

Figure 2B

<https://hgserver1.amc.nl/cgi-bin/r2/main.cgi>

Dataset: [GSE62564](#)

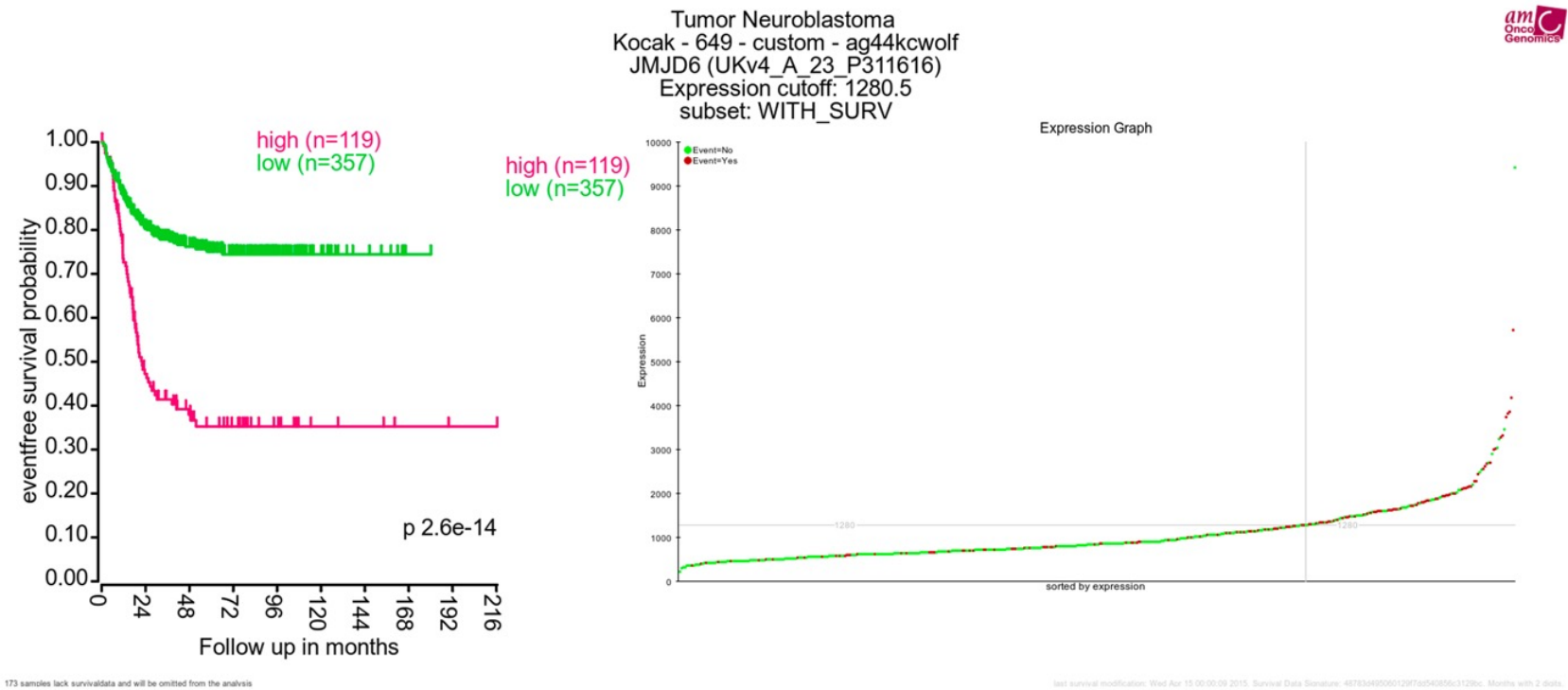
