

Introduction To Python Programming

II: How to Use It

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Today:

- **Use a python module**
 - Example: Generate random values
- **Install a Python package as a user**
 - Often useful when installing software
- **Exercise: Read a file, filter and output another file**
 - Reading and writing files
 - splitting lines and finding bits of information
- **Create a Slurm job and submit from a Python script**
 - Create text file from template ("here" document)
 - run system commands from inside a script

Use a Python module

We will use the random module

```
import random          # import the random module
random.randint(0,10)   # a random integer in [0, 10]
2

import random as rn   # give module a shorter name
rn.randint(0,10)
9

from random import randint # import a module function
randint(0,10)            # directly
6

from random import *      # import all functions - dangerous
```

Make your own Python module

- Modules are just Python files
 - You can organize and import your own code

main.py

```
# use mymodule

import mymodule as mm
print(mm.double(2))
```

mymodule.py

```
# mymodule example

def double(x):
    return x*2
```

Use modules to:

- Organize your program

Split up a large program into modules with related code

Makes organization and debugging easier

- Reuse useful code

Example: data analysis and graphing code

Build a library of useful tools for interactive analysis of your data sets

Python packages

- Python Packages: a directory with
 - One or more module files, and
 - An `__init__.py` file
 - It can be empty
 - It's run when you load the package

```
packagename/
  ├── __init__.py
  ├── module1.py
  └── module2.py
    ...
    . . .
```

You don't often write packages yourself

- Useful for large projects
- When you want to share a package in public

Install a Python package

Problem: You want to install a program in Deigo.

The instructions tell you to run:

`pip install <some package>`

or you download it, unpack, then run:

`python setup.py install`

But **this fails** with an error about not having permission

What do you do?

Install a Python package

1. Load our Python module

Load python 3:

```
$ module load python/3.7.3
```

Use 'pip3' and 'python3'

Use our python modules,
never the system Python

Install a Python package

1. Load our Python module

Load python 3:

```
$ module load python/3.7.3
```

Use 'pip3' and 'python3'

Use our python modules,
never the system Python

Packages are at PyPi:

<https://pypi.org/>

Over 300 000 (!) projects

Anything you need may already
be available there

Install a Python package in your home

2. Install locally using the '--user' option

for pip3:

```
$ pip3 install --user <package>
```

for setup.py:

```
$ python3 setup.py --user
```

This installs the package into your home.

Programs go to:

~/.local/bin

packages go to:

~/.local/lib/python3.7/site-packages



this depends on your Python version

Install a Python package in your home

2. Install locally using the '--user' option

Example:

```
$ pip3 install --user cli-weather
```

```
$ ~/.local/bin/cli-weather city Naha -f
```

This installs the package into your home.

Programs go to:

~/.local/bin

packages go to:

~/.local/lib/python3.7/site-packages



this depends on your Python version

Install a Python package in /apps

You can choose the installation directory with PYTHONUSERBASE:

```
$ export PYTHONUSERBASE="/apps/unit/UnitU"  
$ pip3 install --user <package>
```

Programs go to:

/apps/unit/UnitU/bin

packages go to:

/apps/unit/UnitU/lib/python3.7/site-packages/

Remember this path



Install a Python package in /apps

3. Tell Python where the local library and binary is

Add the path to site-packages to PYTHONPATH:

```
export PYTHONPATH="/apps/unit/UnitU/lib/python3.7/site-packages:$PYTHONPATH"
```

Add the path to bin to PATH

```
export PATH="/apps/UnitU/bin:$PATH"
```

Install a Python package in /apps

3. Tell Python where the local library is

Add the path to site-packages to PYTHONPATH:

```
export PYTHONPATH="/apps/unit/UnitU/lib/python3.7/site-packages:$PYTHONPATH"
```

set PYTHONPATH to

our new path

separate paths
with ':'

followed by whatever is
already in PYTHONPATH

export makes PYTHONPATH available to the entire
shell environment

Exercise:

Convert a file from one format to another

We will convert a FASTA file from one format to another

- "Fasta" is a bioinformatics sequence file format
- This kind of problem shows up in any field; the specific file format here is not important.

Exercise:

Convert a file from one format to another

We will convert a FASTA file from one format to another

- "Fasta" is a bioinformatics sequence file format
- This kind of problem shows up in any field; the specific file format here is not important.

