Computer Lab 8: Land snow cover (part 2)

Climate Data Analysis, ATS 301, Fall 2018

As before, we start by specifing that we want plots to be displayed inside the Jupyter Notebook, and load the modules we'll need.

```
In [ ]: %matplotlib inline
   import numpy as np
   import matplotlib.pyplot as plt

# Set default figure size
   # This is so I don't have to keep specifying figsize later.
   plt.rcParams['figure.figsize'] = (10.0, 8.0)

import xarray as xr
   import cartopy.crs as ccrs
   from scipy import stats
   import math
```

Read in data, exclude ocean points, and points that never have snow.

```
In []: snow_file='/data/ATS_301/Data/nhsce_v01r01_19661004_20181001.nc'
    dsnow = xr.open_dataset(snow_file)
    snow = dsnow.snow_cover_extent.where(dsnow.land > 0)
    new_snow = snow.where(np.any(snow > 0,axis=0))
```

The reference for this dataset is:

Robinson, David A., Estilow, Thomas W., and NOAA CDR Program (2012):NOAA Climate Date Record (CDR) of Northern Hemisphere (NH) Snow Cover Extent (SCE), Version 1. NOAA National Climatic Data Center. doi:10.7289/V5N014G9 [accessed 10/19/2018].

Verify that this looks like what we expect:

Grid-point statistics

Our goal here is to determine whether two points have the same mean, standard deviation, etc.

Take a quick look at the time-average plot to identify some particular grid points to explore.

```
In [ ]: time_average_snow_new.plot()
In [ ]: # Compare with the plot with all land points, just to see the effect of omitting
# the snow-free points.
time_average_snow=snow.mean(dim='time')
time_average_snow.plot()
```

Pick a few points. Choose two near each other, and one further away with a different time average. Points with averages near the middle of the colorbar will work best, since they are will have more variation.

Calculate the annual mean for this single point.

Save these values, for later use.

stats.describe doesn't include the standard deviation and standard error of the mean, so we use different functions for these.

Next, calculate the mean, variance, std dev, and standard error yourself, using only np.sum, np.size, math.sqrt, and math operations. Verify that they match what python gets.

Note, you may have to use the float function (e.g., float (mean1)) to get your values to print correctly.

```
In [ ]: ## Now, calculate the mean, variance, and std dev yourself
    mean1= XXXXXX
    var1= XXXXXXX
    st_dev1= XXXXXXXX
    st_err1=XXXXXXXX
In [ ]: # ↔
```

Plot the histogram. You may want to play around with the number of bins it uses.

```
In [ ]: plt.hist(snow_1, density=True,bins='auto')
# bins='auto' or bins=14
```

Now, what would a normal distribution with this mean and std dev look like?

The stats.norm.rvs function samples size times from a normal distribution with mean loc and standard deviation scale.

```
In []: r = stats.norm.rvs(loc=m1,scale=sd1,size=n1)
    print(stats.describe(r))

# Plot our data and the sampled normal data.
    plt.hist((snow_1,r), density=True, bins='auto',color=('b','g'))

plt.show() # used here so that we don't get all the histogram output.
```

Re-run the above cell a few times. Notice how much the distribution can vary.

This is an indication that the limited number of years we have will increase the uncertainty in our statistics, as we will see soon.

Do two distributions have the same means?

Does your grid point data have the same mean as the normal distribution?

They should, since we are specifying the mean value for the distribution. The question is, can we prove it? Or rather, can we disprove that they have the same means?

[Some code is repeated here so it is all in one place.]

```
In [ ]: # Calculate the annual avereages for each year for this single point.
          snow 1=snow[:,47,30].groupby('time.year').mean()
          # Statistics for this point.
         n1, (min1, max1), m1, v1, s1, k1 = stats.describe(snow_1)
          sd1=stats.tstd(snow_1) # standard deviation (unbiased)
         sem1=stats.sem(snow 1) # Standard error
          # Samples from a normal distribution using same mean and std dev
         r = stats.norm.rvs(loc=m1,scale=sd1,size=n1)
         norm n, (norm min, norm max), norm m, norm v, norm s, norm k = stats.describe(r)
          norm std=stats.tstd(r)
         norm sem=stats.sem(r)
          # Create a table comparing the two
         print('
                              Snow Normal')
         print('Mean: {0:.3f} {1:.3f}'.format(m1, norm_m))
print('Var: {0:.4f} {1:.4f}'.format(v1, norm_v))
print('Std: {0:.3f} {1:.3f}'.format(sd1, norm_std))
print('SEM: {0:.3f} {1:.3f}'.format(sem1, norm_sem))
                               {0:.2f} {1:.2f}'.format(s1, norm_s))
{0:.2f} {1:.2f}'.format(k1. norm k))
          print('Skew:
         print('Kurt:
                                             {1:.2f}'.format(k1, norm_k))
                               {0:.2f}
          # Plot our data and the sampled normal data.
          plt.hist((snow_1,r), density=True, bins='auto',color=('b','g'))
          plt.show() # used here so that we don't get all the histogram output.
```

```
In [ ]: #--
```

The **means** and **standard** errors give us an idea of the significance of the difference of the means. More formally, we can do statistical tests to check this.

