

# Project

October 15, 2024

## 1 №2

### 1.1 ,

#### 1.1.1 -02

```
[38]: #jupyter nbconvert --to markdown Project.ipynb
```

### 1.2

```
[39]: import numpy as np
import matplotlib.pyplot as plt
from scipy import stats
```

## 2 №1

### 2.1

```
[40]: exp_sample_100 = np.random.exponential(scale=1, size=100)
exp_sample_1000 = np.random.exponential(scale=1, size=1000)

norm_sample_100 = np.random.normal(loc=0, scale=1, size=100)
norm_sample_1000 = np.random.normal(loc=0, scale=1, size=1000)
```

### 2.2

- 1.
- 2.

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2. , ..

#### 2.2.1 numpy

```
[41]: def calculate_mean_variance(sample, distribution_name):
    sample_mean = np.mean(sample)
    sample_variance = np.var(sample, ddof=1)
```

```

if distribution_name == "exponential":
    theoretical_mean = 1
    theoretical_variance = 1
elif distribution_name == "normal":
    theoretical_mean = 0
    theoretical_variance = 1

print(f"                ({distribution_name}): {theoretical_mean}")
print(f"                ({distribution_name}, n={len(sample)}): {sample_mean:.4f}")
print(f"                ({distribution_name}): {theoretical_variance}")
print(f"                ({distribution_name}, n={len(sample)}): {sample_variance:.4f}")
print("")

```

## 2.3

```

[42]: calculate_mean_variance(exp_sample_100, "exponential")
      calculate_mean_variance(exp_sample_1000, "exponential")
      calculate_mean_variance(norm_sample_100, "normal")
      calculate_mean_variance(norm_sample_1000, "normal")

```

```

(exponential): 1
(exponential, n=100): 0.9101
(exponential): 1
(exponential, n=100): 0.5744

(exponential): 1
(exponential, n=1000): 0.9969
(exponential): 1
(exponential, n=1000): 0.9619

(normal): 0
(normal, n=100): -0.0815
(normal): 1
(normal, n=100): 1.2893

(normal): 0
(normal, n=1000): 0.0251
(normal): 1
(normal, n=1000): 0.9782

```

## 2.4

```
[43]: def calculate_quantiles(sample, distribution_name):
        sample_quantiles = np.quantile(sample, [0.5, 0.99])

        if distribution_name == "exponential":
            theoretical_quantiles = stats.expon.ppf([0.5, 0.99], scale=1)
        elif distribution_name == "normal":
            theoretical_quantiles = stats.norm.ppf([0.5, 0.99], loc=0, scale=1)

        print(f"                ({distribution_name}, n={len(sample)}):␣
↪{sample_quantiles}")
        print(f"                ({distribution_name}): {theoretical_quantiles}")
        print("\n")
```

```
[44]: calculate_quantiles(exp_sample_100, "exponential")
        calculate_quantiles(exp_sample_1000, "exponential")
        calculate_quantiles(norm_sample_100, "normal")
        calculate_quantiles(norm_sample_1000, "normal")
```

```
(exponential, n=100): [0.79599225  3.10469233]
(exponential): [0.69314718  4.60517019]
```

```
(exponential, n=1000): [0.69396014  4.1181586 ]
(exponential): [0.69314718  4.60517019]
```

```
(normal, n=100): [-0.11287732  2.37344646]
(normal): [0.          2.32634787]
```

```
(normal, n=1000): [-1.7143188e-03  2.2685730e+00]
(normal): [0.          2.32634787]
```

