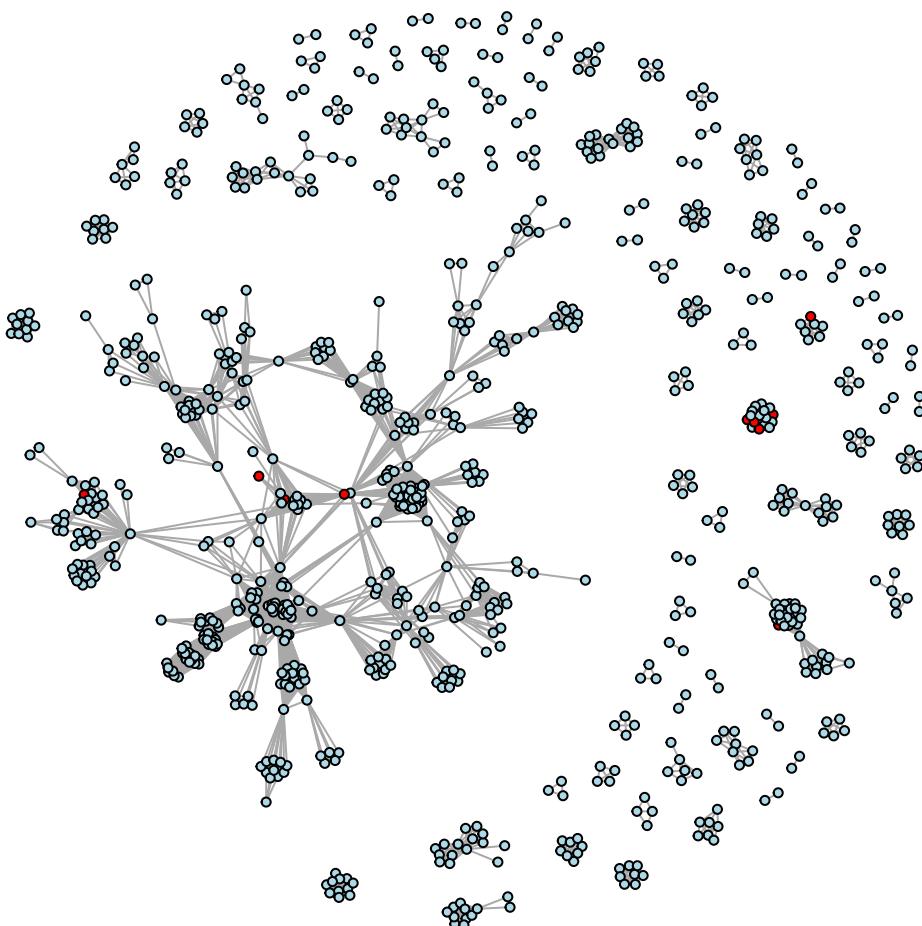


# Co-functional (EcoCyc)

Average path length =4.1098. Number of vertexes =885

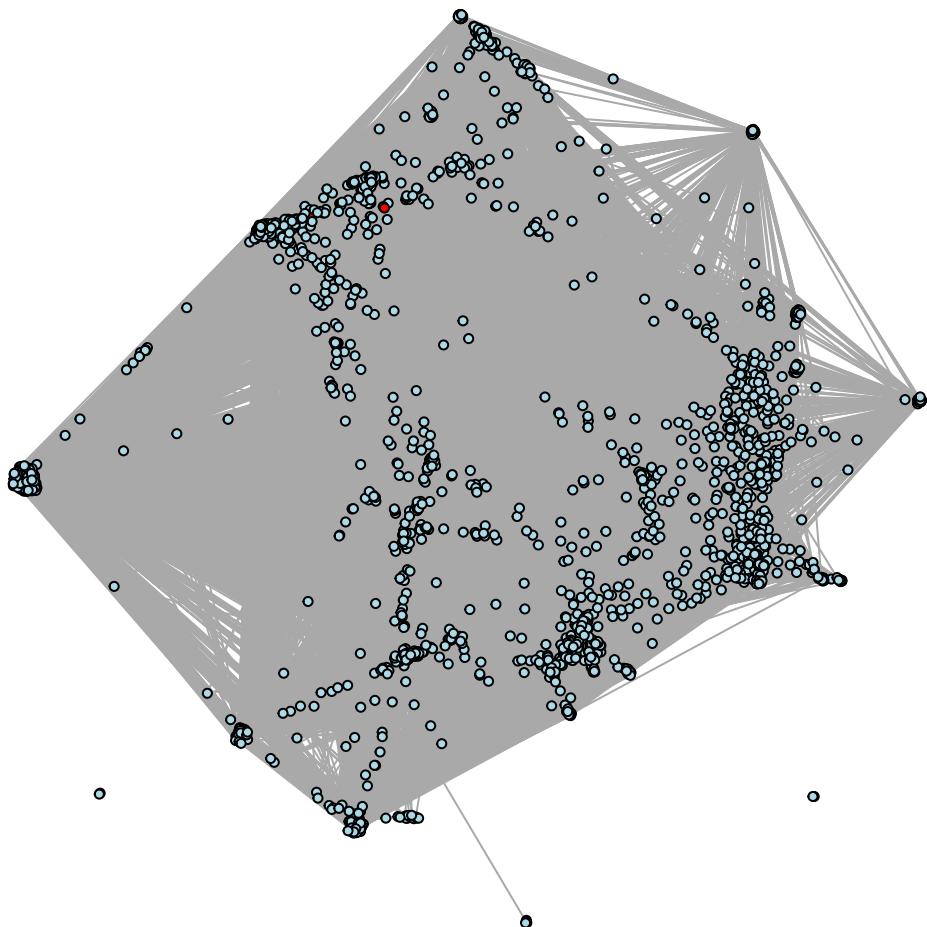
