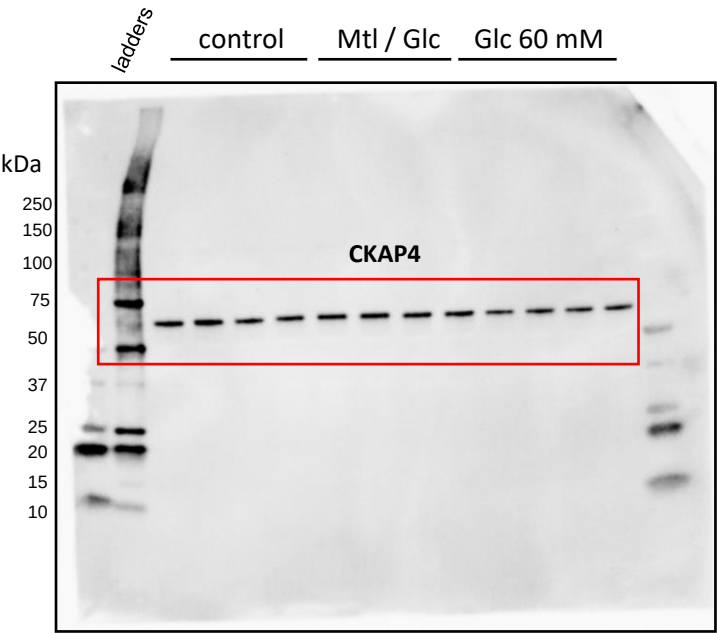
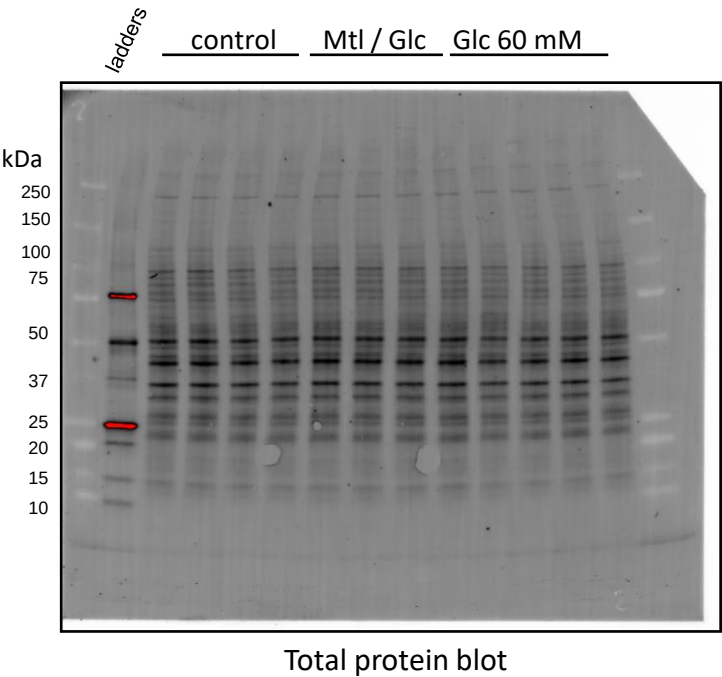
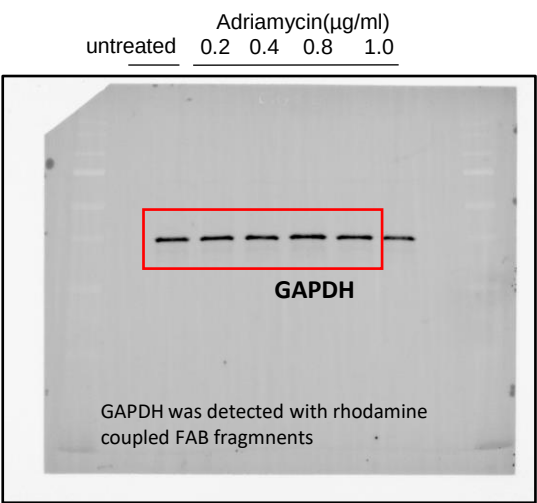
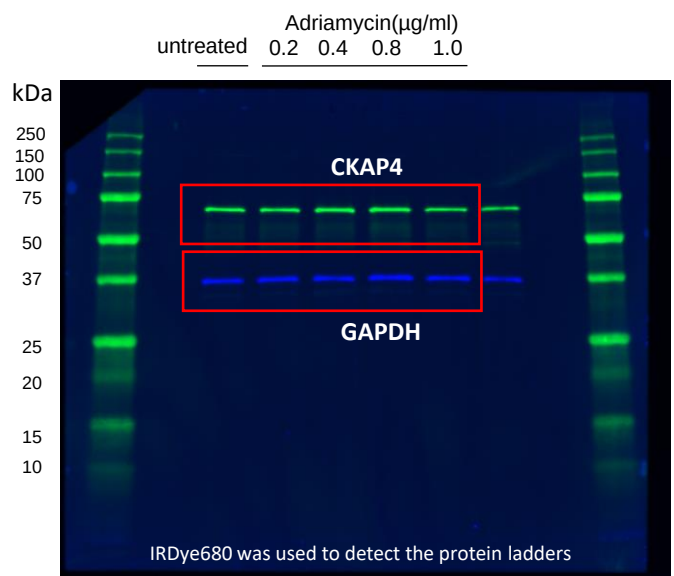
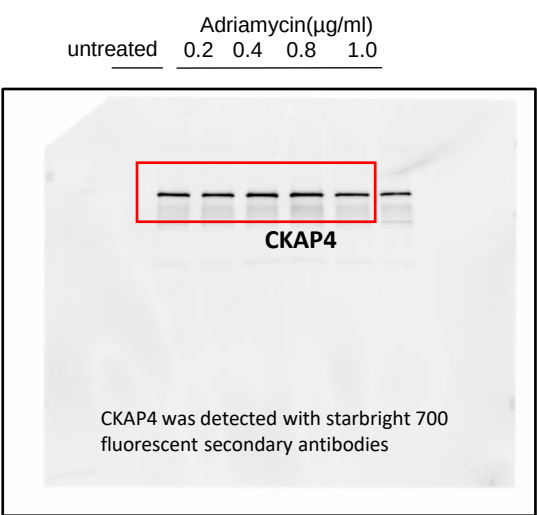
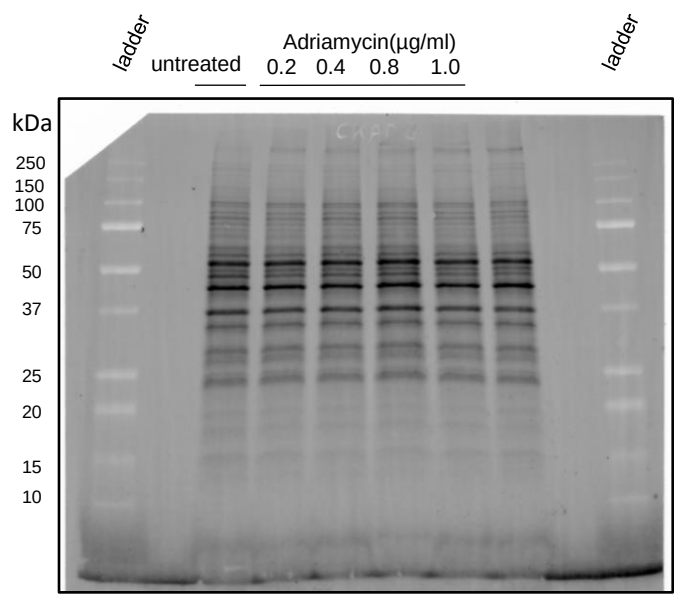


