## Problem Set 1

Submit your solution via NTULearn, with one Python file per problem. Problems labeled like this—(5\* marks)—are optional for undergraduates, and can be attempted for a bonus of 1/5 the indicated marks; for graduate students, these problems are compulsory and receive full marks. Your code should follow good programming style, with clear comments. Plots should be labeled clearly.

## 0. GAUSSIAN ELIMINATION

In this problem, you will write and test an implementation of the *Gaussian elimination* algorithm, which solves linear systems of equations of the form

$$A\vec{x} = \vec{b},\tag{0}$$

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where A is a square matrix, and  $\vec{x}$  and  $\vec{b}$  are vectors. The goal is to find  $\vec{x}$  given A and  $\vec{b}$ . See  $\langle \text{http://spike.spms.ntu.edu.sg/wiki/Gaussian_elimination} \rangle$  for a description of the algorithm. We will split the implementation into several pieces.

(a) (2 marks) Write a function to perform the row reduction phase of the algorithm:

def row_reduce(A, b, n):		
Inputs		
A	A 2D array specifying a square matrix $A$ .	
b	A 1D array specifying a vector $\vec{b}$ .	
n	The pivot row index (an integer).	

This function has no return values, and directly modifies the contents of the A and b arrays, in order to perform "row reduction". This means modifying A and b so that all the elements underneath A[n,n] become zero. The procedure is as follows:

For each 
$$m > n$$
, 
$$\begin{cases} \text{(i)} & b_m \to b_m - \left(\frac{A_{mn}}{A_{nn}}\right) b_n \\ \text{(ii)} & A_{mk} \to A_{mk} - \left(\frac{A_{mn}}{A_{nn}}\right) A_{nk}, \end{cases}$$

where k runs over all the columns of A.

You should test this code. For example, try the example given in the course notes:

This should print:

```
[[ 1. 2. 3.]
[ 0. -4. -7.]
[ 0. 2. -4.]]
[ 3. -5. -2.]
```

(b) (2 marks) Write a function to perform pivoting:

<pre>def pivot(A, b, n):</pre>		
Inputs		
A	A 2D array specifying a square matrix $A$ .	
b	A 1D array specifying a vector $\vec{b}$ .	
n	The pivot row index (an integer).	

This function has no return values, and directly modifies the contents of the A and b arrays. It performs "pivoting", which consists of the following steps:

- (i) Look through rows  $m \geq n$ , and find the row with the largest value of  $|A_{mn}|$ .
- (ii) If  $m \neq n$  was the row round in (i), then swap rows m and n in the matrix A; also, swap elements m and n in the vector  $\vec{b}$ .

Again, be sure to test your code for correctness.

(c) (3 marks) Write a function to perform Gaussian elimination:

def gauss_eliminate(A, b):		
Inputs		
A	A 2D array specifying a square matrix $A$ . Will not be altered.	
b	A 1D array specifying a vector $\vec{b}$ . Will not be altered.	
Return value		
х	A 1D array containing the solution to $A\vec{x} = \vec{b}$ .	

This should use the functions written in (a) and (b) as subroutines, as well as code for performing the back-substitution part of the Gaussian elimination algorithm.

Note that the gauss\_eliminate function is *not* supposed to alter the contents of the input arrays A and b, unlike the functions written in (a) and (b). Use copy appropriately to accomplish this.

(d) (3 marks) Write a function, gauss\_eliminate\_profile(), to measure and report the performance of the Gaussian elimination algorithm. This function should measure the time taken to run Gaussian elimination,  $t_N$ , versus the problem size N. You are free to choose an approprate range for N, such as  $10 \lesssim N \lesssim 500$ . Performance times can be measured using Python's time.perf\_counter function.

The function should then show a "log-log" plot of  $t_N$  versus N (e.g., using the plt.loglog function). Within the same figure, plot the performance graphs for two different solvers: (i) the gauss\_elimination solver you wrote in part (c), and (ii) the scipy.linalg.solve function, which is Scipy's own implementation of Gaussian elimination.

Furthermore, for each graph, compute and show the trend-line for the least-squares fit

$$\log(t) \approx p \log(N) + q.$$

(To get cleaner results, you might want to fit to a subset of the data points; if so, indicate this choice clearly in code comments.) Make separate fits for the gauss\_eliminate and scipy.linalg.solve results. You can use polyfit for the linear fitting. Label the two graphs clearly, and show the fitted value of p in the trend-line labels.

Discuss in code comments: What is the significance of the values of p and q found?

