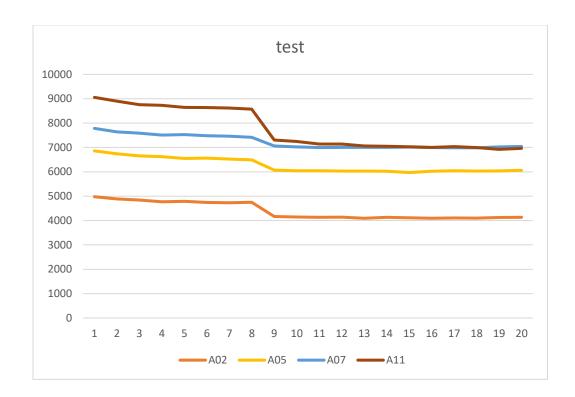
221 images from excel

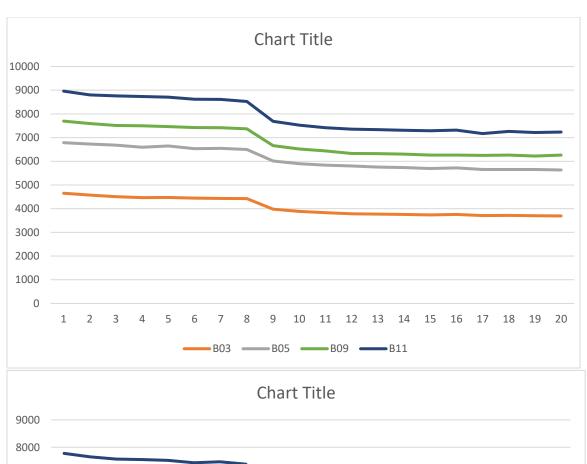
#Same group different concentrations (2,3,4,5 uM ACMA)