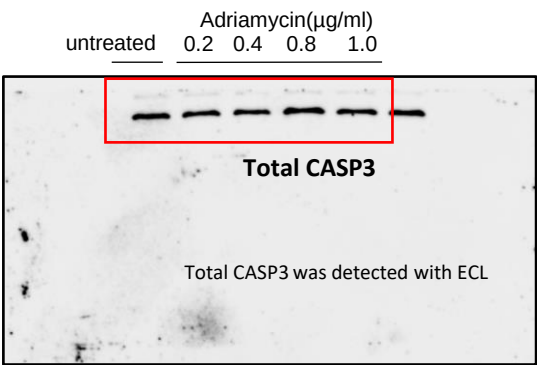
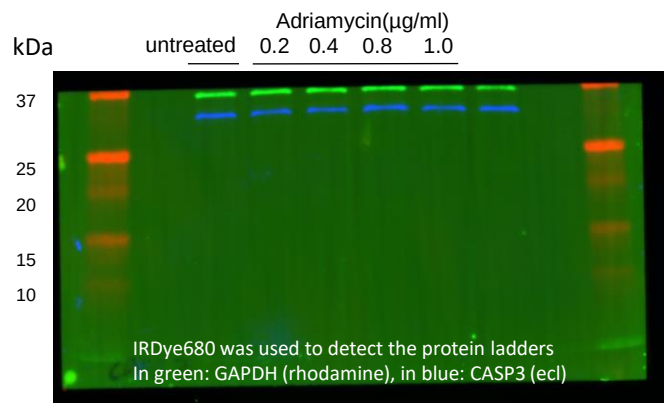
Full unedited blots for Figure 1E



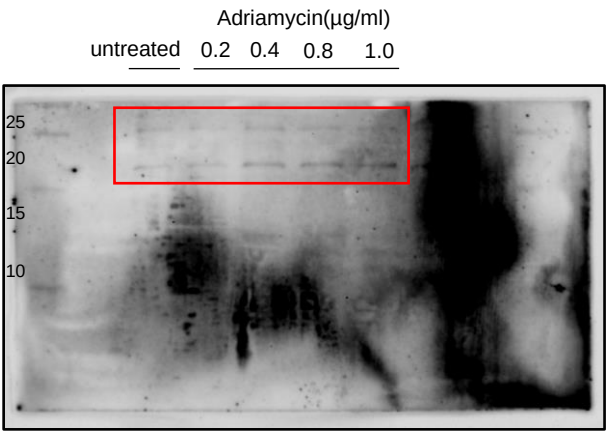
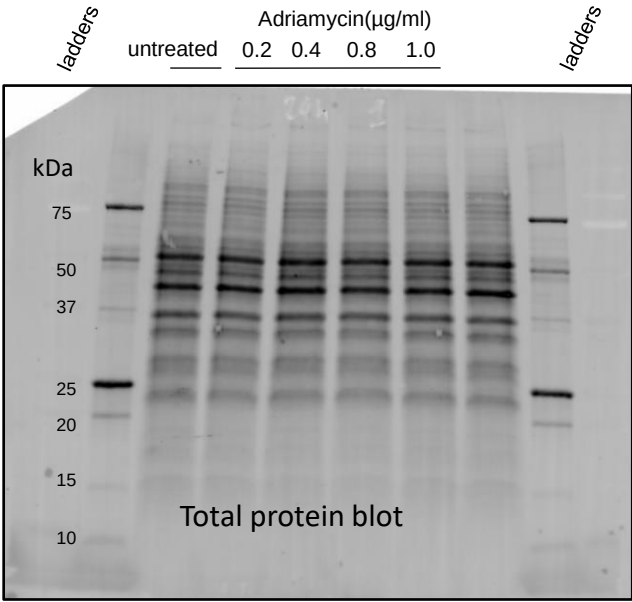
Full unedited blots for Figure 1F, part 1