Here, we're just adding a t-test to the previous code cell.

```
In [ ]: # Calculate the annual averages for each year for this single point.
        snow 1=snow[:,47,30].groupby('time.year').mean()
        # Statistics for this point.
        n1, (min1, max1), m1, v1, s1, k1 = stats.describe(snow 1)
        sdl=stats.tstd(snow 1) # standard deviation (unbiased)
        sem1=stats.sem(snow 1) # Standard error
        # Samples from a normal distribution using same mean and std dev
        r = stats.norm.rvs(loc=m1,scale=sd1,size=n1)
        norm n, (norm min, norm max), norm m, norm v, norm s, norm k = stats.describe(r)
        norm std=stats.tstd(r)
        norm sem=stats.sem(r)
        # Create a table comparing the two
        print('
                                    Normal')
                           Snow
                       {0:.4f}
{0:.3f}
{0:.3f}
        print('Mean:
                                    {1:.3f}'.format(m1, norm_m))
                                    {1:.4f}'.format(v1, norm_v))
        print('Var:
                          {0:.3f} {1:.3f}'.format(sd1, norm_std))
{0:.3f} {1:.3f}'.format(sem1, norm_sem))
        print('Std:
        print('SEM:
                                    {1:.2f}'.format(s1, norm_s))
        print('Skew:
                           {0:.2f}
        print('Kurt:
                           {0:.2f}
                                        {1:.2f}'.format(k1, norm k))
        # Plot our data and the sampled normal data.
        plt.hist((snow_1,r), density=True, bins='auto',color=('b','g'))
        plt.show() # used here so that we don't get all the histogram output.
        # Student's t-test
        print()
        print(stats.ttest_ind(snow_1,r,equal_var=False))
```

Small p-values indicate that the difference in means is unlikely to be larger than what we obtained here due just to chance.

Large p-values indicate that it's not unlikely you'd get the same difference in sample means from the same distribution by change.

Based on the p-values, we cannot reject the null hypothesis that the mean are consistent.

Repeating the comparison using a different sample of the normal distribution

Try running the above cell again. Do the statistics and results of the Student's t-test change? Does the p-value ever indicate a statistically significant (at the 0.05 level) difference? Should it?

For your homework, you will re-sample from the normal distribution a number of times (~20). Record the t-statistics and p-values. It's easiest to do this using a **for loop** (see below). However, you can also repeat the calculation manually, recording the p values and t statistics each time. Put the t-statistics and p-values into a table.

```
In []: df = n1+n1-2
    x = np.linspace(stats.t.ppf(0.01, df),stats.t.ppf(0.99, df), 100)
    plt.plot(x, stats.t.pdf(x, df))

# For your homework, you can either use a for loop to generate these values 20 times,
    # or run the previous cell 20 times and record the t-statistics and p values.

# Copy this line 20 times and replace with the t-statistics for your data.
    plt.axvline(x=-0.3)
    plt.axvline(x=0.19541631165695444)
```

Comparing two normal distributions

The data we have is probably not really a normal distribution. What happens if we compare one random normal distribution to another?

By comparing two sets of normal distributions samples (which we know for sure either are or are not sampled from the same normal distribution population), we can get a feel for how well we can expect the student's t-test to work.

Repeat the above. With what sort of frequency do you get a result that is "statistically significant"?

Now, increase the *number* of values in each sample (and re-run the above cell). What happens?

Finally, create two different normal distributions and see how well they compare to each other.

```
In [ ]: # Create two _different_ normal distributions and see how well they compare to eac
h other.

r1 = stats.norm.rvs(loc=ml+sd1/2,scale=sd1,size=n1)
r2 = stats.norm.rvs(loc=m1,scale=sd1,size=n1)

print(stats.describe(r1))
print(stats.describe(r2))

print(stats.ttest_ind(r1,r2,equal_var=False))

plt.hist((r1,r2), density=True, bins=20, color=('g','b'))
plt.show()
```

Even though we know they are different, the test doesn't always come back "significant." Increase the number of data points (i.e. "years") until you usually get a "significant" result.

