# Python Code for QSS Chapter 7: Uncertainty

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First Printing

```
[]: import pandas as pd
import numpy as np
from scipy import stats
import matplotlib.pyplot as plt
import seaborn as sns
```

#### Section 7.1: Estimation

#### Section 7.1.1: Unbiasedness and Consistency

```
[]: # simulation parameters
n = 100 # sample size
mu0 = 0 # mean of Y_i(0)
sd0 = 1 # standard deviation of Y_i(0)
mu1 = 1 # mean of Y_i(1)
sd1 = 1 # standard deviation of Y_i(1)

# generate a sample
Y0 = stats.norm.rvs(size=n, loc=mu0, scale=sd0)
Y1 = stats.norm.rvs(size=n, loc=mu1, scale=sd1)
tau = Y1 - Y0 # individual treatment effect
# true value of the sample average treatment effect
SATE = tau.mean()
SATE
```

#### []: 1.1324239959603015

```
[]: # repeatedly conduct randomized controlled trials
sims = 5000 # repeat 5,000 times, we could do more
diff_means = np.zeros(sims) # container
sample_vector = np.concatenate((np.ones(int(n/2)), np.zeros(int(n/2))))

for i in range(sims):
    # randomize the treatment by sampling of a vector of 0's and 1's
    treat = np.random.choice(sample_vector, size=n, replace=False)
    # difference-in-means
    diff_means[i] = Y1[treat==1].mean() - Y0[treat==0].mean()
```

```
# estimation of error for SATE
     est_error = diff_means - SATE
     est_error.mean()
[]: 0.002340959712900045
[]: pd.Series(est_error).describe().round(5)
[]: count
              5000.00000
    mean
                 0.00234
                 0.12593
     std
    min
                -0.43821
    25%
                -0.08159
    50%
                 0.00175
    75%
                 0.08941
                 0.48091
    max
     dtype: float64
[]: # PATE simulation
     PATE = mu1 - mu0
     diff_means = np.zeros(sims)
     for i in range(sims):
         # generate a sample for each simulation
         Y0 = stats.norm.rvs(size=n, loc=mu0, scale=sd0)
         Y1 = stats.norm.rvs(size=n, loc=mu1, scale=sd1)
         treat = np.random.choice(sample_vector, size=n, replace=False)
         diff_means[i] = Y1[treat==1].mean() - Y0[treat==0].mean()
     # estimation error for PATE
     est_error = diff_means - PATE
     # unbiased
     est_error.mean()
[]: 0.0029577230122792265
[]: pd.Series(est_error).describe().round(5)
              5000.00000
[]: count
                 0.00296
    mean
    std
                 0.19731
    min
                -0.76042
    25%
                -0.13340
    50%
                 0.00249
    75%
                 0.14015
                 0.70650
    max
```

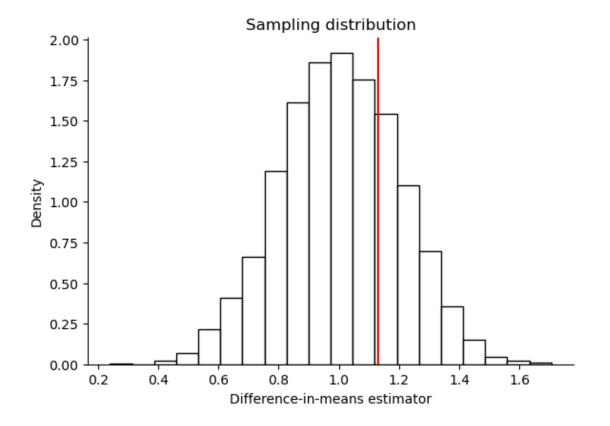
dtype: float64

# Section 7.1.2: Standard Error

```
[]: sns.displot(
    diff_means, stat='density', color='white', edgecolor='black',
    height=4, aspect=1.5, bins=20
).set(title='Sampling distribution', xlabel='Difference-in-means estimator')

plt.axvline(SATE, color='red') # true value of SATE
```

[]: <matplotlib.lines.Line2D at 0x28c69242380>



```
[ ]: diff_means.std(ddof=1)
[ ]: 0.19730602332768601
[ ]: np.sqrt(((diff_means - SATE)**2).mean())
```

