Evolutionary Genetics

LV 25600-01 | Lecture with exercises | 4KP

How to approach new software?





HS20 | UniBS | JCW

Google

nucleotide sequence length application linux

KaKs Calculator - kaks-calculator - A software package for ... code.google.com/p/kaks-calculator/wiki/KaKs_Calculator

If you use other Linux/Unix OS, you have to compile the program in the source codes ... Let us assume that the number of lengths between two DNA sequences ...

GenSkew - visualization of nucleotide skew in genome sequences genskew.csb.univie.ac.at/ - Cached

GenSkew is an **application** for computing and plotting **nucleotide** skew data. ... unlimited sequence length are available in the GenSkew Java Application. ... Operating Systems: Every OS that runs Java2 (tested on Linux, Windows, MacOS X) ...

Software/list - SEQwiki - SEQanswers

seqanswers.com/wiki/Software/list - Cached - Similar

21 Nov 2012 ... Linux. Alcovna, ALgorithms for COmparing and Visualizing Non Assembled data ... Arf, arf is a genetic analysis program for sequencing data. to obtain all sequences that overlaps to construct the a "full length" sequence. ... BFCounter, BFCounter is a program for counting k-mers in DNA sequence data.

DNA sequence Reverse and Complement Tool Free Bioinformatics ...

www.cellbiol.com/scripts/.../dna_sequence_reverse_complement.php - Cached With this tool you can reverse a DNA sequence, complement a DNA sequence or reverse ... A web application written in Python by Andrea Cabibbo ... web applications, open source, linux, strider, biology news, bioinformatics, biology software, ... Q

Freeware (from "free" and "software") is computer software that is available for use at no cost or for an optional fee (donation).



The **GNU General Public License** (GNU GPL or simply GPL) is the most widely used free software license.

Commercial software, or less commonly, **payware**, is computer software that is produced for sale or that serves commercial purposes.

The term **shareware** (also known as trialware or demoware) refers to proprietary software that is provided to users without payment on a trial basis and is often limited by any combination of functionality, availability or convenience.

Input sequence(s) (or upload file):	?
Upload File	
Run Sequence length:	

Input sequence(s) (or upload file):	?
>Seq1	
ATGGC ATGCG	
>Seq2	
TGACA TTNNN	
Upload File	
Run Sequence length: 11,11	



- What should you know before you download?
- What should you do before you start analysing your data?
- Where can you get help?
- What do you need for the publication / report?

What should you know **before** you download?

- requirements
- type (e.g. freeware)
- in- and output file format
- last update / update history
- literature search
- what language is used

What should you do **before** you start analysing your data?

- avoid installation errors
- understand installation warnings
- run tests simple tests
- explore limitations
- check in- and output format
- speed

Where can you get **help**?

- manual(s) > RFM
- help options (e.g. help button, -help, --h)
- README file(s)
- log file(s)
- user forum
- author(s)

What do you need for the **publication / report** ?

- literature reference or web link
- version
- parameters (better: command line)

Bioinformatics - Introduction

OPEN O ACCESS Freely available online

PLOS COMPUTATIONAL BIOLOGY

Editorial

Ten Simple Rules for Getting Help from Online Scientific Communities

Giovanni M. Dall'Olio¹*, Jacopo Marino², Michael Schubert³, Kevin L. Keys¹, Melanie I. Stefan⁴, Colin S. Gillespie⁵, Pierre Poulain^{6,7,8}, Khader Shameer^{9,10}, Robert Sugar³, Brandon M. Invergo¹, Lars J. Jensen¹¹, Jaume Bertranpetit¹, Hafid Laayouni¹

Netiquette

- Rule 1. Do Not Be Afraid to Ask a Question
- Rule 2. State the Question Clearly
- Rule 3. New to a Mailing List? Learn the Established Customs before Posting
- Rule 4. Do Not Ask What Has Already Been Answered
- Rule 5. Always Use a Good Title
- Rule 6. Do Your Homework before Posting
- Rule 7. Proofread your Post and Write in Correct English
- Rule 8. Be Courteous to Other Forum Members
- Rule 9. Remember That the Archive of Your Discussion Can Be Useful to Other People
- Rule 10. Give Back to the Community



entry

Euclid's algorithm for the greatest

An algorithm is a **step-by-step procedure** for solving a problem or accomplishing some end especially by a computer. Procedure that produces the answer to a question or the solution to a problem in **a finite number of steps**. An algorithm that produces a yes or no answer is called a decision procedure; one that leads to a solution is a computation procedure. A mathematical formula and the instructions in a computer program are examples of algorithms.

Dynamic Programming is a method for solving complex problems by breaking it down into a collection of simpler subproblems. In order to solve a given problem using a dynamic programming approach, we need to solve different parts of the problem (subproblems), then combine the solutions of the subproblems to reach an overall solution.





Pseudocode is an informal high-level description of the operating principle of a computer program or other algorithm. It uses the structural conventions of a programming language, but is **intended for human reading** rather than machine reading.



1.. If student's grade is greater than or equal to 20

Print "passed"

else

Print "failed"

