

exercicio-06-giuliavieira

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UNIVERSIDADE FEDERAL DE MINAS GERAIS INSTITUTO DE CIÊNCIAS EXATAS
GRADUAÇÃO EM CIÊNCIA DA COMPUTAÇÃO DISCIPLINA: Introdução a Física Estatística e Computacional

ALUNA: Giulia Monteiro Silva Gomes Vieira MATRICULA: 2016006492

EXERCÍCIO AVALIATIVO 06: Propagação de doenças

```
[1]: import numpy as np
import matplotlib.pyplot as plt
import matplotlib.colors as mcolors
from matplotlib.animation import FuncAnimation
from PIL import Image
import imageio
import os
```

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[2]: #HEALTHY = 0
#INFECTED = 1
#RECOVERED = 2
```

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[3]: def initialize_network(size):
    network = np.zeros((size, size), dtype=int)
    infected_individual = np.random.randint(0, size**2)
    network.flat[infected_individual] = 1
    return network
```

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[4]: def sir_simulation(size, p_infection, p_recovery, num_steps):
    network = initialize_network(size)
    history = []

    for step in range(num_steps):
        new_network = network.copy()

        for i in range(size):
            for j in range(size):
                if network[i, j] == 0: # Indivíduo suscetível
                    neighbors = [
                        network[(i - 1) % size, j],
                        network[(i + 1) % size, j],
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        network[i, (j - 1) % size],
        network[i, (j + 1) % size]
    ]

    # Regras do modelo SIR
    if 1 in neighbors and np.random.rand() < p_infection:
        new_network[i, j] = 1 # infecta individuo
    elif network[i, j] == 1: # Indivíduo infectado
        if np.random.rand() < p_recovery:
            new_network[i, j] = 2 # recupera individuo

    network = new_network
    history.append(network.copy())

    return np.array(history)

```

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[5]: # Parâmetros da simulação
size = 20
p_infection_values = [0.1, 0.3, 0.5] # Escolha livremente
p_recovery_values = [0.05, 0.1, 0.2] # Escolha livremente
num_simulations = 10
num_steps = 50

cm = plt.get_cmap('Set2')

```

01. Médias das populações

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[11]: for p_infection in p_infection_values:

    fig, axes = plt.subplots(1, 3, figsize=(12, 6))
    fig.suptitle(f'Médias da população para p_infection = {p_infection}',
        ↪fontsize=15),

    output_directory = "sir_last_simulations"
    os.makedirs(output_directory, exist_ok=True)

    for i, p_recovery in enumerate(p_recovery_values):
        average_populations = np.zeros((num_simulations, 3))

        history = []
        for sim in range(num_simulations):
            history = sir_simulation(size, p_infection, p_recovery, num_steps)
            final_state = history[-1]

            s_population = np.sum(final_state == 0)
            i_population = np.sum(final_state == 1)
            r_population = np.sum(final_state == 2)

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        average_populations[sim] = [s_population, i_population,
↪r_population]

    fig_anim, ax_anim = plt.subplots(figsize=(6, 4))
    cmap = mcolors.ListedColormap(['white', cm(1), cm(2)])
    bounds = [0, 1, 2, 3]
    norm = mcolors.BoundaryNorm(bounds, cmap.N)
    frames = []

    for frame in range(len(history)):
        ax_anim.clear()
        ax_anim.imshow(history[frame], cmap=cmap, norm=norm)
        ax_anim.set_title(f'Time: {frame}')
        fig_anim.canvas.draw()

        # Convert figure to PIL Image and append to frames
        image = Image.frombytes('RGB', fig_anim.canvas.get_width_height(),
↪fig_anim.canvas.tostring_rgb())
        frames.append(image)

    # Save frames as animated GIF
    gif_filename = f'sir_simulation_{p_infection}_{p_recovery}-last.gif'
    gif_path = os.path.join(output_directory, gif_filename)
    imageio.mimsave(gif_path, frames, duration=0.5)

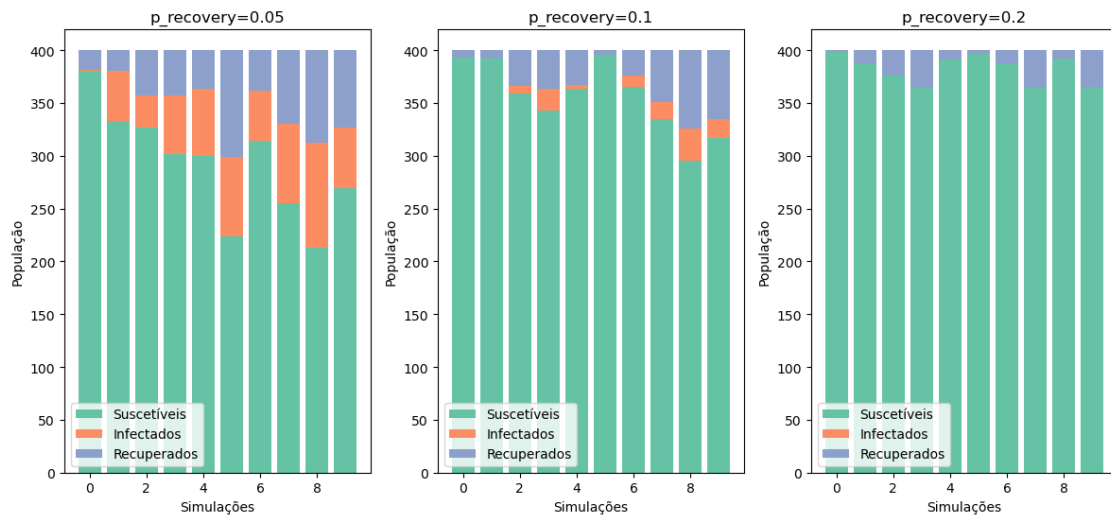
    plt.close(fig_anim)