The same membrane used for CKAP4 and GAPDH was cut at 37 kDa and used for the detection of total CASP 3 (31 kDa)



Full unedited blots for Figure 1F, part 2



Cleaved CASP3

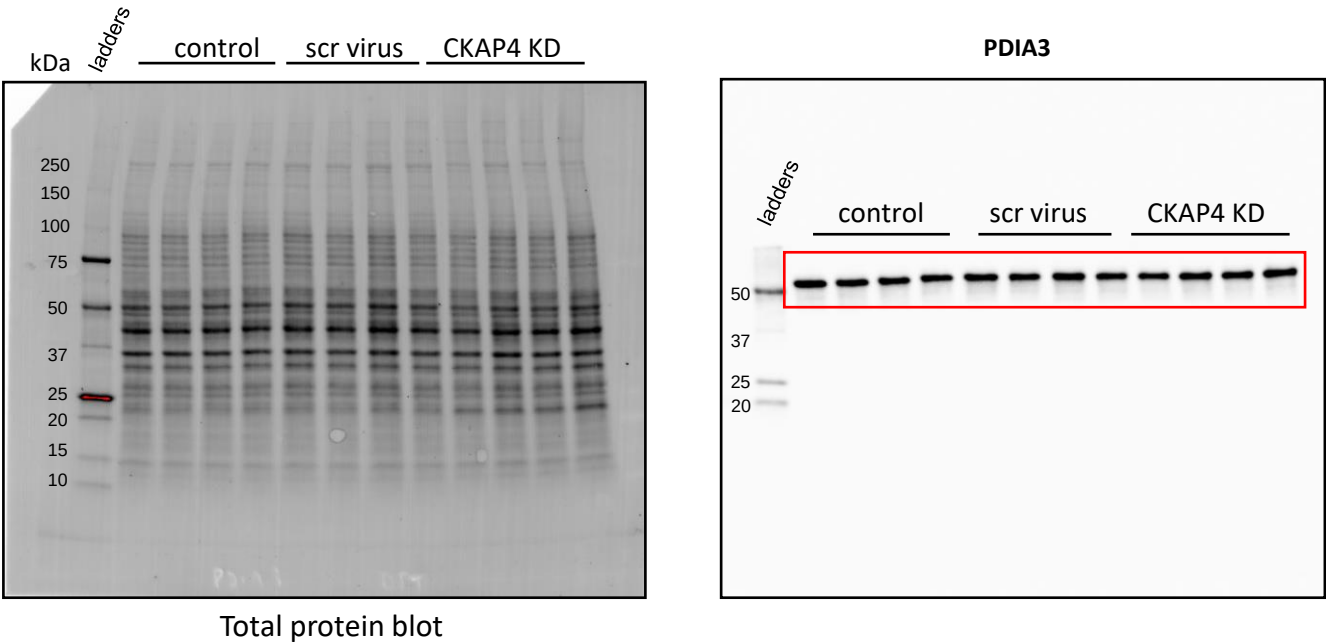
The membrane was cut at 25 Kda

Full unedited representative blots for Figure 3B

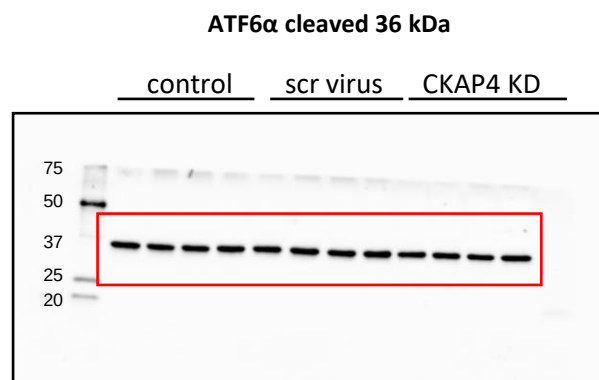
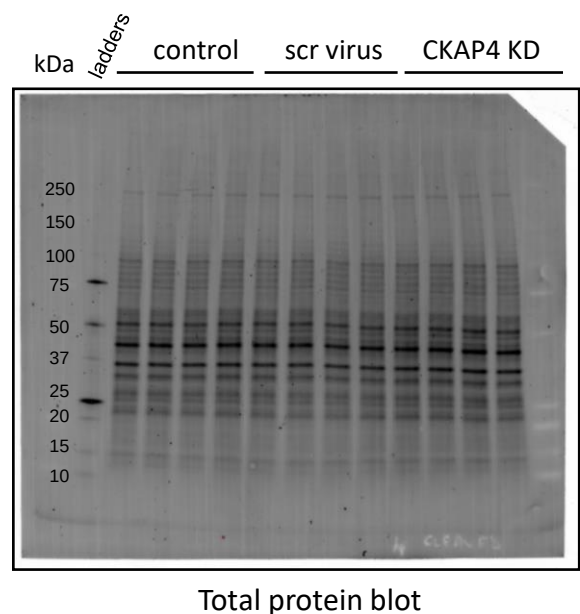
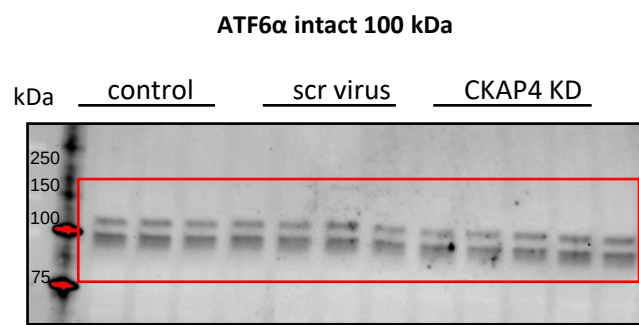
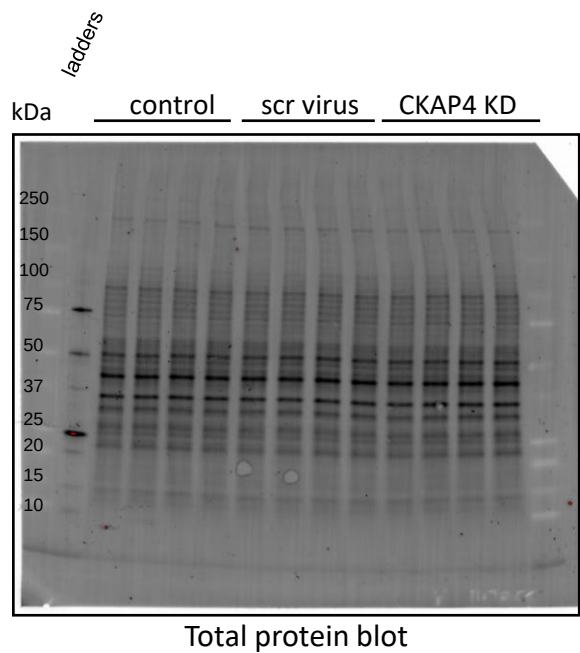