## 2.5

!

```
[45]: def plot_histogram(sample, distribution_name, n):
        plt.figure(figsize=(8,6))
        plt.hist(sample, bins=30, density=True, alpha=0.6, color="skyblue",␣
↪label="                ")

        x = np.linspace(min(sample), max(sample), 1000)
        if distribution_name == "exponential":
```

```

pdf = stats.expon.pdf(x, scale=1)
elif distribution_name == "normal":
    pdf = stats.norm.pdf(x, loc=0, scale=1)
plt.plot(x, pdf, "r", label="")

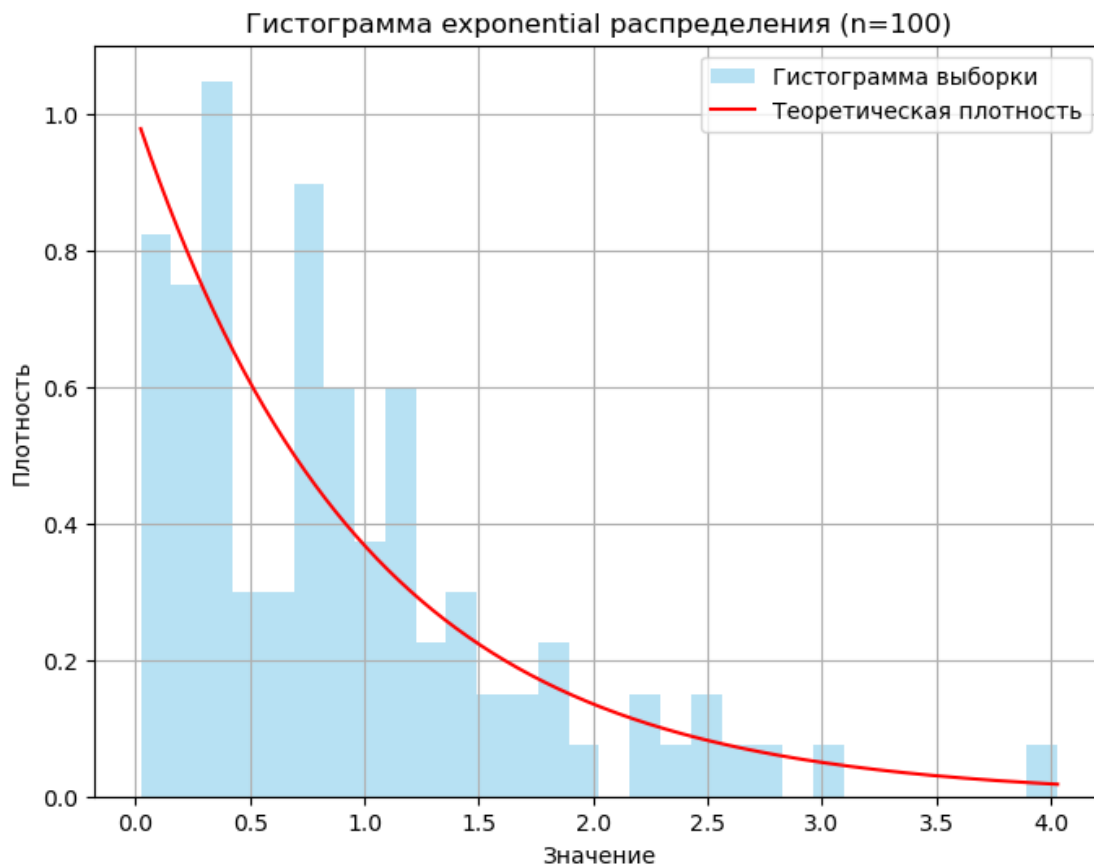
plt.title(f"      {distribution_name}      (n={n})")
plt.xlabel("      ")
plt.ylabel("      ")
plt.legend()
plt.grid(True)
plt.show()

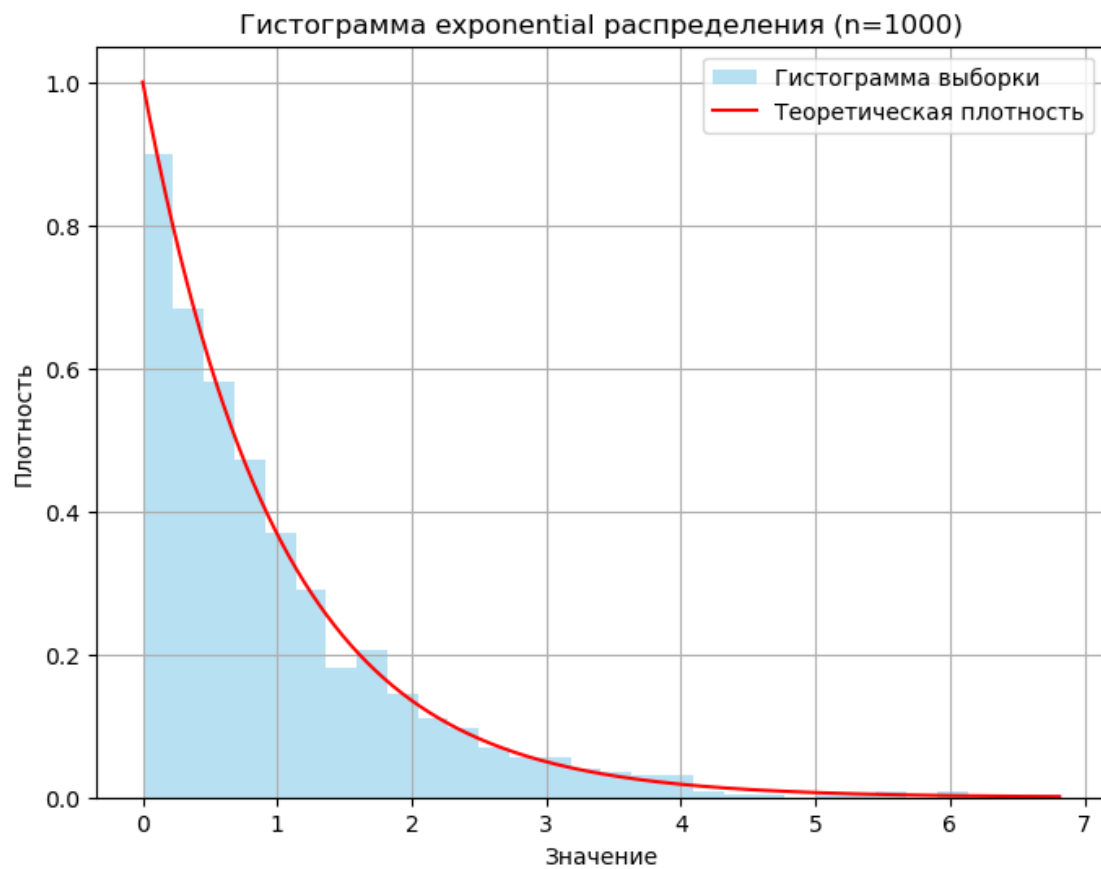
```

