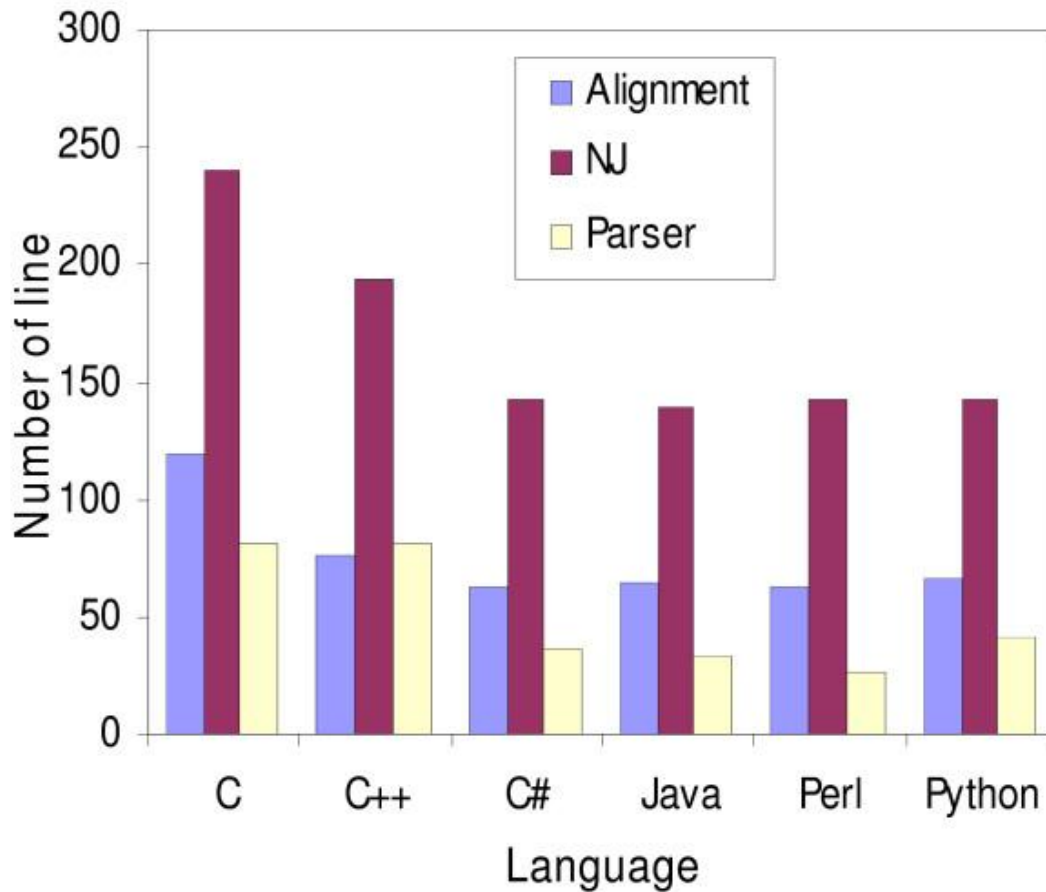
# Comparison of different programming languages for bioinformatics



**Memory usage comparison of the Neighbor-Joining and global alignment programs.** Memory usage comparison for the Neighbor-Joining and global alignment programs implemented in C, C++, C#, Java, Perl and Python. The programs were run on a Linux platform.



# Comparison of different programming languages for bioinformatics



**M Number of lines for each program.** Number of lines for the global alignment, BLAST parser and Neighbor-Joining programs implemented in C, C++, C#, Java, Perl and Python.





# Course goals

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- ④ Understand C programming language in general
  - ④ Learn how to write good code
  - ④ Learn how to work together
  - ④ Practice, practice and practice...
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# Course contents

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- Introduction to C programming language
- \*Programming skills other than languages
  - Debugging tools
  - Using open source communities
  - Keeping projects documented and manageable
  - Source code managing
  - Parallel computing
  - Understanding the hardware
  - Value your time

\*Dudley JT, Butte AJ (2009) A Quick Guide for Developing Effective Bioinformatics Programming Skills. *PLoS Comput Biol* 5(12): e1000589.