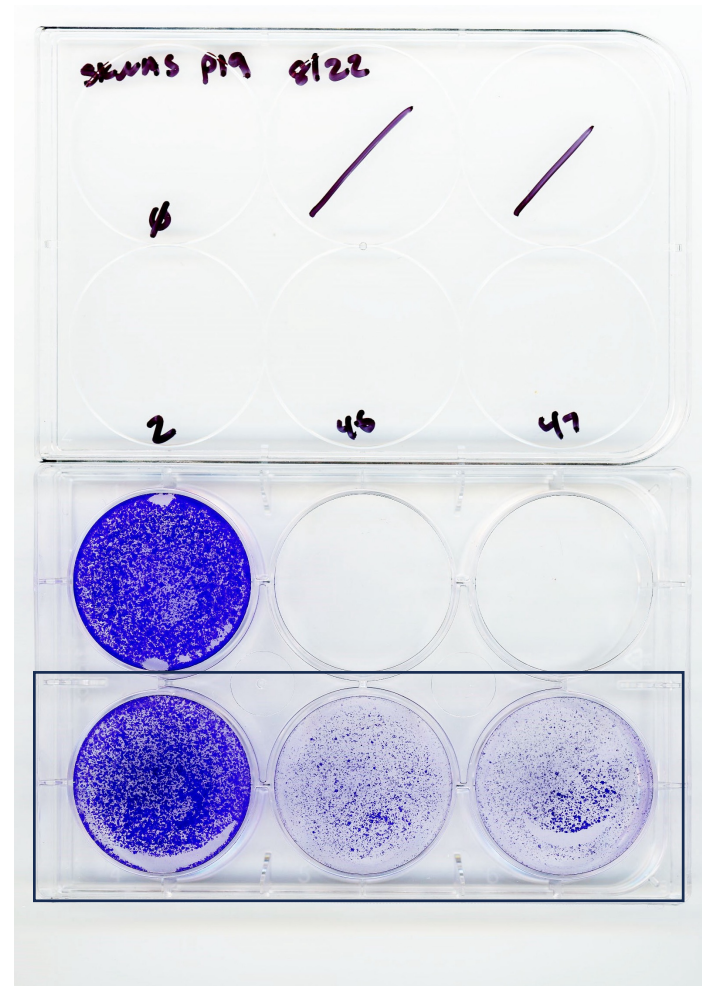
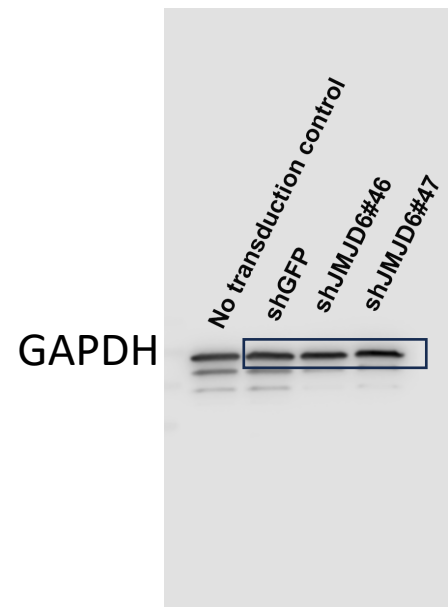
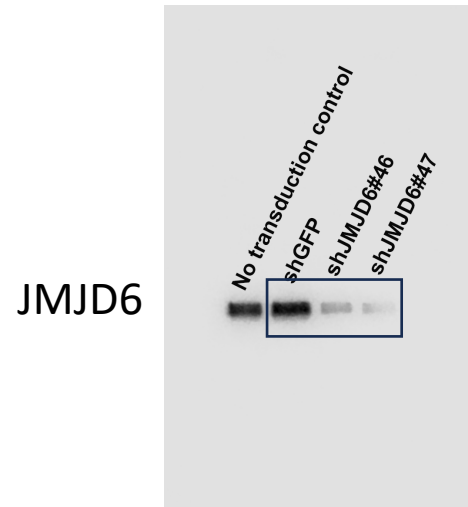