Number of targets =10. Number of drugs =16



# EN: all networks

Average path length =2.8575. Number of vertexes =4099

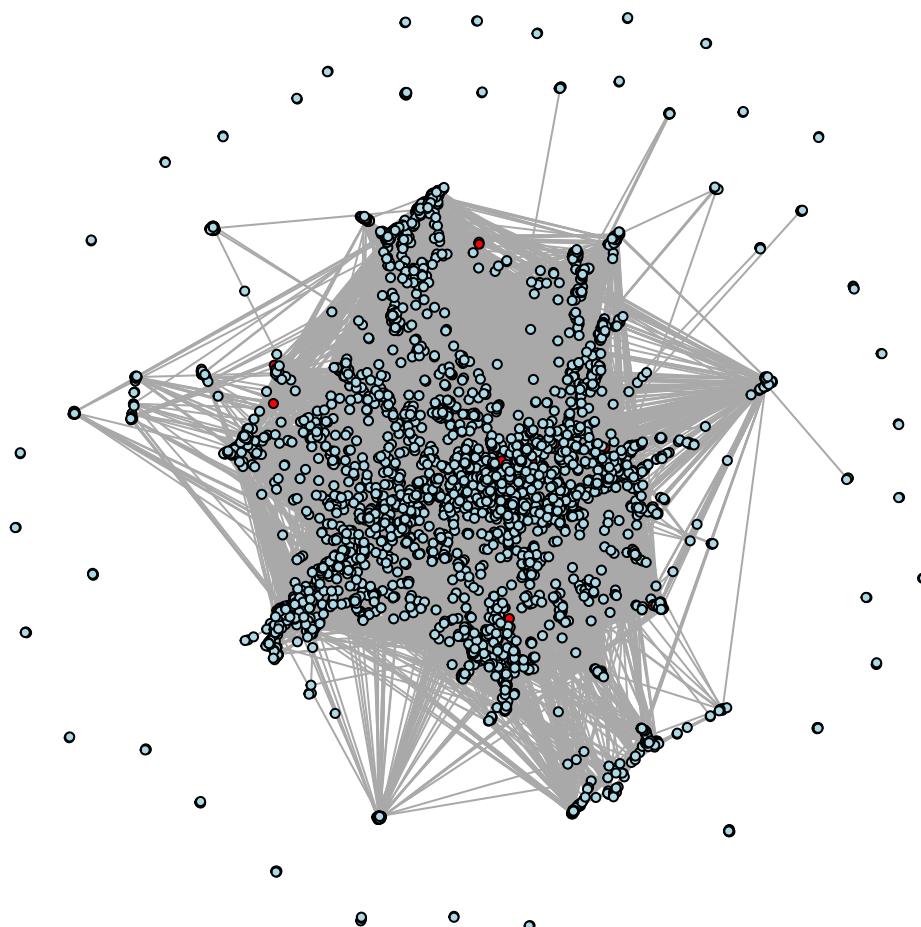
Number of targets =25. Number of drugs =36



