Empowering Biologists to Decode Omics Data: The Genekitr R Package and Web Server

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Supplementary Figure and Tables

Table S1. GO CC enrichment analysis result without simplification method.
Table S2. GO CC enrichment analysis result after simplification.
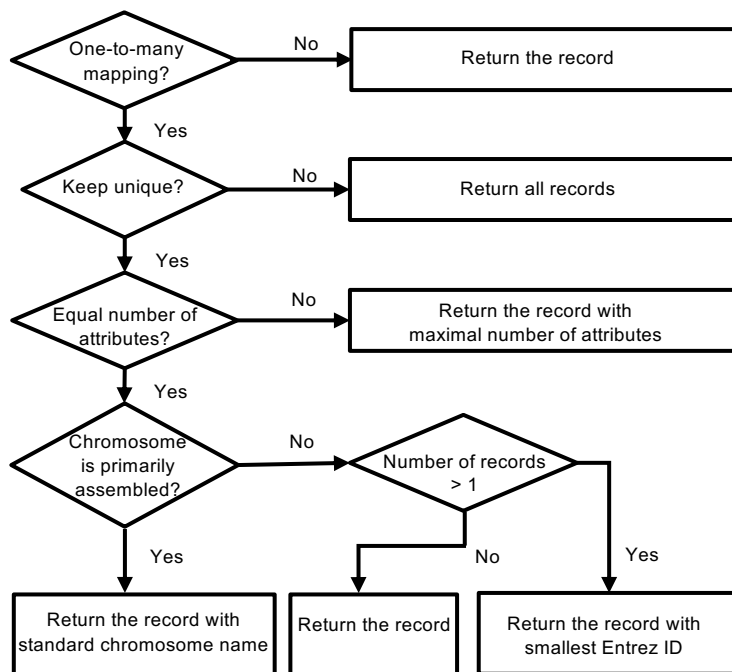

Fig. S1. Flowchart of one-to-many mapping rules for gene information retrieval.