Python I

Arjan van der Velde

arjan.vandervelde@umassmed.edu

https://sites.google.com/site/gsbsbootstrappers/courses-workshops/python

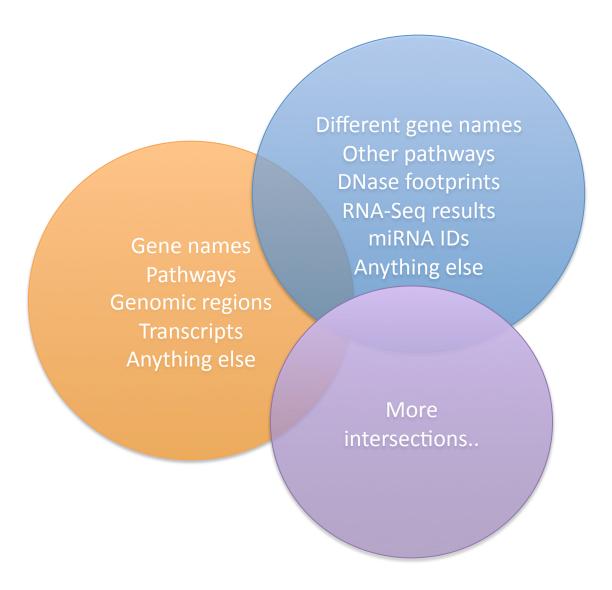
http://bioinfo.umassmed.edu/bootstrappers/bootstrappers-courses/python1/index.html



Sessions:

Tuesday, December 2nd, 5pm-7pm Thursday, December 4th, 5pm-7pm Tuesday, December 9th, 5pm-7pm Thursday, December 11th, 5pm-7pm **Amp III S6-102**

Basic Bioinformatics

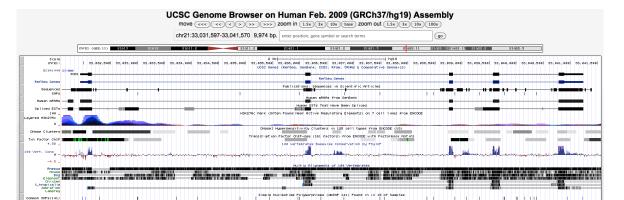


Examples

- Find pathways overlapping with upregulated genes
- Find promoters overlapping with ChIP-Seq reads
- Find upregulated genes that are novel (that don't overlap) relative to some other treatment
- Find primers that don't overlap with high GCcontent regions of the genome

Basic Bioinformatics

chr9	4974730 4974744	CEBPB 14.6061	+	6.11e-06	GATTGCGCCACTGC	-1 K562	CEBPB	ENCFF00	2CLR	0
chr13 /	25037026 We sec	25037040 mlends	CEBPB	14.8182 -	4.55e-06	TATTGCACCACAGC	-1	K562	CEBPB	ENCF
chr2	204064308 favors	204064322 ing th	CEBPB	14.6061 -	6.11e-06	GATTGCGCCACTGC	-1	K562	CEBPB	ENCF
chr9	129127385	129127399	CEBPB	14.5758 -	6.38e-06	GATTGCACCACTGC	-1	K562	CEBPB	ENCF
chr1	147043983 Thu	147043997	CEBPB	14.6061 +	6.11e-06	GATTGCGCCACTGC	-1	K562	CEBPB	ENCF
chr12ov	98781472	98781486	CEBPB	13.9394 -	1.34e-05	GATTGCAAAACTCC	-1	K562	CEBPB	ENCF
chr7	7157279 7157293	CEBPB 14.6061	+	6.11e-06	GATTGCGCCACTGC	-1 K562	CEBPB	ENCFF00	2CLR	0
chr19	17969839	17969853	CEBPB	14.0758 -	1.15e-05	GATTGCGCCACTGT	-1	K562	CEBPB	ENCF
chr2ing	238986128	238986142	CEBPB	14.3788 -	8.21e-06	TGTTGCATCATACT	32.2695	23804255	9	K562
chr4ing	72514304 /0	72514318	CEBPB	13.6212 +	1.82e-05	TGTTGCATCATTTC	-1	K562	CEBPB	ENCF
chr21lo	45973113	45973127	CEBPB	13.6818 -	1.73e-05	GATTGCTCAACCTC	-1	K562	CEBPB	ENCF
chr5esu	62157235	62157249	CEBPB	14.1818 -	1.03e-05	GCTTGCACCATCTC	-1	K562	CEBPB	ENCF
chr11el	129734538	129734552	CEBPB	13.3182 -	2.36e-05	GGTTGCATCATTGT	-1	K562	CEBPB	ENCF
chr4out	154492332 Nov	21544923466 EST	CEBPB	14.1667 +	1.05e-05	CGTTGCATAATCTT	-1	K562	CEBPB	ENCF
chr3sul	42644802	42644816	CEBPB	14.5758 +	6.38e-06	GATTGCACCACTGC	-1	K562	CEBPB	ENCF
chr11	27165617 oresult	27165631 va\$ svn	CEBPB	14.6061 +	6.11e-06	GATTGCGCCACTGC	-1	K562	CEBPB	ENCF
chr12	22628393 oresult	22628407 vas pvt	CEBPB	14.5455 -	6.62e-06	TATTGCACCATACA	-1	K562	CEBPB	ENCF
chr2	171532292 thon 2	171532306 onda 2	CEBPB	614.5758d+fault.	6.38e-06014, 15	GATTGCACCACTGC	-1	K562	CEBPB	ENCF
chr6	13491393800 4.2	134913952 Tnc. h	CEBPB 57	713.4242	2.17e-05	TATTGCATAATAAA	-1	K562	CEBPB	ENCF
							-			