# Similar genomic context

Average path length =3.957. Number of vertexes =3568

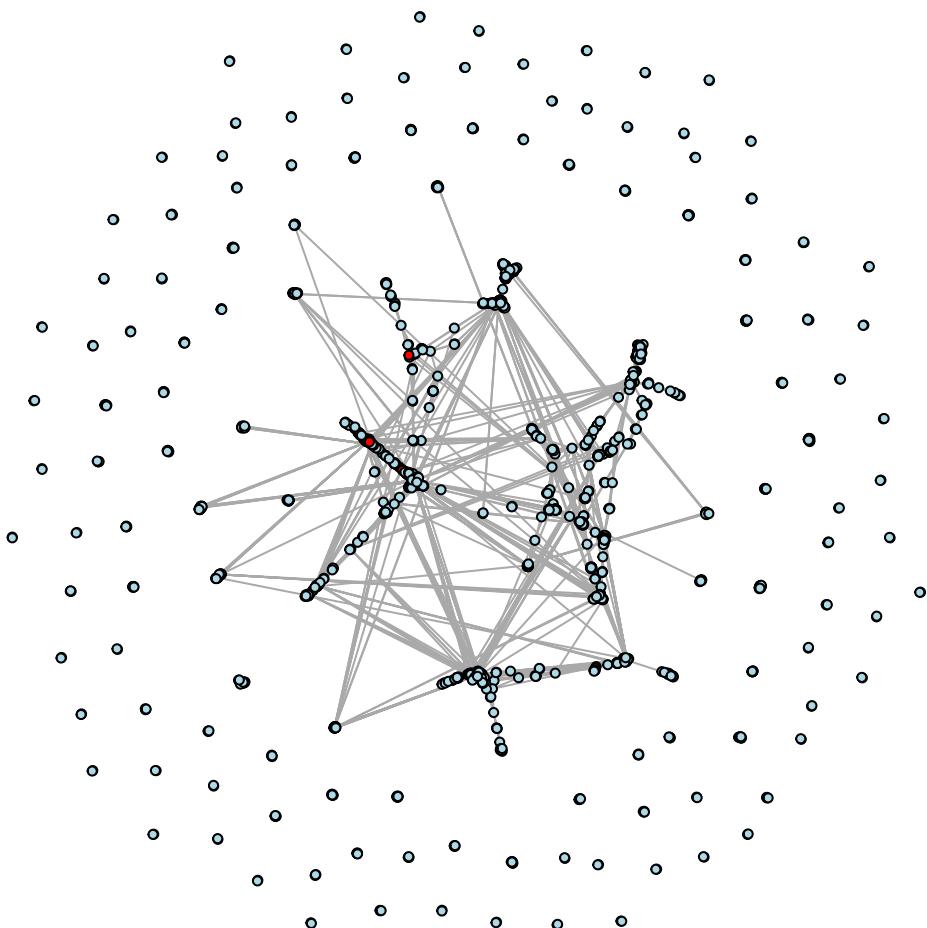
Number of targets =25. Number of drugs =36



