String grid-connected inverter

Remove redundant interaction pairs (AB-BA): By default, the links files encode interactions in both directions (AB and BA). However, since functional and physical interactions are symmetric, ...

In that case, you could simply enlarge the network (e.g. clicking the "+" button or modifying the String parameters at the bottom of the page) and relaunch the clustering. ...

Gordon et al., SARS-CoV-2 protein interaction map (Nature 2020) All 332 virus-interacting human proteins are marked for emphasis in this version of STRING; The color-code indicates the ...

Upload a complete species proteome to STRING, and we"ll generate its interaction network and predict protein functions, including Gene Ontology terms and KEGG pathways. Once ...

STRING API STRING has an application programming interface (API) which enables you to get the data without using the graphical user interface of the web page. The ...

STRING is a Core Data Resource as designated by Global Biodata Coalitionand ELIXIR.

STRING is a database of known and predicted protein-protein interactions. The interactions include direct (physical) and indirect (functional) associations; they stem from computational ...

Web: https://stanfashion.pl