Objectives

- Learn the logical structure of programming languages
- Open, analyze and close large data files
- By the end, write simple programs to calculate statistics, compare large data files to each other and extract relevant data
- Know how to apply programming to simple large scale tasks

Python is a great place to start

- The command are like natural English
- Many modules exist that enable common tasks, such as statistics, graphing, etc.
- Python is an open language and a large community exists that supports and continuously improves the language
- Python is a good language for text processing and therefore for many Bioinformatics tasks

Course Layout

Session I

- Intro
- What is programming?
- Introduction iPython Notebooks
- Variables and data types, simple conversions
- Expressions and conditionals

Session II

- Lists and dictionaries
- Simple iteration
- Splitting and sorting
- Functions

Course Layout

Session III

- Scripting
- Reading and writing files
- How are (text) files represented?
- Delimiters, column based text data
- Data transformations

Session IV

- Sample problems
- Using existing code (modules)
- Loading modules
- Read a genome?
- How about sequence data?

Course Layout

Session V

Learning by doing

- Programming is very practical
- It's a craft
- Short explanations (5-10 min) followed by a set of instructive exercises (10-30 min)
- 3-5 blocks each session

What is Python?

- Python is a programming language that came about in the 90s and has become very popular over the past ten years
- There is a solid community and there are many online resources
- Currently the are two main branches of Python, version 2 and version 3. Python 3 is not fully backward compatible and 2.7 is still the most widely used version
- Python is an interpreted language; programs written in the Python language are run by the Python interpreter (called python)
- Python commands can be typed directly into the interpreter but are usually saved in a file (a script) that is then read and executed by python.
- In the first two sessions we will be using iPython Notebook to interact with python

Getting help

- Python comes with a built-in help system, called pydoc, as well full documentation online.
- https://docs.python.org/2.7/reference/
- https://docs.python.org/2.7/library/

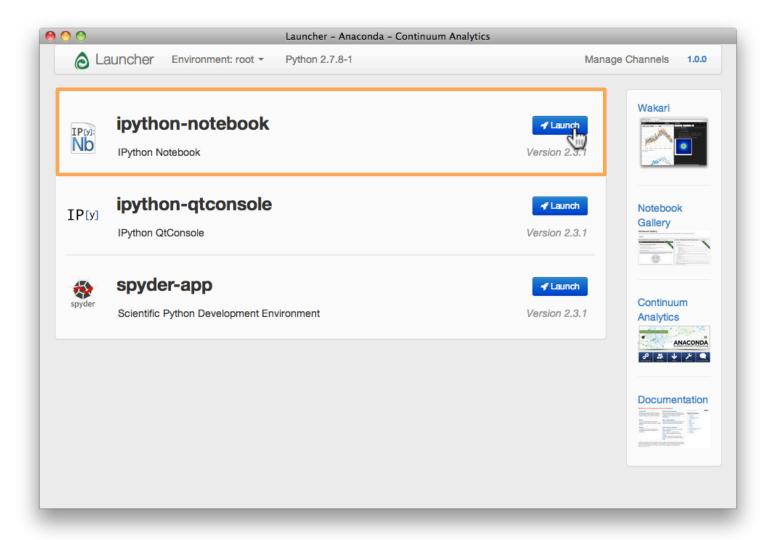
Troubleshooting errors

- In programming, most things don't work the first time
- By breaking a complex statement into parts, you can often see where the programming language didn't understand you
- Example: 2 / (4 + 6) * 4 should give 0.8 but it doesn't

```
4 + 6
```