For Figure 3B, we have decided to use GAPDH as normalization factor instead of the total lane signal. Even when the lanes are equally loaded in terms of μg of total protein, protein content was broadly different when comparing the three cell types investigated (different proteomes). Thus, we decide against using total lane normalization (only) in this case.

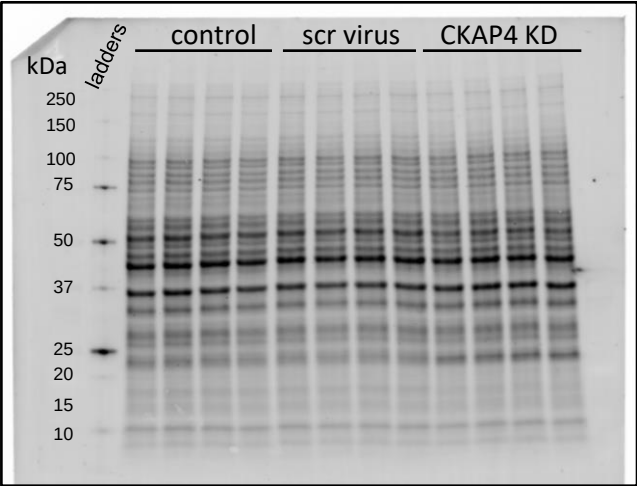
Full unedited blots for Figure 5B



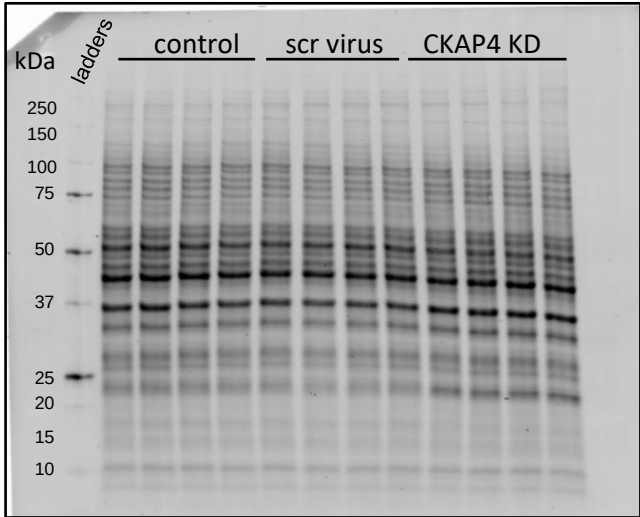
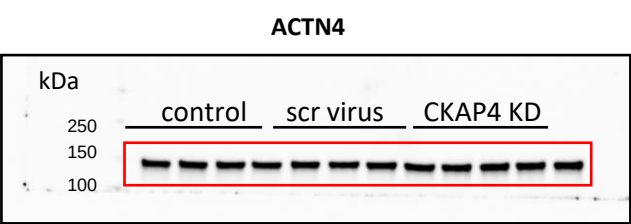
Full unedited blots for Figure 5C



