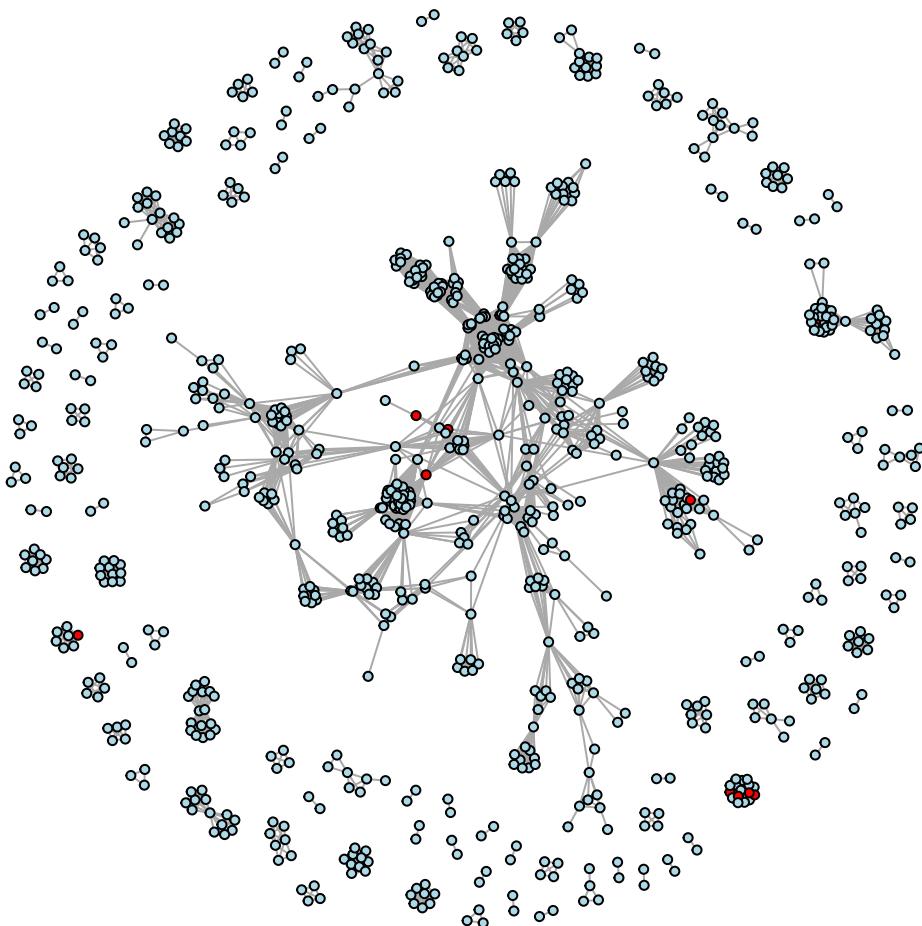


# Co-functional(EcoCyc)

Average path length = 4.1098.Number of vertexes = 885

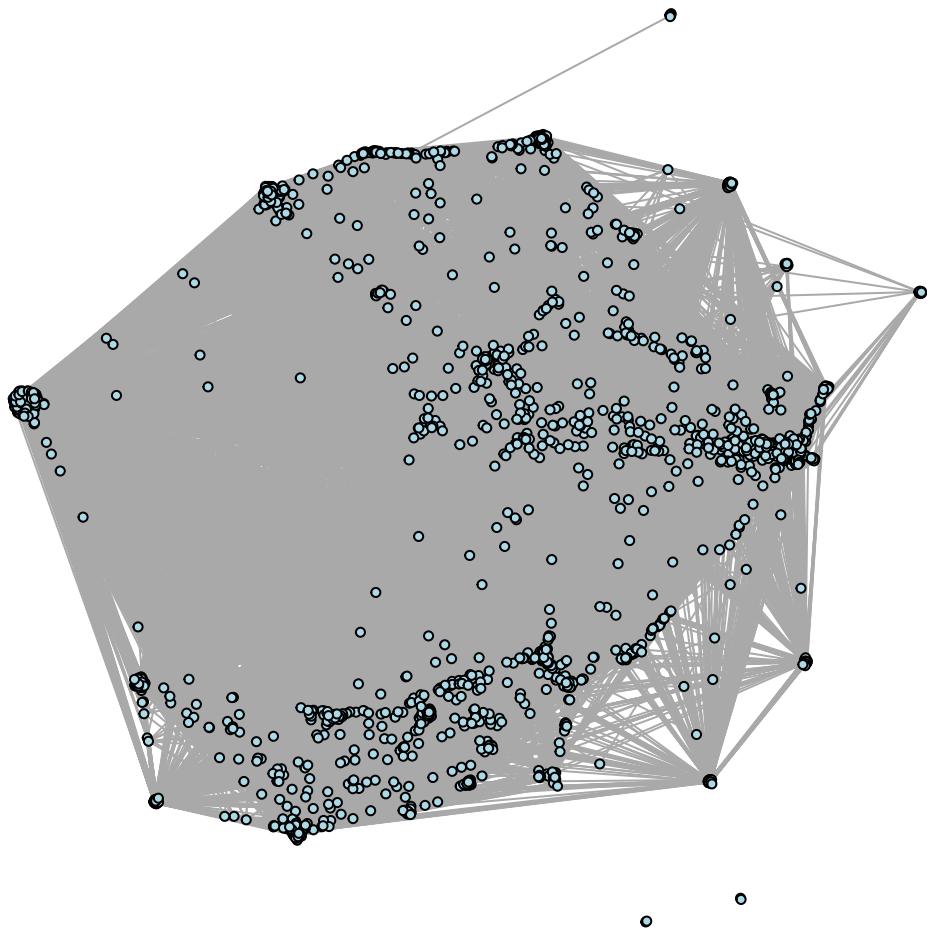