Compare annual means from two grid points

Finally, we are ready to compare snow data at different points.

```
In [ ]: # Use your own point here.
        snow_2=snow[:,47,31].groupby('time.year').mean()
        n2, (min2, max2), m2, v2, s2, k2 = stats.describe(snow_2)
        sd2=stats.tstd(snow_2) # standard deviation (unbiased)
                                # Standard error
        sem2=stats.sem(snow_2)
        # Create a table comparing the two
                          Snow 1
                                   Snow 2')
        print('
        print('Mean:
                          {0:.3f}
                                     {1:.3f}'.format(m1, m2))
                                   {1:.4f}'.format(v1, v2))
        print('Var:
                          {0:.4f}
                                    {1:.3f}'.format(sd1, sd2))
        print('Std:
                          {0:.3f}
                                    {1:.3f}'.format(sem1, sem2))
        print('SEM:
                          {0:.3f}
        print('Skew:
                          {0:.2f}
                                      {1:.2f}'.format(s1, s2))
        print('Kurt:
                          {0:.2f}
                                      {1:.2f}'.format(k1, k2))
        # Student's t-test
        print()
        print(stats.ttest_ind(snow_1,snow_2,equal_var=False))
        print(stats.ttest_rel(snow_1,snow_2))
        plt.hist((snow 1,snow 2), density=True, bins='auto', color=('b', 'g'))
        plt.show()
```

For my two points, I can reject the null hypothesis that these point have the same statistics. (You may get a different result, depending on the points you choose.)

```
In [ ]: # Pick another point and compare to the first.↔
```

For these two points, I can reject the null hypothesis that these point have the same statistics. (You may get a different result, depending on the points you choose.)

```
In [ ]: # Test points 2 and 3 \leftrightarrow
```

The difference in these two means is not statistically significant according to either test. We cannot reject the null hypothesis.

Create a nice table summarizing these three points.

Using Loops (AKA Making the computer do all the work)

Ok, we've calculated the t-statistics and p-values for a few sample distributions. But we'd like to do 20-50.

Here's where a **loop** is useful. One thing that computers are great at is repeating actions. We can repeat commands over and over (and over) by placing the commands we want to repeat inside a **for loop**. Here's an example that prints "Hello World" 10 times:

```
for i in range(10):
    print("Hello, World")
```

The __indentation__ (the spaces in front of the `print` command) is very important here. This is how the command interpreter knows which commands are within the loop!

The variable i is an **index**. range(10) is a simple way of creating a **list** of 10 values, from 0 to 9 (i.e., 0, 1, 2, 3, 4, 5, 6, 7, 8, 9). Remember that Python uses an index of 0 for the first value in a list. If you are used to languages that start with a 1, this behavior will probably cause some bugs in your code at some point, so be aware of the different indexing conventions.

The **for loop** runs the code inside the loop once for every value in the list. Within the loop, the variable i has whatever the current value from the list is. That is, the first time the body of the loop is evaluated, i has a value of 0. The second time i is 1. for i in range(10) is just a way of saying we will evaluate the statements within the loop 10 times.

Try this simple loop. To see how the index (i) changes, we'll output during every **iteration** of the loop. *Notice that the two indented lines are repeated, while the last command is not.*

```
In [ ]: print('before loop')
    for i in range(10):
        print("Hello, World")
        print('The current value of i is ',i)
    print('after loop')
```

As you wrote this code, you may have noticed that Jupyter Notebook automatically indents forward, but you have to manually delete spaces when you return to the previous indentation level.

You don't have to stick to indicies of 0, 1, 2, etc. You can also use any list (such as a list of solar constants) to loop over. The commands within the for loop will be repeated once for every index in the list.

Note: do not change the value of your index (i.e., 'i' above) within the loop. Bad stuff will happen!

```
In [ ]: print('before loop')
    for solar_constant in np.linspace(300,400,11):
        print("Hello, Earth")
        print('The current value of the solar constant is ',solar_constant)
    print('after loop')
```

So, rather than running the following cell multiple times, being sure to record the values:

and then using those values to make a plot:

```
In []: df = n1+n1-2
    x = np.linspace(stats.t.ppf(0.01, df), stats.t.ppf(0.99, df), 100)
    plt.plot(x, stats.t.pdf(x, df))

# For your homework, you can either use a for loop to generate these values 20 times,
# or run the previous cell 20 times and record the t-statistics and p values.

# Copy this line 20 times and replace with the t-statistics for your data.
    plt.axvline(x=-0.3)
    plt.axvline(x=0.19541631165695444)
```

We can do this all in one cell:

```
In []: df = n1+n1-2
x = np.linspace(stats.t.ppf(0.01, df), stats.t.ppf(0.99, df), 100)
plt.plot(x, stats.t.pdf(x, df))

for i in range(20):
    r = stats.norm.rvs(loc=m1, scale=sd1, size=n1)
    print(stats.ttest_ind(snow_1, r, equal_var=False))
    t_stat, p_val=stats.ttest_ind(snow_1, r, equal_var=False)
    plt.axvline(x=t_stat)
```

```
In [ ]:
```