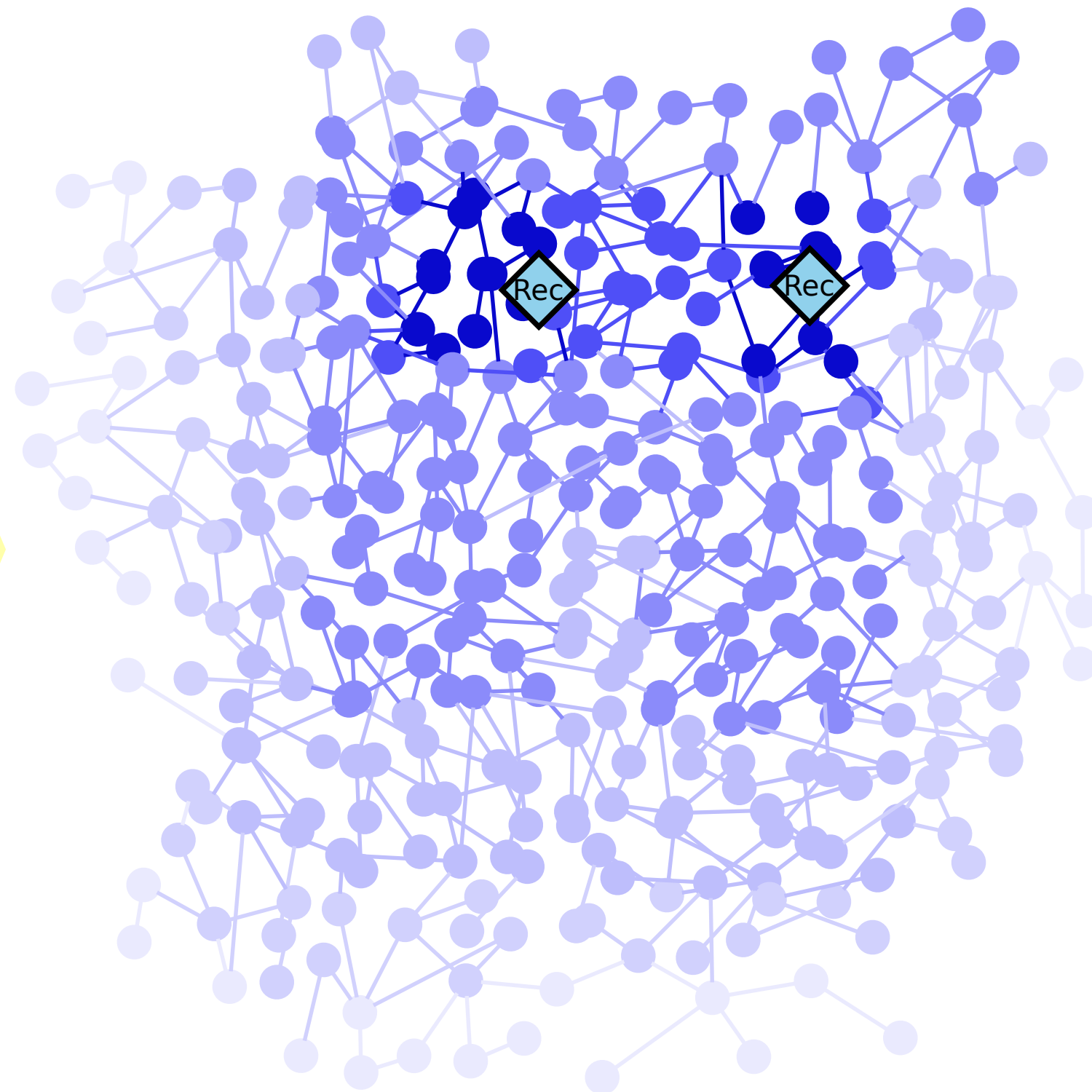


Human Interactome

PRFlux

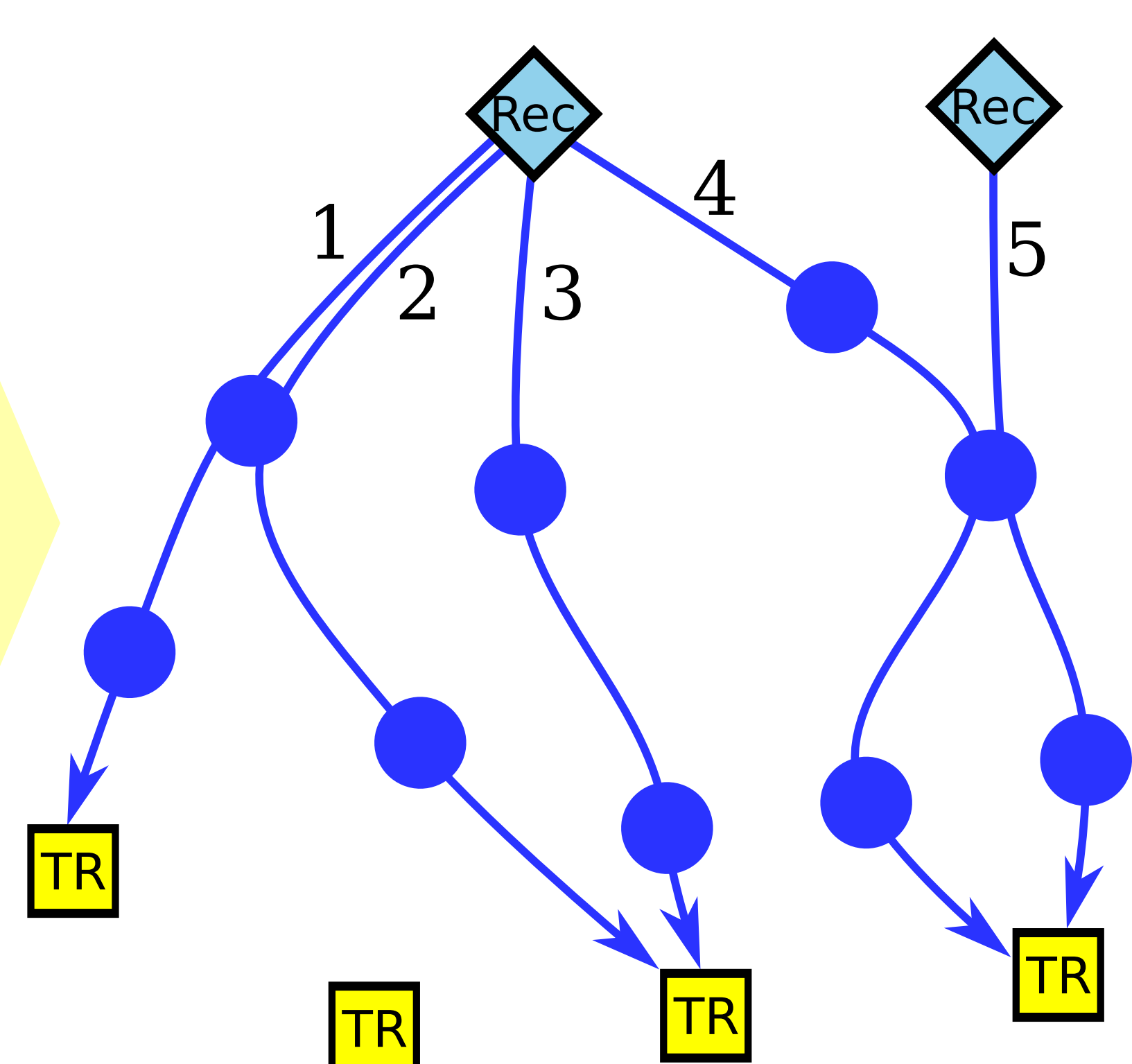
Random walk
to compute
node visitation
probabilities and
edge flux scores



Weighted Interactome

KSP

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



Computed Pathway