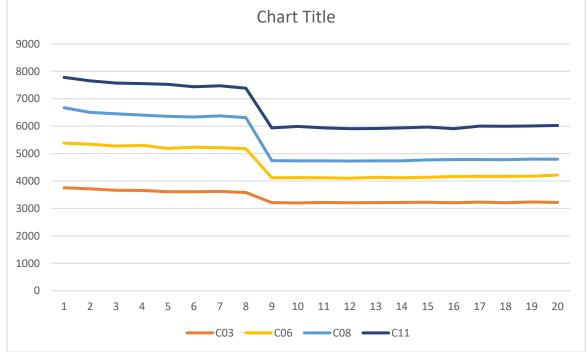
A02,A05,A07,A11

B03, B05, B09, B11

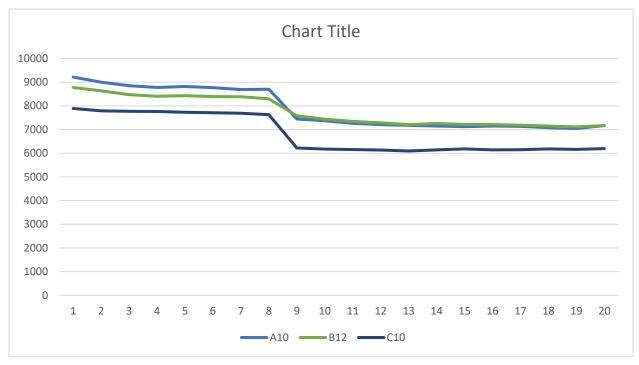
C03, C06, C08, C11

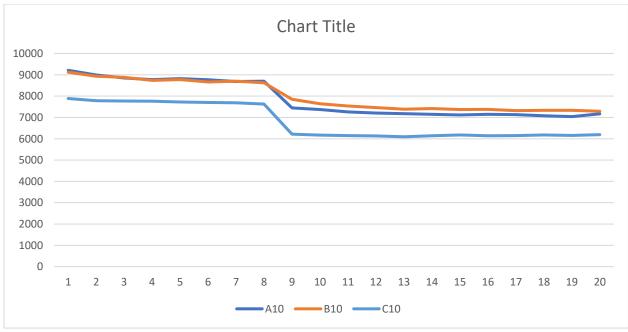




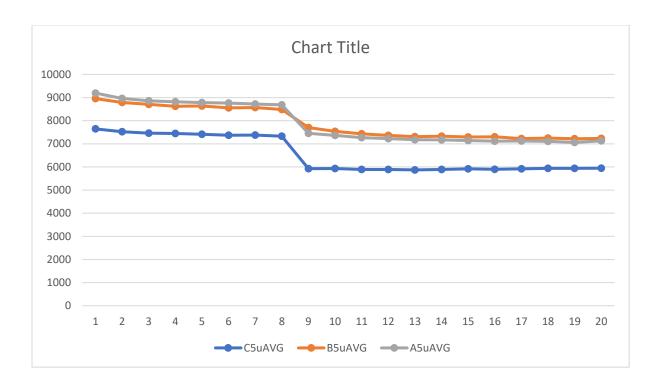


Same concentrations different group

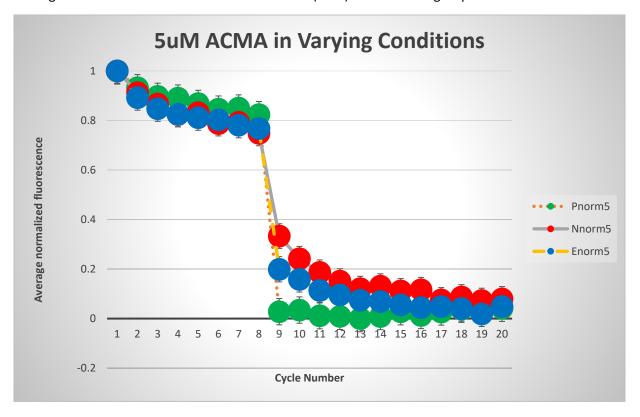


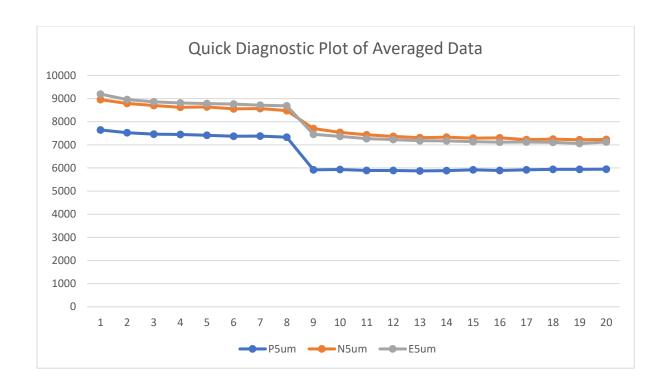