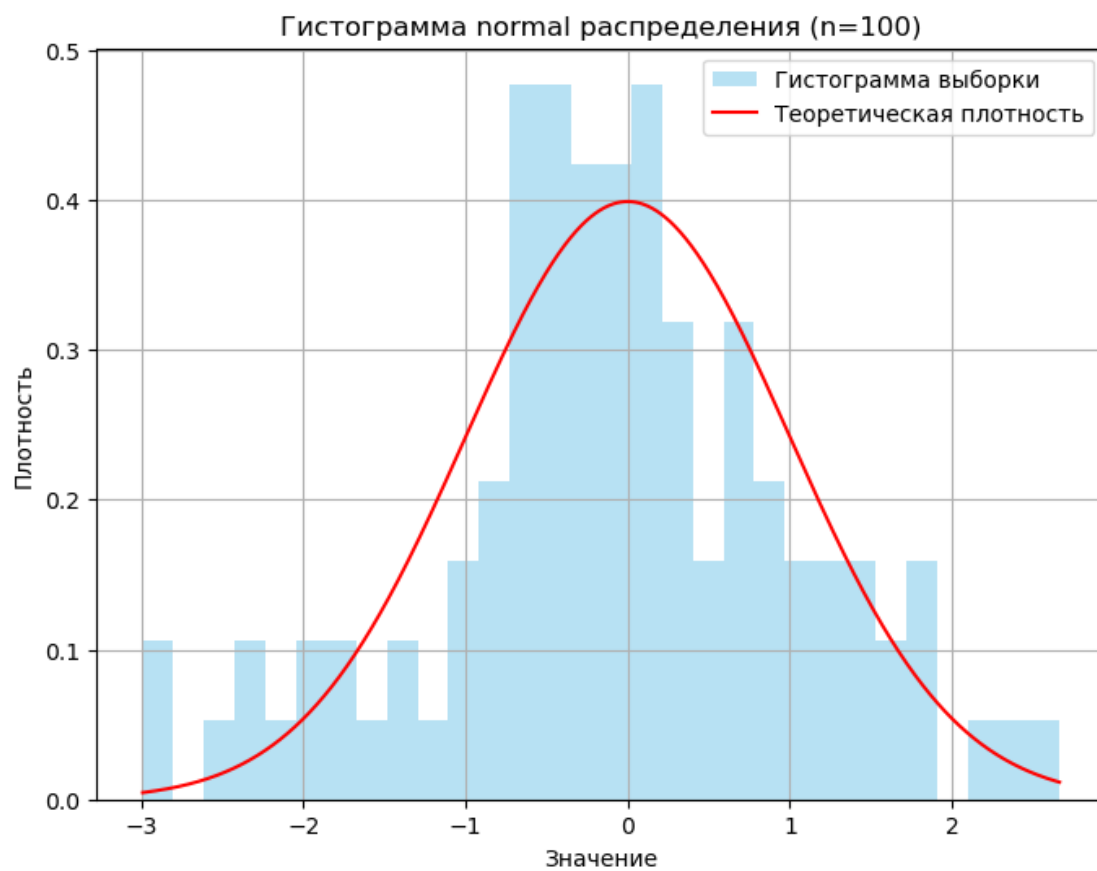
```

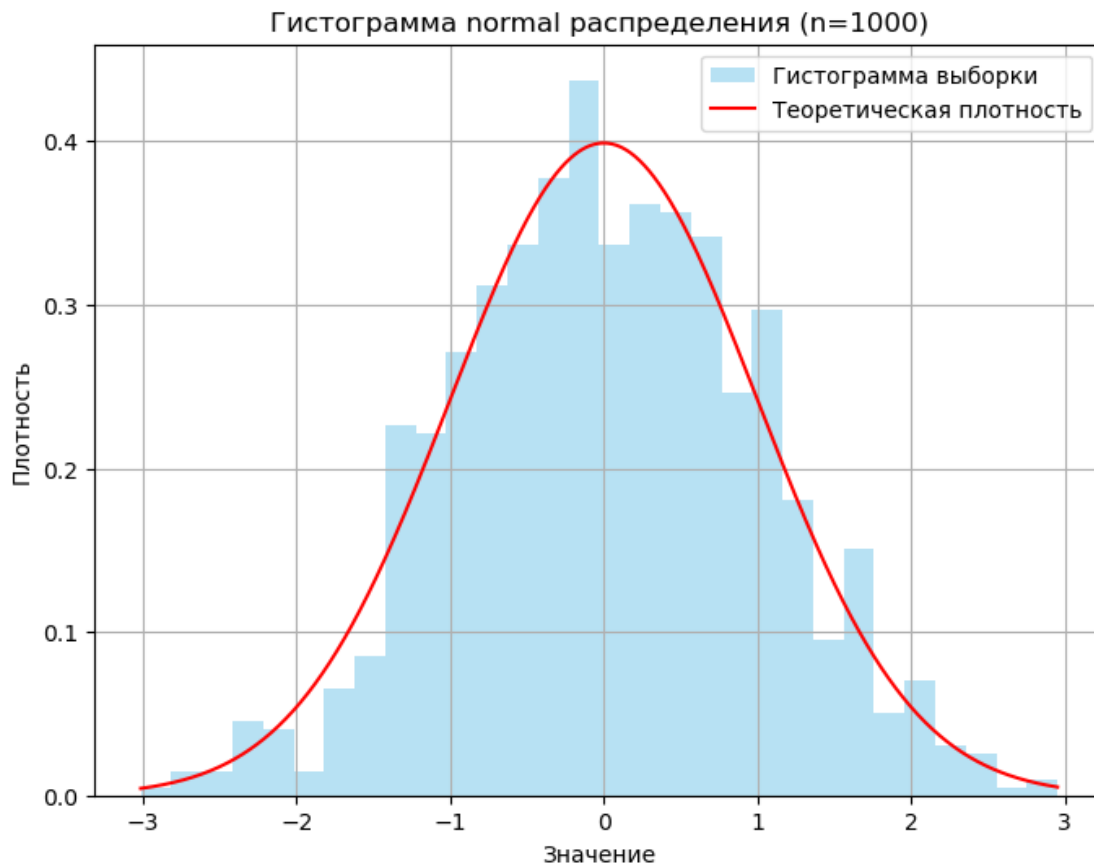
[46]: plot_histogram(exp_sample_100, "exponential", 100)
      plot_histogram(exp_sample_1000, "exponential", 1000)
      plot_histogram(norm_sample_100, "normal", 100)
      plot_histogram(norm_sample_1000, "normal", 1000)

