How to be a Computational Physicist after you've taken all the classes

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Caveats ...

- This is a personal list
- This is not comprehensive (backups? reg-ex? debugging? database management?)
- This is a non-essential list

1 How to present

- 2 How to learn
- 3 How to ask for help
- 4 How to manage
- 5 How to navigate
- 6 Putting it together

LaTeX



- LaTeX is NOT WYSIWYG
- LaTeX forces you to focus on content
- LaTeX is Portable
- LaTeX is Pretty!
- LaTeX is Free!

Table Table

Typesetting - Line breaks and hyphenations

`Oh. I've had such a curious dream!' said Alice, and she told her sister, as well as she could remember them. all these strange Adventures of hers that you have just been reading about; and when she had finished, her sister kissed her, and said, `It was a curious dream, dear, certainly: but now run in to your tea; it's getting late.' So Alice got up and ran off, thinking while she ran, as well she might, what a wonderful dream it had been.

Figure: Microsoft Office

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Figure: LATEX



Wang-Landau Density of States Calculation in Crowded Protein Environments

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Introduction

The Wang-Landau scheme, a recent development in the calculation of thermodynamic quantities, is used to sample the entire density of states efficiently for a biological system. We show examples of lattice proteins in crowded environments and the calculation of a two parameter density of states using a multi-pass algorithm.

Model

Wang-Landau

Wang-Landau sampling [4] is a generic algorithm to calculate 0, the density of state, of a given system. Once It has been calculated to the desired accuracy, all thermodynamic quantifies follow from it as S = hin 0. The WL algorithm attempts to find the Ω bardively, by generating a flat histogram in energy space. Given a moveset, the WL accestence rate from confourtion A to B is:

$$P(A \rightarrow B) = \min \left(1, \frac{\Omega(E_A)n_{B \rightarrow A}/n_B}{\Omega(E_B)n_{A \rightarrow B}/n_A}\right)$$

Where the factor $\left(\frac{2\pi a_{1}/2a_{2}}{a_{1}-a_{1}/2a_{2}}\right)$ accounts for movesets that are not reversible in one step [5] and $\Omega(E)$ is the currently measured density of states at energy E. As the process brantse, the density of states is updated as $\Omega(A) \to \Omega(A)$ where f is a modification. Inside that monotonically decreases during the simulation.

Lattice Peptides

We use a coarse grained model; proteins are represented as a bead sequence of amino acid residues and restricted to move on a lattice.

- . A length N: Number of amino acids.
- A conformation represented by an ordered sequence S_i ∈ {1,...,20}, i = 1,...N of amino acids where each amino acid type is indexed by an integer corresponding to the twenty natural amino acids found in proteins.
- A conformation c_i ∈ C_N denoting the location of each residue on the lattice.

Movesets

We use two movisets, pull movies [3] and bord-redridging [1]. The combination of these moves has the advertage of small energy changes per move, which allows WL to focus down on the rare low-energy configurations. The bond-redridging allows the system to sample the low-energy states in a more efficient manner.

Mean-Field Crowding

Crowding is implemented by modeling the physical process of excluded volume [2]. As the peptide folds we imagine the environment to favor compact conformations so we extend the physical volume of the protein by k lattice units (k = 1 in this study).



Figure 1: Examples of 2D and 3D lattice proteins. The excluded volume zone is shown schematically with transparent yellow spheres.

Multi-pass Wang-Landau

One it has been calculated for a suitable parameter (usually energy), the density of states can be reased to estimate other cbservables. Since a converged VU, algorithm will visit each state with equal probability, we can sample a different observable energy without having to worry about reaching all low-energy status. This approach is particularly well-studied if a correlation exists between *E* and the observable. Results are shown for $(12, R_{\rm ob})$, the o parameter density of states for energy and radius of gyration.



Figure 2: Specific Heat Capacity curves for a 3D lattice homopolymer with L = 3i at various crowing parameters. When the crowing reflect is absent there are two mething points, one for the collapse from an unfolded chain to motion globule, then to the native state. As the environment becomes more crowided, the first transition excernse less pronounced.



Figure 3: Specific Hast Capacity curves for a 2D lattice homophymer 1— is all various convelling parameters shown as a contour map. A horizontal taxos across the graph is the specific heat at constant II. The effect of higher crowder concentration (as seen in the 3D case) is also observed here. Having the complete density of states allows smooth curves to be shown at any temperature which utrither cabustion.