Number of targets = 10.Number of drugs = 16



# EN:allnetworks

Average path length = 2.8575.Number of vertexes = 4099

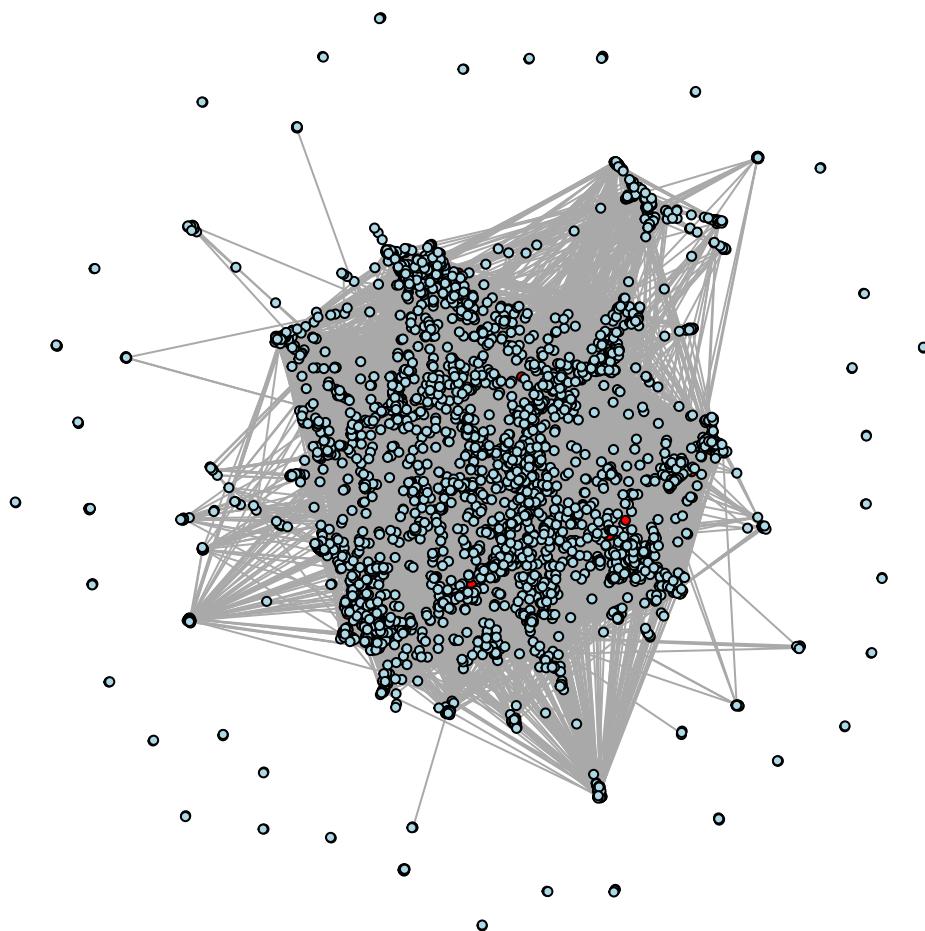
Number of targets = 25.Number of drugs = 36



