**Related work :**

KEGG pathway could make compared with gene Ontol-ogy (GO),2a way database to gene set improvement Investigation. Try will be an accumulation from claiming controlled vocabularies for gene func-tions sorted out for three ontologies: living process, cellu-lar component, Furthermore atomic capacity. Large portions genomes are annotated with try terms Eventually Tom's perusing group keeping efforts, Along these lines that GO enrichment dissection camwood make performed around Different gene sets. For example, a improvement Investigation figures try terms that areover-represented to a provided for set about mankind's genes, empowering itsbiological elucidation. In contrast, KEGG pathway map-ping displays extra data over how genes orgene results cooperation done pathways, yet the scope o fgenes is lesquerella far reaching over try. The BRITE database, which is an metaphysics database, was introduced should supplement pathway and to extend the coverage for genes to KEGG mapping. The module data-base might have been included Similarly as a endeavor will computerize utilitarian inter-pretation. Unfortunately, however, these two databases have not been well utilized