[]: 0.235973296665256

```
[]: # PATE simulation with standard error
     sims = 5000
     diff_means = np.zeros(sims)
     se = np.zeros(sims)
     for i in range(sims):
         # generate a sample for each simulation
         Y0 = stats.norm.rvs(size=n, loc=mu0, scale=sd0)
         Y1 = stats.norm.rvs(size=n, loc=mu1, scale=sd1)
         # randomize treatment by sampling the vector of 0's and 1's created above
         treat = np.random.choice(sample vector, size=n, replace=False)
         diff_means[i] = Y1[treat==1].mean() - Y0[treat==0].mean()
         se[i] = (np.sqrt(Y1[treat==1].var(ddof=1) / (n/2) +
                          Y0[treat==0].var(ddof=1) / (n/2))
     diff_means.std(ddof=1)
[]: 0.19686379792583616
[]: se.mean()
[]: 0.1996177467877422
    Section 7.1.3: Confidence Intervals
[]: n = 1000 \# sample size
     x_bar = 0.6 # point estimate
     s_e = np.sqrt(x_bar * (1-x_bar) / n) # standard error
     # 99% confidence intervals; display as a tuple
     ((x_bar - stats.norm.ppf(0.995) * s_e).round(5),
      (x_bar + stats.norm.ppf(0.995) * s_e).round(5))
[]: (0.5601, 0.6399)
[]: # 95% confidence intervals
     ((x_bar - stats.norm.ppf(0.975) * s_e).round(5),
      (x_bar + stats.norm.ppf(0.975) * s_e).round(5))
[]: (0.56964, 0.63036)
[]: # 90% confidence intervals
     ((x_bar - stats.norm.ppf(0.95) * s_e).round(5),
      (x_bar + stats.norm.ppf(0.95) * s_e).round(5))
[]: (0.57452, 0.62548)
```

```
[]: # empty container matrices for 2 sets of confidence intervals
    ci95 = np.zeros(sims*2).reshape(sims, 2)
    ci90 = np.zeros(sims*2).reshape(sims, 2)

# 95 percent confidence intervals
    ci95[:,0] = diff_means - stats.norm.ppf(0.975) * se # lower limit
    ci95[:,1] = diff_means + stats.norm.ppf(0.975) * se # upper limit

# 90 percent confidence intervals
    ci90[:,0] = diff_means - stats.norm.ppf(0.95) * se # lower limit
    ci90[:,1] = diff_means + stats.norm.ppf(0.95) * se # upper limit

# coverage rate for 95% confidence interval
    ((ci95[:,0] <= 1) & (ci95[:,1] >= 1)).mean()
```

```
[]: # coverage rate for 90% confidence interval ((ci90[:,0] <= 1) & (ci90[:,1] >= 1)).mean()
```

[]: 0.8982

```
[]: p = 0.6 \# true parameter value
     n = np.array([50, 100, 1000]) # 3 sample sizes to be examined
     alpha = 0.05
     sims = 5000 # number of simulations
     results = np.zeros(len(n)) # a container for results
     for i in range(len(n)):
         ci results = np.zeros(sims) # a container for whether CI contains truth
         # loop for repeated hypothetical survey sampling
         for j in range(sims):
             data = stats.binom.rvs(n=1, p=p, size=n[i]) # simple random sampling
             x_bar = data.mean() # sample proportion as an estimate
             s_e = np.sqrt(x_bar * (1-x_bar) / n[i]) # standard errors
             ci_lower = x_bar - stats.norm.ppf(1-alpha/2) * s_e
             ci_upper = x_bar + stats.norm.ppf(1-alpha/2) * s_e
             ci_results[j] = (p >= ci_lower) & (p <= ci_upper)</pre>
         # proportion of CIs that contain the true value
         results[i] = ci_results.mean()
     results
```

[]: array([0.9366, 0.9448, 0.942])

## Section 7.1.4: Margin of Error and Sample Size Calculation in Polls

```
[]: MoE = np.array([0.01, 0.03, 0.05]) # the desired margin of error
    p = np.arange(0.01, 1, 0.01)
    n = 1.96**2 * p * (1-p) / MoE[0]**2
    n2 = 1.96**2 * p * (1-p) / MoE[1]**2
    n3 = 1.96**2 * p * (1-p) / MoE[2]**2

fig, ax = plt.subplots(figsize=(6,4))

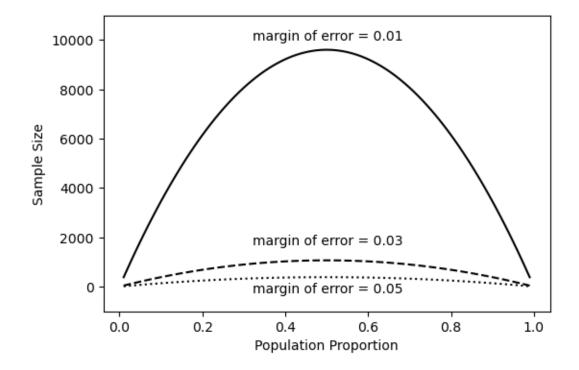
sns.lineplot(x=p, y=n, ax=ax, color='black').set(
    ylim=(-1000, 11000), xlabel='Population Proportion', ylabel='Sample Size')

sns.lineplot(x=p, y=n2, ax=ax, color='black', linestyle='--')

sns.lineplot(x=p, y=n3, ax=ax, color='black', linestyle='--')

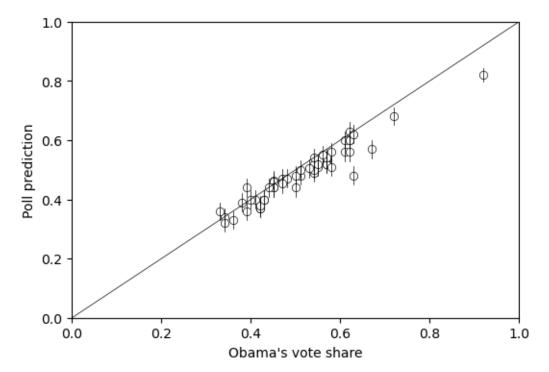
# Add text labels
    ax.text(0.32, 10000, 'margin of error = 0.01', fontsize=10)
    ax.text(0.32, 1700, 'margin of error = 0.03', fontsize=10)
    ax.text(0.32, -250, 'margin of error = 0.05', fontsize=10)
```