# Co-functional (GO-BP)

Average path length =4.7648. Number of vertexes =1474

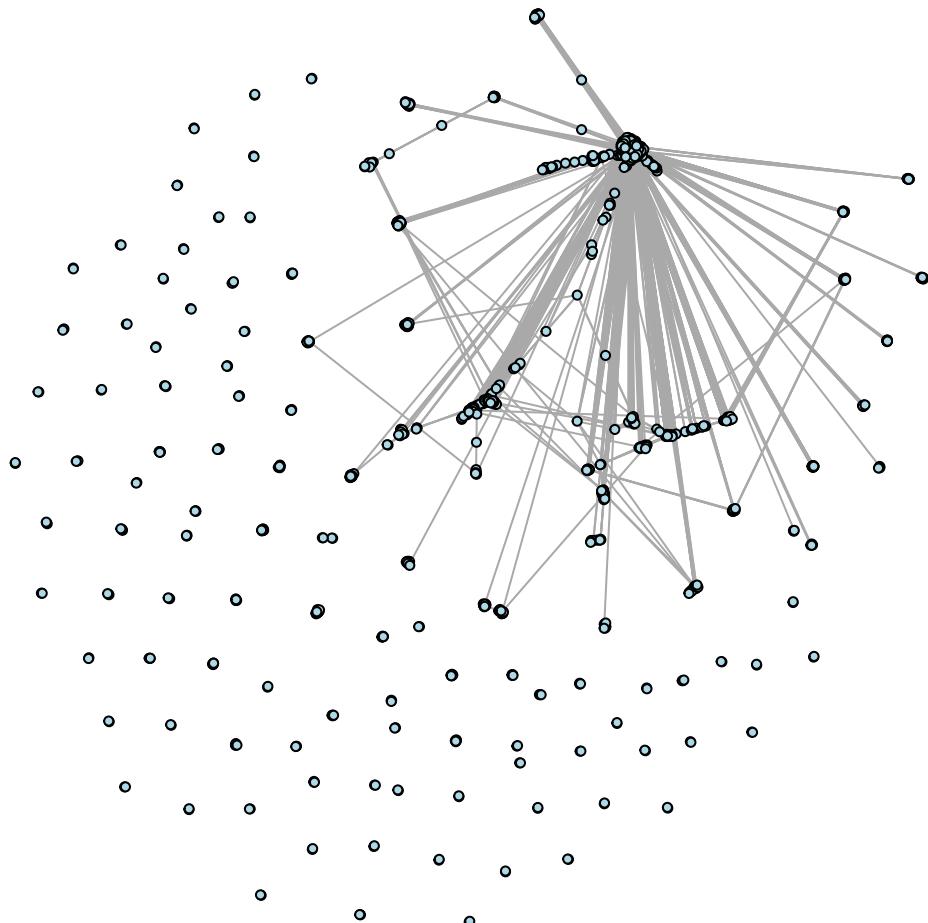
Number of targets =17. Number of drugs =31



# Co-citation

Average path length =3.6296. Number of vertexes =2296

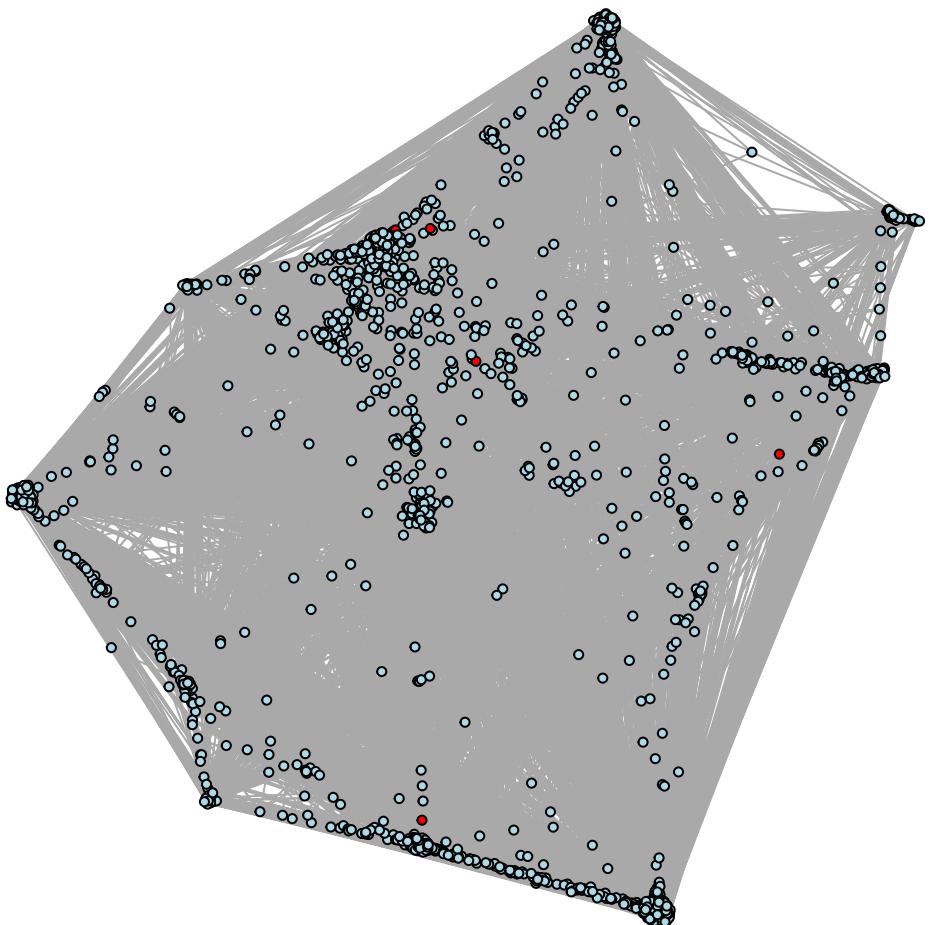
Number of targets =12. Number of drugs =23