```









## 2.6

```
[47]: def plot_cdf(sample_100, sample_1000, distribution_name):
    sorted_sample_100 = np.sort(sample_100)
    sorted_sample_1000 = np.sort(sample_1000)

    ecdf_100 = np.arange(1, len(sorted_sample_100)+1) / len(sorted_sample_100)
    ecdf_1000 = np.arange(1, len(sorted_sample_1000)+1) /
    ↪ len(sorted_sample_1000)

    plt.figure(figsize=(8,6))
    plt.step(sorted_sample_100, ecdf_100, where="post", label="
    ↪ (n=100)")
    plt.step(sorted_sample_1000, ecdf_1000, where="post", label="
    ↪ (n=1000)")

    x = np.linspace(min(np.min(sorted_sample_100), np.min(sorted_sample_1000)),
    ↪ max(np.max(sorted_sample_100), np.max(sorted_sample_1000)),
    ↪ 1000)
```

```

if distribution_name == "exponential":
    cdf = stats.expon.cdf(x, scale=1)
elif distribution_name == "normal":
    cdf = stats.norm.cdf(x, loc=0, scale=1)
plt.plot(x, cdf, "r", label="")

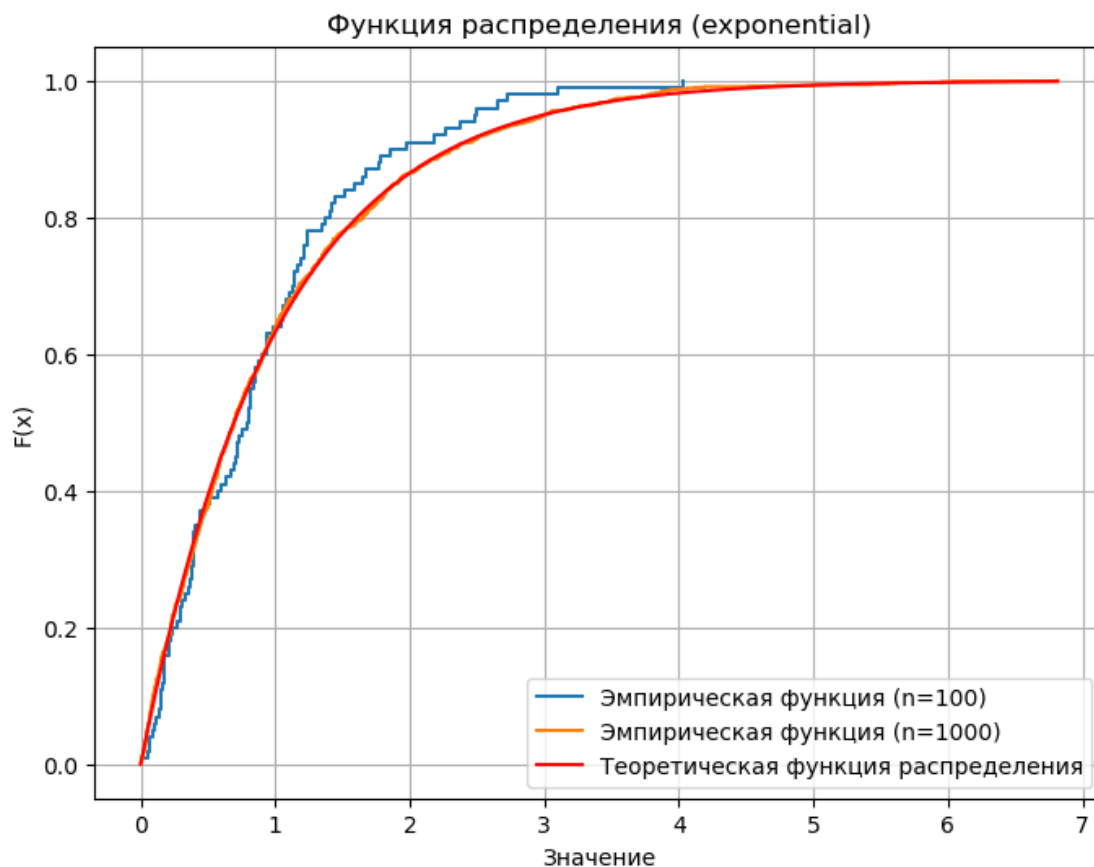
plt.title(f"({distribution_name})")
plt.xlabel("")
plt.ylabel("F(x)")
plt.legend()
plt.grid(True)
plt.show()

