Random walk to compute node visitation probabilities and edge flux scores
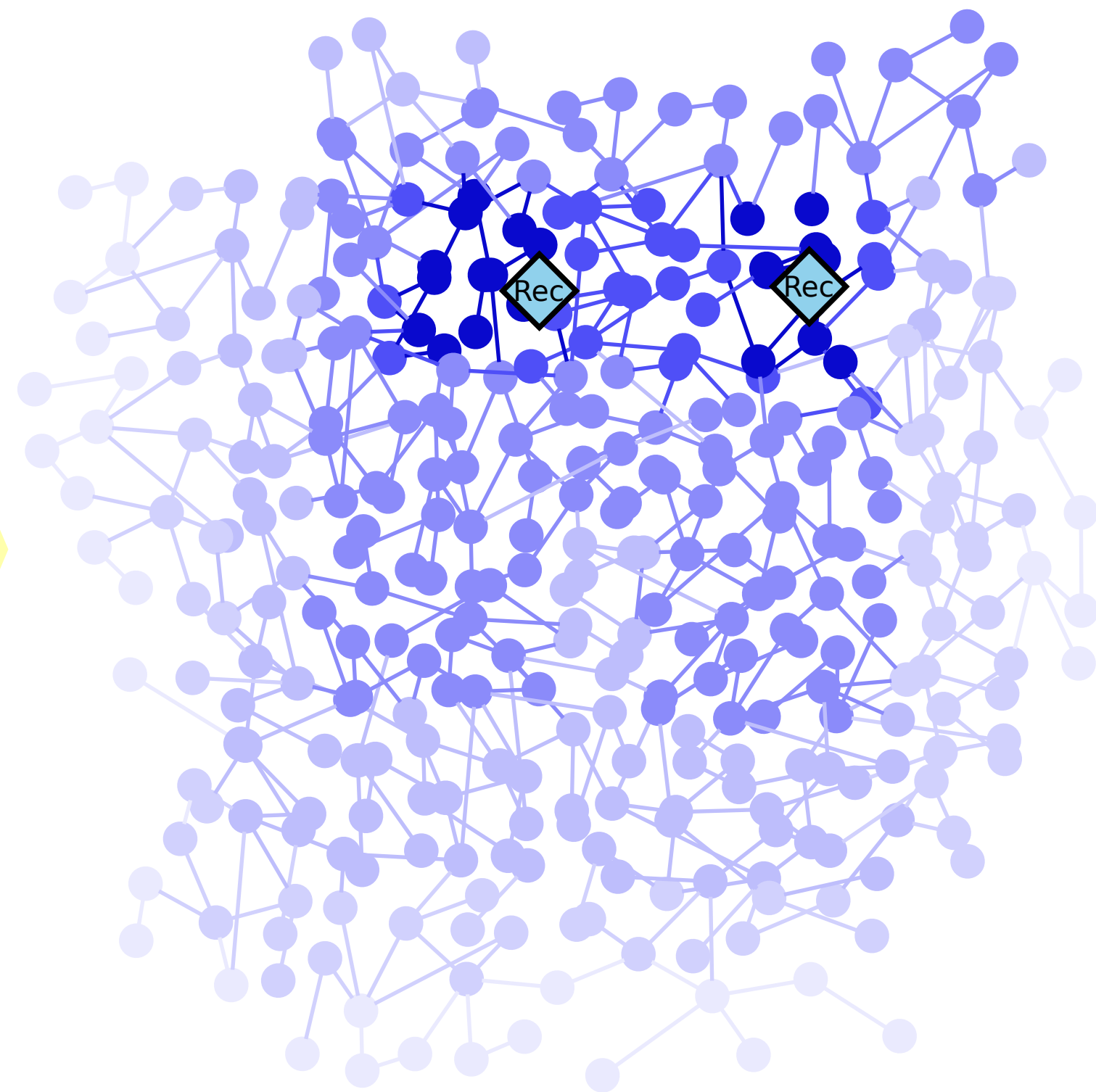
Human Interactome

Compute k paths with the largest edge flux scores from any receptor to any TR