Homework 2: Lyman-break Galaxies in R Due October 2, 11:59 pm

Before reading the rest of this handout, please check these links, which are also posted under "Before Th 9/17" on Canvas, as background:

• Lyman data background video playlist:

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https://mediaspace.wisc.edu/playlist/dedicated/1_gu7iqxze
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• Lyman data background links:

http://pages.stat.wisc.edu/~jgillett/605/data/lyman

This homework is a first step in our search for an undiscovered, gravitationally lensed, high-redshift Lyman-break galaxy. It requires you to search through 100 spectra for those that look like cB58, the target Lyman-break galaxy. Later in the semester, we will use the tools that we develop in this homework, along with distributed computing tools, to search a much larger collection of spectra.

Over the course of this homework, you will

- become familiar with our first computing tools, including the VM/Linux, and a powerful editor, either emacs or vim. If you are tempted to solve HW2 without these tools, note that we will need them soon when we do remote large-scale computing, so please take this opportunity to learn them.
- practice creating, selecting and applying statistical methods to a research problem for which there isn't a known right answer and for which trial-and-error is required. This is preparation for the project phase of the course in which students do original work.
- become familiar with the Lyman-break galaxy data from the Sloan Digital Sky Survey (SDSS), in which we will do a large search later, doing a small-scale search in this homework.

1 Overview

Before we dive in, here are some high-level instructions. Details are provided in the sections below.

- 1. Our high-level goal is to find the 3 SDSS spectra closest to cB58 by a distance measure of your choice. We will work together to develop measures.
- 2. When we show our best spectra to Christy (the astronomer), here are some ways she will decide whether it is a Lyman-break galaxy. Look at http://pages.stat.wisc.edu/~jgillett/605/2/exampleMatch.pdf, which shows the low-noise target spectrum of cB58 in red and the SDSS noisy spectrum of the same cB58 in blue. Note the presense of:
 - the Lyman-alpha absorption trough, around index 800
 - the Lyman-alpha emission peak just right of the trough, around 900
 - $\bullet\,$ metal abosorption lines right of the peak, in red (and blue) from about 950 to $2600\,$
- 3. We will consider integer redshifts (shift 1, shift 2, shift 3, ...) of wavelengths in attempting to match cB58.
- 4. For each of the 3 closest spectra, make a graph showing that spectrum aligned with cB58. If graphing only one spectrum, we would use x=wavelength, y=flux. However, in light of red shifting and the need to graph two aligned spectra at a time, we can abandon wavelength and instead use x = shifted wavelength, or, more simply, just x = index (that is, 1 to the length of the flux vector). Each of your graphs should show the alignment, or lack of alignment, as well as exampleMatch.pdf. Different graph designs are ok if they make deciding "Is this a nice alignment?" easy.
- 5. Report on what you tried, what worked, and what didn't.

2 Lyman-breaks data: Details

Detailed instructions on this assignment follow. Please make sure you have read and understood the links at the beginning of this handout before proceeding.

First, we need to get the data.

- 1. On your VM, create a new directory, ~/Desktop/2.
- 2. From the command line (either the terminal or an emacs shell buffer):
 - (a) Get to the right directory: cd ~/Desktop/2

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- (b) Download the spectra data (21 MB): wget http://www.stat.wisc.edu/~jgillett/605/2/data.tar
- (c) Unpack the data by running: tar xvf data.tar (tar="tape archive", xvf="extract verbose the file ..."). That will extract 100 .fits files in a new subdirectory, data. Each file contains a noisy SDSS spectrum of an astronomical object.
- 3. Download cB58_Lyman_break.fit (34 KB), the spectrum of the known Lymanbreak galaxy, cB58:

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wget http://www.stat.wisc.edu/~jgillett/605/2/cB58_Lyman_break.fit
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It is a high-quality spectrum from a bigger telescope that is relatively noise-free.