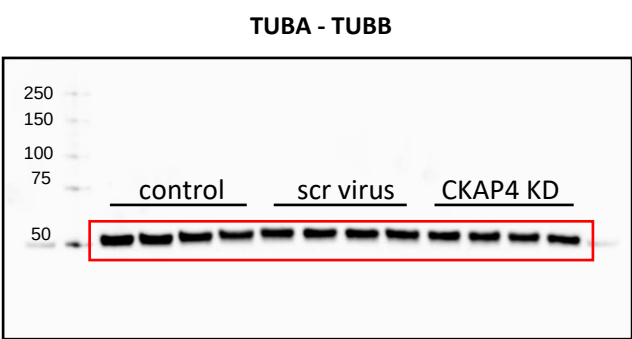
Full unedited blots for Figure 6A



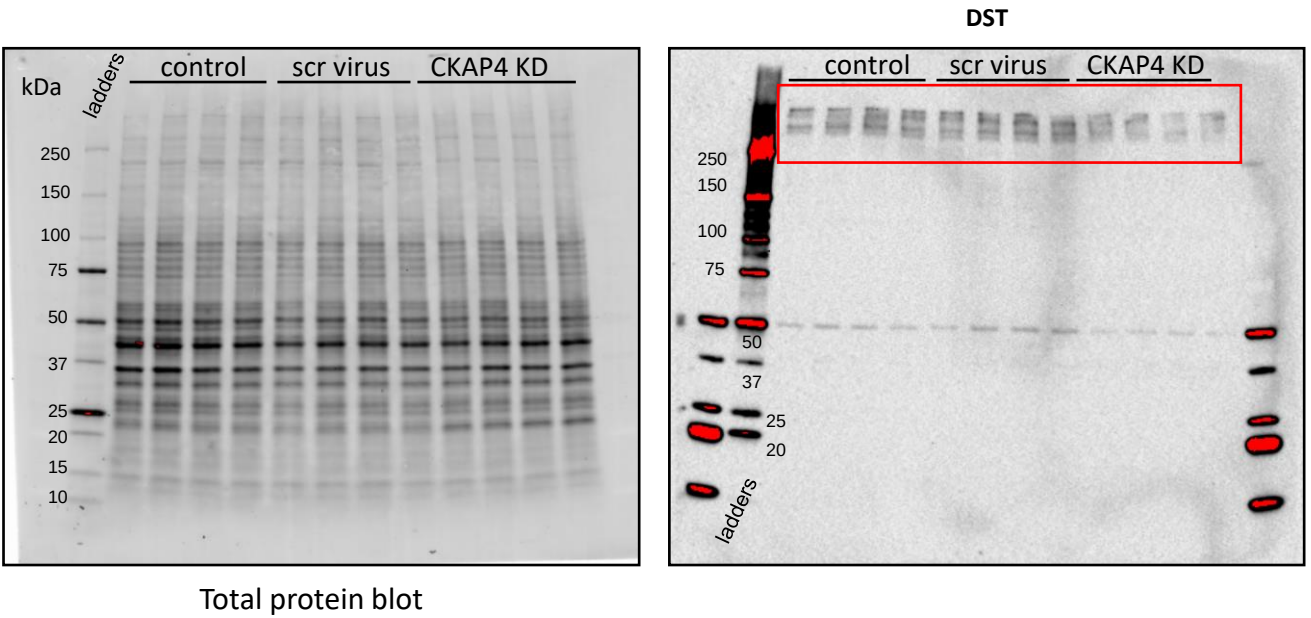
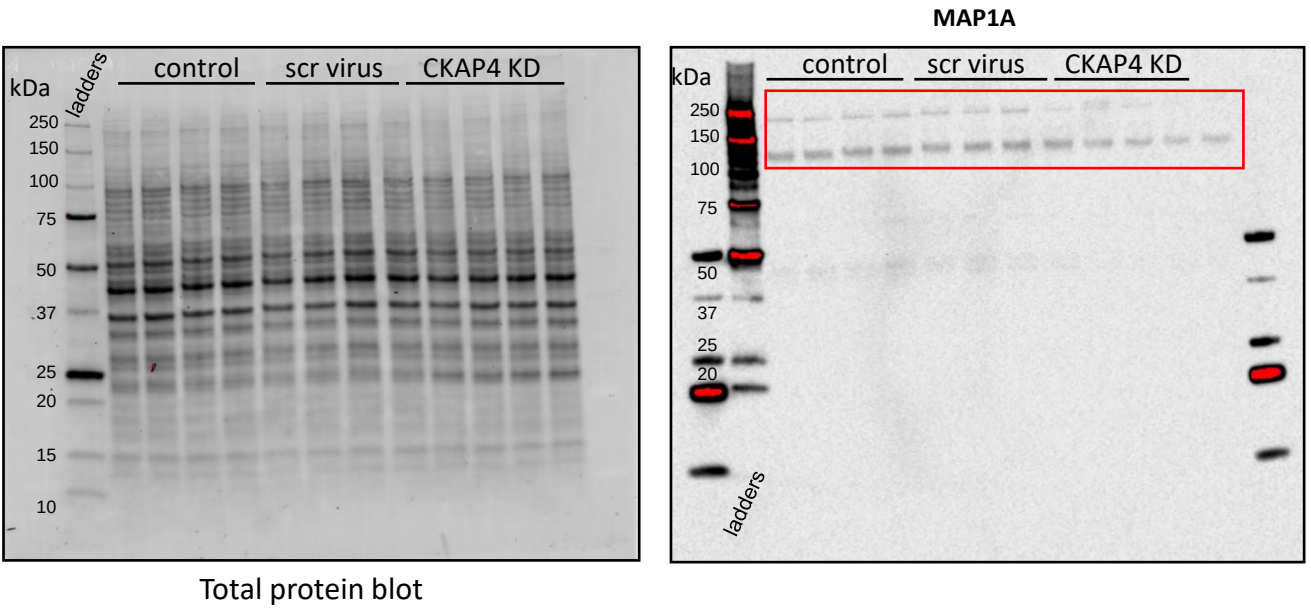
Total protein blot