# Course organization

- Course introduction ( Week 1)
  - Code editor: Emacs
- Part I: Introduction to C programming language (Week 1 - 12)
  - Chapter 1: Overall Introduction (Week 1-4)
    - C
    - Unix/Linux
  - Chapter 2: Types, operators and expressions (Week 5)
  - Chapter 3: Control flow (Week 6)
  - Chapter 4: Functions and program structure (Week 7-8)
  - Chapter 5: Pointers and arrays (Week 9)
  - Chapter 6: Structures (Week 10)
  - Chapter 7: Input and Output (Week 11)
- Part II: Skills others than programming languages (Week 12- 14)
  - Debugging tools (Week 12-13)
  - Keeping projects documented and manageable (Week 14)
  - Source code managing (Week 14)
- Part III: Reports from the battle field (student forum) (Week 15 – 16)



## Course organization (2)

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- Thursday 14:00-15:40 (Every Week)
  - 生物药楼4号楼-225, 生信实验室
- Tuesday 8:00-9:40 (Even weeks)
  - 生物药楼4号楼-225, 生信实验室



**Course features :**  
**this is a practice course and**  
**you need lots and lots of**  
**practices**



# Prerequisites

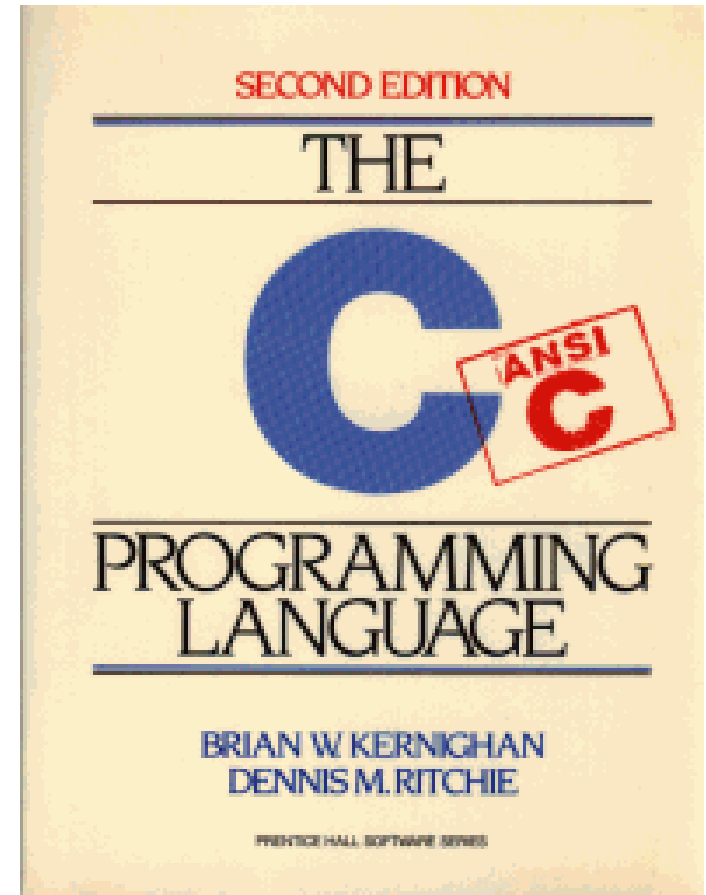
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- Mathematics (no)
- Computer Science ( no)
- Biology ( no)



# Text Book

The C Programming Language, Second Edition  
by Brian W. Kernighan and  
Dennis M. Ritchie. Prentice  
Hall, Inc., 1988.





# References

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- Text Editors

- Emacs

- tutorial: <http://www.gnu.org/software/emacs/tour/>

- Manual:

- <http://www.gnu.org/software/emacs/manual/emacs.pdf>

- Debug Tools

- GDB

- Document: <http://www.gnu.org/software/gdb/documentation/>





# Grading

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- Homework 50%
- Projects 30%
  - Design and implementation of a bioinformatics tool
- Presentation 20%



# 作业规定

- 作业允许合作，但是必须注明各人的贡献
- 作业报告必须用自己的语言独立完成
- 严禁抄袭
  - 抄袭者：不及格(F)
  - 被抄袭者：成绩降一级 ( $A \rightarrow B, B \rightarrow C, C \rightarrow D, D \rightarrow F$ )



# Course website

- ④ <http://cbb.sjtu.edu.cn/~ccwei//pub/courses/2014/plb/plb.php>
- ④ If you have any question, send me an email at: [ccwei@sjtu.edu.cn](mailto:ccwei@sjtu.edu.cn)