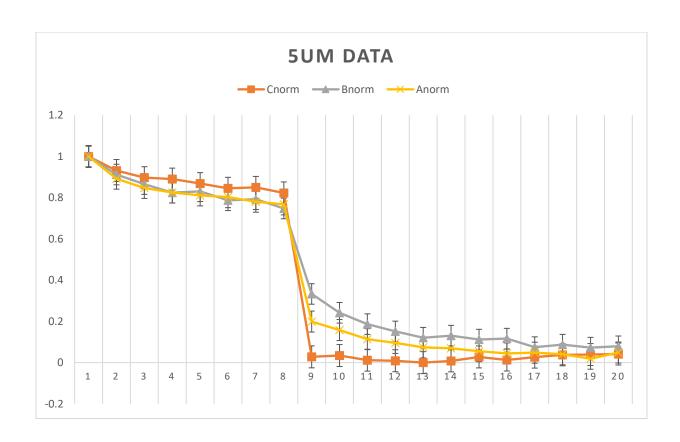
A10,B10,C10

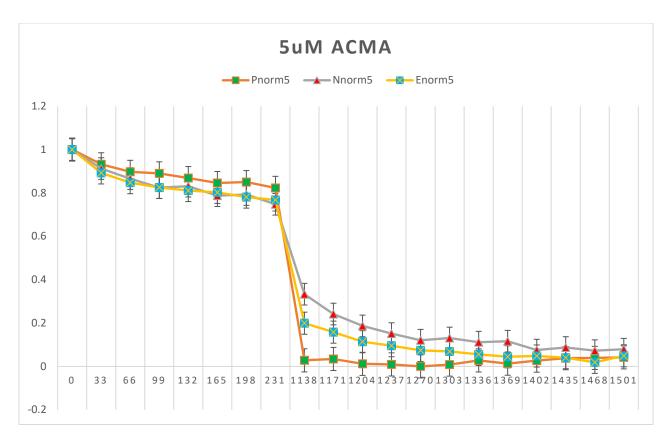


Average Normalized fluor for same concentration (5uM) and different groups





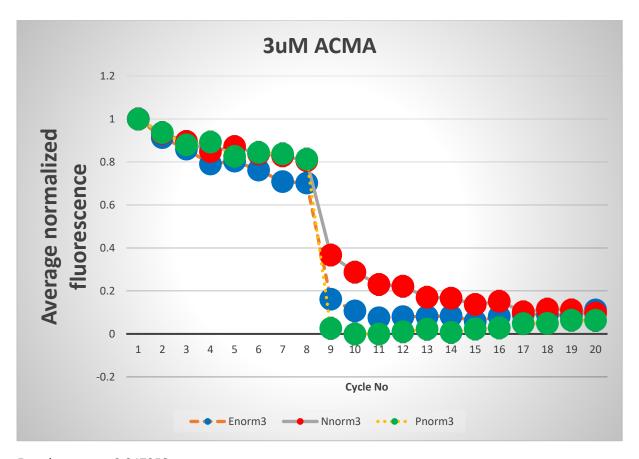




Positive Control= Value+-SEM (0.05325)

Negative Control=Value+-SEM(0.049876)

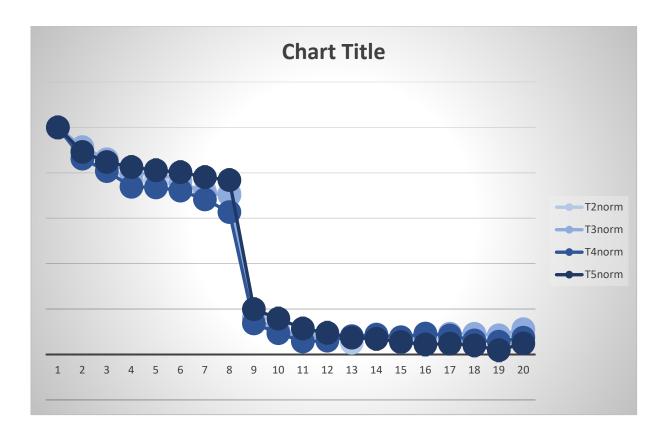
Experiment=Value+-SEM(0.05091)