[]: Text(0.32, -250, 'margin of error = 0.05')



```
[]: # election and polling results, by state
     pres08 = pd.read_csv('pres08.csv')
     polls08 = pd.read_csv('polls08.csv')
     # convert to a date object
     polls08['middate'] = pd.to_datetime(polls08['middate'])
     # number of days to the election
     from datetime import datetime
     election_day = datetime.strptime('2008-11-04', '%Y-%m-%d')
     polls08['days_to_election'] = (election_day - polls08['middate']).dt.days
     # extract unique state names which the loop will iterate through
     st_names = polls08['state'].unique()
     # create an empty 51 X 3 placeholder Data Frame
     poll_pred = pd.DataFrame(np.zeros(51*3).reshape(51, 3), index=st_names)
     # loop across the 50 states plus DC
     for i in range(len(st_names)):
         # subset the ith state
         state_data = polls08[polls08['state'] == st_names[i]]
         # further subset the latest polls within the state
         latest = (state_data['days_to_election']==
                   state_data['days_to_election'].min())
         # compute the mean of the latest polls and store it
         poll_pred.iloc[i, 0] = state_data['Obama'][latest].mean() / 100
     # upper and lower confidence limits
     n = 1000 \# sample size
     alpha = 0.05
     se = np.sqrt(poll_pred.iloc[:,0] * (1-poll_pred.iloc[:,0]) / n) # standard error
     poll_pred.iloc[:,1] = poll_pred.iloc[:,0] - stats.norm.ppf(1-alpha/2) * se
     poll_pred.iloc[:,2] = poll_pred.iloc[:,0] + stats.norm.ppf(1-alpha/2) * se
[]: # plot the results
     fig, ax = plt.subplots(figsize=(6,4))
     sns.scatterplot(
         x = pres08['Obama'] / 100, y = poll_pred.iloc[:,0].reset_index(drop=True),
         ax=ax, color='white', edgecolor='black'
     ).set(xlabel="Obama's vote share", ylabel='Poll prediction',
           xlim=(0, 1), ylim=(0, 1))
     ax.axline((0, 0), slope=1, color='black', linewidth=0.5)
     # adding 95% confidence intervals for each state
```

```
for i in range(len(st_names)):
    ax.plot(
        [pres08['Obama'][i] / 100] * 2,
        [poll_pred.iloc[i,1], poll_pred.iloc[i,2]],
        color='black', linewidth=0.5
)
```



```
[]: # proportion of confidence intervals that contain the election day outcome # reset index: can only compare identically-labeled Series objects

((poll_pred.iloc[:,1].reset_index(drop=True) <= pres08['Obama'] / 100) & (poll_pred.iloc[:,2].reset_index(drop=True) >= pres08['Obama'] / 100)).mean()
```

```
[]: # bias bias=(poll_pred.iloc[:,0].reset_index(drop=True) - pres08['Obama']/100).mean() bias
```

#### []: -0.026797385620915028

```
[]: # bias corrected estimate
poll_bias = poll_pred.iloc[:,0] - bias
# bias corrected standard error
```

```
se_bias = np.sqrt(poll_bias * (1-poll_bias) / n)

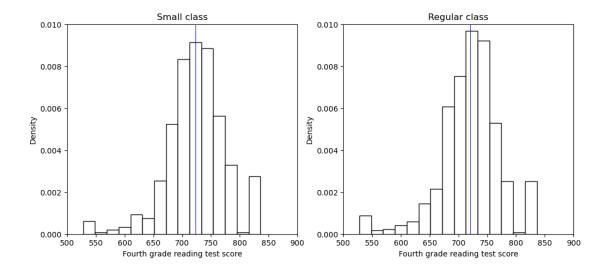
# bias corrected confidence intervals
ci_bias_lower = poll_bias - stats.norm.ppf(1-alpha/2) * se_bias
ci_bias_upper = poll_bias + stats.norm.ppf(1-alpha/2) * se_bias

# proportion of bias corrected CIs that contain election day outcome
((ci_bias_lower.reset_index(drop=True) <= pres08['Obama'] / 100) &
    (ci_bias_upper.reset_index(drop=True) >= pres08['Obama'] / 100)).mean()
```