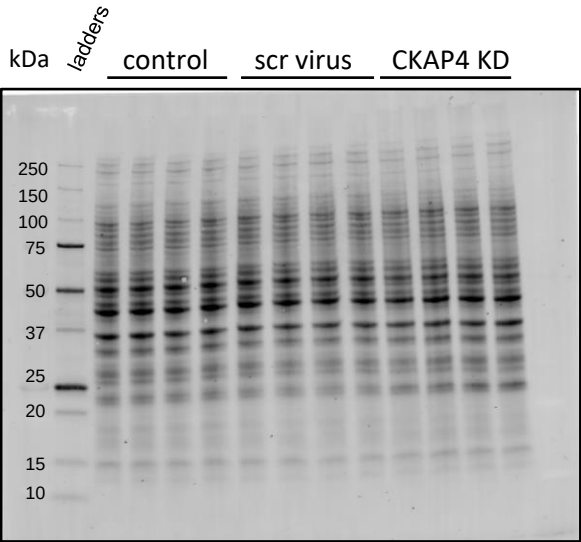
Total protein blot



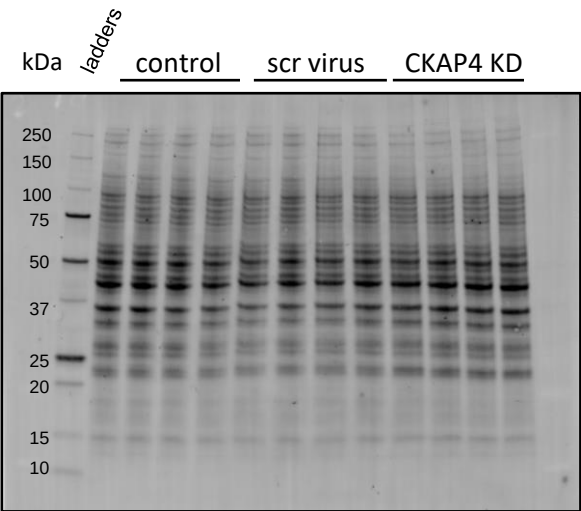
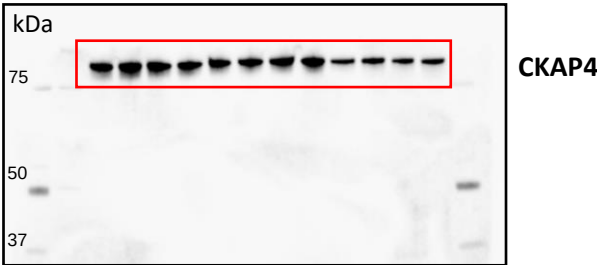
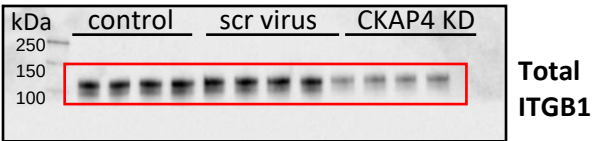
Full unedited blots for Figure 7D



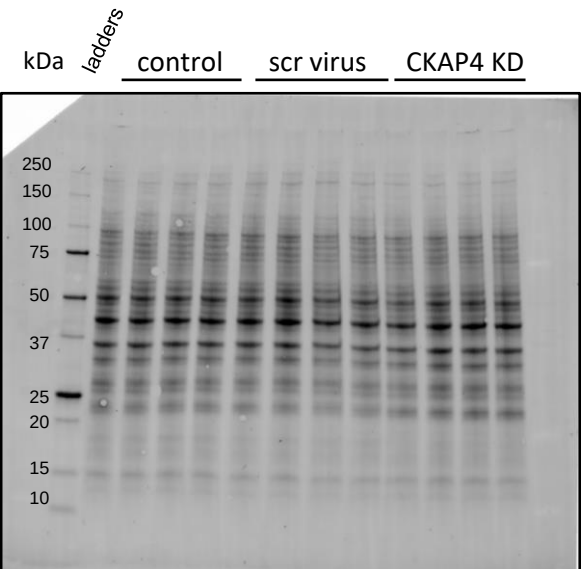
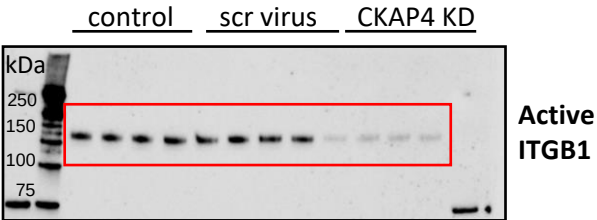
Full unedited blots for Figure 8B, part 1



