```
In []: import numpy as np
    import pandas as pd
    import scipy
    from icecream import ic
    from matplotlib import pyplot as plt
    import seaborn as sns
    from tqdm import tqdm
    from scipy.stats import chi2_contingency, chi2
    import time
```

# **Excercise 2**

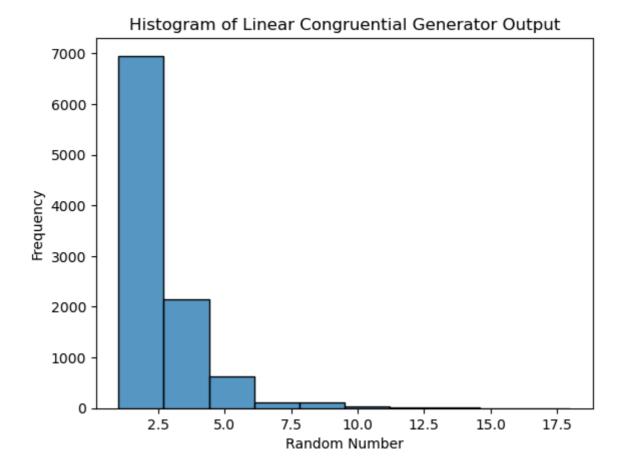
1. Choose a value for the probability parameter p in the geometric distribution and simulate 10,000 outcomes. You can experiment with a small, moderate and large value if you like.

```
In []: # Number of outcomes
N = 10_000

# Probability p
p = 0.45

x = np.random.geometric(p=p, size=N)

In []: # Plot the histogram
sns.histplot(x, bins=10)
plt.xlabel('Random Number')
plt.ylabel('Frequency')
plt.title('Histogram of Linear Congruential Generator Output')
plt.show()
```



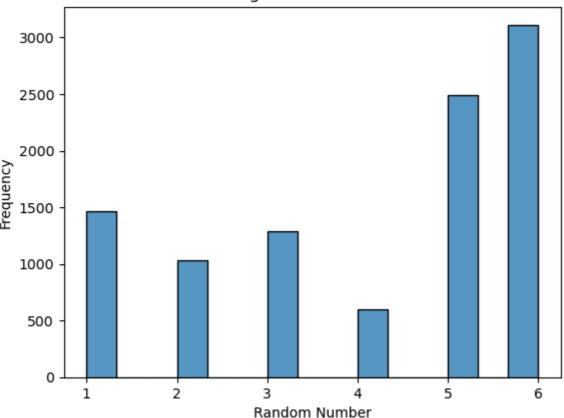
### 2. Simulate the 6 point distribution with given probabilities

First we simulate it using numpy's build in RNG as a reference

```
In []: # Define the outcomes and their probabilities
   outcomes = [1, 2, 3, 4, 5, 6]
   p = [7/48, 5/48, 1/8, 1/16, 1/4, 5/16]

# Simulate 10,000 outcomes
   start_time = time.time()
   numpy_distribution = np.random.choice(outcomes, N, p=p)
   end_time = time.time()
   numpy_time = end_time - start_time
In []: # Plot the histogram
   sns.histplot(numpy_distribution)
   plt.xlabel('Random Number')
   plt.ylabel('Frequency')
   plt.title('Histogram of Distribution')
   plt.show()
```

#### Histogram of Distribution



### 2.1 Crude method

We generate the accumulated probabilities first

```
In []: x = np.random.rand(10000)
    outcomes = [1, 2, 3, 4, 5, 6]
    probabilities = [7/48, 5/48, 1/8, 1/16, 1/4, 5/16]
    accumulated_probabilities = [sum(probabilities[0:i+1]) for i in range(len
```

We then loop through the probabilities for each uniformly distributed U to simulate a roll with probabilities equal to probabilities = [7/48, 5/48, 1/8, 1/16, 1/4, 5/16]

```
In []: # We make a list of counts
   outcomes = [1, 2, 3, 4, 5, 6]
   rolls = []

start_time = time.time()
   for _ in tqdm(range(10000)):
        U = np.random.uniform()
        for i, prob in enumerate(accumulated_probabilities):
            if U <= prob:
                rolls.append(outcomes[i])
                break
   end_time = time.time()
   crude_time = end_time - start_time

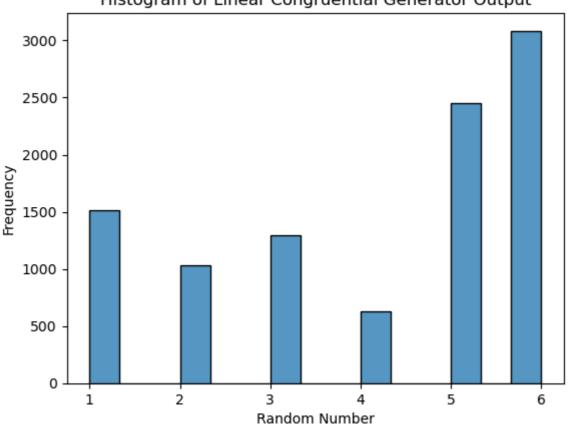
crude_method = rolls

# Plot the histogram</pre>
```

```
sns.histplot(rolls)
plt.xlabel('Random Number')
plt.ylabel('Frequency')
plt.title('Histogram of Linear Congruential Generator Output')
plt.show()

100%| 10000/10000 [00:00<00:00, 675911.95it/s]</pre>
```