# Similar genomic context

Average path length = 3.957.Number of vertexes = 3568

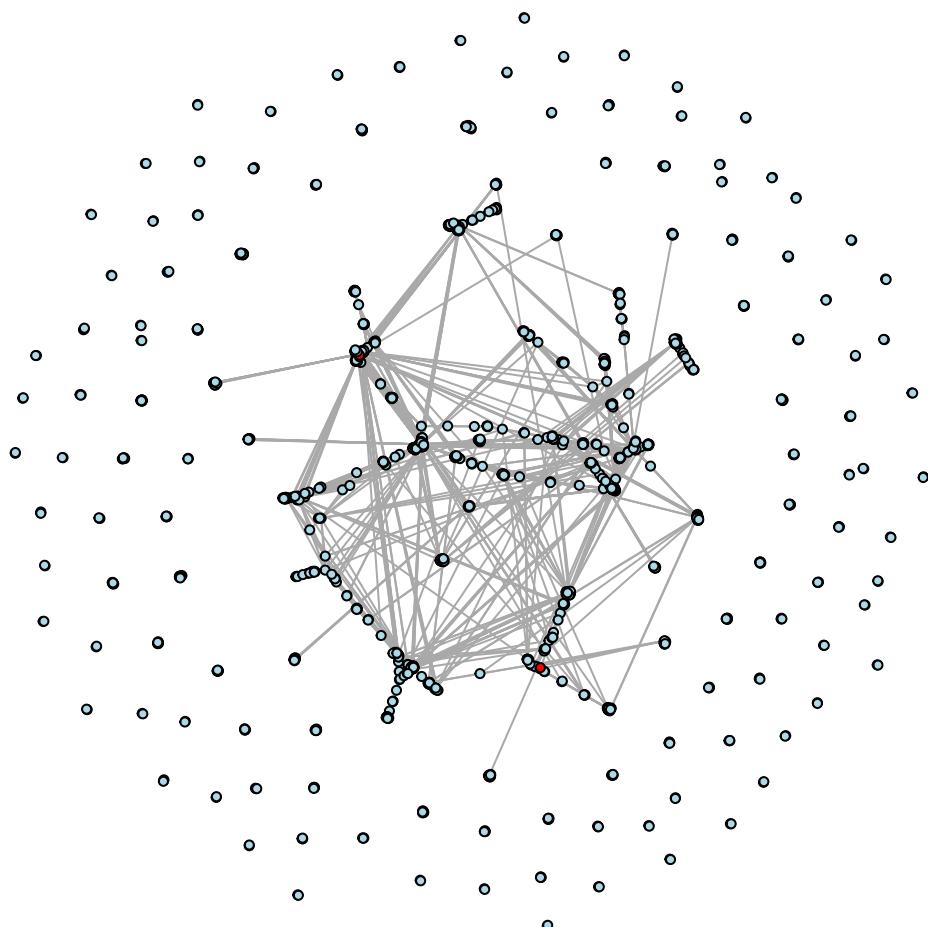
Number of targets = 25.Number of drugs = 36



