Post Delta Network Refactoring

# Motivation

* Now that we have finished and tested the delta network functionality, let’s refactor some code on the client side. We know for sure that we won’t need a separate tab for the delta network, so we can remove all code that attempts to distinguish controllers between delta and non-delta
* Even though the server side code that handles the R output is for the most part unique to each R script, we should still turn parts of it into functions as a way of minimizing the lines of code in the server.js file.

# Features to Add

## Small

* Multiple gene selection from autocomplete
* Prevent zooming out too far or too close into the graph
* Gene locator multiple genes
* Perhaps update to latest cytoscape version
* Add redirect if users don’t add /app to url
* Add genecard to the table headers

## Large

* Community Detection
* Filtering on Interaction Explorer
* Separate data from layouts