```

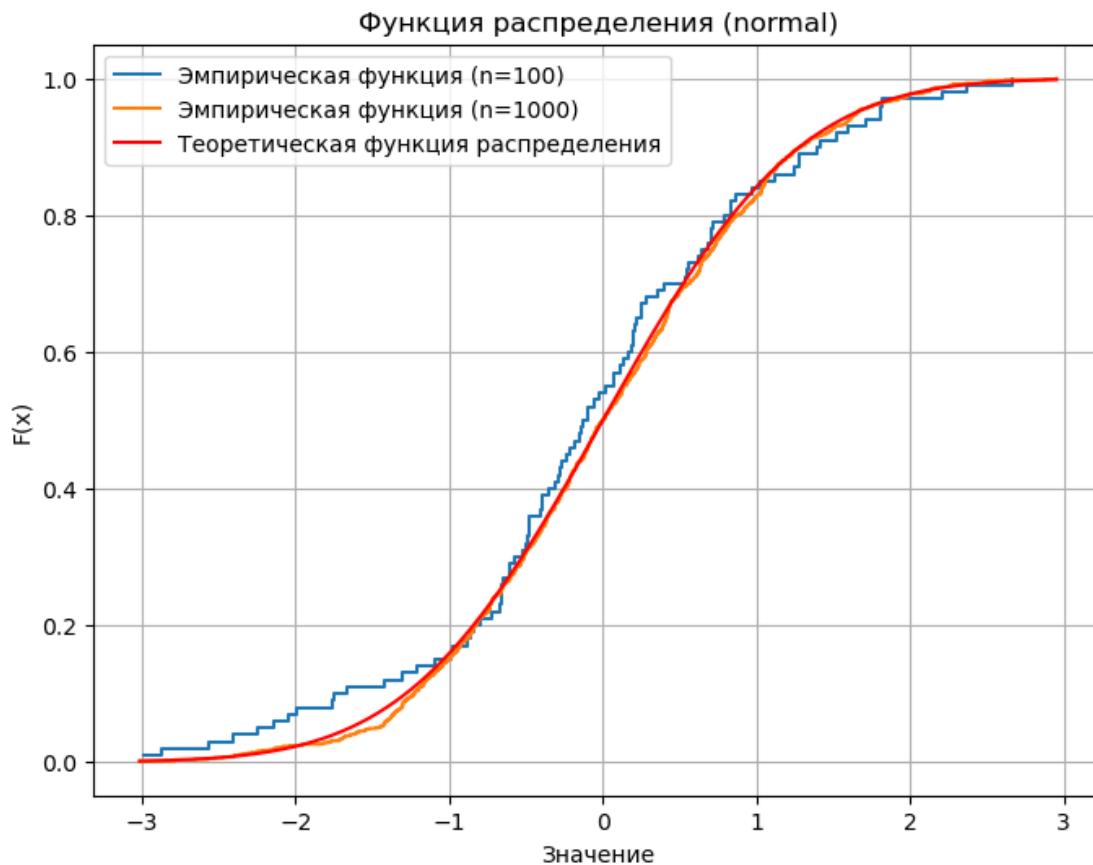
```

[48]: plot_cdf(exp_sample_100, exp_sample_1000, "exponential")
      plot_cdf(norm_sample_100, norm_sample_1000, "normal")