# Co-functional (GO-BP)

Average path length = 4.7648.Number of vertexes = 1474

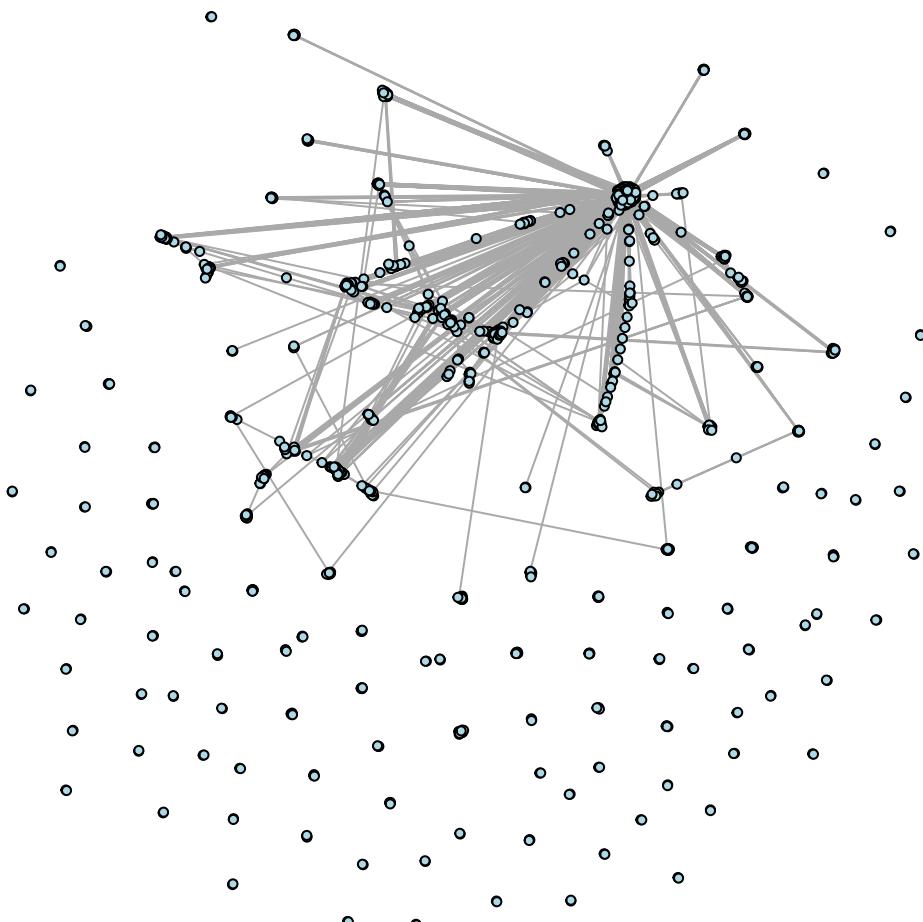
Number of targets = 17.Number of drugs = 31



# Co-citation

Average path length = 3.6296.Number of vertexes = 2296

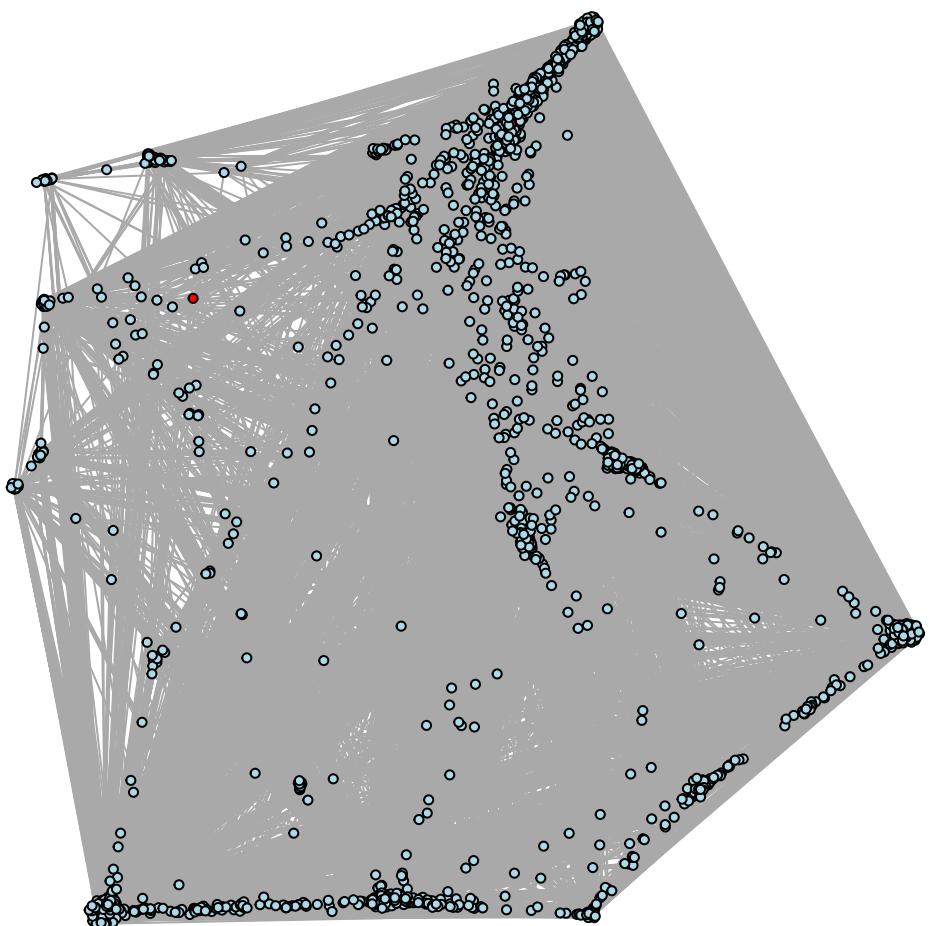
Number of targets = 12.Number of drugs = 23



