

Additional file 4.1. – File with supplemental figures in pdf format. Additional_file_4.1

Figure S4.1. – Database schema for data retrieved from KEGG.

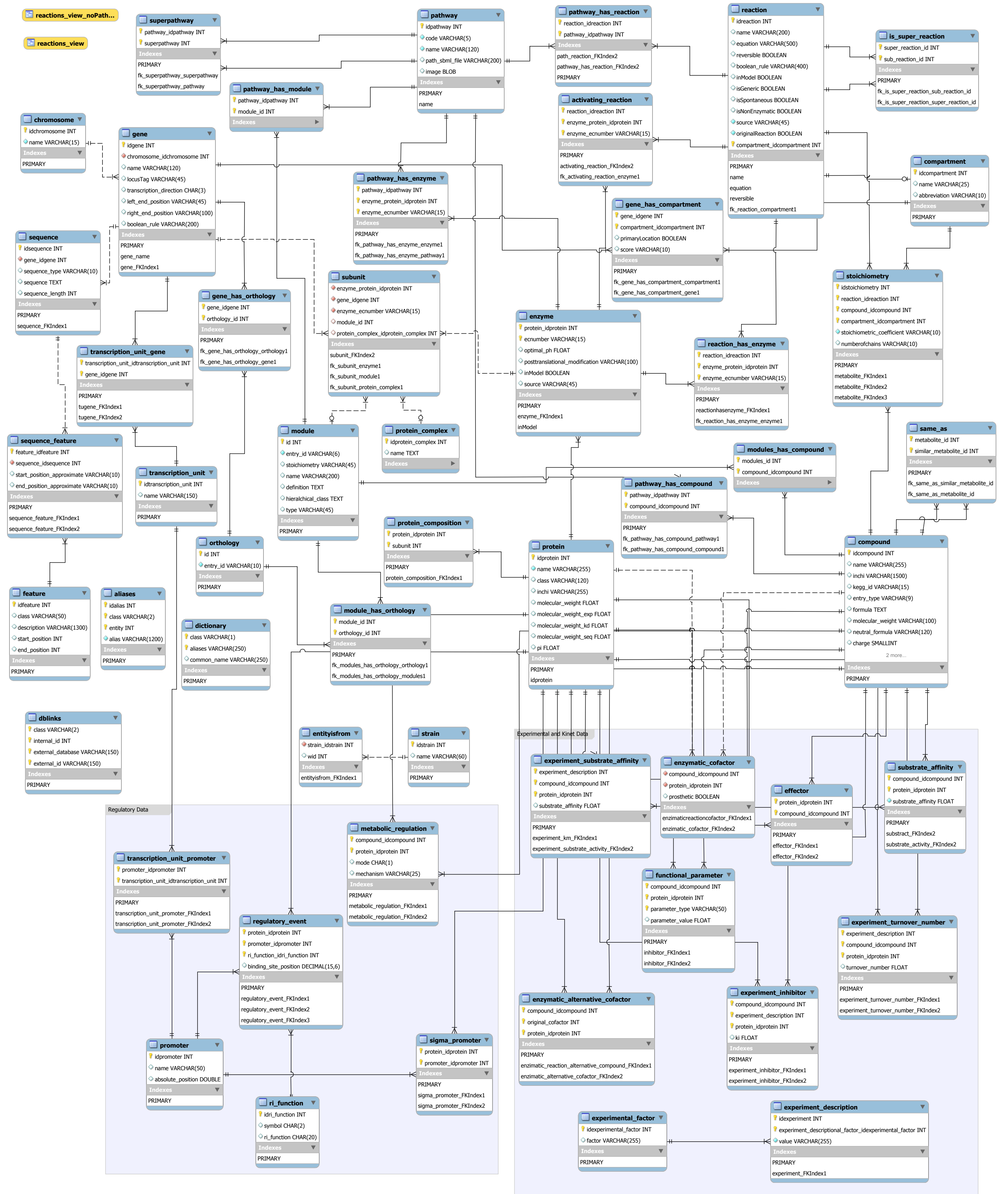
Figure S4.2. - Database schema for data retrieved from remote homology alignments for the genome annotation.