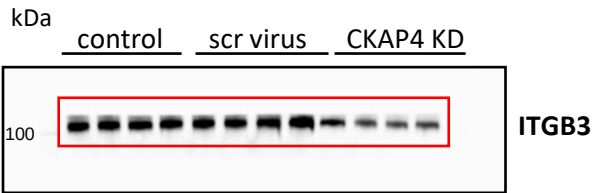
Total protein blot



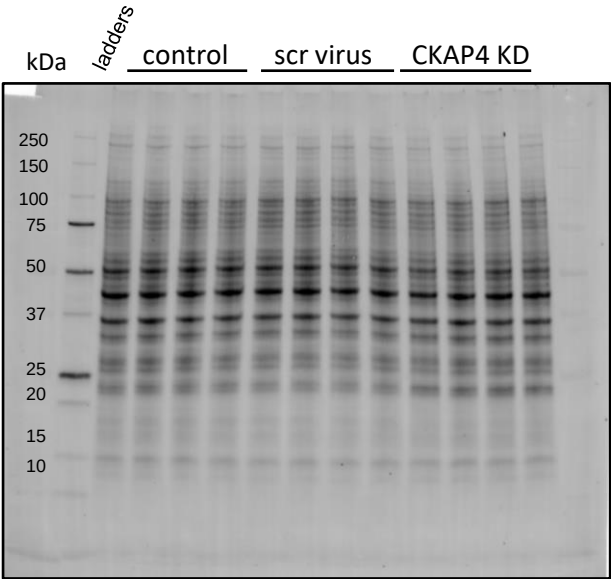
Total protein blot



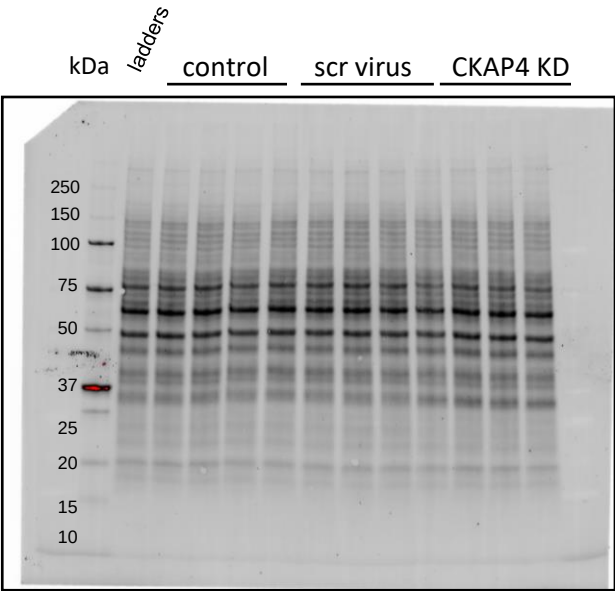
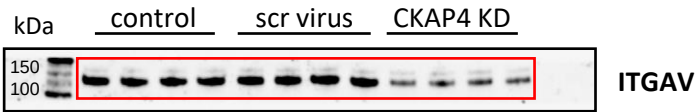
Total protein blot



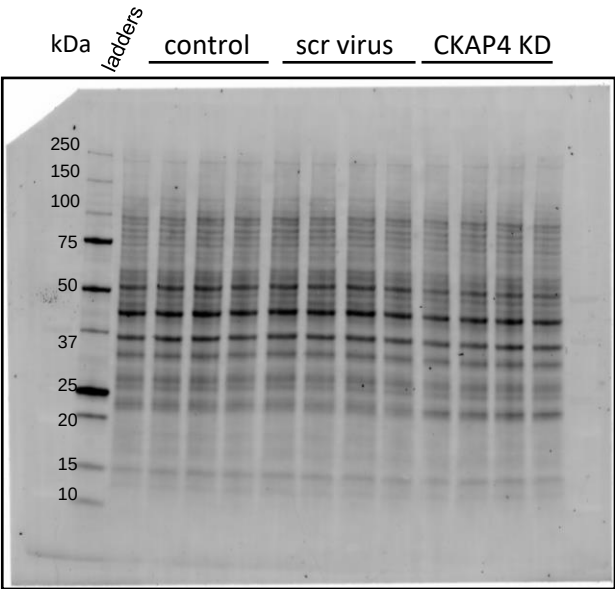
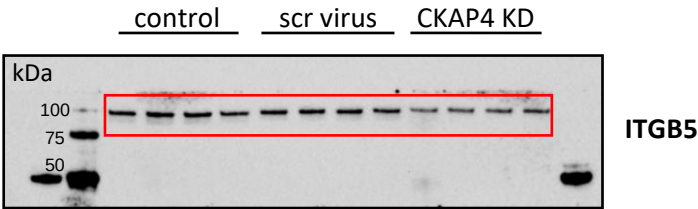
Full unedited blots for Figure 8B, part 2