# Co-expression

Average path length = 3.1348.Number of vertexes = 4039

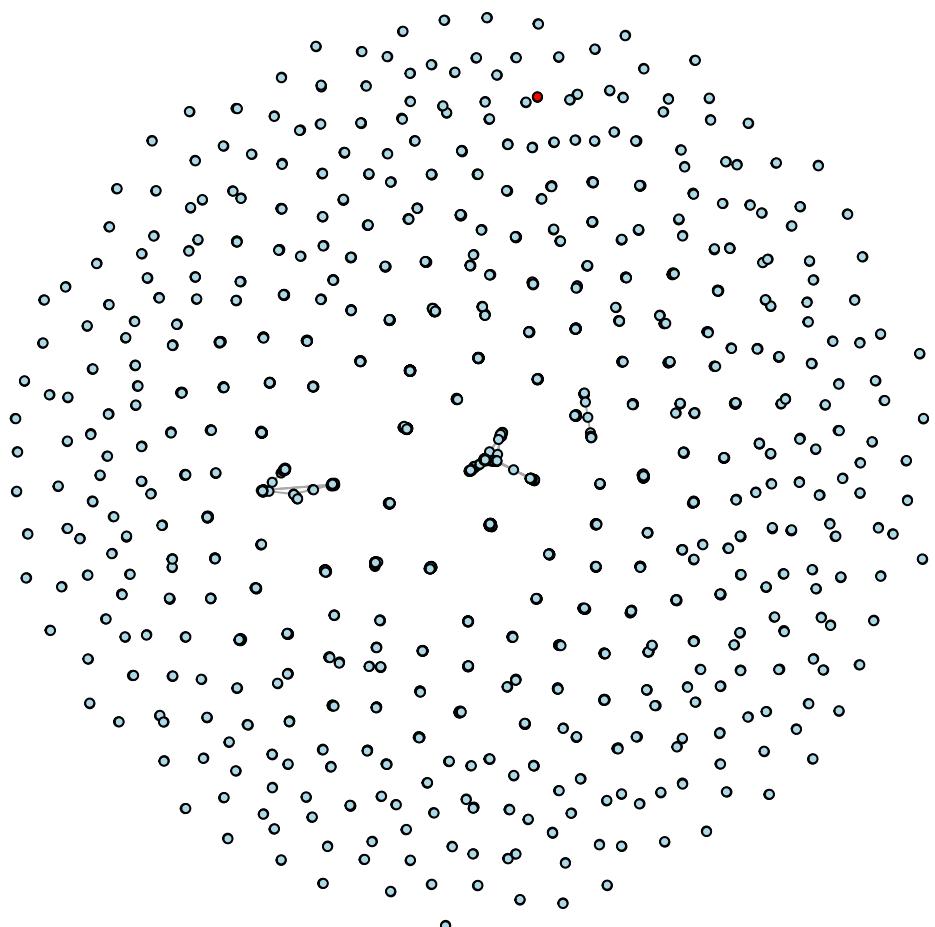
Number of targets = 25.Number of drugs = 36



