

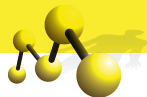
File Parsing

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Programming for Scientists

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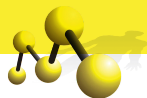


What's a File?

A sequence of bytes.

(and meta-data).

File Format Examples: FASTA format



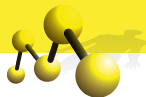
> qqq

```
ACTTTGTTATATATACTATCTGTATTTTC
CTGGGTGAGAGAGTGGTTGAGAGGGGGAA
CCCCCAACCACATTTCCCCACACCCCCTG
ACTTTCCTATATGTCCATTTTTTATAATC
```

> ppp

```
TTTTTGGTATCTATTTTCCACTCATTCTTTAT
TACCCCAGTCATCACAAAACACACACAACC
ATTATCTCTAATATATAATTTTACCTTT
```

Parsing FASTA

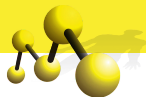


```
sequences = []
curseq = ''
for line in file('input.fsa'):
    if line[0] == '>':
        sequences.append(curseq)
    else:
        curseq += line.strip()
sequences.append(curseq)
```

File Format Examples (II): GenBank



LOCUS SCU49845 5028 bp DNA PLN 21-JUN-1999
DEFINITION Saccharomyces cerevisiae TCP1-beta gene, partial cds, and A
(AXL2) and Rev7p (REV7) genes, complete cds.
ACCESSION U49845
VERSION U49845.1 GI:1293613
KEYWORDS .
SOURCE Saccharomyces cerevisiae (baker's yeast)
ORGANISM Saccharomyces cerevisiae
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
...
ORIGIN
1 gatcctccat atacaacggt atctccacct cagggtttaga tctcaacaac ggaaccattg
61 ccgacatgag acagttaggt atcgtcgaga gttacaagct aaaacgagca gtagtcagct
121 ctgcatctga agccgctgaa gttctactaa ggggtggataa catcatccgt gcaagaccaa
181 gaaccgccaa tagacaacat atgtaacata tttaggatat acctcgaaaa taataaacgg
241 ccacactctc attattataa ttagaacaac aacgcacaaa ttatccacta tataattcaa

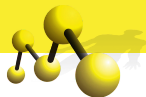


ASCII

- 65: A
- 66: B
- ...
- 48: 0
- 49: 1
- ...
- ...

127 code points taken.

Two File Format Classes

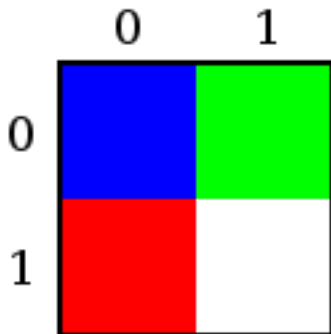
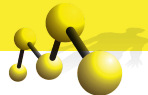


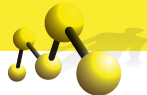
- Text files
- Non-text files (binary files)

Size	Hex Value	Value	Meaning
2	42 4D	"BM"	Magic Number (66, 77)
4	46 00 00 00	70 Bytes	Size of Bitmap
2	00 00	Unused	Application Specific
2	00 00	Unused	Application Specific
4	36 00 00 00	54 bytes	The offset of data.
4	28 00 00 00	40 bytes	Size of header.
4	02 00 00 00	2 pixels	The width in pixels
4	02 00 00 00	2 pixels	The height in pixels
2	01 00	1 plane	Number of color planes.
2	18 00	24 bits	The bits/pixel.
4	00 00 00 00	0	No compression used
4	10 00 00 00	16 bytes	The size of the raw BMP data
4	13 0B 00 00	2,835 pixels/m	The horizontal resolution
4	13 0B 00 00	2,835 pixels/m	The vertical resolution
4	00 00 00 00	0	Number of colors in the palette
4	00 00 00 00	0	Means all colors are important

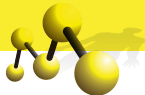
Size	Hex Value	Value	Meaning
3	00 00 FF	0 0 255	Red, Pixel (0,1)
3	FF FF FF	255 255 255	White, Pixel (1,1)
2	00 00	0	Padding for 4 bytes/row
3	FF 00 00	255 0 0	Blue, Pixel (0,0)
3	00 FF 00	0 255 0	Green, Pixel (1,0)
2	00 00	0	Padding for 4 bytes/row

(Wikipedia)





When possible, prefer text formats.
They are simpler.