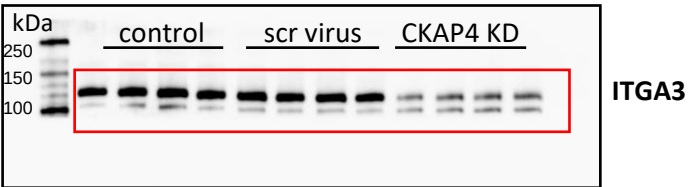
Total protein blot



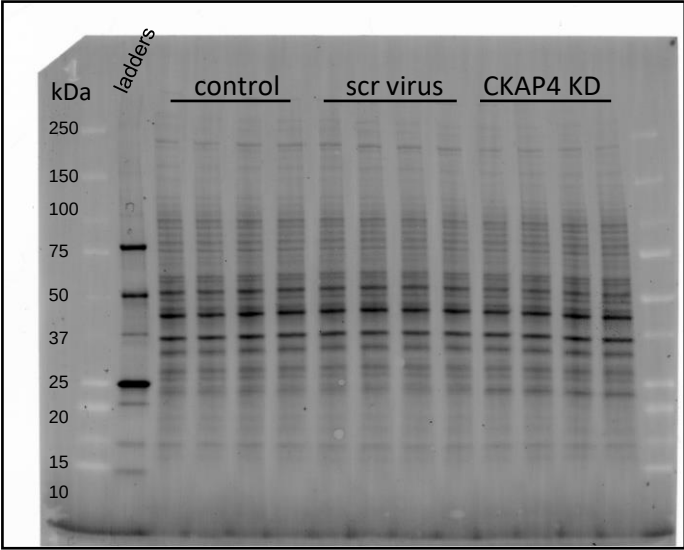
Total protein blot



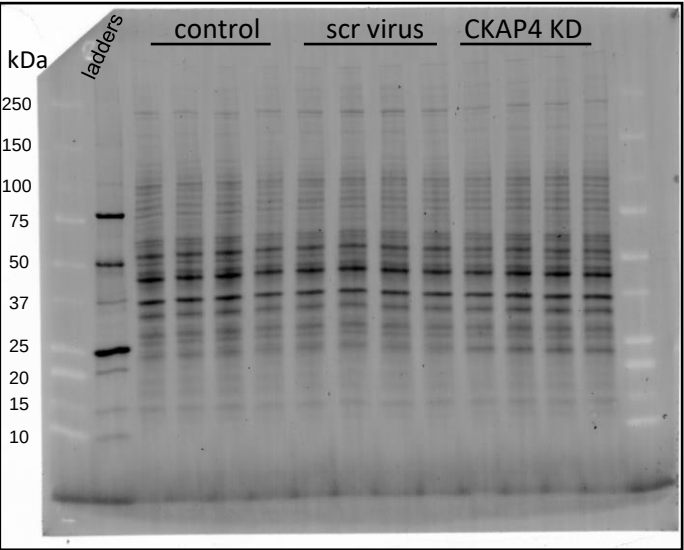
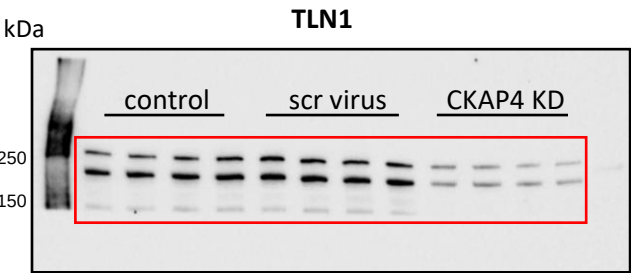
Total protein blot



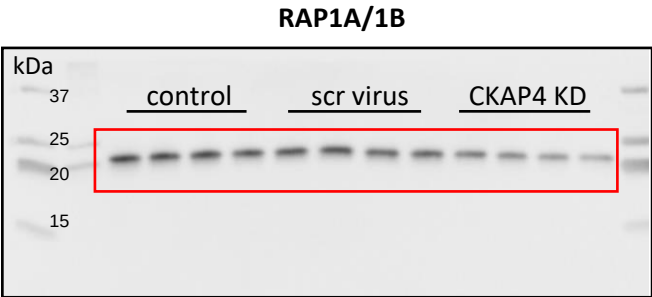
Full unedited blots for Figure 8B, part 3



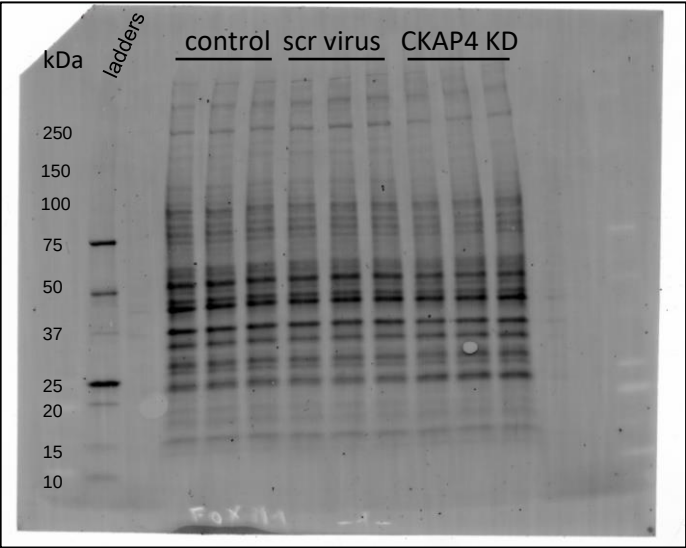
Total protein blot



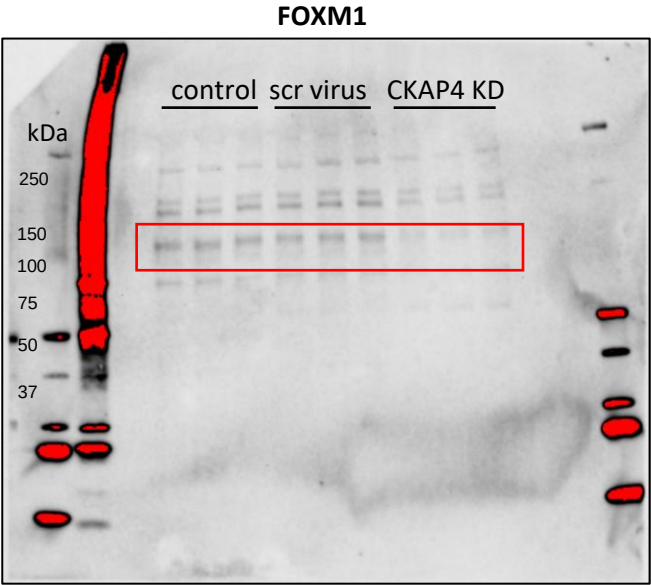
Total protein blot



Full unedited blots for Figure 8D

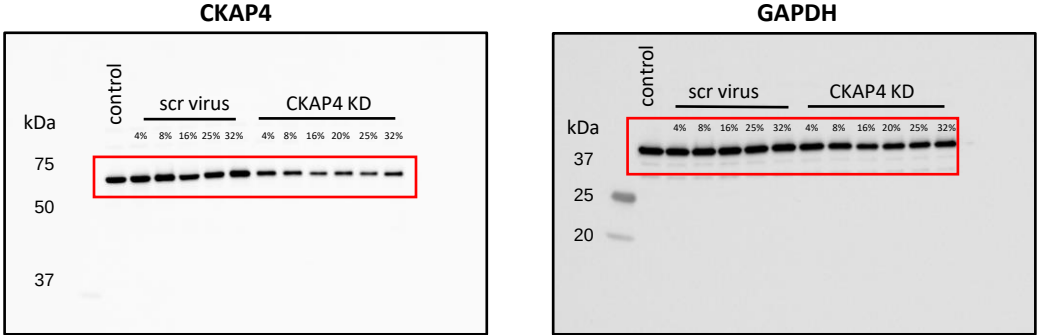


Total protein blot



Full unedited blots for Supplemental Figure 1

A



B