# Co-expression

Average path length =3.1348. Number of vertexes =4039

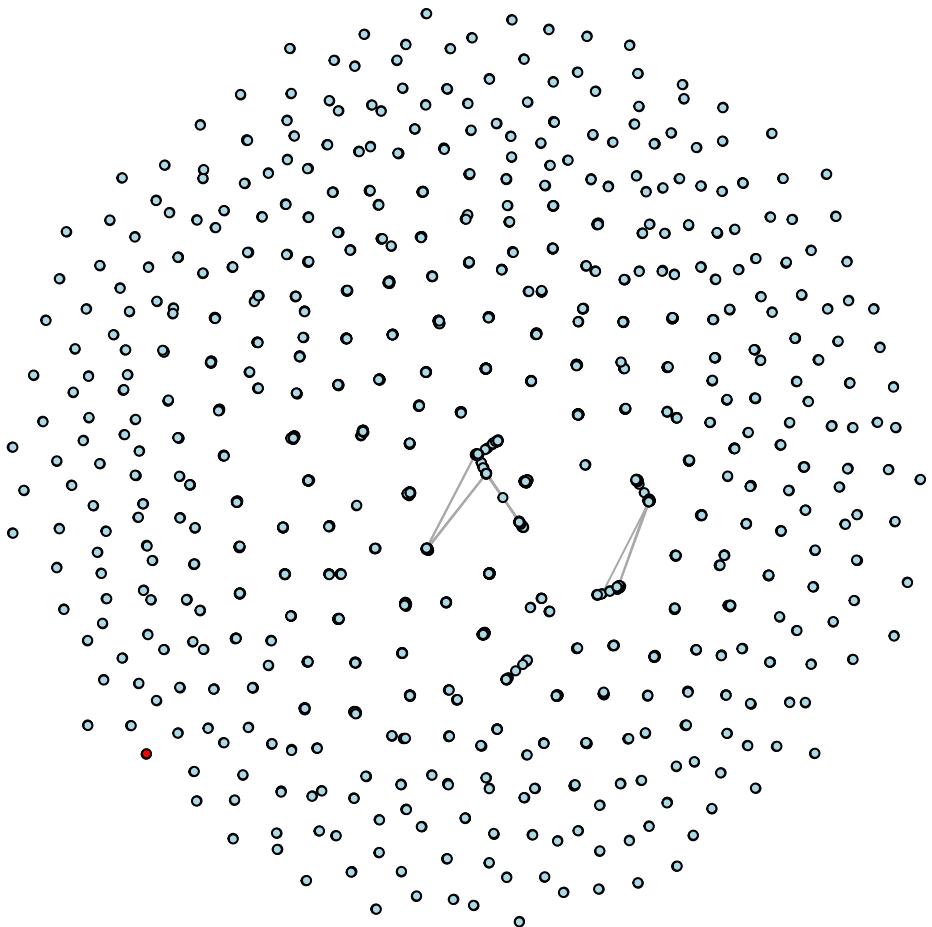
Number of targets =25. Number of drugs =36