```







## 2.7

```
[49]: def plot_density(sample_100, sample_1000, distribution_name):
    from scipy.stats import gaussian_kde

    density_100 = gaussian_kde(sample_100)
    density_1000 = gaussian_kde(sample_1000)

    x = np.linspace(min(np.min(sample_100), np.min(sample_1000)),
                    max(np.max(sample_100), np.max(sample_1000)), 1000)

    plt.figure(figsize=(8,6))
    plt.plot(x, density_100(x), label="n=100")
    plt.plot(x, density_1000(x), label="n=1000")

    if distribution_name == "exponential":
        pdf = stats.expon.pdf(x, scale=1)
```

```

elif distribution_name == "normal":
    pdf = stats.norm.pdf(x, loc=0, scale=1)
    plt.plot(x, pdf, 'r', label="")

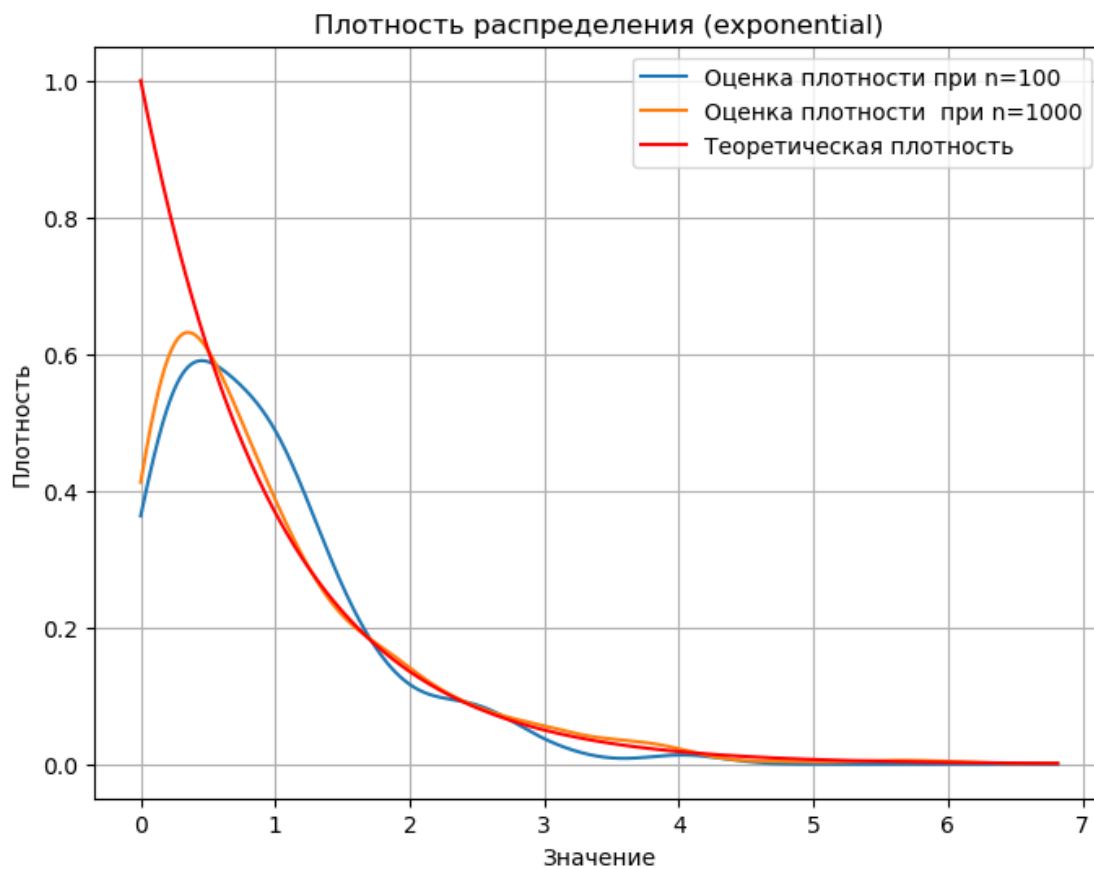
plt.title(f"({distribution_name})")
plt.xlabel("")
plt.ylabel("")
plt.legend()
plt.grid(True)
plt.show()

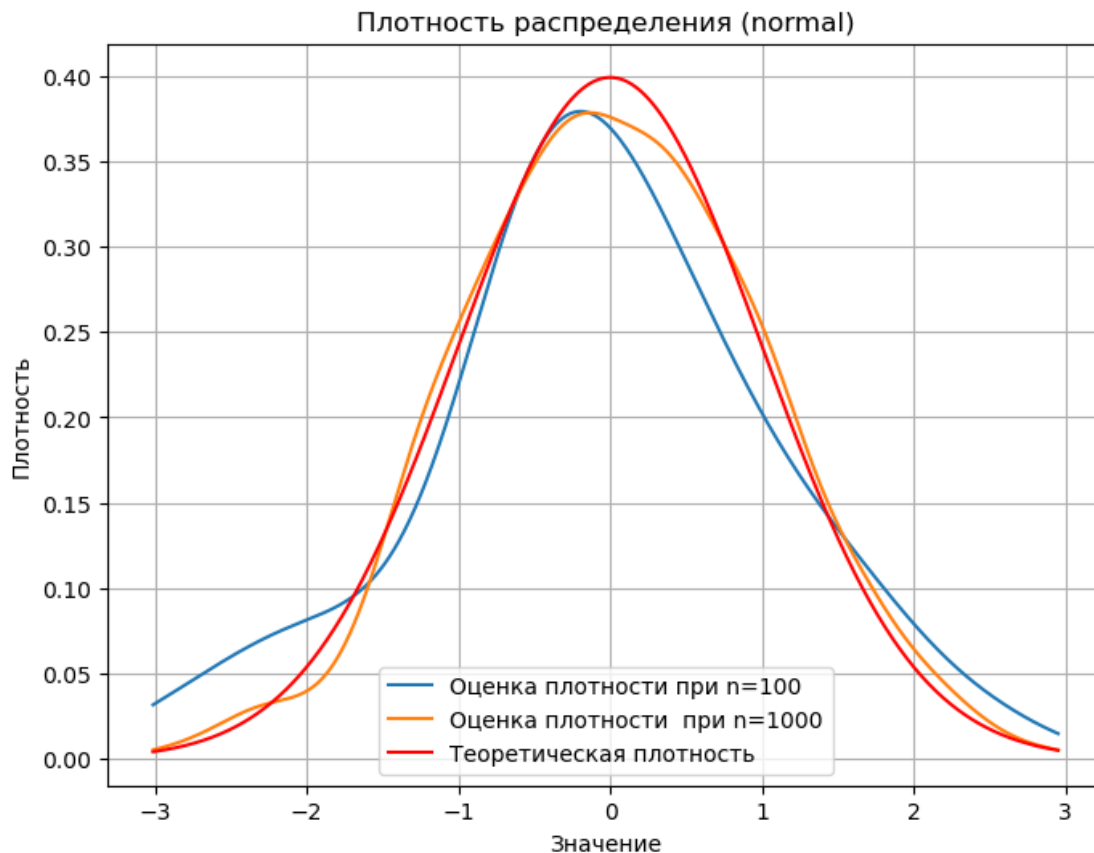
```

```

[50]: plot_density(exp_sample_100, exp_sample_1000, "exponential")
      plot_density(norm_sample_100, norm_sample_1000, "normal")