- Unix: LF (line feed)
- Windows: CRLF (carriage return, line feed)
- (Old Mac OS: CR)

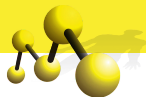
The extra carriage returns will often show up as ^M in Unix.
Some unix text files will show up as a single ultra-long line on Windows.

What About International Characters?

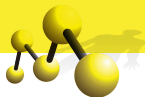


- Such as á or ç?
- Or μ ?
- Or Asian characters?
- Or —?

It's a mess!



- Traditional (latin-1,latin-9,latin-15,...)
- Unicode (16-bits, or 32-bits)
- UTF-8



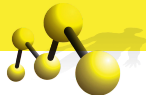
Unicode

Use 16 bits for (almost) all possible possible characters.

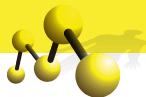
Use 32 bits for all possible characters.

Byte Order

If you have a 2 byte number, which byte do you write first?



Emerging standard (at some levels).



Emerging standard (at some levels).

You still see errors

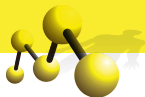
I saw this in an ATM receipt the other day
confirmação (which is confirmação in UTF-8, but printed out in
Latin-15)



- Text format is not only for text
- Your Python files are text files
- Anything can be represented as a text-file

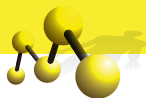


- Easier to parse
- Easier to debug
- Easier to transfer from one machine to the next
- Larger files (but you can compress them)



- Comma- or TAB- separated files are flexible
- Can be used to transfer from and to spreadsheet software
- Unfortunately, these are often not 100% well specified

Example: SAM Format



@HD VN:1.0 SO:unsorted

@SQ SN: LN:9732

@PG ID:bowtie2 PN:bowtie2 VN:2.0.0-beta7

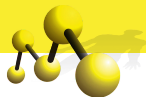
FC5:1101:1441:2131	4	*	0	0	*	*	0	0	TNTCTATT
--------------------	---	---	---	---	---	---	---	---	----------

FC5:1101:1371:2227	4	*	0	0	*	*	0	0	TTGTCTCT
--------------------	---	---	---	---	---	---	---	---	----------

FC5:1101:1650:2171	4	*	0	0	*	*	0	0	GTGTAATT
--------------------	---	---	---	---	---	---	---	---	----------

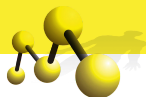
FC5:1101:1599:2183	4	*	0	0	*	*	0	0	GTATATAC
--------------------	---	---	---	---	---	---	---	---	----------

FC5:1101:1539:2201	4	*	0	0	*	*	0	0	CCTTCTCA
--------------------	---	---	---	---	---	---	---	---	----------



FC5:1101:1441:2131\t4\t*\t0\t0...

Parsing This File



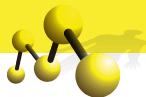
```
ifile = open('input.sam')
matched = []
nreads = 0
for line in ifile:
    if line[0] == '@':
        continue
    nreads += 1
    tokens = line.strip().split('\t')
    if tokens[1] != '4': # Four is *unmatched*
        matched.append( (tokens[0], tokens[2], int(tokens[3])

print 'Matched {0} reads out of {1} ({2:.2%})'.format(len(m
```



- Use UTF-8
- Prefer text-based formats (use generic compression on top)

Homework I



- Download FastQ file from course webpage
- Plot (for each base pair position) average quality & std. dev.
- Write a sequence trimmer

Homework I