# Co-occurrence of prot. domains

Average path length = 2.7359. Number of vertexes = 2283

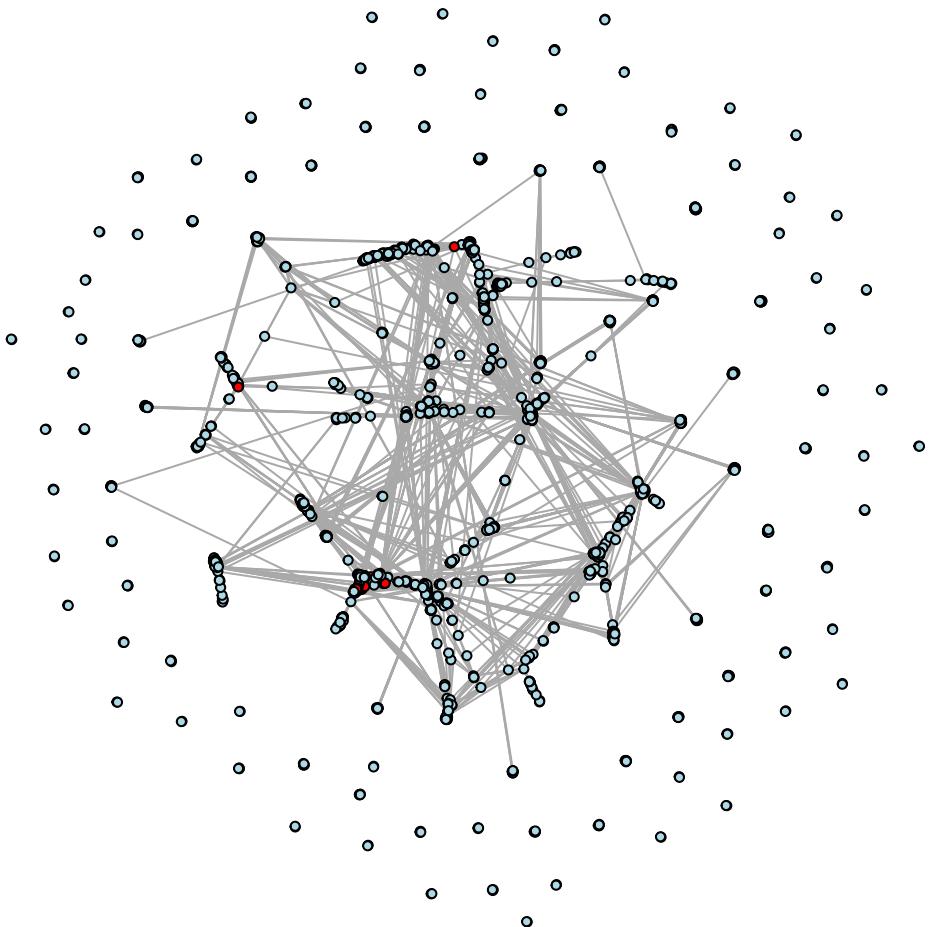
Number of targets = 6. Number of drugs = 16



# Co-functional (EcoCyc/GO-BP)

Average path length = 4.8203.Number of vertexes = 1835

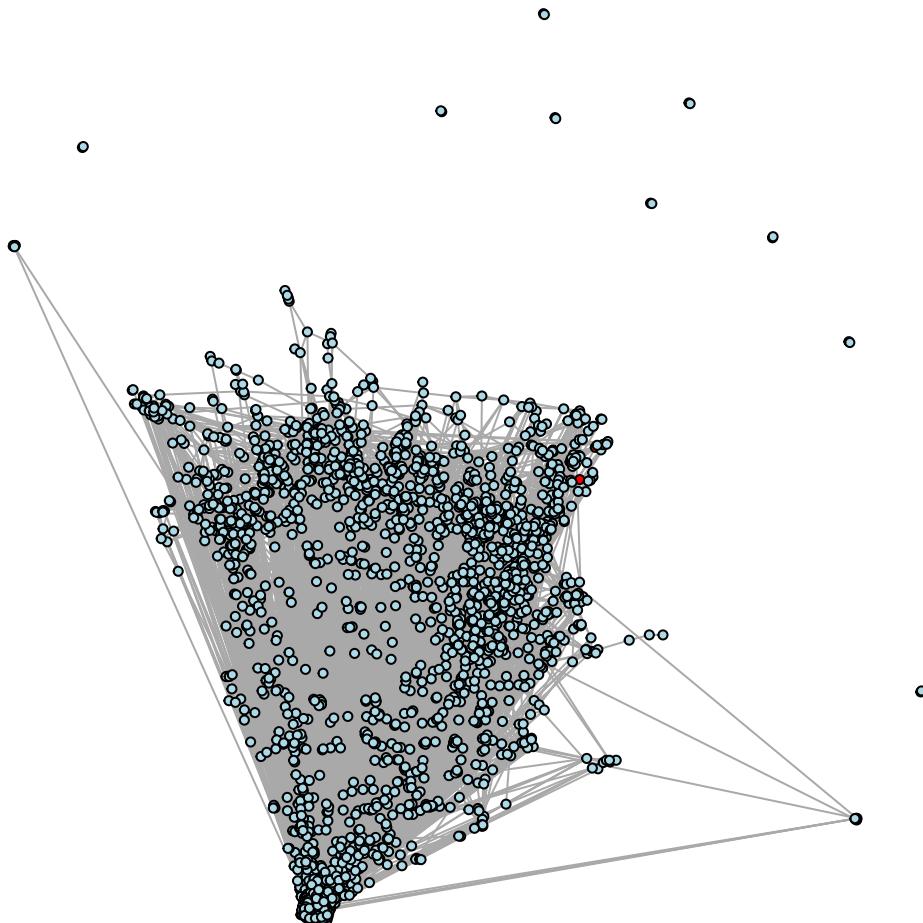
Number of targets = 20.Number of drugs = 31



