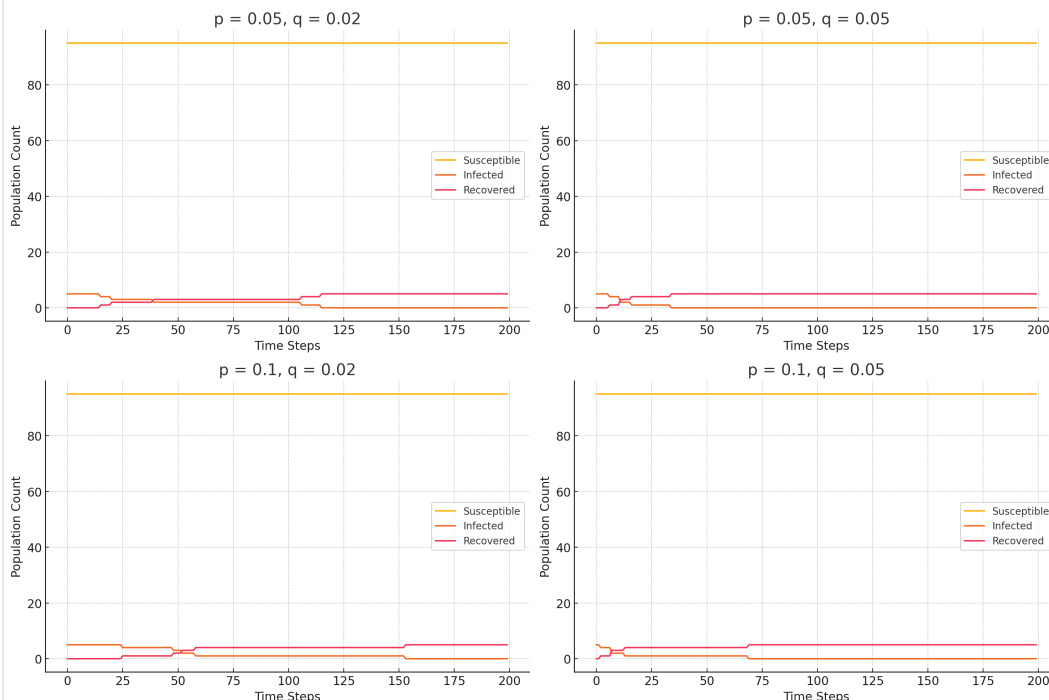




yes, try to implement in python



P = 0.1, Q = 0.05



Here's the simulation output, showing the infection dynamics over time with different combinations of infection probability p and recovery probability q . Each subplot corresponds to a specific p and q configuration, tracking the populations of susceptible, infected, and recovered agents across the 200 time steps.

This setup provides insights into how different infection and recovery rates impact the progression of the infection within the population. Let me know if you would like further analysis or adjustments! [\[>-\]](#)

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