#### Section 7.1.5: Analysis of Randomized Controlled Trials

```
[]: STAR = pd.read csv('STAR.csv')
     fig, axs = plt.subplots(1, 2, figsize=(12,5))
     sns.histplot(
         STAR['g4reading'][STAR.classtype==1], stat = 'density', ax=axs[0],
         color='white', edgecolor='black', bins=15
     ).set(ylim=(0, 0.01), xlim=(500, 900), title='Small class',
           xlabel='Fourth grade reading test score')
     axs[0].axvline(STAR['g4reading'][STAR.classtype==1].mean(),
                    color='blue', linewidth=0.75)
     sns.histplot(
         STAR['g4reading'][STAR.classtype==2], stat = 'density', ax=axs[1],
         color='white', edgecolor='black', bins=15
     ).set(ylim=(0, 0.01), xlim=(500, 900), title='Regular class',
           xlabel='Fourth grade reading test score')
     axs[1].axvline(STAR['g4reading'][STAR.classtype==2].mean(),
                     color='blue', linewidth=0.75)
```

[]: <matplotlib.lines.Line2D at 0x28c69241fc0>



```
[]: # estimate and standard error for small class size

n_small = (STAR['classtype']==1 & STAR['g4reading'].notnull()).sum()

est_small = STAR['g4reading'][STAR.classtype==1].mean()

se_small = STAR['g4reading'][STAR.classtype==1].std() / np.sqrt(n_small)

est_small, se_small
```

[]: (723.3911845730028, 1.9130122952458233)

[]: (719.88995215311, 1.8388496908502467)

[]: (719.6417493723386, 727.1406197736669)

```
ci_regular
[]: (716.2858729860609, 723.4940313201591)
[]: # difference in means estimator
     ate_est = est_small - est_regular
     ate_est
[ ]: 3.5012324198927445
[]: # standard error and 95% confidence interval
     ate_se = np.sqrt(se_small**2 + se_regular**2)
     ate_se
[]: 2.653485298112982
[]: ate_ci = (ate_est - stats.norm.ppf(1-alpha/2) * ate_se,
               ate_est + stats.norm.ppf(1-alpha/2) * ate_se)
     ate_ci
[]: (-1.699503197915229, 8.701968037700718)
    Section 7.1.6: Analysis Based on Student's t-Distribution
[]: # 95% CI for small class
     (est_small - stats.t.ppf(0.975, df=n_small-1) * se_small,
      est_small + stats.t.ppf(0.975, df=n_small-1) * se_small)
[]: (719.635479522832, 727.1468896231735)
[]: # 95% CI based on the central limit theorem
     ci small
[]: (719.6417493723386, 727.1406197736669)
[]: # 95% CI for regular class
     (est_regular - stats.t.ppf(0.975, df=n_regular-1) * se_regular,
      est_regular + stats.t.ppf(0.975, df=n_regular-1) * se_regular)
[]: (716.2806412822123, 723.4992630240077)
[]: # 95% CI based on the central limit theorem
     ci_regular
[]: (716.2858729860609, 723.4940313201591)
[]: test_result = stats.ttest_ind(
        STAR['g4reading'][STAR.classtype==1],
```

```
STAR['g4reading'][STAR.classtype==2],
    # override default equal_var=True; False is Welch t-test
    equal_var=False,
    # override default nan_policy='propagate'
    nan_policy='omit')

# we can extract results from the test_result object
test_result.pvalue
```

```
[]: # find the confidence interval
ci = test_result.confidence_interval(confidence_level=0.95)

# print summary of results
print(
    'Welch Two Sample t-test\n'
    f"t-stat: {test_result.statistic:.4f}\n"
    f"p-value: {test_result.pvalue:.4f}\n"
    f"df: {test_result.df:.1f}\n"
    f"95% confidence interval: ({ci[0]:.4f}, {ci[1]:.4f})"
)
```

```
Welch Two Sample t-test
t-stat: 1.3195
p-value: 0.1872
df: 1541.2
95% confidence interval: (-1.7036, 8.7061)
```