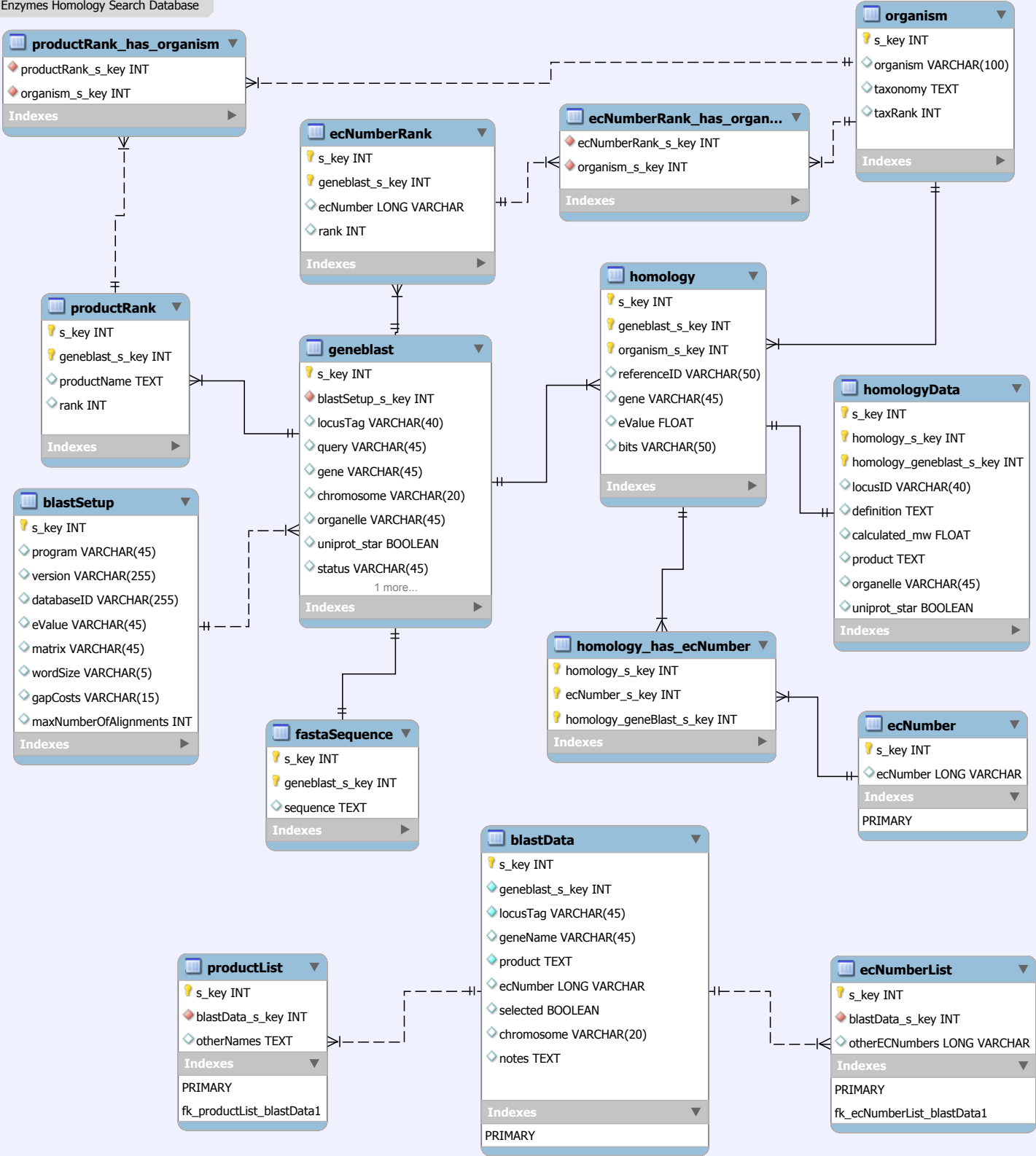
Figure S4.3. - Database schema for storing data on the local similarity alignments between the genome and TCDB.

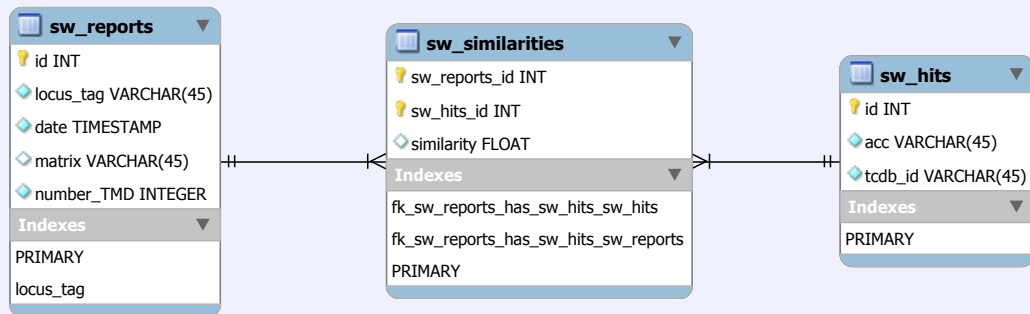
Figure S4.4. - Database schema for the transport reactions generation database.

