Figure S4. Network plot showing the relationship between tissue enriched and group enriched genes in their respective expressed tissues. Red nodes represent the number of tissue enriched genes in each tissue. Orange nodes show the number of genes that are group enriched in combinations of up to five different tissues and with a minimum of three genes in each combination. This results in a total number of grey tissue nodes of 31 as smooth muscle has no tissue enriched genes and no combinations with more than three group enriched genes. The size of the orange and red nodes is related to the number of genes enriched in a particular combination of tissues.