# Section 7.2: Hypothesis Testing

#### Section 7.2.1: Tea-Testing Experiment

```
[]: from math import comb

# truth: enumerate the number of assignment combinations
true = np.array(
        [comb(4,0) * comb(4,4),
        comb(4,1) * comb(4,3),
        comb(4,2) * comb(4,2),
        comb(4,3) * comb(4,1),
        comb(4,4) * comb(4,0)]
)

true
```

```
[]: array([1, 16, 36, 16, 1])
```

```
true = pd.Series(true / true.sum(), index=[0,2,4,6,8])
     true
[]: 0
         0.014286
         0.228571
     4
         0.514286
         0.228571
         0.014286
     dtype: float64
[]: # simulations
     sims=1000
     # lady's quess: M stands for 'Milk first', T stands for 'Tea first'
     guess=np.array(['M', 'T', 'T', 'M', 'M', 'T', 'T', 'M'])
     sample_vector=np.array(['T']*4 + ['M']*4)
     correct=pd.Series(np.zeros(sims)) # place holder for number of correct guesses
     for i in range(sims):
         # randomize which cups get Milk/Tea first
         cups=np.random.choice(sample_vector, size=len(sample_vector), replace=False)
         correct[i]=(guess==cups).sum() # number of correct guesses
     # estimated probability for each number of correct quesses
     correct.value_counts(normalize=True).sort_index()
[]: 0.0
           0.015
    2.0
           0.254
    4.0
           0.493
     6.0
           0.225
     8.0
           0.013
     Name: proportion, dtype: float64
[]: # comparison with analytical answers; the differences are small
     correct.value_counts(normalize=True).sort_index() - true
[]: 0.0
           0.000714
    2.0
         0.025429
     4.0
         -0.021286
     6.0
         -0.003571
     8.0
         -0.001286
     dtype: float64
```

[]: # compute probability: divide it by the total number of events

#### Section 7.2.2: The General Framework

```
[]: # all correct
    x = pd.DataFrame(\{'M': [4,0], 'T': [0,4]\}, index=['M','T'])
    # six correct
    y = pd.DataFrame(\{'M': [3,1], 'T': [1,3]\}, index=['M','T'])
Г1:
       M T
       4 0
    M
    T 0 4
[ ]: y
Г1:
       Μ
    M 3 1
    T 1 3
[]: # one-sided test for 8 correct guesses
    fisher_one=stats.fisher_exact(x.values, alternative='greater')
    print(
        "Fisher's Exact Test for Count Data: One-Sided\n"
        f"P-value: {fisher one.pvalue:.5f}"
    )
    Fisher's Exact Test for Count Data: One-Sided
    P-value: 0.01429
[]: # two-sided test for 6 correct guesses
    fisher_two=stats.fisher_exact(y.values)
    print(
         "Fisher's Exact Test for Count Data: Two-Sided\n"
        f"P-value: {fisher two.pvalue:.5f}"
    )
    Fisher's Exact Test for Count Data: Two-Sided
    P-value: 0.48571
    Section 7.2.3: One-Sample Tests
[ ]: n = 1018
    x bar = 550 / n
    se = np.sqrt(0.5 * 0.5 / n) # standard deviation of sampling distribution
     # upper red area in the figure
    upper = 1 - stats.norm.cdf(x_bar, loc=0.5, scale=se)
```

```
# lower red area in the figure; identical to the upper area
    lower = stats.norm.cdf(0.5 - (x_bar - 0.5), loc=0.5, scale=se)
     # two-sided p-value
    upper + lower
[]: 0.010168663287718531
[]: 2 * upper
[]: 0.010168663287718482
[]: # one-sided p-value
    upper
[]: 0.005084331643859241
[]: z_score = (x_bar - 0.5) / se
    z score
[]: 2.5700404773096097
[]: # one-sided p-value
    1 - stats.norm.cdf(z_score)
[]: 0.005084331643859241
[]: # two-sided p-value
    2 * (1 - stats.norm.cdf(z_score))
[]: 0.010168663287718482
[]: # 99% confidence interval contains 0.5
     (x_bar - stats.norm.ppf(0.995) * se, x_bar + stats.norm.ppf(0.995) * se)
[]: (0.499909283428347, 0.58064081480348)
[]: # 95% confidence interval does not contain 0.5
     (x_bar - stats.norm.ppf(0.975) * se, x_bar + stats.norm.ppf(0.975) * se)
[]: (0.5095604956138589, 0.5709896026179682)
[]: from statsmodels.stats.proportion import proportions_ztest, proportion_confint
    # no continuity correction to get the same p-value as above
    stat, pval = proportions_ztest(count=550, nobs=n, value=0.5, prop_var=0.5)
    ci = proportion_confint(count=550, nobs=n)
```

```
print(
    'One-sample z-test without continuity correction\n'
    'Alternative hypothesis: true p is not equal to 0.5\n'
    f"Sample proportion: {x_bar:.4f}\n"
    f"Test statistic: {stat:.4f}\n"
    f"P-value: {pval:.4f}\n"
    f"95% confidence interval: ({ci[0]:.4f}, {ci[1]:.4f})"
)
```

```
One-sample z-test without continuity correction Alternative hypothesis: true p is not equal to 0.5 Sample proportion: 0.5403
Test statistic: 2.5700
P-value: 0.0102
95% confidence interval: (0.5097, 0.5709)
```

