-So, may have fourth PC3 cavin1 exosome mir data, raw counts, unsure about how well it will integrate with the current data. So exclude the horrible replicate, put that one in its place. Identify the differentially expressed miRNAs. Then assess through qPCR.

- Can look at the different markers for the different vesicle types. Eg EPHA2 and 4F2(SLC3A2). To find if the miRNAs and proteins are exported separately depending on vesicle with the different markers.

- How does it get in there based on the different markers? IE find correlation or interaction between marker and protein that might be the interaction partner for the miRNAs. Verify by experimental data: eg co-ip localization, knockout?

Background or biological significance?