- (e) (2 marks) Modify your Gaussian elimination code to detect the case where A is non-invertible, and generate an error using Python's raise statement. Hint: you should not do this by calculating det(A); why?
- (f) (4\* marks) Modify your Gaussian elimination code so that it handles the case where x is a 2D array, representing an  $N \times M$  matrix. The code should still work as usual if x is a 1D array representing a vector.

Then, write a function <code>gauss\_eliminate\_profile2()</code>, to profile the performance of matrix inversion, comparing your Gaussian elimination code to <code>scipy.linalg.inv</code>.

## 1. THE HARPER MODEL

The Harper model is a theoretical quantum system consisting of a particle existing along a discrete one-dimensional "chain". The chain consists of a set of discrete points labeled by n = 0, 1, ..., N - 1. A quantum state is described by a complex vector,

$$\psi = \begin{vmatrix} \psi_0 \\ \vdots \\ \psi_{N-1} \end{vmatrix}, \text{ where } \sum_{n=0}^{N-1} |\psi_n|^2 = 1.$$

For each n, the complex number  $\psi_n$  gives the "quantum wavefunction" at position n; the probability to observe the particle at that position is  $|\psi_n|^2$ .

An "energy state" is a quantum state that satisfies the eigenvalue problem

$$H\psi = E\psi$$
,

where  $E \in \mathbb{R}$  is the energy and H is an  $N \times N$  matrix called the "Hamiltonian". The Hamiltonian has the form

$$H = \begin{bmatrix} V_0 & 1 \\ 1 & V_1 & \ddots \\ & \ddots & \ddots & 1 \\ & & 1 & V_{N-1} \end{bmatrix}, \text{ where } V_n = W \cos(2\pi\alpha n + \phi).$$

The diagonal entries  $V_n$  represent the potential along the 1D chain. This potential varies sinusoidally, with amplitude W and phase parameters  $\alpha$  and  $\phi$ . If  $\alpha$  is irrational, the potential function is "aperiodic" (i.e., it does not repeat with n).

## (a) (2 marks) Write the following function:

	def harper_hamiltonian(N, W, phi, alpha):			
Inputs				
N	The Hamiltonian size (an integer).			
W	The amplitude of the potential function (a number).			
phi	The parameter $\phi$ in the potential function (a number).			
alpha	The parameter $\alpha$ in the potential function (a number).			
Return value				
Н	The Harper model's Hamiltonian matrix (a 2D array).			

(b) (2 marks) Write the following function:

<pre>def harper_levels_plot(N=199, W=2., phi=0.):</pre>		
Inputs		
N	The Hamiltonian size (an integer).	
W	The amplitude of the potential function (a number).	
phi	The parameter $\phi$ in the potential function (a number).	

This function should plot the "energy-level diagram" of E versus  $\alpha$ : i.e., for each value of  $\alpha$  on the horizontal axis, plot the discrete energies E (the eigenvalues of H) on the vertical.

(c) (2 marks) Write harper\_wavefunction\_demo(N=199, alpha=1.618034, phi=0.), which plots the probability density  $|\psi_n|^2$  for the ground state (the energy state with lowest energy), versus position n. Do this for several choices of W, in the range  $1.2 \lesssim W \lesssim 2.5$ . Label all plots clearly; you may use subplots for clarity.

Discuss in code comments: How does the behavior of  $|\psi_n|^2$  change with W?

(d) (2 marks) The "inverse participation ratio" (IPR) of a wavefunction is the quantity

$$IPR[\psi] = \sum_{n} |\psi_n|^4.$$

It is a crude measure of whether a wavefunction is "localized" (i.e., concentrated near a few points) or "extended" (i.e., spread out over the whole chain). Write a function  $harper_ipr_demo(N=199, alpha=1.618034, phi=0.)$ , which plots the mean IPR of the Harper model's energy states, versus W.

Discuss in code comments: Is the IPR large or small for localized states? Estimate the critical W where localization begins to occur.

(e) (6\* marks) We now consider the  $\phi$  parameter. Write a function harper\_phi\_demo(), which plots the energy levels versus  $\phi \in [-\pi, \pi]$ . Choose appropriate values for the other parameters, but let  $\alpha$  be irrational. You should find that most of the energies cluster into "bands". However, there exist some states lying outside the bands. In a separate subplot or figure, plot the probability density for one or more of these "out-of-band" states. In the energy level plot, use markers to indicate the states you are plotting.

Discuss in code comments: How do the in-band and out-of-band states differ? How does this phenomenon vary with  $\alpha$ ?