The continuity correction factor subtracts 0.5 from the sample proportion before computing the test statistic. The correction factor is not built into the one-sample z-tests available in Python. However, we can build a function to implement the correction using the hypothesis testing logic we have developed.

```
[]: # Define a function to implement the continuity correction factor
     def proportions_ztest_correct(count, nobs, value, conf_level=0.95):
         # compute the p-value
         prop = count / nobs
         correction = 0.5 / nobs # Yates' continuity correction
         adjusted_prop = prop - correction
         se_null = np.sqrt(value * (1-value) / nobs) # SE under the null hypothesis
         z_score = np.abs(adjusted_prop-value) / se_null
         # assume a two-tailed test, but we could generalize this
         pval = 2 * (1 - stats.norm.cdf(z_score))
         # compute the confidence interval
         se_sample = np.sqrt(adjusted_prop * (1-adjusted_prop) / nobs)
         alpha = 1-conf level
         ci_lower = adjusted_prop - stats.norm.ppf(1-alpha/2) * se_sample
         ci_upper = adjusted_prop + stats.norm.ppf(1-alpha/2) * se_sample
         conf_print = int(conf_level * 100)
         print(
             'One-sample z-test with continuity correction\n'
             f"Alternative hypothesis: true p is not equal to {value}\n"
            f"Sample proportion: {prop:.4f}\n"
             f"Test statistic: {z_score:.4f}\n"
             f"P-value: {pval:.4f}\n"
             f"{conf_print}% confidence interval: ({ci_lower:.4f}, {ci_upper:.4f})"
```

```
[]: proportions_ztest_correct(count=550, nobs=n, value=0.5)
    One-sample z-test with continuity correction
    Alternative hypothesis: true p is not equal to 0.5
    Sample proportion: 0.5403
    Test statistic: 2.5387
    P-value: 0.0111
    95% confidence interval: (0.5092, 0.5704)
[]:|proportions_ztest_correct(count=550, nobs=n, value=0.5, conf_level=0.99)
    One-sample z-test with continuity correction
    Alternative hypothesis: true p is not equal to 0.5
    Sample proportion: 0.5403
    Test statistic: 2.5387
    P-value: 0.0111
    99% confidence interval: (0.4995, 0.5800)
[]: # two-sided one-sample t-test
     star_ttest = stats.ttest_1samp(STAR.g4reading, popmean=710, nan_policy='omit')
     ci lower = star ttest.confidence interval()[0]
     ci_upper = star_ttest.confidence_interval()[1]
     print(
         'One-sample t-test\n'
         'Alternative hypothesis: true mean is not equal to 710\n'
         f"Sample mean: {STAR.g4reading.mean():.3f}\n"
         f"t-statistic: {star_ttest.statistic:.3f}\n"
         f"df: {star ttest.df}\n"
         f"P-value: {star_ttest.pvalue:.5f}\n"
         f"95% confidence interval: ({ci_lower:.3f}, {ci_upper:.3f})"
     )
    One-sample t-test
    Alternative hypothesis: true mean is not equal to 710
    Sample mean: 721.248
    t-statistic: 10.407
    df: 2352
    P-value: 0.00000
    95% confidence interval: (719.128, 723.367)
```

# Section 7.2.4: Two-Sample Tests

```
[]: # one-sided p-value
     stats.norm.cdf(-np.abs(ate_est), loc=0, scale=ate_se)
[]: 0.09350361332918433
[]: # two-sided p-value
     2 * stats.norm.cdf(-np.abs(ate_est), loc=0, scale=ate_se)
[]: 0.18700722665836866
[]: # testing the null of zero average treatment effect
     ttest = stats.ttest_ind(STAR.g4reading[STAR.classtype==1],
                             STAR.g4reading[STAR.classtype==2],
                             equal_var=False, nan_policy='omit')
     ci_lower = ttest.confidence_interval()[0]
     ci_upper = ttest.confidence_interval()[1]
     print(
         'Welch Two Sample t-test\n'
         'Alternative hypothesis: true difference in means is not equal to 0\n'
        f"Sample mean difference: {ate est:.3f}\n"
        f"t-statistic: {ttest.statistic:.3f}\n"
        f"df: {ttest.df:.1f}\n"
        f"P-value: {ttest.pvalue:.5f}\n"
        f"95% confidence interval: ({ci_lower:.3f}, {ci_upper:.3f})"
     )
    Welch Two Sample t-test
    Alternative hypothesis: true difference in means is not equal to 0
    Sample mean difference: 3.501
    t-statistic: 1.319
    df: 1541.2
    P-value: 0.18720
    95% confidence interval: (-1.704, 8.706)
[]: resume = pd.read_csv('resume.csv')
     # organize the data in a cross-tab
     x = pd.crosstab(resume.race, resume.call)
     x
[]: call
     race
    black 2278 157
     white 2200 235
```

```
[]: # one-sided test with continuity correction factor
     result = stats.chi2_contingency(x)
     print(
         '2-sample test for equality of proportions with continuity correction\n'
         'Alternative hypothesis: greater\n'
         f"X-squared: {result.statistic:.3f}\n"
         f"P-value: {result.pvalue/2}\n"
     )
    2-sample test for equality of proportions with continuity correction
    Alternative hypothesis: greater
    X-squared: 16.449
    P-value: 2.4987891949816276e-05
[]: # sample size
    n0 = (resume.race=='black').sum()
     n1 = (resume.race=='white').sum()
     # sample proportions
     p = resume['call'].mean() # overall
     p0 = resume['call'][resume.race=='black'].mean()
     p1 = resume['call'][resume.race=='white'].mean()
     # point estimate
     est = p1 - p0
     est
[ ]: 0.032032854209445585
[]: # standard error
     se = np.sqrt(p * (1 - p) * (1 / n0 + 1 / n1))
[]: 0.007796894036170457
[]: # z-statistic
     zstat = est / se
     zstat
[]: 4.108412152434346
[]: # one-sided p-value
     stats.norm.cdf(-abs(zstat))
```