Copy the session data files:

```
cp -r /apps/share/training/Python/session2 .
```

Exercise:

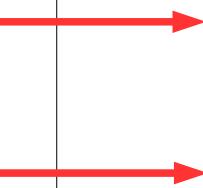
Convert a file from one format to another

Input file:

```
>NODE_130_length_9368_cov_2.522454  
GCGCCGTTTTCATA...  
>NODE_357_length_6506_cov_2.493338  
GCCTCACCTGTGGAA...
```

Output file:

```
>sequence_number_1_[cov=2.52]  
GCGCCGTTTTCATA...  
>sequence_number_2_[cov=2.49]  
GCCTCACCTGTGGAA...
```



the gene data is just copied right over

Copy the session data files:

```
cp -r /apps/share/training/Python/session2 .
```

Exercise:

Convert a file from one format to another

Input file:

```
>NODE_130_length_9368_cov_2.522454
GCGCCGTTTTCATA...
>NODE_357_length_6506_cov_2.493338
GCCTCACCTGTGGAA...
```

begins with '>'

>NODE_130_length_9368_cov_2.522454

covariance

stuff separated by '_'

Output file:

```
>sequence_number_1_[cov=2.52]
GCGCCGTTTTCATA...
>sequence_number_2_[cov=2.49]
GCCTCACCTGTGGAA...
```

same covariance,
rounded to 2 decimals

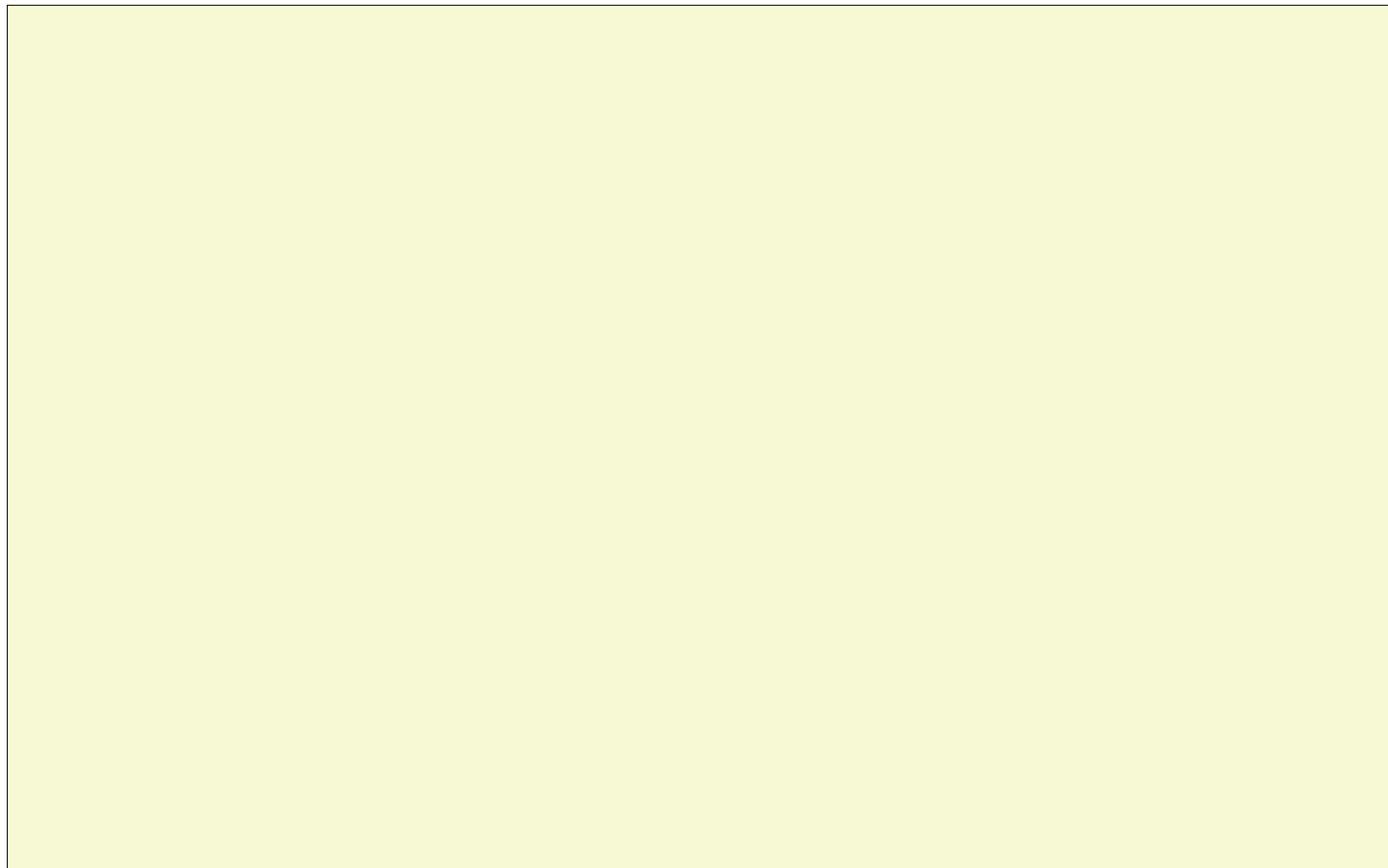
>sequence_number_1_[cov=2.52]