Future work

Recently, we have introduced realistic potential energy functions to bother capture the actual interactions between armino acid readures (4-bed and MJ matrix (not shown)). The peruitimate goal is to study protein aggregation; ufitmately we would like to calculate the full density of states of two or more lattice proteins and their biolism cooperatively in the oresence of enrowless.

References

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- [2] T. Hoppe and J. Yuan. Entropic flows, crowding effects, and stability of asymmetric proteins. Phys. Rev. E, 80(1):011404, July 2009.
- [3] N. Lesh, M. Mitzenmacher, and S. Whitesides. A complete and effective more set for simplified protein folding. In *Proceedings of the seventh annual International conference on Research in computational molecular biology*, pages 185–195, Berlin, Germany, 2003. ACM.
- [4] F. Wang and D. P. Landau. Determining the density of states for classical atstitutical models: A random welk algorithm to produce a flat histogram. *Physical Review E*, 64(5):355101, Oct. 2001.
- [5] T. Wast and D. P. Landau. Versatile approach to access the low temperative thermodynamics of lattice polymers and proteins. *Phys. Rev. Lett.*, 102(17):178101-4. May 2009.

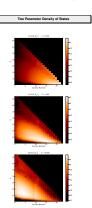


Figure 4: Probability $V(T, H) = \frac{0.16 \times 10^{-10} (mm^{-1} {\rm cm})^2}{100}$ as uncreased at 20 lattice handwards of the start of the

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Sample LaTeX code

```
Adapted from : 
http://amath.colorado.edu/documentation/LaTeX/basics/example.html
```

```
\documentclass[12pt]{article}
\title{Mv Sample \LaTeX{} Document}
\author{Travis Hoppe}
\begin{document}
\maketitle
                                  % automatic title!
This is (very) short primer to LaTeX
\section{Formulae: inline vs. displayed}
I insert an inline formula by surrounding it with a pair of
single \ symbols: what is x = 3 \times 5?
For a \emph{displayed} formula, use double-\$
before and after --- include no blank lines!
$$\mu^{\alpha+3} + (\alpha^{\beta}+\theta {\gamma}+\delta+\zeta)$$
Use the \emph{equation} environment to get numbered formulae, e.g.,
\begin{equation}
    v {i+1} = x {i}^{2n} - \sart{5}x {i-1}^{n} + \sart{x {i-2}^7} -1
\end{equation}
\begin{equation}
    \frac{\partial u}{\partial t} + \nabla^{4}u + \nabla^{2}u +
        frac12
                   | nab|a u|^{2} = c^{2}
\end{equation}
\end{document}
                           % End of document.
```

Sample LaTeX code: output

My Sample $\LaTeX\ensuremath{\mathsf{TE}}\ensuremath{\mathsf{X}}$ Document

Travis Hoppe

March 11, 2010

This is (very) short primer to LaTeX

1 Formulae; inline vs. displayed

I insert an inline formula by surrounding it with a pair of single \$ symbols; what is $x = 3 \times 5$? For a *displayed* formula, use double-\$ before and after — include no blank lines!

$$\mu^{\alpha+3} + (\alpha^{\beta} + \theta_{\gamma} + \delta + \zeta)$$

Use the equation environment to get numbered formulae, e.g.,

$$y_{i+1} = x_i^{2n} - \sqrt{5}x_{i-1}^n + \sqrt{x_{i-2}^7} - 1 \tag{1}$$

$$\frac{\partial u}{\partial t} + \nabla^4 u + \nabla^2 u + \frac{1}{2} |\nabla u|^2 = c^2 \tag{2}$$

Project Euler

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Often we are faced with learning a new programming language (or some new library from a known one).

- Textbooks: Will give a good detailed instruction to the language. Often too comprehensive - will detail many features irrelevant to you
- Verbal instruction (that's me!): Allows immediate feedback with questions. Less of a hands-on approach, the information is given but rarely self-analyzed
- Practice : Programing is as much of an art as it is a science by far best way to learn

Choosing good material is hard - after all you don't know the language in the first place!

Project Euler

Time and time again, you will encounter the same problems in a different setting.



- Project Euler is a great collection of problems that require three elements needed for physics, mathematics, critical thinking and tight coding practices.
- Once you solve a problem you are given access to a forum where you can see answers from other users. These answers span the gamut of languages from C++, Python, Ruby, Perl, Assembly, Scheme, Delphi, etc...
- Often your solution is not as clever as others use them to learn!