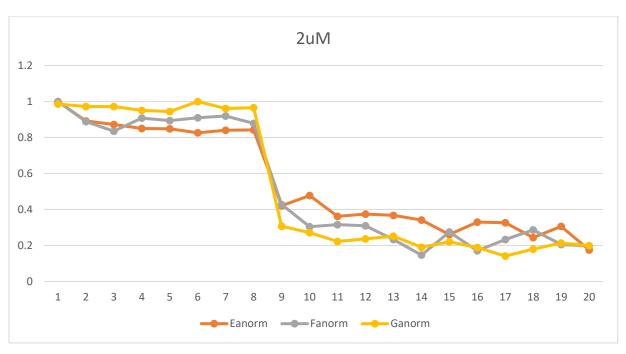


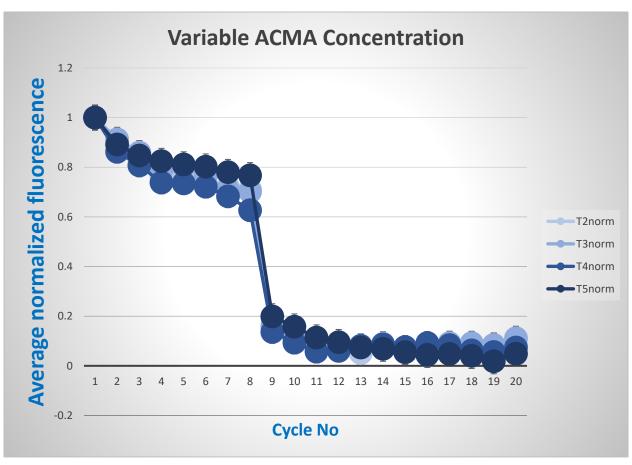
E =value +- sem 0.047958

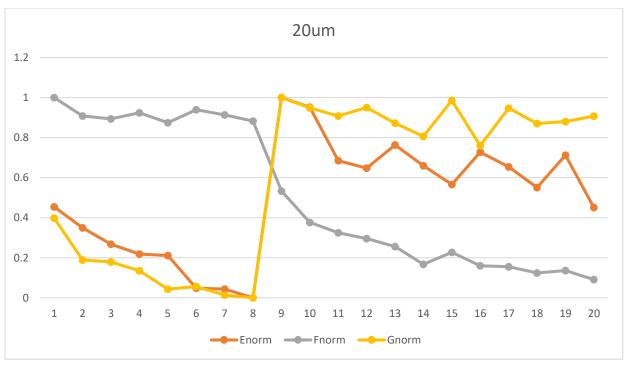
Nc =value +- sem 0.050838

Pc =value +- sem 0.05148

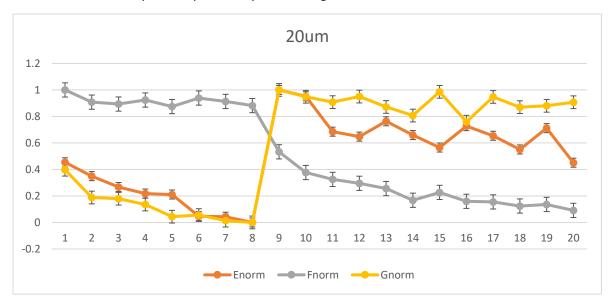