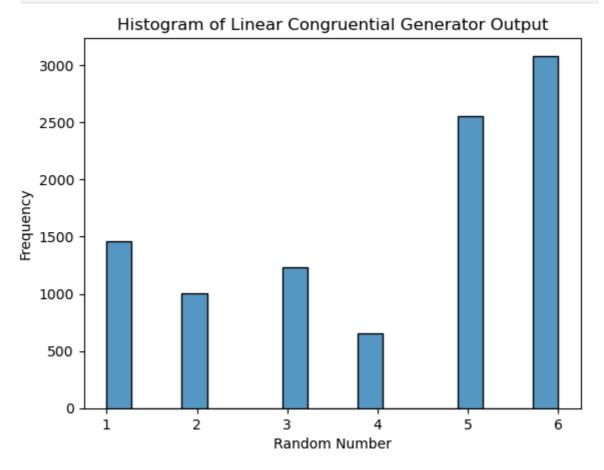
Histogram of Linear Congruential Generator Output



## 2.2 Rejection method

```
In [ ]:
        outcomes = [1, 2, 3, 4, 5, 6]
        probabilities = [7/48, 5/48, 1/8, 1/16, 1/4, 5/16]
        # We determine minimum value for C in order to accept as many rolls as po
        c = np.max(probabilities)
        rolls = []
        start_time = time.time()
        while len(rolls) < 10000:</pre>
            U1 = np.random.uniform()
            U2 = np.random.uniform()
            I = np.floor(6 * U1) + 1
            if U2 <= probabilities[int(I-1)] / c:</pre>
                 rolls.append(int(I))
        end_time = time.time()
        rejection_time = end_time - start_time
        rejection_method = rolls
        # Plot the histogram
        sns.histplot(rolls)
        plt.xlabel('Random Number')
        plt.ylabel('Frequency')
```

```
plt.title('Histogram of Linear Congruential Generator Output')
plt.show()
```

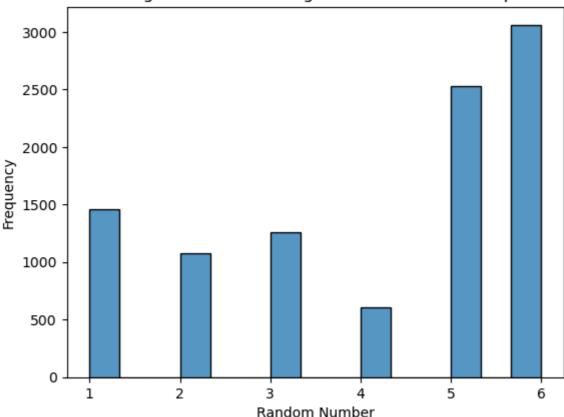


## 2.3 Alias method

```
In [ ]: import numpy as np
        from tqdm import tqdm
        import matplotlib.pyplot as plt
        import seaborn as sns
        p = [7/48, 5/48, 1/8, 1/16, 1/4, 5/16]
        L = [1, 2, 3, 4, 5, 6]
        F = np.array(p) * n # We scale the probabilities
        alias = np.zeros(n, dtype=int) # Alias
        prob = np.zeros(n)
        G = [] \# Big
        S = [] # Small
        start_time = time.time()
        for i, f in enumerate(F):
            if f >= 1.0:
                G.append(i)
            else:
                S.append(i)
        while G and S:
            i = G.pop()
                        # Big index
            j = S.pop() # Small index
```

```
# Add indexes where scaled probabilities F is over 1 to the alias ind
            #at the locations where the scaled probabilities are smaller
            alias[j] = i
            # Add the scaled probability for the smaller indexes in the smaller i
            prob[j] = F[j]
            F[i] = F[i] - (1 - F[j])
            if F[i] >= 1.0:
                G.append(i)
            else:
                S.append(i)
        # For remaining elements in G and S
        while G:
            i = G.pop()
            prob[i] = 1.0
        while S:
            i = S.pop()
            prob[i] = 1.0
        # List to store rolls
        rolls = []
        # Draw rolls using the loop structure
        for _ in tqdm(range(10000)):
            a = True
            while a:
                U1 = np.random.uniform()
                I = int(np.floor(6 * U1) + 1) # Choose a random number from the
                U2 = np.random.rand() # Generate a uniform random number
                if U2 < prob[I-1]:
                     rolls.append(I) # Append the index, +1 for 1-based indexing
                else:
                    rolls.append(alias[I-1] + 1) # Append the alias, +1 for 1-ba
                a = False # Exit the while loop
        end_time = time.time()
        alias_time = end_time - start_time
        alias_method = rolls
                      | 10000/10000 [00:00<00:00, 526896.14it/s]
       100%||
       100%
                      1 10000/10000 [00:00<00:00, 526896.14it/s]</pre>
In [ ]: # Plot the results
        sns.histplot(rolls)
        plt.xlabel('Random Number')
        plt.ylabel('Frequency')
        plt.title('Histogram of Linear Congruential Generator Output')
        plt.show()
```