# Co-occurrence of prot. domains

Average path length =2.7359. Number of vertexes =2283

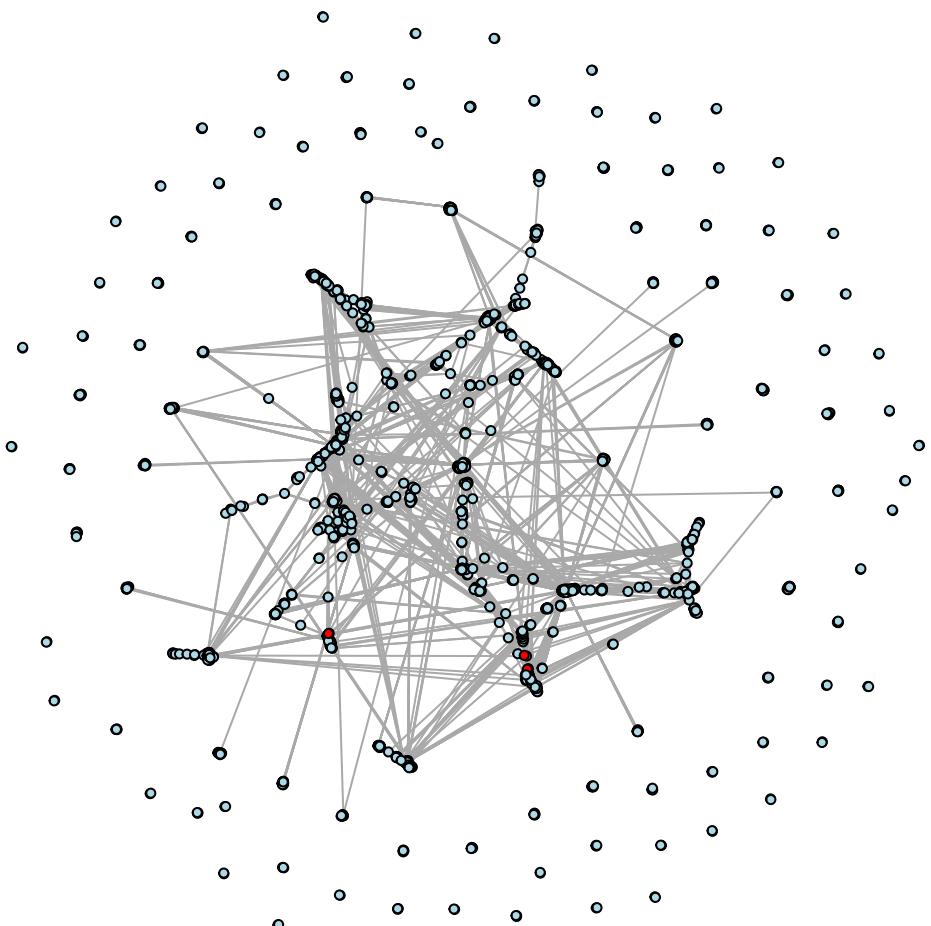
Number of targets =6. Number of drugs =16



# Co-functional (EcoCyc/GO-BP)

Average path length =4.8203. Number of vertexes =1835

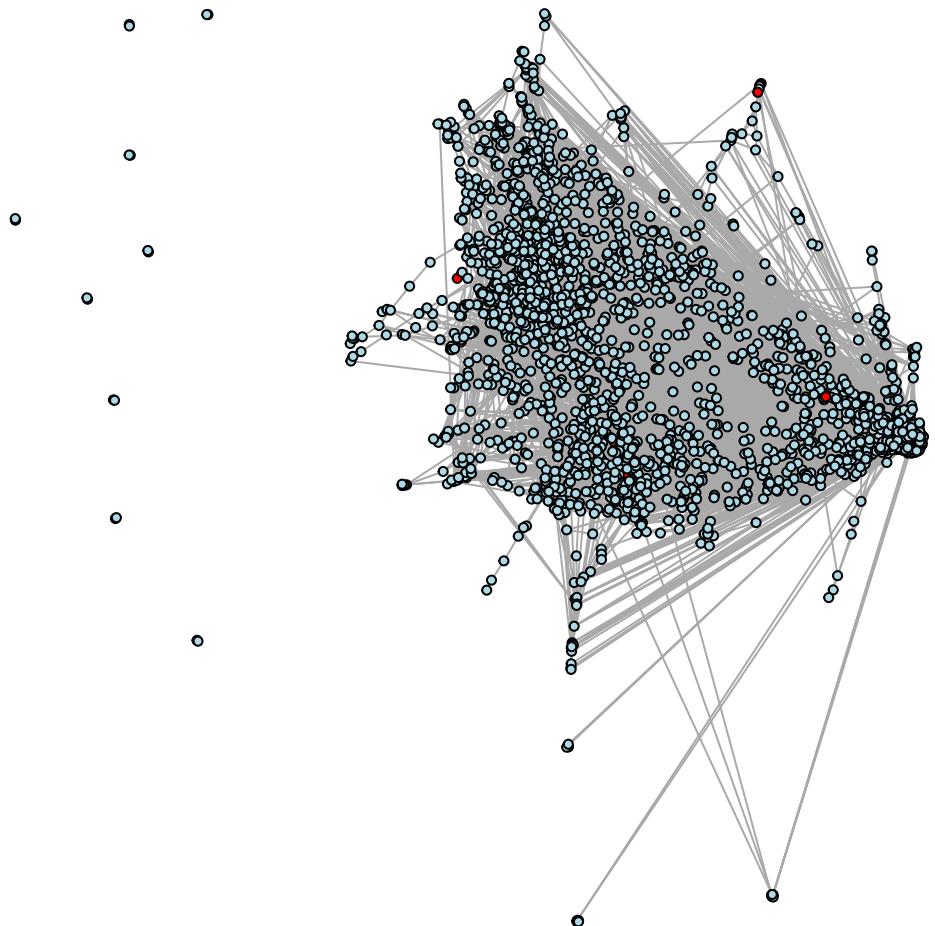
Number of targets =20. Number of drugs =31