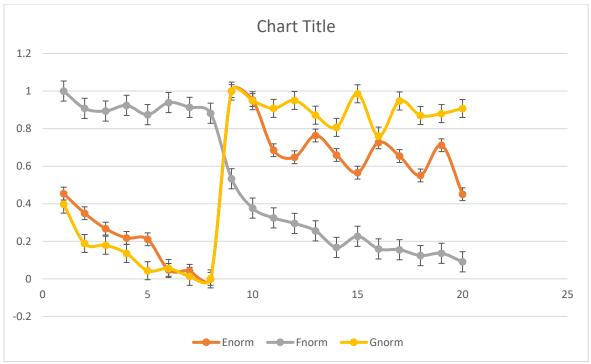


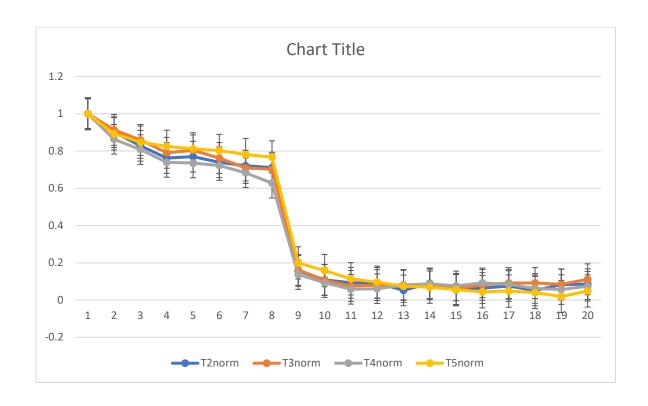


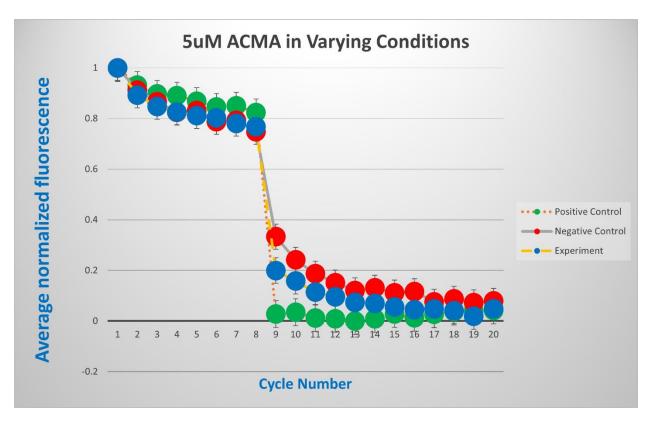


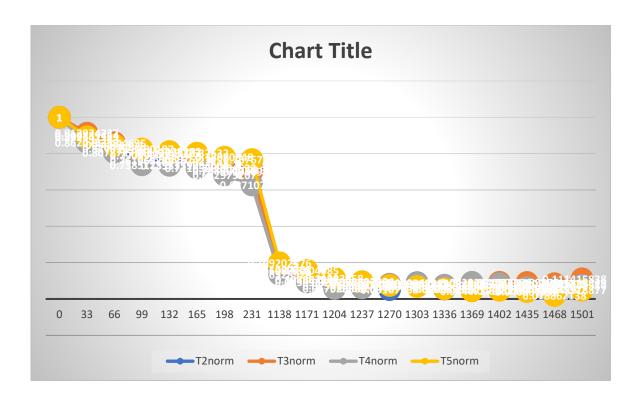
Normalization of noisy data is potentially a bad thing to do.