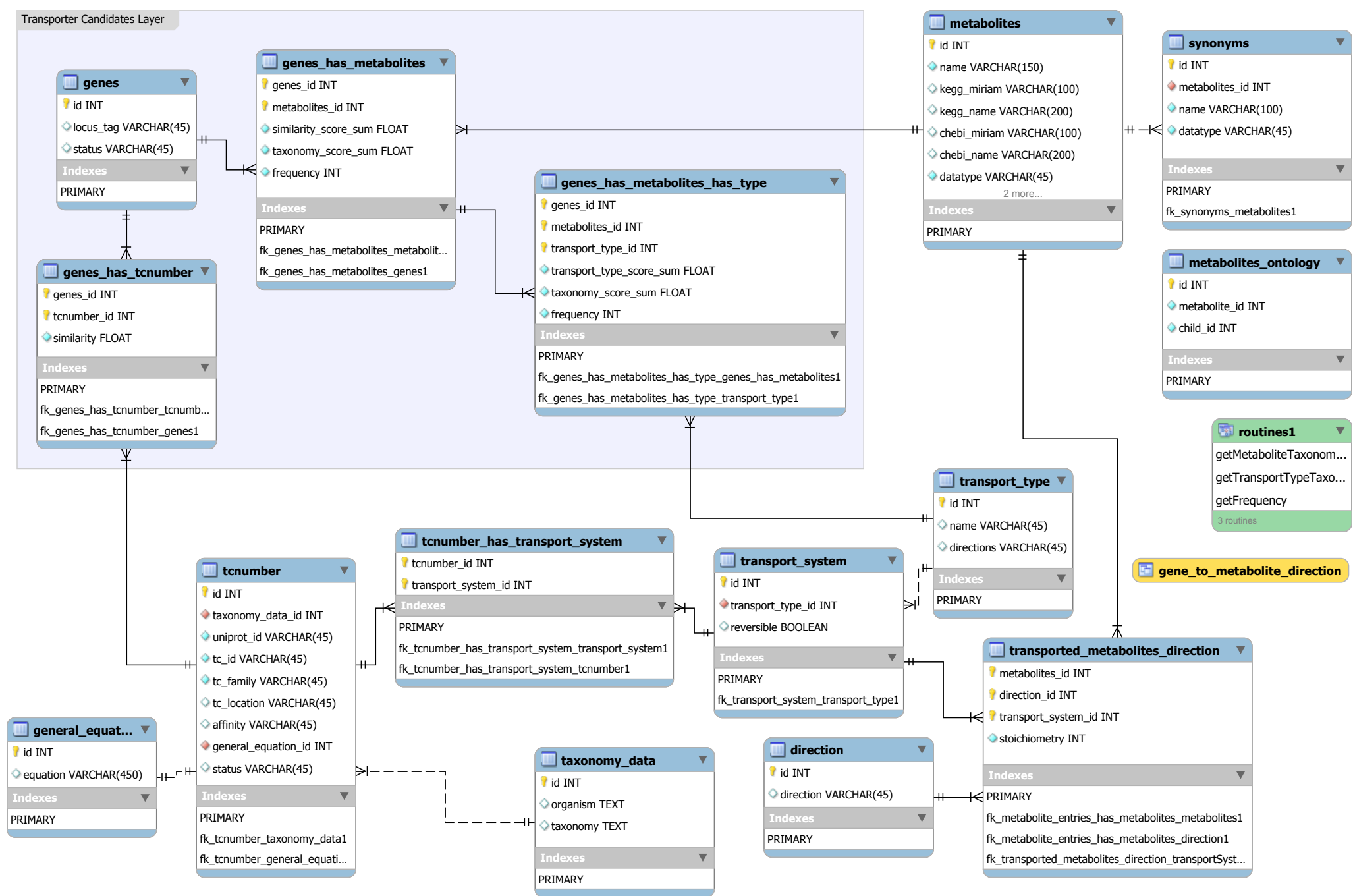
Figure S4.5. - Database schema for the compartments prediction database.

Figure S4.6 - KEGG Pathway map with enzymes and reactions coloured by *merlin* 2.0.

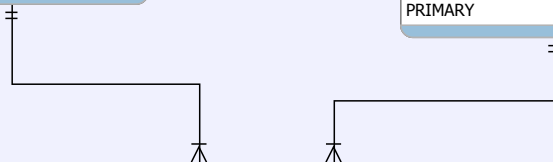
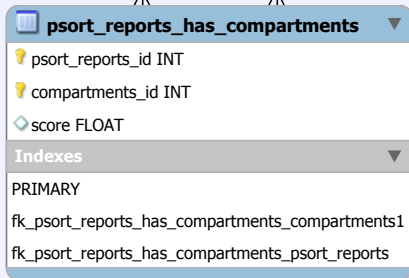
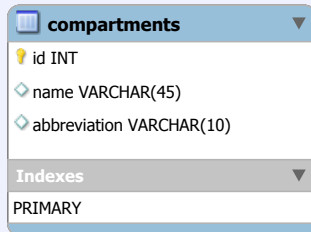
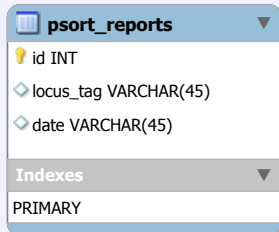








Compartments Prediction



CITRATE CYCLE (TCA CYCLE)