# High-throughput PPI

Average path length =4.144. Number of vertexes =3209

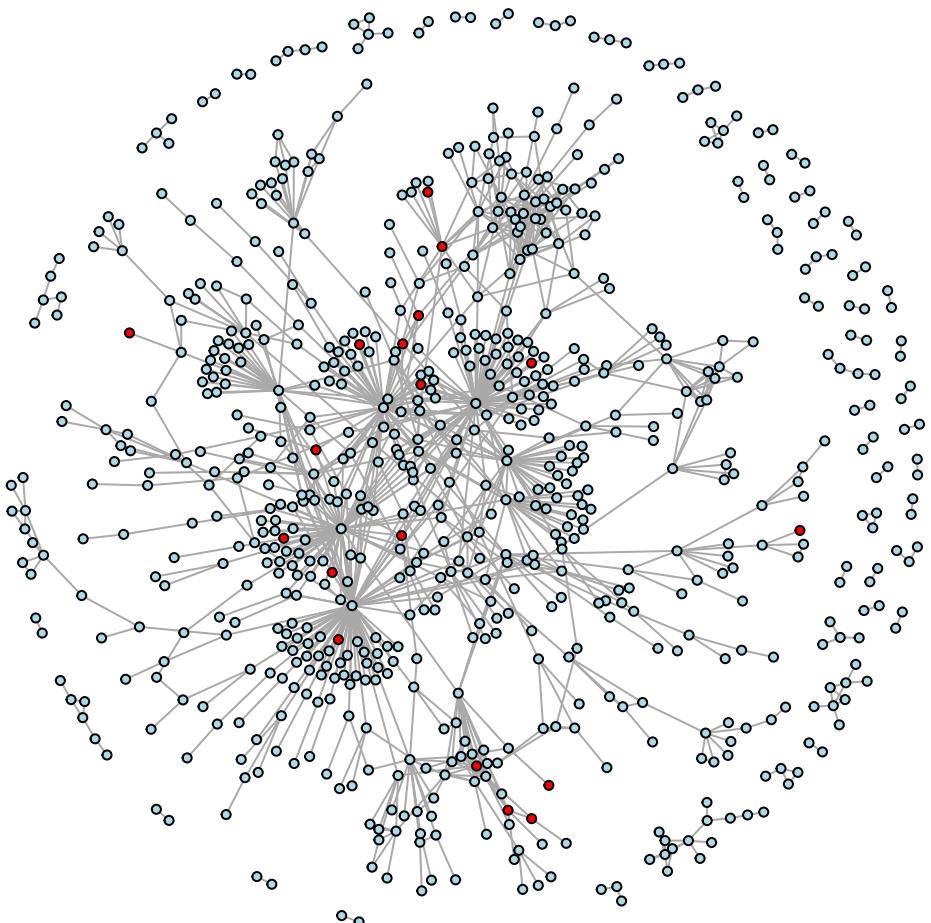
Number of targets =25. Number of drugs =36



# 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