[]: 1.9919434187925383e-05

```
print(
    '2-sample test for equality of proportions without continuity correction\n'
    'Alternative hypothesis: greater\n'
    f"X-squared: {result_uncorrected.statistic:.3f}\n"
    f"P-value: {result_uncorrected.pvalue/2}"
)
```

2-sample test for equality of proportions without continuity correction Alternative hypothesis: greater X-squared: 16.879

P-value: 1.991943418792538e-05

#### Section 7.2.5: Pitfalls of Hypothesis Testing

# Section 7.2.6: Power Analysis

```
[]: # set the parameters
n = 250
p_star = 0.48 # data generating process
p = 0.5 # null value
alpha = 0.05

# critical value
cr_value = stats.norm.ppf(1-alpha/2)

# standard errors under the hypothetical data generating process
se_star = np.sqrt(p_star * (1 - p_star) / n)

# standard error under the null
se = np.sqrt(p * (1 - p) / n)

# power
(stats.norm.cdf(p - cr_value * se, loc=p_star, scale=se_star) +
1 - stats.norm.cdf(p + cr_value * se, loc=p_star, scale=se_star))
```

#### []: 0.09673113765989816

```
[]: # parameters
n1 = 500
n0 = 500
p1_star = 0.05
p0_star = 0.1

# overall call back rate as a weighted average
p = (n1 * p1_star + n0 * p0_star) / (n1 + n0)
# standard error under the null
```

```
se = np.sqrt(p * (1 - p) * (1 / n1 + 1 / n0))
     # standard error under the hypothetical data generating process
     se_star = np.sqrt(p1_star * (1 - p1_star) / n1 + p0_star * (1 - p0_star) / n0)
     (stats.norm.cdf(-cr_value * se, loc=(p1_star-p0_star), scale=se_star) +
     1 - stats.norm.cdf(cr_value * se, loc=(p1_star-p0_star), scale=se_star))
[]: 0.8522799668094607
[]: from statsmodels.stats.proportion import (power_proportions_2indep,
                                               samplesize_proportions_2indep_onetail)
     power = power_proportions_2indep(
        diff=(p0_star - p1_star), prop2=p1_star, alpha=0.05, nobs1=n1
     print(power)
    power = 0.8522799668094605
    p_pooled = 0.07500000000000001
    std_null = 0.37249161064378356
    std_alt = 0.37080992435478316
    nobs1 = 500
    nobs2 = 500
    nobs ratio = 1
    alpha = 0.05
[]: samplesize_proportions_2indep_onetail(
        diff=(p0_star - p1_star), prop2=p1_star, alpha=0.05, power=0.9
     )
[]: 581.082053834476
[]: from statsmodels.stats.power import TTestPower, TTestIndPower
     TTestPower().solve power(effect size=0.25, nobs=100, alpha=0.05)
[]: 0.696980269099517
[]: TTestPower().solve_power(effect_size=0.25, power=0.9, alpha=0.05)
[]: 170.05107691102737
[]: TTestIndPower().solve_power(effect_size=0.25, power=0.9, alpha=0.05,
                                 alternative='larger')
```