- 4. Start R, either from the terminal or by starting an R buffer in emacs. To download software for reading the spectra, run install.packages("FITSio") (reply "y" to the personal library questions; select any of the CRAN mirrors).
- 5. Run require("FITSio") to load FITSio into the current R session. (Or use library("FITSio") if you prefer that interface for loading a package.) Now you can use the function readFrameFromFITS() to read a .fits file (https://en.wikipedia.org/wiki/FITS) containing a spectrum into a data frame. It has these columns:
 - (a) flux is light intensity at a given wavelength. I think it is theoretically nonnegative, but with noise it can be negative.
 - (b) loglam is log(x=wavelength, base=10), so wavelength = 10^{loglam}.
 - (c) ivar is 1/sigma_i^2, where sigma_i^2 is an estimated variance of the i-th flux.
 - (d) and_mask is 0 for a good observation. We could exclude data with nonzero and_mask. (Each spectrum is constructed from several observations. Masks flag problems in the data. and_mask indicates whether there is bad data at this (wavelength, flux) pixel in all observations.)
 - (e) **or_mask** is 0 for a good observation. We can ignore it. (It says there is bad data at this pixel in at least one observation.)
 - (f) wdisp is the resolution of the spectrograph at that wavelength. We can ignore it.
 - (g) sky is the spectrum of the sky, mostly already subtracted out from flux. We can ignore it. (Some search algorithms may need to mask out +/-5 pixels around 5577 Angstroms to avoid spurious matches to a strong emission line due to Earth's atmosphere.)
 - (h) model is SDSS's hypothesis of the true spectrum, shifted to the redshift of the object. We can ignore it. (Well, one possible search criterion is to seek each spectrum that is closer to cB58 than to the spectrum's model.)

Note: readFrameFromFITS() can also read cB58_Lyman_break.fit, a .fit file that contains only two columns, FLUX and LOGLAM.

To get started, run readFrameFromFITS("cB58_Lyman_break.fit") to see the lownoise target data for cB58. Then, to see the noisy SDSS data for cB58, run readFrameFromFITS("data/spec-1353-53083-0579.fits").

3 Developing distance measures

To explore and develop ideas for a distance measure, you must participate in the Canvas Discussion at https://canvas.wisc.edu/courses/220494/discussion_topics/761673. Make at least three posts between 9/22/20 and 10/1/20:

- one post proposing or developing an idea
- one asking a helpful question about an idea
- one giving a constructive criticism of an idea

We want to simulate, via Canvas, the brainstorming that goes on in a research-anddevelopment meeting of a motivated team.

We are exploring! Some ideas may work, some may fail. Please don't worry that your idea might get shot down: if you articulate a reasonable thought, probably ten other students (and teachers) share it, and getting it out advances the conversation. If it gets shot down, fine! Then we can shoot back (kindly) or move to the next idea. Teachers will award a bonus point or two to a few of the best posts, and mistaken ideas may be among these best posts.

Here are examples of ideas from past semesters:

- Find the correlation, r, between cB58 and a spectrum, for all spectra and all redshifts; then use largest r.
- For a target $y_1, ..., y_n$ and spectrum $s_1, ..., s_n$, find difference vectors, $d_i = y_{i+1} y_i$ and $e_i = s_{i+1} - s_i$, and then find spectra with minimum difference in difference vectors d and e.
- Use a chi-squared statistic to get the distance from cB58.

Implement a search from the Canvas Discussion. A successful search should find at least one persuasive match in its top 3 spectra, namely data/spec-1353-53083-0579.fits, since that is the the SDSS noisy spectrum of cB58. Some hints:

• Consider the and_mask and ivar variables for some distance measures. Note that cB58 has about 2000 points, but the other spectra have about 5000 each. Consider all possible alignments, from 1-2000, 2-2001, 3-2002, through 3001-5000.

• (If your search is successful, please ignore this bullet point.) If you cannot get an idea from the Discussion to work, briefly report on the failure, including your failed distance code. Then you can implement the following simple idea. Mostly forget about astronomy and recognize that the x-coordinate wavelengths are not relevant (because we'll redshift them) and the y-coordinates (flux) are relevant. So we have one vector of y-coordinates for cB58, the known Lyman break galaxy. We have another 100 vectors of y-coordinates (in the data directory). Which of the 100 vectors is most like the cB58 vector?

