SUPPLEMENTARY INFORMATION

Use of Graph Database for the Integration of Heterogeneous Biological Data

Byoung-Ha Yoon^{1,2}, Seon-Kyu Kim¹, Seon-Young Kim^{1,2}*

¹Personalized Genomic Medicine Research Center, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Daejeon 34141, Korea, ²Department of Functional Genomics, University of Science and Technology (UST), Daejeon 34113, Korea

http://www.genominfo.org/src/sm/gni-15-19-s001.pdf.

Supplementary Fig. 1

Neo4j Cypher query

MATCH (drug)-[r:Drug_Gene_Interaction {role:'increases^expression'}]-> (gene:Gene {GeneSymbol:'BRCA1'})-[r1]->(disease) WHERE disease.DiseaseName =~ '.*[Cc](ancer|ANCER).*' RETURN drug, r, gene, r1, disease

Supplementary Figure 1. Cypher query to retrieve the relationships among genes, drugs, and diseases that increased the expression of the BRCA1 gene.