# High-throughput PPI

Average path length = 4.144.Number of vertexes = 3209

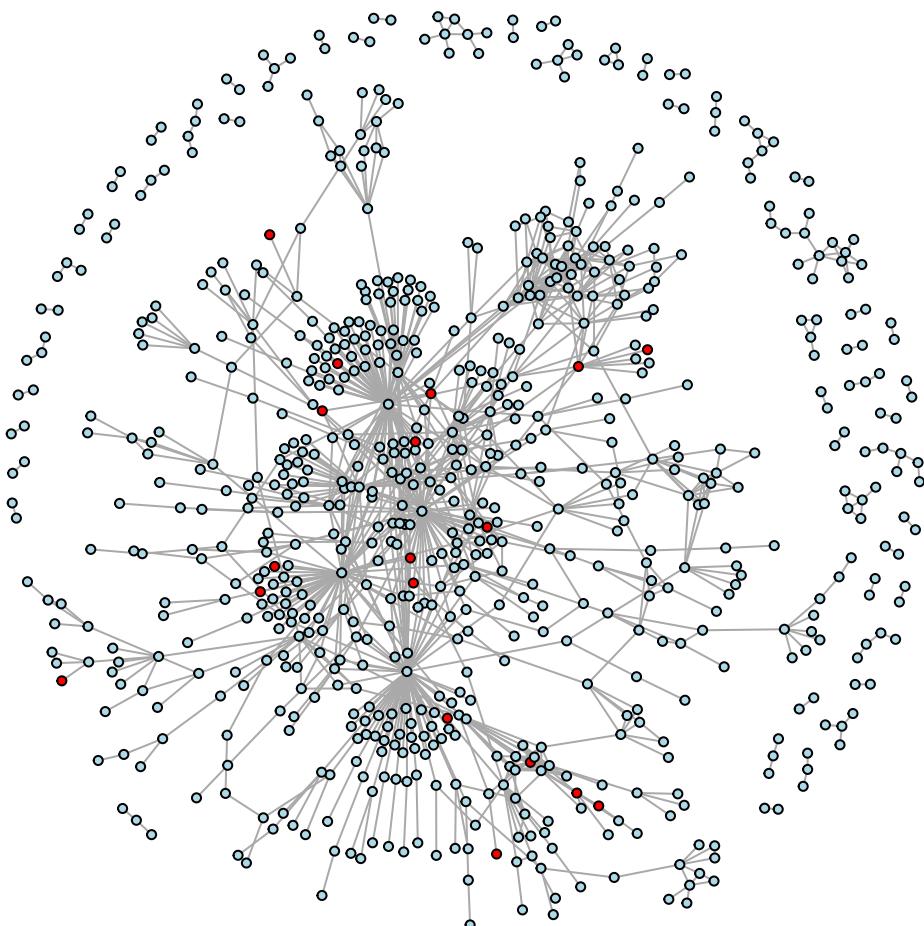
Number of targets = 25.Number of drugs = 36



# Small/medium-scale PPI

Average path length = 4.854. Number of vertexes = 764

Number of targets = 18. Number of drugs = 31



# Similar phylogenetic profiles

Average path length = 3.2474.Number of vertexes = 1817

Number of targets = 15.Number of drugs = 26

