



Figure 4 : Endocytosis and receptor mediated endocytosis are barriers to reprogramming

Using a combination of gene interaction network analysis and gene ontology analysis we identified genes central to endocytosis and receptor mediated endocytosis among the top 5% significance level ($p=0.0502$) TRA-1-81+ screen hits. Receptor mediated endocytosis frequently requires ubiquitination of the receptor and/or proximal surface proteins to target them for degradation. Screen hits MARCH3 ($p=0.0152$), RNF40 ($p=0.0044$), and NEDD4 ($p=0.0172$) have all been shown to mediate receptor ubiquitination and subsequent targeting for endocytosis [1-5]. Once targeted for proteolysis, transport vesicle formation requires RABEP1 ($p=0.0262$) and EHD2 ($p=0.0262$) which are screen hits. Early endosome formation involves formation of coated pits comprised of Clathrin (CLTA $p=0.0046$) as well as WDFY1 ($p=0.0222$), both top screen hits. Multiple screen hits were identified as vesicle surface proteins mediating late endosome-lysosome fusion (MCOLN1 $p=0.0108$, PIK3R4 $p=0.023$, VPS25 $p=0.015$, HSPA8 $p=0.0242$, SCARB2 $p=0.014$). Lysosomal lipases and proteases such as LIPA ($p=0.0394$), GM2A ($p=0.035$), and LGMN ($p=0.0006$) were screen hits as well as PIK3C3 ($p=0.0166$) which mediates lysosomal protein shuttling into the lysosome. Lastly, the lysosomal protein SLC17A5 ($p=0.0192$) and DRAM1 ($p=0.044$), a lysosome surface membrane protein associated with autophagy, were also screen hits.

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