    # Plot population evolution
    axes[i].bar(np.arange(num_simulations), average_populations[:, 0],
↪color = cm(0), label='Suscetíveis')
    axes[i].bar(np.arange(num_simulations), average_populations[:, 1],
↪bottom=average_populations[:, 0], color = cm(1), label='Infectados')
    axes[i].bar(np.arange(num_simulations), average_populations[:, 2],
↪bottom=average_populations[:, 0] + average_populations[:, 1], color = cm(2),
↪label='Recuperados')
    axes[i].set_xlabel('Simulações')
    axes[i].set_ylabel('População')
    axes[i].set_title(f'p_recovery={p_recovery}')
    axes[i].legend(loc='best')

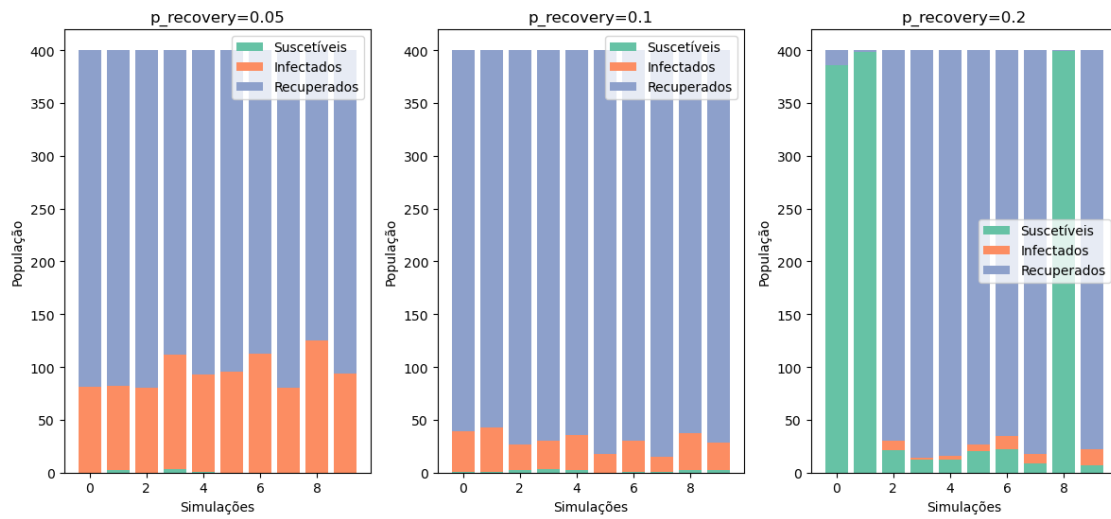
plt.tight_layout()
plt.show()

```

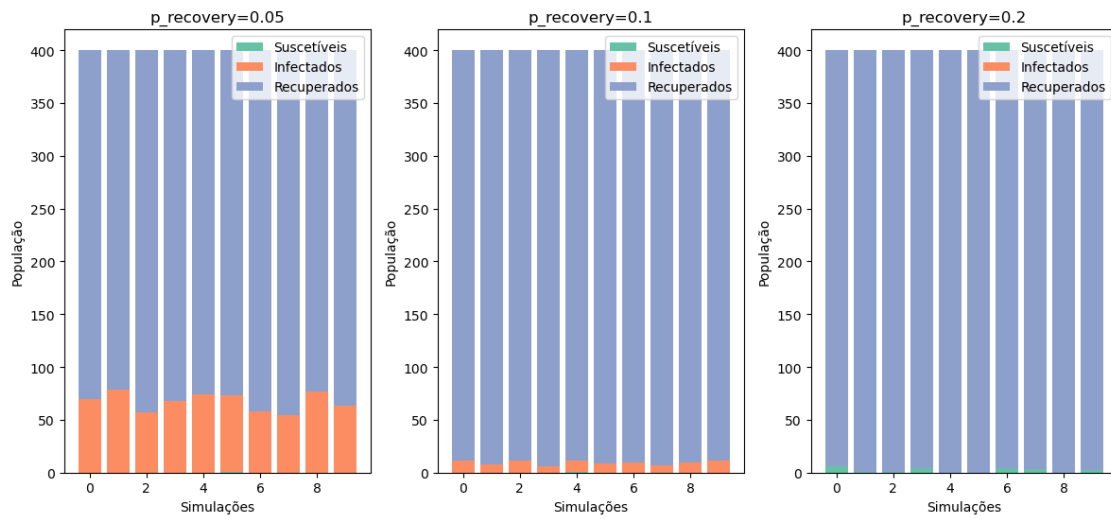
Médias da população para $p_{\text{infection}} = 0.1$



Médias da população para $p_{\text{infection}} = 0.3$



Médias da população para $p_{\text{infection}} = 0.5$



02. Comportamento geográfico na rede gifs podem ser encontrados aqui:
<https://www.dropbox.com/scl/fo/qyq3g55v1hkd3ixvdeu98/h?rlkey=7wl12o37z2b7g6295zcfaf1y7&dl=0>