fixed text

counter increases
by one

Exercise:

Convert a file from one format to another



← Our program

Exercise:

Convert a file from one format to another

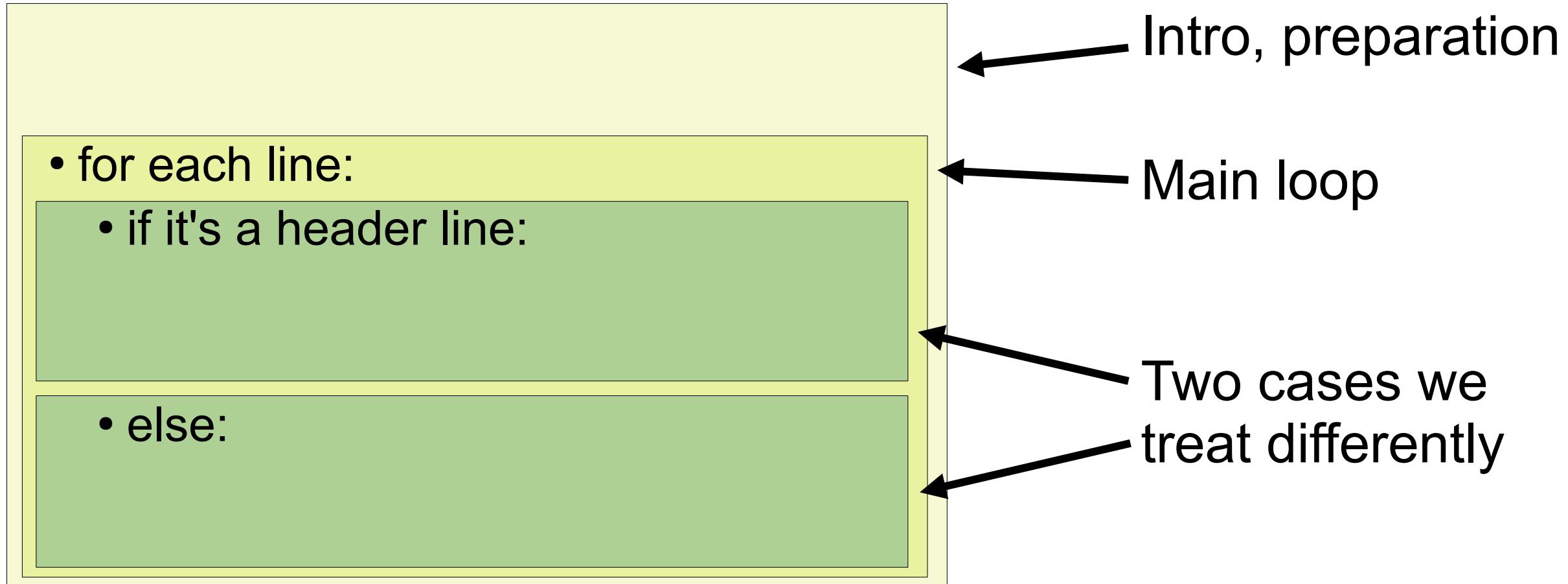
- for each line:

Intro, preparation

Main loop

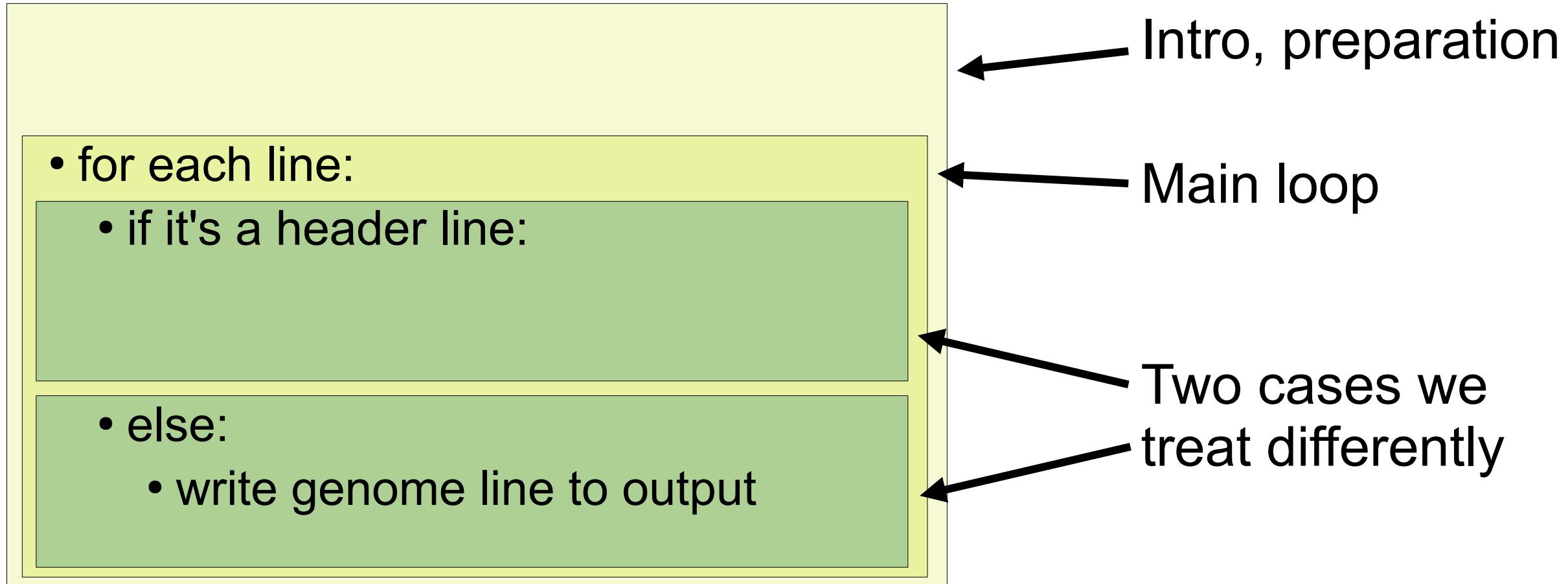
Exercise:

Convert a file from one format to another



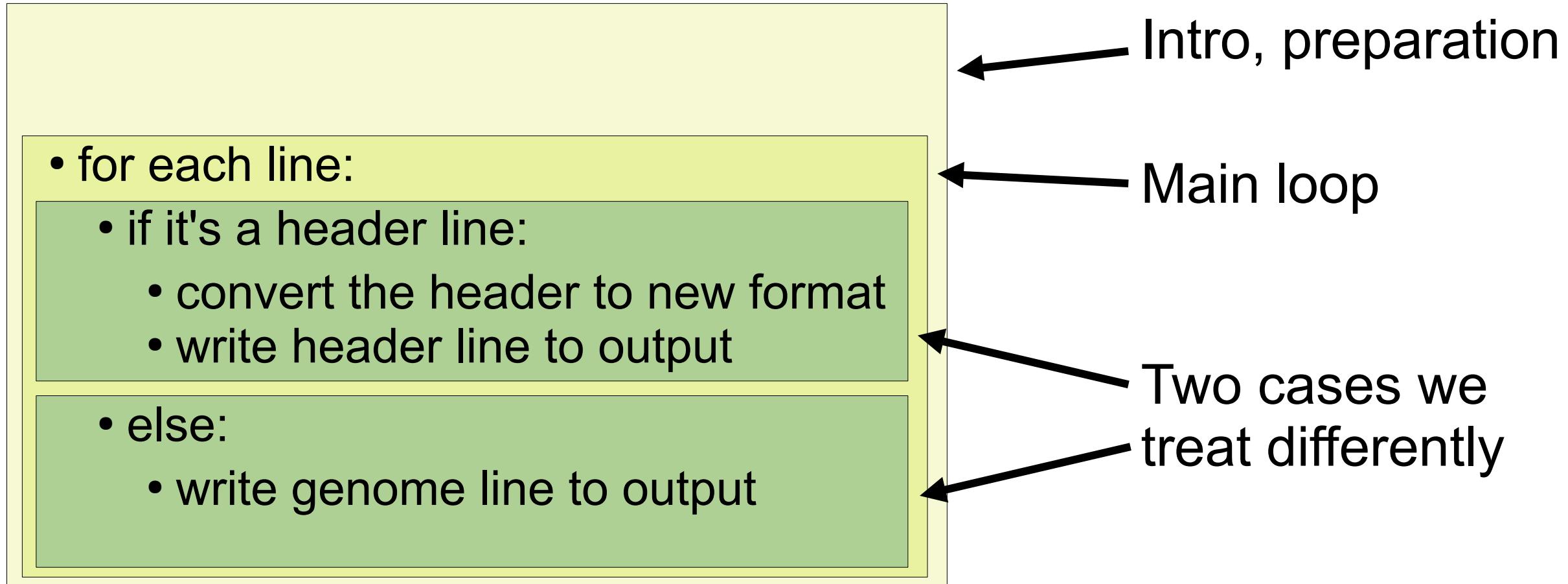
Exercise:

Convert a file from one format to another



Exercise:

Convert a file from one format to another



Exercise:

Convert a file from one format to another

- get input and output file names
- open input and output files

- for each line:

- if it's a header line:
 - convert the header to new format
 - write header line to output

- else:
 - write genome line to output

Intro, preparation

Main loop

Two cases we
treat differently

Exercise:

Convert a file from one format to another

- get input and output file names
- open input and output files
- for each line:
 - if it's a header line:
 - convert the header to new format
 - write header line to output
 - else:
 - write genome line to output

Questions: How do we ...

- convert the header?
- open files?
- read and write from files?
- get our filenames?

begins with '>' covariance same covariance, rounded to 2 decimals

>NODE_130_length_9368_cov_2.522454

stuff separated by '_' fixed text >sequence_number_1_[cov=2.52] index increases by one

Convert the header: write a function

- take as input an index value "index", and input header text line "line"
- output a new header line (using "return")
- see `string.split()` - split line into a list
 - `split()` splits a line between spaces
 - `split('_')` splits a line between '_'

```
def myfunction(a,b,c):  
    ...  
    return result  
  
"string: {} bipp".format(value)
```

begins with '>'

covariance

>NODE_130_length_9368_cov_2.522454

stuff separated by '_'

same covariance,
rounded to 2 decimals

>sequence_number_1_[cov=2.52]

fixed text

index increases
by one

Convert the header: write a function

- take as input an index value "index", and input header text line "line"

- output a new header line (using "return")
- see `string.split()` - split line into a list
 - `split()` splits a line between spaces
 - `split('_')` splits a line between '_'

```
def convert_header(index, line):  
  
    split_line = line.split('_')  
    cov = float(split_line[-1])  
    return '>sequence_number_{}_[cov={:.2f}]\n'.format(index, cov)
```

Exercise:

Convert a file from one format to another

- Opening files: use the "with" syntax

```
with open('filename', 'r') as f:          # open filename for reading
    <do stuff with filehandle f>

# open two files at once:

with open('file1', 'r') as fin, open('file2', 'w') as fout:
    <read from fin, write to fout>
```

Exercise:

Convert a file from one format to another

- read and write from files:

```
with open('filename', 'r') as fin:          # open filename for reading
    all_lines = fin.readlines()                # read all lines in file

with open('filename', 'r') as fin:          # open filename for reading
    for line in fin:
        <do something with each line>

with open('filename', 'w') as fout:          # write a line or a list
    fout.writelines(lines)                  # of lines
```

Exercise:

Convert a file from one format to another

Our main loop:

```
infile = 'inputfile.fa'  
outfile = 'output.txt'  
  
with open(infile, 'r') as fin, open(outfile, 'w') as fout:  
  
    for line in fin:
```

Exercise:

Convert a file from one format to another

Our main loop:

```
infile = 'inputfile.fa'
outfile = 'output.txt'

with open(infile, 'r') as fin, open(outfile, 'w') as fout:

    for line in fin:
        if line[0] == '>':

            else:
```

Exercise:

Convert a file from one format to another

Our main loop:

```
infile = 'inputfile.fa'
outfile = 'output.txt'

with open(infile, 'r') as fin, open(outfile, 'w') as fout:

    for line in fin:
        if line[0] == '>':

            else:
                fout.writelines(line)
```

Exercise:

Convert a file from one format to another

Our main loop:

```
infile = 'inputfile.fa'
outfile = 'output.txt'

with open(infile, 'r') as fin, open(outfile, 'w') as fout:

    for line in fin:
        if line[0] == '>':
            outline = convert_header(index, line)
            fout.writelines(outline)
        else:
            fout.writelines(line)
```

Exercise:

Convert a file from one format to another

Our main loop:

```
infile = 'inputfile.fa'
outfile = 'output.txt'
index = 1

with open(infile, 'r') as fin, open(outfile, 'w') as fout:

    for line in fin:
        if line[0] == '>':
            outline = convert_header(index, line)
            fout.writelines(outline)
            index += 1
        else:
            fout.writelines(line)
```

Exercise:

Convert a file from one format to another

Get the input and output files: use the 'sys' module:

```
import sys

infile = sys.argv[1]
outfile = sys.argv[2]
```

Run like:

```
$ python3 convfasta.py inputfile.fa output.txt
```

Exercise:

Convert a file from one format to another

Make our program a "real" program

1. Tell linux that python3 knows what to do with it:

```
#!/bin/env python3
import sys
...
```

Run like:

```
$ ./convfasta.py input.fa out.txt
```

2. Make it executable:

```
$ chmod +x convfasta.py
```

Generate and run a Slurm script from Python

Take a look at this Slurm script:

```
#!/bin/bash

#SBATCH -t 1:00
#SBATCH --mem=2G

tail mylittlefile.txt
```

What if we wanted to choose the input file when we submit?

Let's write a Python script that:

- 1) gets the filename
- 2) creates a multi-line string with this Slurm script and the filename filled in
- 3) saves it to a file
- 4) Runs it with sbatch

we already know how to do these!

Generate and run a Slurm script from Python

Remember **multi-line strings** and format substitutions:

```
s='''#!/bin/bash

#SBATCH -t 1:00
#SBATCH --mem=2G

tail {}'''

s.format(...)
```

Run a command:

```
import subprocess as sb

# just run a command
sb.run(['sbatch', 'myscript.slurm'])

# run command, get the output
ret=sb.run(..., stdout=sb.PIPE,
           universal_newlines=True)

# output as a list
ret.stdout.split()
```

The final script

```
import sys
import subprocess as sb
sfile = sys.argv[1]                      # Get first argument

script=''#!/bin/bash
#SBATCH -t 10:00
#SBATCH --mem=2G
tail {}
'''

with open('myscript.slurm', 'w') as f:
    f.writelines(script.format(sfile))      # substitute and write

sb.run(['sbatch', 'myscript.slurm'])       # start job
```

The final script - get the job ID

```
import sys
import subprocess as sb
sfile = sys.argv[1]                      # Get first argument

script=''#!/bin/bash
#SBATCH -t 10:00
#SBATCH --mem=2G
tail {}
'''

with open('myscript.slurm', 'w') as f:
    f.writelines(script.format(sfile))    # substitute and write

ret = sb.run(['sbatch', 'myscript.slurm'],
            stdout=sb.PIPE, universal_newlines=True)
jobid = ret.stdout.split()[3]
print(jobid)
```

Generate and run a Slurm script from Python

WARNING!!!!

Don't use this to submit lots and lots of jobs!!!!

you can break Slurm for everyone
and we may need to kill your jobs if you do

Always use Slurm Array jobs for this

Next Session

Let's get Scientific!

- Numpy and Scipy
 - The workhorses for all scientific Python programming
- Matplotlib
 - Plot your data