ABSTRACT

Method of processing protein peptide data and system

The invention provides a method of processing protein peptide data obtained from healthy or pathological samples for analysis, comprising the steps of: providing a list of peptide sequences and associated auxiliary information representing an input data set; compiling from the input data set a new peptide sequence list by removing peptide sequence redundancy in the peptide sequence list, said new peptide sequence list representing a peptide data set; and grouping together members of the peptide data set originating from the same protein thus generating a protein data set.

Fig 1