```





### 3 №2

100, 1000 10000  
10 30.

#### 3.1 2

```
[51]: def generate_points(sample_size, width=10, height=30):
    x1 = np.random.uniform(0, width, sample_size)
    y1 = np.random.uniform(0, height, sample_size)
    x2 = np.random.uniform(0, width, sample_size)
    y2 = np.random.uniform(0, height, sample_size)
    distances = np.sqrt((x1 - x2)**2 + (y1 - y2)**2)
    return distances
```

```
[52]: sizes = [100, 1000, 10000]
distance_samples = {}

for size in sizes:
    distances = generate_points(size, width=10, height=30)
    distance_samples[size] = distances
```

```
[55]: from scipy.stats import gaussian_kde
def plot_cdf(samples_dict):
    plt.figure(figsize=(10,6))
    for size, distances in samples_dict.items():
        sorted_distances = np.sort(distances)
        ecdf = np.arange(1, len(sorted_distances)+1) / len(sorted_distances)
        plt.step(sorted_distances, ecdf, where="post", label=f"n={size}")

    plt.title("CDF of distances")
    plt.xlabel("distance")
    plt.ylabel("F(x)")
    plt.legend()
    plt.grid(True)
    plt.show()

def plot_pdf(samples_dict):
    plt.figure(figsize=(10,6))
    x_min = min([min(distances) for distances in samples_dict.values()])
    x_max = max([max(distances) for distances in samples_dict.values()])
    x = np.linspace(x_min, x_max, 1000)

    for size, distances in samples_dict.items():
        density = gaussian_kde(distances)
        plt.plot(x, density(x), label=f'n={size}')

    plt.title("PDF of distances")
    plt.xlabel("distance")
    plt.ylabel("density")
    plt.legend()
    plt.grid(True)
    plt.show()
```

```
[56]: plot_cdf(distance_samples)
plot_pdf(distance_samples)
```