For two vectors of y-coordinates A and B, the simplest measure I can think of is the Euclidean distance $(A, B) = \sqrt{\sum_{i=1}^{n} (A_i - B_i)^2}$. (Using absolute values instead of squaring seems equally simple.)

For each of cB58 and the other spectrum, I first standardized y-coordinates by subtracting their mean and dividing by their standard deviation. I figured subtracting the mean allows aligning spectra of objects with different brightness (height on the y-axis).

I divided by the standard deviation (that is, I multiplied by a constant) because I wanted to force graphical alignment of cB58 and spec-1353-53083-0579.fits. Christy, the astronomer, said two objects with the same luminosity at different distances would have brightness at each wavelength differ by the ratio of their distances squared, so multiplying by a constant allows aligning objects at different distances.

This approach put spec-1353-53083-0579.fits in the top 3.

4 Submission Instructions

Please turn in these four files:

- 0 There's no file for this item (0), but don't forget to write the three Discussion posts mentioned above.
- 1 hw2.R, which implements your search
 - Please put this code as its first line: rm(list=ls()) This will ensure that your script does not rely on variables defined only in your current R Console session.
 - Include your name and primaryEmail@wisc.edu email address.
 - Before turning in hw2.R, run source("hw2.R") to ensure that your script works. We will run this in a directory that contains cB58_Lyman_break.fit and the "data" directory when grading.
 - Your hw2.R should read cB58_Lyman_break.fit and the 100 data/spec-....fits files (from data.tar, and one at a time, in a loop). It should write a file called hw2.csv whose 100 lines have the format distance,spectrumID,i, where
 - distance is your measure of the distance from this spectrum to cB58

- spectrumID is the spectrum ID, e.g., spec-1353-53083-0579.fits
- i is the index in the spectrum at which your alignment with cB58 begins

These lines should be sorted by increasing distance. As an example, a possible line is 1032, spec-1353-53083-0579.fits, 519, which says "the object in spec-1353-53083-0579.fits has a distance 1032 from cB58 when red-shifted by 519."

- 2 hw2.csv, the record of your search and the output of running source("hw2.R")
- 3 hw2.Rmd, which is your report (you may write this in RStudio on your local machine or on your VM):
 - Include your name and primaryEmail@wisc.edu email address.
 - Include a brief introduction describing what you tried, what worked, and what didn't. In particular,
 - Describe the measure you chose and why you chose it.
 - Mention difficulties you had.
 - Your hw2.Rmd should read your hw2.csv file and make three graphs, showing cB58 aligned with each of the top three spectra from your search in hw2.R. Include a legend with each graph identifying cB58 and the other spectrum.
 - Knit your hw2.Rmd to make hw2.html.
- 4 hw2.html, the output of hw2.Rmd

To turn in these files only, please make a new directory, e.g., ~/Desktop/turnIn/2. Copy the four required files there. Go to its parent directory, ~/Desktop/turnIn. In the shell, run tar cvf 2.tar 2 ("tape archive c=create, v=verbose, f=file to create 2.tar from the 2 directory") to create the file 2.tar from your 2 directory. It contains your four files.

Upload 2.tar as Canvas's HW2_Lyman100 assignment. Great work! You're all set!

5 Additional Notes and Helpful Tips

Here are some notes regarding this assignment and some common errors:

- Do not use setwd(), as it will fail if we do not have your directory when we run source("HW2.R"); similarly, use relative paths, not absolute paths in your code.
- Do not change file and directory names.
- Make sure variables are defined by including this as your first line: rm(list=ls())
- Make sure FITSio is loaded within your script. Please don't install and load packages we didn't indicate (unless you have permission).
- Here is a test:

- 1. Download your submitted .tar file.
- 2. Extract it in a new directory (tar xvf 2.tar test_HW2)
- 3. Use source("HW2.R") in R, or Rscript HW2.R from the shell, to confirm that
 it works.