Figure S3. Heat map showing the leading edge subsets (LES) of pathways in the 'nucleocytoplasmic transport' cluster. The heat map fill color indicates the pairwise comparison/s in which each LES protein was implicated.
Figure S4. Heat map showing the leading edge subsets (LES) of pathways in the ‘regulation of gene expression’ cluster. The heat map fill color indicates the pairwise comparison/s in which each LES protein was implicated.
Figure S5. Heat map showing the leading edge subsets (LES) of pathways in the 'ion and vascular homeostasis', 'signal transmission', and 'solute transport' clusters. The heat map fill color indicates the pairwise comparison/s in which each LES protein was implicated.