[]: 274.72216286128617

# Section 7.3: Linear Regression Model with Uncertainty

#### Section 7.3.1: Linear Regression as a Generative Model

```
[]: import statsmodels.formula.api as smf
     minwage = pd.read_csv('minwage.csv')
     # compute proportion of full employment before minimum wage increase
     minwage['fullPropBefore'] = minwage['fullBefore'] / (
         minwage['fullBefore'] + minwage['partBefore']
     )
     # same thing after minimum wage increase
     minwage['fullPropAfter'] = minwage['fullAfter'] / (
         minwage['fullAfter'] + minwage['partAfter']
     # an indicator for NJ: 1 if it's located in NJ and 0 if in PA
     minwage['NJ'] = np.where(minwage['location']=='PA', 0, 1)
     minwage_model = 'fullPropAfter ~ -1 + NJ + fullPropBefore + wageBefore + chain'
     fit_minwage = smf.ols(minwage_model, data=minwage).fit()
     # regression result
     fit_minwage.params
[]: chain[burgerking]
                         -0.115625
     chain[kfc]
                         -0.150800
     chain[roys]
                         -0.206386
     chain[wendys]
                        -0.220133
    NJ
                         0.054220
     fullPropBefore
                         0.168794
     wageBefore
                          0.081334
     dtype: float64
[]: minwage model1 = 'fullPropAfter ~ NJ + fullPropBefore + wageBefore + chain'
     fit_minwage1 = smf.ols(minwage_model1, data=minwage).fit()
     fit_minwage1.params
[]: Intercept
                       -0.115625
     chain[T.kfc]
                       -0.035175
     chain[T.roys]
                       -0.090761
     chain[T.wendys]
                       -0.104507
    NJ
                        0.054220
```

fullPropBefore 0.168794 wageBefore 0.081334

dtype: float64

```
[]: fit_minwage.predict(minwage.iloc[[0]])
```

[]: 0 0.270937 dtype: float64

```
[]: fit_minwage1.predict(minwage.iloc[[0]])
```

[]: 0 0.270937 dtype: float64

# Section 7.3.2: Unbiasedness of Estimated Coefficients

```
[]: women = pd.read_csv('women.csv')

fit_women = smf.ols('water ~ reserved', data=women).fit()

print(fit_women.summary())
```

# OLS Regression Results

Dep. Variable:	water	R-squared:	0.017
Model:	OLS	Adj. R-squared:	0.014
Method:	Least Squares	F-statistic:	5.493
Date:	Wed, 15 Nov 2023	Prob (F-statistic):	0.0197
Time:	23:23:31	Log-Likelihood:	-1586.1
No. Observations:	322	AIC:	3176.
Df Residuals:	320	BIC:	3184.

Df Model: 1
Covariance Type: nonrobust

=========	========					
	coef	std err	t	P> t	[0.025	0.975]
Intercept	14.7383	2.286	6.446	0.000	10.240	19.236
reserved	9.2524	3.948	2.344	0.020	1.486	17.019
=========						
Omnibus:		398.10	04 Durbin	n-Watson:		1.990
Prob(Omnibus	3):	0.00	00 Jarque	e-Bera (JB):		26829.354
Skew:		5.69	90 Prob(	JB):		0.00
Kurtosis:		46.24	46 Cond.	No.		2.41
========	========		========			=======

# Notes:

<sup>[1]</sup> Standard Errors assume that the covariance matrix of the errors is correctly specified.

# []: # 95% confidence intervals fit\_women.conf\_int().rename(columns={0:'2.5%', 1:'97.5%'})

[]: 2.5% 97.5% Intercept 10.240240 19.236395 reserved 1.485608 17.019238

# [ ]

fullPr Least	.======	ion Results ====================================			
Least	-	R-squared:			
fullPropAfter OLS Least Squares Wed, 15 Nov 2023 23:23:31		Adj. R-squared: F-statistic: Prob (F-statistic): Log-Likelihood:		0.070 0.054 4.401 0.000264 0.84929	
no	358 351 6 onrobust	AIC: BIC:		12.30 39.47	
coef	std err	t	P> t	[0.025	
0.1156	0.179	-0.646	0.518	-0.467	
0.1508	0.183	-0.824	0.411	-0.511	
0.2064	0.187	-1.105	0.270	-0.574	
0.2201	0.188	-1.168	0.243	-0.591	
0.0542	0.033	1.633	0.103	-0.011	
0.1688	0.057	2.981	0.003	0.057	
0.0813	0.039		0.037	0.005	
15.383 Durbin-Watson:			1.973		
	0.000 0.501	_	(JB):	16.262	
	0.2064 0.2201 0.0542 0.1688 0.0813	0.2064 0.187 0.2201 0.188 0.0542 0.033 0.1688 0.057 0.0813 0.039	0.2064 0.187 -1.105 0.2201 0.188 -1.168 0.0542 0.033 1.633 0.1688 0.057 2.981 0.0813 0.039 2.090	0.2064 0.187 -1.105 0.270 0.2201 0.188 -1.168 0.243 0.0542 0.033 1.633 0.103 0.1688 0.057 2.981 0.003 0.0813 0.039 2.090 0.037  15.383 Durbin-Watson: 0.000 Jarque-Bera (JB):	

Notes:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

```
[]: # confidence interval just for the 'NJ' variable fit_minwage.conf_int().rename(columns={0:'2.5%', 1:'97.5%'}).loc['NJ']
```

[]: 2.5% -0.011093 97.5% 0.119533

Name: NJ, dtype: float64

# Section 7.3.3: Standard Errors of Estimated Coefficients

In Progress