Stack Overflow

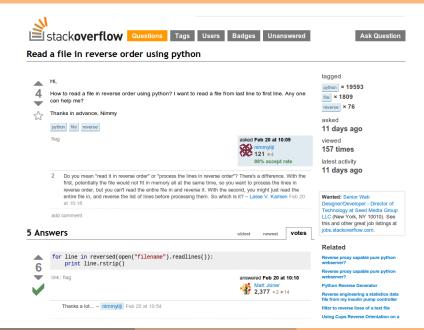
- The typical response to a dumb (not naive) question is to Google it (JFGI).
- There are times when Google fails this is often because you don't know how to ask the right question in the first place!
- Similar to looking up how to spell a word in a dictionary when you don't know how to spell it!

When Google fails and Wikipedia is not specific enough - turn to the most helpful programming message board created:



The site leverages the communication of a form, while encouraging participation through points. In short, it's the ultimate nerd video game.

Stack Overflow: How it works



BitBucket

Source Code Repository (BitBucket)



- Small projects : 100 lines of code
- Large projects : (Linux kernel 2.6.32 12m LOC) (Windows Server 2003 50m LOC)
- Impossible to manage with one central location for the code, graphics, UI, etc...
- Solution: Code repository, allow pieces to be checked out when needed

Unix

man cat just sounds funny

Learning to navigate across your system is akin to learning to use the mouse. Is it necessary?

Commands to know

- locate find files by name
- ssh OpenSSH SSH client (remote login program)
- scp secure copy (remote file copy program)
- grep print lines matching a pattern
- awk pattern scanning and text processing language
- man an interface to the on-line reference manuals
- history GNU History Library
- cat, head, tail concatenate files and print on the standard output
- chmod chown change file owner and group
- top display Linux tasks
- **ps, pkill** look up or signal processes based on name and other attributes

Putting it together

Lit-Py

Lit-Py Input

```
# {\Large Mandbrot Set} \\
# The Mandelbrot set $M$ is defined by a family of complex quadratic polynomials
$P_c:\mathbb C\to\mathbb C$ given by: $P_c: z\to z^2 + c$ where $c$ is a complex parameter.
For each c, one considers the behavior of the sequence (0, P, c(0), P, c(P, c(0))), P, c(P, c(P, c(0))).
ldots obtained by iterated function P_c(z) starting at critical point z = 0, which either escapes
to infinity or stays within a disk of some finite radius. The Mandelbrot set is defined as the set of all
points $c$ such that the above sequence does not escape to infinity.\\
# \textbf{Create a grid}
from pylab import *
X = linspace(-1.5, .8, 200)
Y = linspace(-1, 1, 200)
XG, YG = meshgrid(X, Y)
G = zeros(XG.shape)
# \textbf{Define the Mandelbrot function}
def MBset(c, z=0):
   for n in xrange(80):
       if abs(z)>2: break
       z = z * * 2 + c
    return n
# Test the function to see if it is working properly
print MBset(1 + .5J)
# \textbf{Compute Mandelbrot set}, note that XG and YG are the grid coordinates
for ix in ndindex(G.shape):
   G[ix] = MBset(XG[ix] + YG[ix]*1.J)
# \textbf{Plot the result}
imshow(G, extent=(-1.5, .8,-1,1), interpolation='nearest')
show()
```

examples/mandelbrot.py

Mandbrot Set

The Mandelbrot set M is defined by a family of complex quadratic polynomials $P_i: C \rightarrow C$ given by $P_i: z \rightarrow z^2 + cwhere is a complex parameter. For each <math>c$, one considers the behavior of the sequence $(0, P_i(0), P_i(P_i(0)), P_i(P_i(0))), \ldots)$ obtained by iterated function $P_i(z)$ starting at critical point z = 0, which either secapes to infinity or stays within a disk of some finite radius. The Mandelbrot set is defined as the set of all points c such that the above sequence does not escape to infinity.

Create a grid

from pylab import *
X = linspace(-1.5, .8, 200)
Y = linspace(-1, 1, 200)
XG,YG = meshgrid(X,Y)
G = zeros(XG.shape)

Define the Mandelbrot function

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def MBset(c, z=0):
    for n in xrange(80):
        if abs(z)>2: break
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    return n
```

Test the function to see if it is working properly

print MBset(1 + .5J)

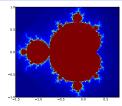
 $\gg 2$

Compute Mandelbrot set, note that XG and YG are the grid coordinates

```
for ix in ndindex(G.shape):
    G[ix] = MBset(XG[ix] + YG[ix]*1J)
```

Plot the result

```
imshow(G, extent=(-1.5, .8,-1,1), interpolation='nearest')
show()
```



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