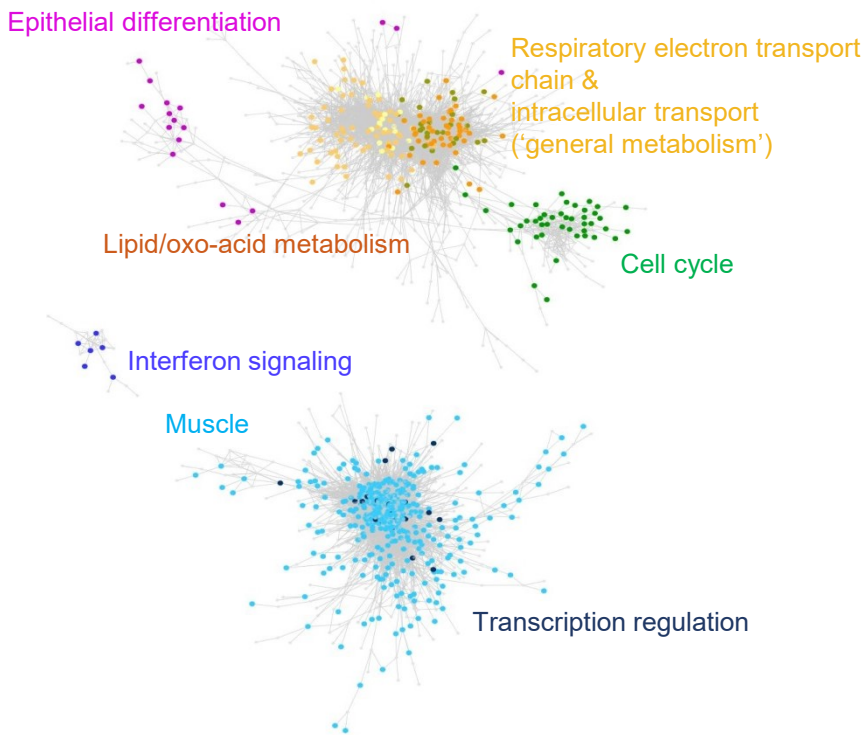


# Figure S3

## NZ along rumen transcriptomic network



**Figure S3: Gene network reproduced using data from the New Zealand CH<sub>4</sub> experiment [1].**

[1] Xiang, R. *et al.* Gene network analysis identifies rumen epithelial cell proliferation, differentiation and metabolic pathways perturbed by diet and correlated with methane production. *Sci Rep* **6** (2016).