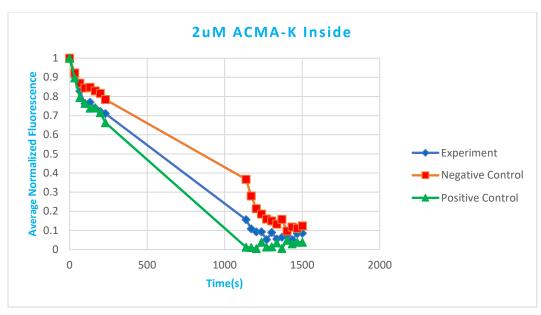


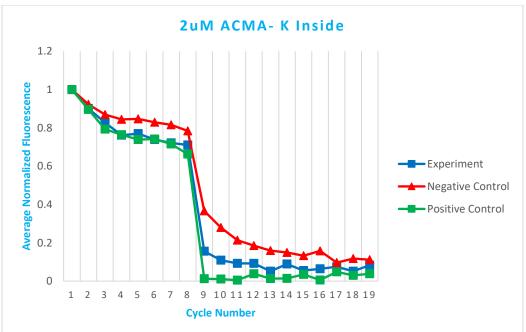








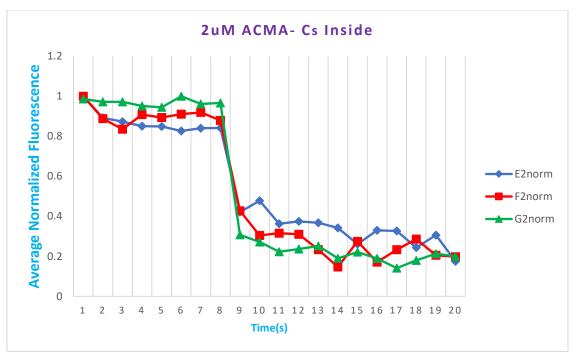


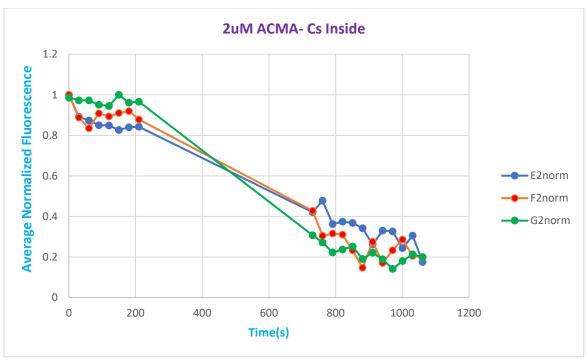


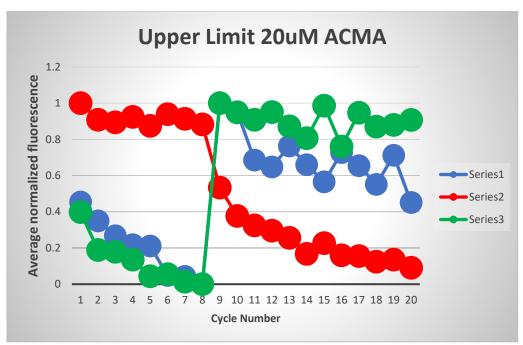
Gain of function experiments(gofs)

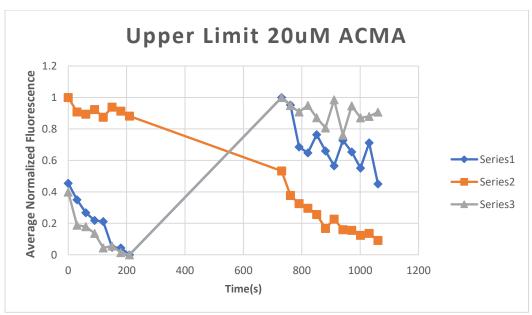
Loss of function experiments (lofs)

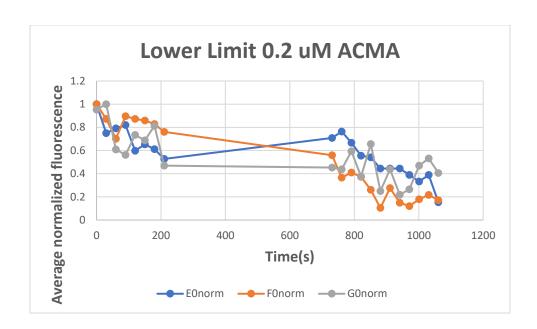
Base recording technique avoiding the recording and stimulating electrode from being together

