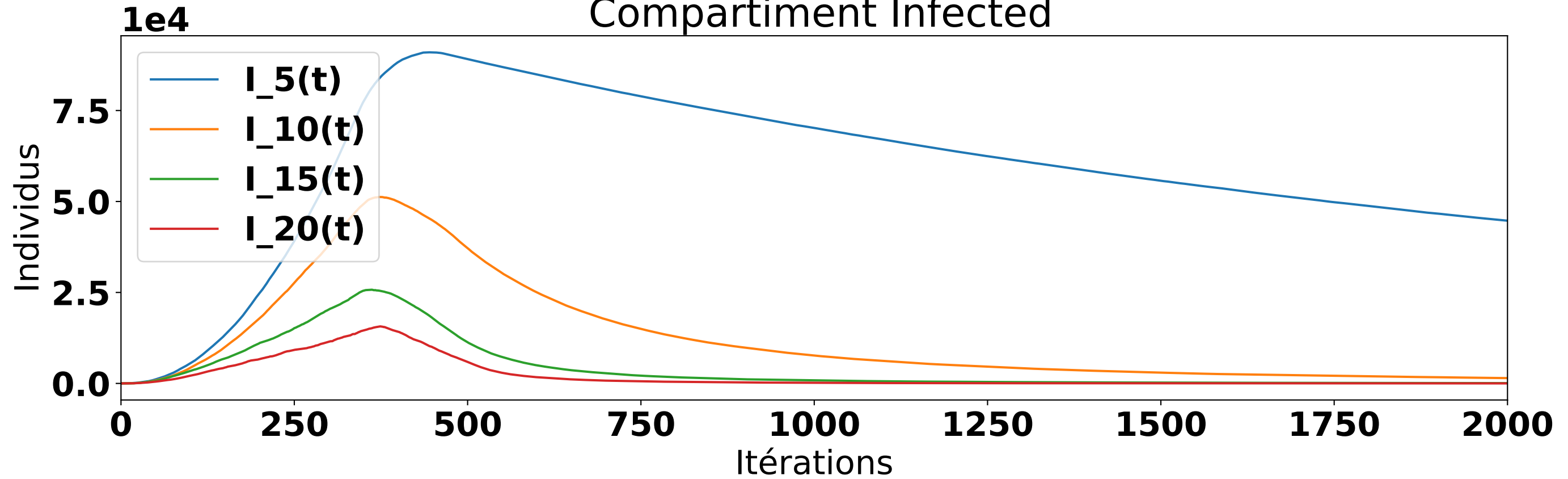
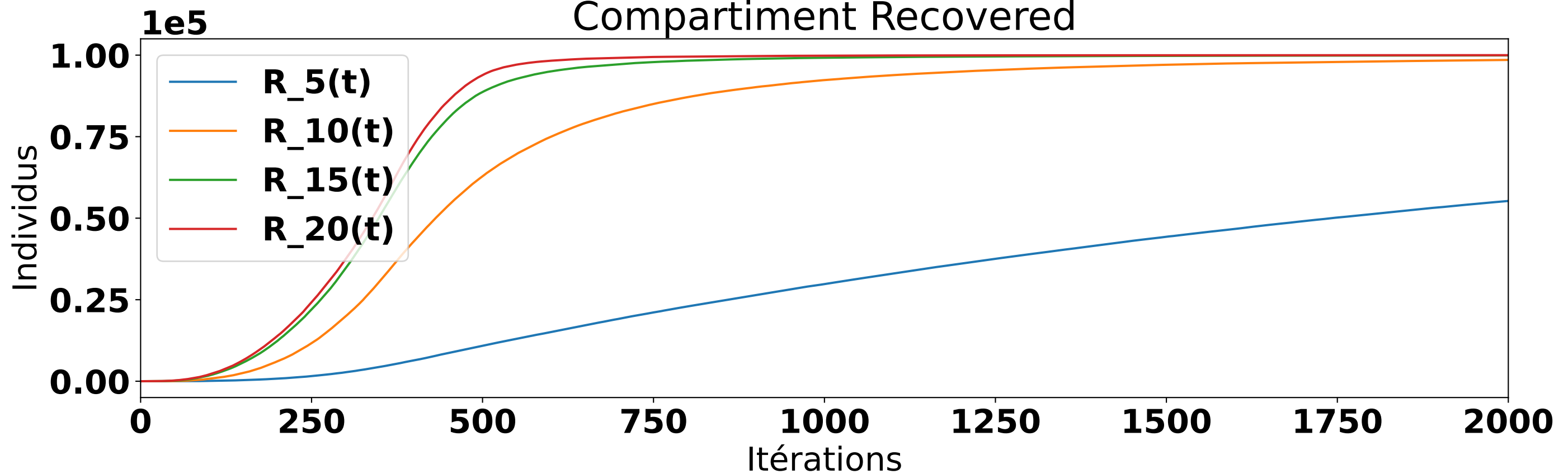


SIR_diversité, distance Hamming 0, densité 1/8, 100000 individus

Compartment Infected



Compartment Recovered



Compartment Susceptible

