* Write Abstract (Elias + Salome)
* Write Project Outline (Salome)
* Change title new colour and font (Lisa)
* Introduction changes (Elias + Salome)
* Quote essential question (Lennart)
* Font change subtitles (Lisa)
* Naming figures with fig.cap=’ ‘ (Salome)
* Add PARP10, ERBB2 and CCND in expression box and whisker plot + one more (Elias)
* Title within sum squares plot + ggplot (make it nicer) (Elias)
* Dataframe with lapply for driver mutation cluster number (Lisa + Elias)
* Wilcoxon test output with lapply to format next to each other (Lisa)
* Reload PIK3CA SST gene functions in csv document (Elias)
* QQ plot titles + colour legend (Elias)
* Correlation coefficient heatmap + QQ plot positioning next to one another (Elias + Lennart)
* F-Test (Lennart)
* References in same format (All)
* Drug/Therapy research -future outcome of SSTs(Elias)