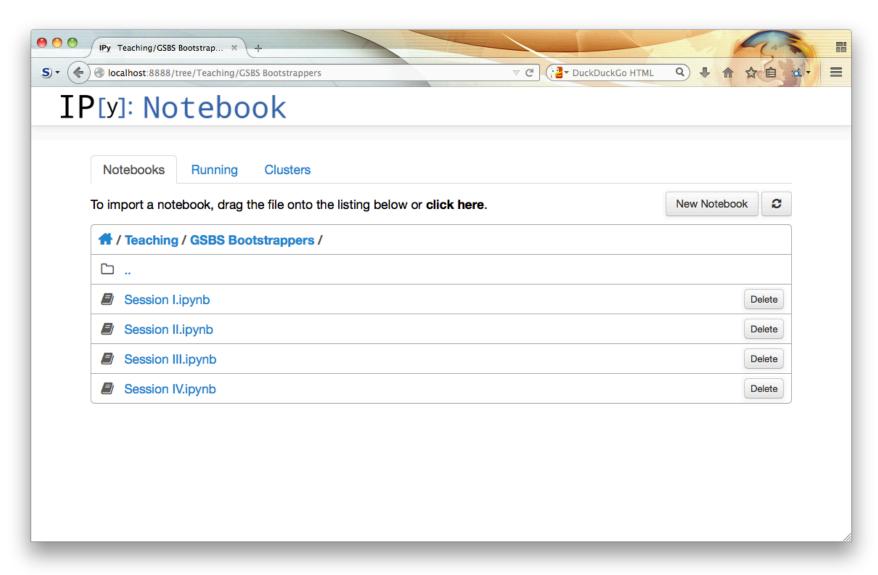
$$0.2*4$$

Let's start





iPython Notebook



Exercises 1

 The goal of these exercises is to get familiar with the Python interpreter and with iPython notebook

Exercises 1 Explanation

- Characters with quotes around them are strings
- Adding two strings together concatenates them
- Numbers without quotes are integers
- Adding two integers together returns the sum of the two integers
- Adding an integer and a string together is not allowed because python doesn't know which type of addition is meant.
- Strings and integers each have certain operations that can be done on them whose results can be built into more complex expressions
- Trying to do math operations on a string or string operations on numbers is an extremely common bug in bioinformatics programs.

Information storage

- All information entered into python has a data type or 'type'
- Simplest data types: str, int, and float (strings, integers, and decimals)
- Examples:
- 'hello' (string, known to python as str)
- 52 (integer, known to python as *int*)
- 35.4 (decimal, known to python as *float*)
- Python automatically guesses each information type and has certain operations it can do on it.
- Information is stored in variables using an "=" sign. From then on, when you type the variable, python retrieves the value.

Things you can do with int, str, and float

Integers:

 Add, subtract, multiply, divide, take exponents, compute remainders, sort numbers by size, anything mathematical, results will always be integers (explains division problem from Exercises 1)

• Floats:

Same as integers, but results will be decimals (python recognizes the decimal point)

Strings:

- In programming, a string is a series of letters. A sentence in English would be an example (python recognizes the quotes).
- Strings can concatenate two strings into a bigger one (like DNA), grab parts of strings, insert extra letters on the end of strings, change parts of a string into something else (like translating cDNA), sort things alphabetically, reverse strings, and much more

Try Exercises 2

Exercises 2 conclusions

- Data types can sometimes be converted between each other, allowing formerly impossible operations to be performed
- When math is done involving both a decimal and an integer, the result will be a decimal
- A variable name can't start with a number or be anything that already has an intrinsic meaning to python (like an '=' sign, the addition sign, the quote symbol, or anything else that python already understands)
- Variables that have been assigned to data take on the properties of the data they have in them (addition of two variables that have numbers assigned is different from addition of two variables holding strings)
- Variables can be reassigned to different values and data types, but can't be used without being initially assigned to a value. Be careful not to confuse strings and variables!

Logic

- Python knows <, >, ==, !=, and other tests
- Strings and numbers can be compared and strings can even be compared to numbers
- Logic operators can be combined with 'and' and with 'or'
- You can even test whether some sequence is found within another sequence.

Conditionals

- Test whether some expression evaluates to True
- Conditional statement starts with 'if' followed by the thing to test followed by a ':' symbol
- Execute conditional commands only when the thing to test is *True*
- Conditional commands are indented under the conditional statement
- First unindented line after the ':' is where normal code resumes (that happens regardless of the conditional)
- If the conditional is not *True*, an '*else*' statement can have its own commands that execute