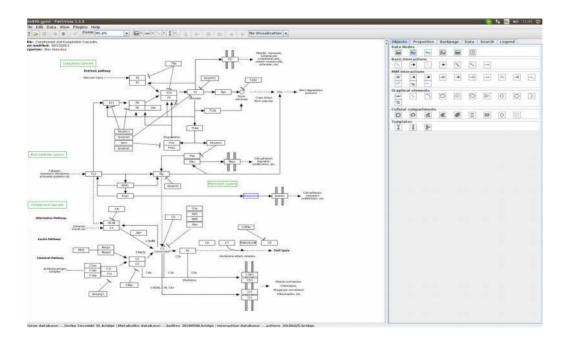
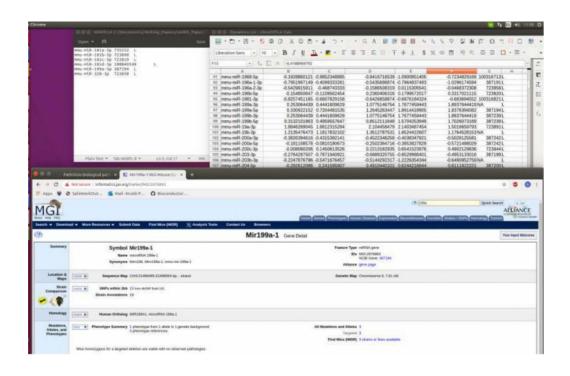
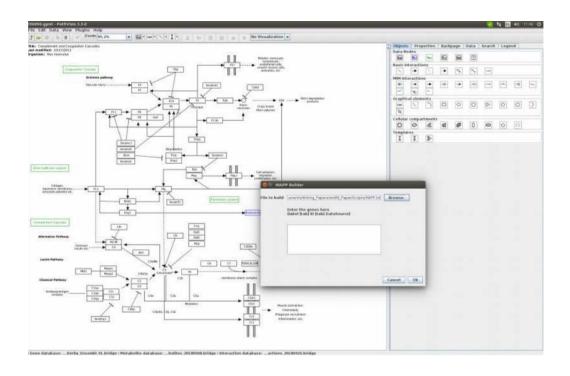
TimiRGeN - Pathvisio GRN creation guide:

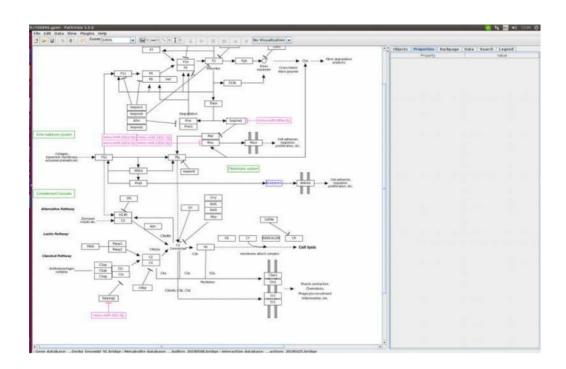
Open selected pathway in Pathvisio. Use the set up instructions found on the pathvisio website. You can download your pathway as a gpml format from wikipathways or use the wikipathways app in pathvisio to import the pathway (1, 2). Can download the wikipathways app if the plugin is unavailable here



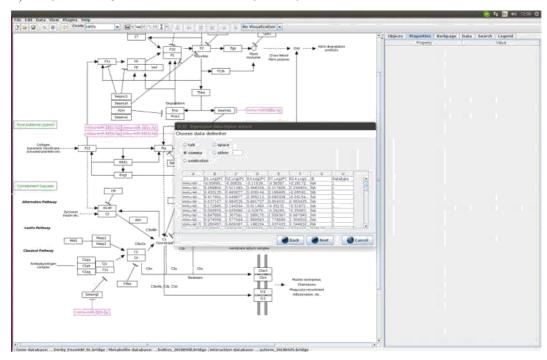


3) Use the MAPP app on Pathvisio to import the MAPP txt file created with TimiRGeN. Can be downloaded from here if plugins are unavailable

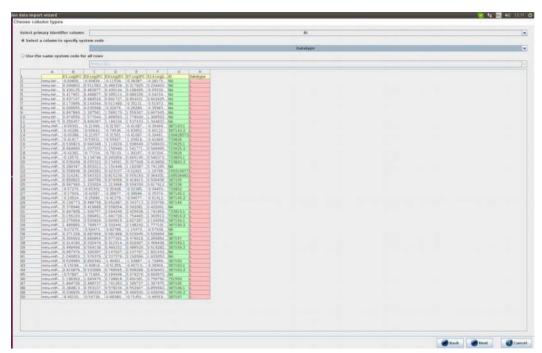




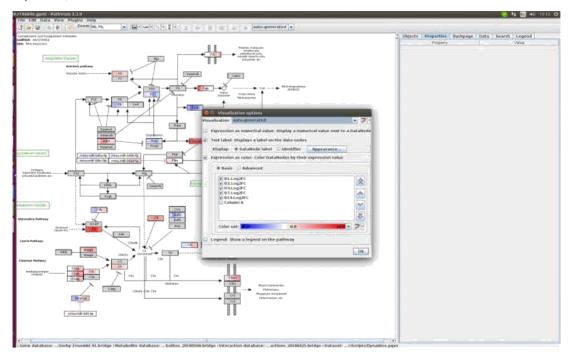
5) Import the dynamics file (Data > Import expression data). Select comma.



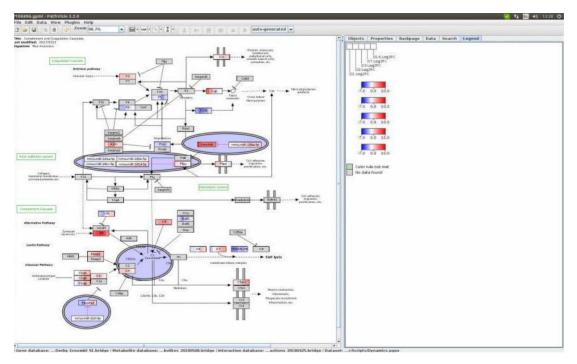
6) Choose Primary identifier column (Green) as ID and column to specify system code as Datatype (red).



7) Select a suitable colour palate to represent the time series. The segments will be shown in an order which matches the Dynamic file. (Data > Visualization options)



8) Isolate key parts of the pathway for bottom-up GRN construction. In this example, the interactions of the miR-mRNA target's can be seen to effect the biogenesis of C3 convertase.



References

- 1) Slenter DN, Kutmon M, Hanspers K, et al. WikiPathways: a multifaceted pathway database bridging metabolomics to other omics research. Nucleic Acids Res. 2018;46(D1):D661D667. doi:10.1093/nar/gkx1064
- 2) van Iersel MP, Kelder T, Pico AR, et al. Presenting and exploring biological pathways with PathVisio. BMC Bioinformatics. 2008;9:399. Published 2008 Sep 25. doi:10.1186/1471-2105-9-399