#### Histogram of Linear Congruential Generator Output



```
In [ ]: def count_frequencies(arr):
            counts = np.bincount(arr)[1:]
            return counts
In [ ]: # Count frequencies for each method
        # Create the perfect_counts array and round it
        perfect_counts = np.round(np.array(p) * 10_000, 0)
        perfect_counts = perfect_counts.astype(int)
        numpy_counts = count_frequencies(numpy_distribution)
        crude_counts = count_frequencies(crude_method)
        rejection_counts = count_frequencies(rejection_method)
        alias_counts = count_frequencies(alias_method)
        # Create a table of frequencies
        table = np.array([perfect_counts, numpy_counts, crude_counts, rejection_c
        columns = ['1', '2', '3', '4', '5', '6']
        methods = ['Perfect Distribution', 'Numpy Distribution', 'Crude Method', '
        df = pd.DataFrame(table, columns=columns, index=methods)
        print(df)
                                1
                                      2
                                             3
                                                  4
                                                        5
                                                              6
       Perfect Distribution 1458
                                   1042
                                         1250
                                               625
                                                     2500 3125
       Numpy Distribution
                             1469
                                   1033
                                         1293
                                               599
                                                     2493
                                                           3113
       Crude Method
                             1436
                                   1060
                                         1244
                                               609
                                                     2501
                                                           3150
```

To compare the distributions we can perform a chi square test

1460

1459

Rejection Method

Alias Method

```
In [ ]: # Chi-Square Test
    chi2_stat, p_val, dof, ex = chi2_contingency(df)
```

1236

1263

657

607

2557

2532

3083

3062

1007

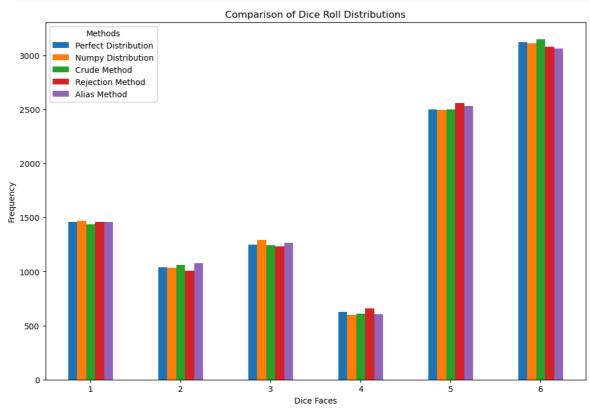
1077

```
print("Chi-Square Statistic:", chi2_stat)
print("p-value:", p_val)
print("Degrees of Freedom:", dof)
```

Chi-Square Statistic: 10.859450719310086

p-value: 0.9497862181882478
Degrees of Freedom: 20

```
In []: # Plot the distributions
    df.T.plot(kind='bar', figsize=(12, 8))
    plt.title('Comparison of Dice Roll Distributions')
    plt.xlabel('Dice Faces')
    plt.ylabel('Frequency')
    plt.xticks(rotation=0)
    plt.legend(title='Methods')
    plt.show()
```



There is no statisticaly significant difference between the distribution methods.

We can compare the time it takes between the different methods

```
In []: time_data = {
    'Method': ['Numpy', 'Crude', 'Rejection', 'Alias'],
    'Elapsed Time (seconds)': [numpy_time, crude_time, rejection_time, al
}

# Convert the dictionary to a DataFrame
df = pd.DataFrame(time_data)
print(df)
```

```
Method Elapsed Time (seconds)

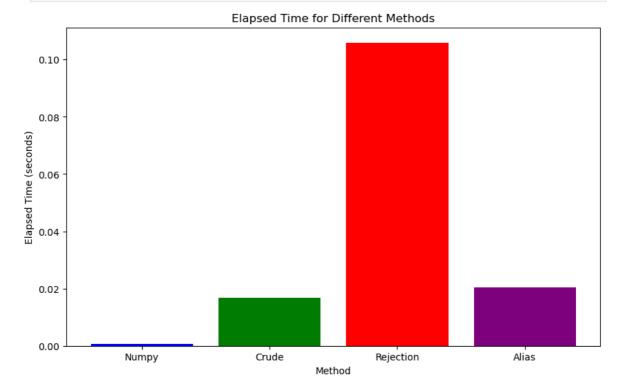
Numpy 0.000540

Crude 0.016685

Rejection 0.105927

Alias 0.020444
```

```
In []: # Plot the times
plt.figure(figsize=(10, 6))
plt.bar(df['Method'], df['Elapsed Time (seconds)'], color=['blue', 'green
plt.xlabel('Method')
plt.ylabel('Elapsed Time (seconds)')
plt.title('Elapsed Time for Different Methods')
plt.show()
```



We notice that there is quite a large difference in time between the methods. Especially we see that the rejection method takes the longest time to run by far. The crude and alias methods are very similar in our examples, and the numpy method is fastest by a large factor