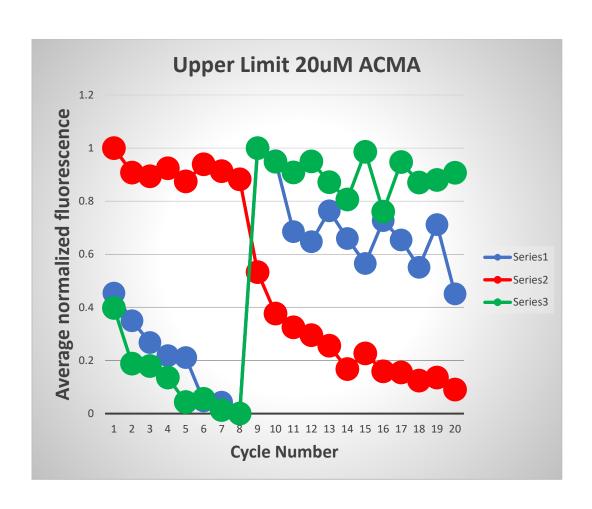


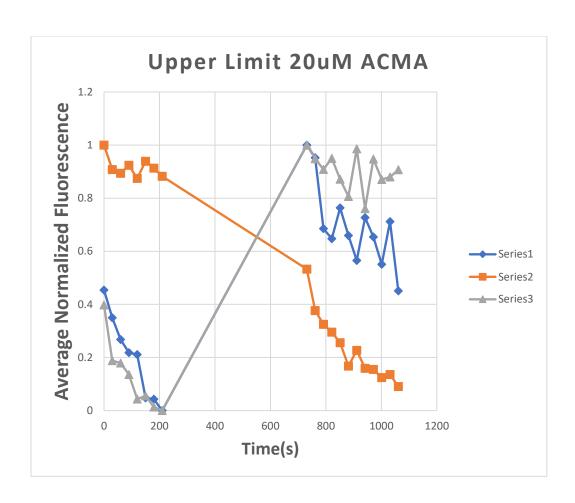


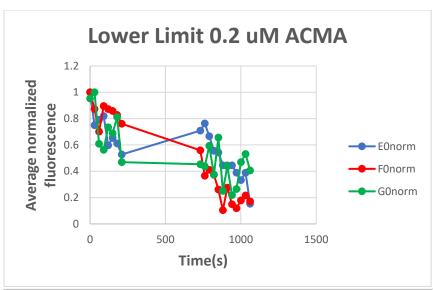


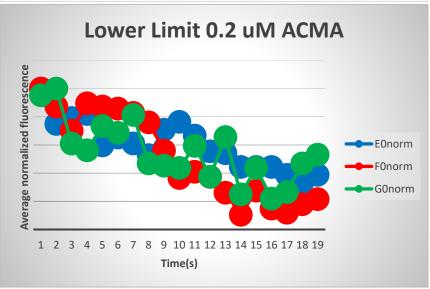


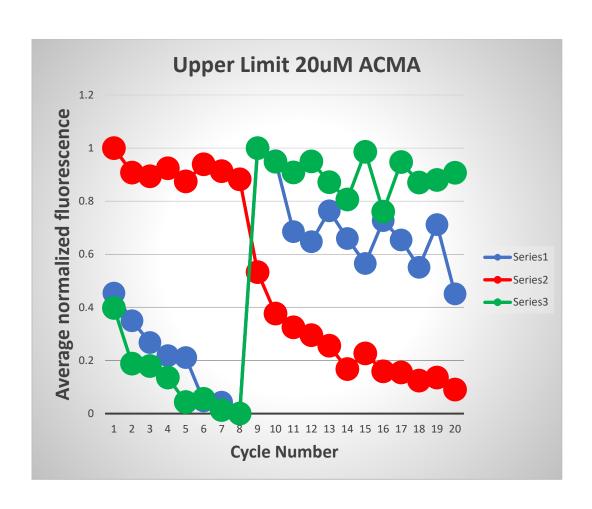












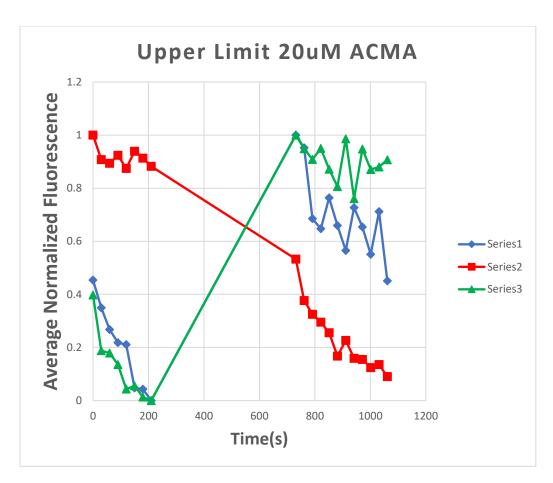


Figure 1: Averaged Normalized Fluorescence (+- SEM) vs Time (s) or Cycle Number, following NavAb flux assay experiments for three groups (Experiment, Negative and positive control). N =40 based on averaged data from triplicate experiments. Image legend

Data set names code to be used in plotting in R with no error bars

Navab221

Navab227

Legend

Figure 1: Averaged Normalized Fluorescence (+- SEM) vs Time (s) or Cycle Number, following NavAb flux assay experiments for three groups (Experiment, Negative and positive control). N = 40 based on averaged data from triplicate experiments. Image legend represents groups correctly. (Source: R Studio)