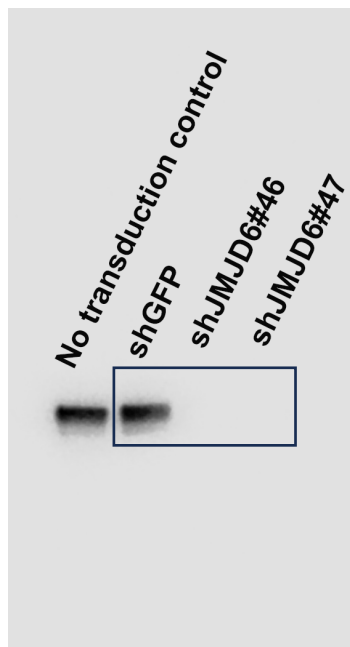
Figure 2C- SKNAS



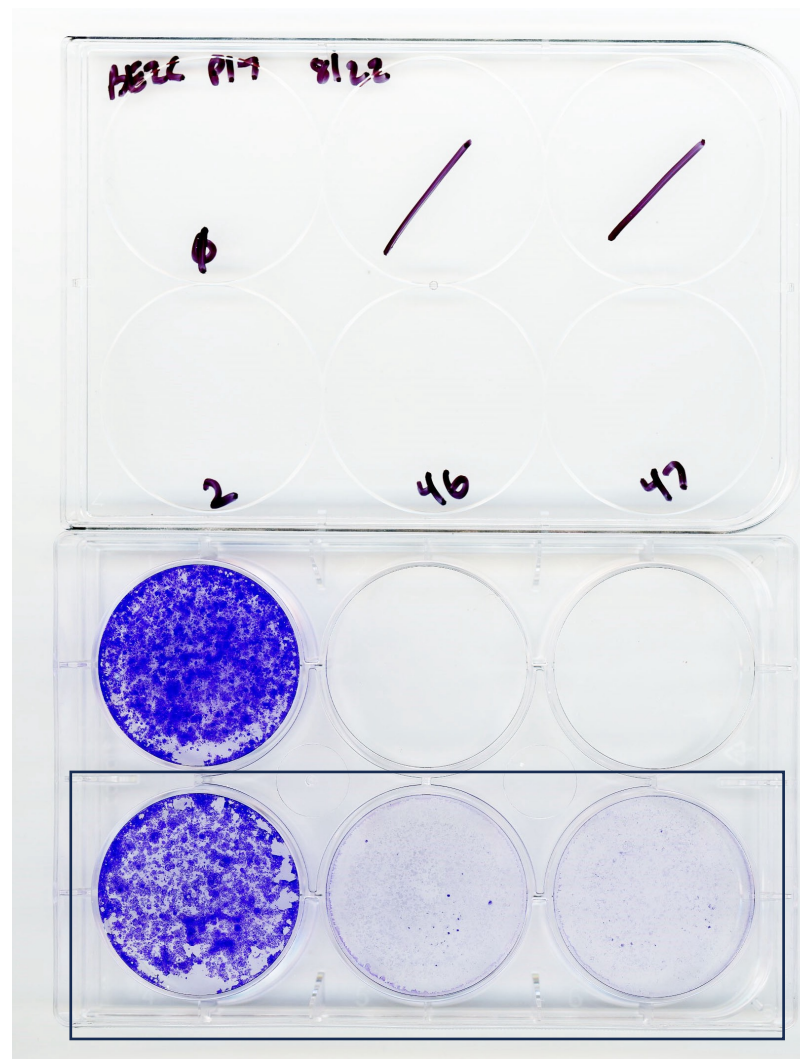
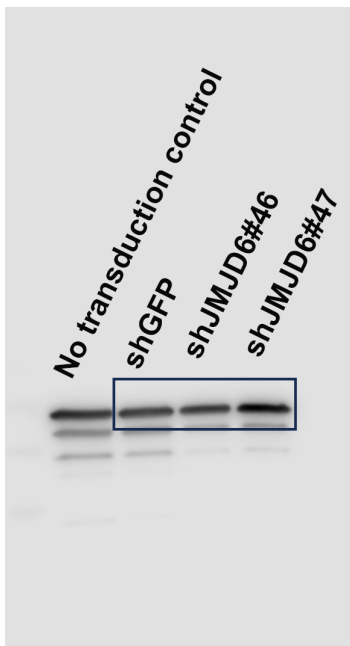
Ø = non transduction control
2 = shGFP
46 = shJMJD6#46
47 = shJMJD6#47

Figure 2C- BE2C

JMJD6



GAPDH



Ø = non transduction control

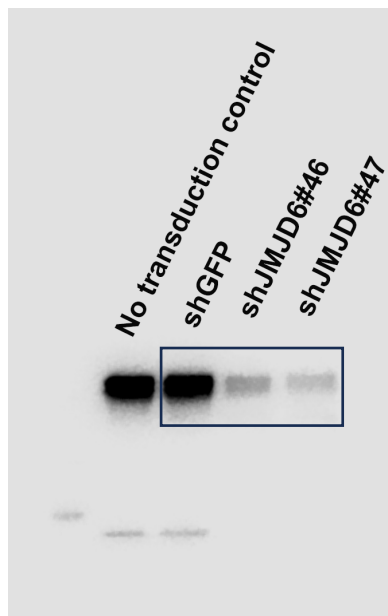
2 = shGFP

46 = shJMJD6#46

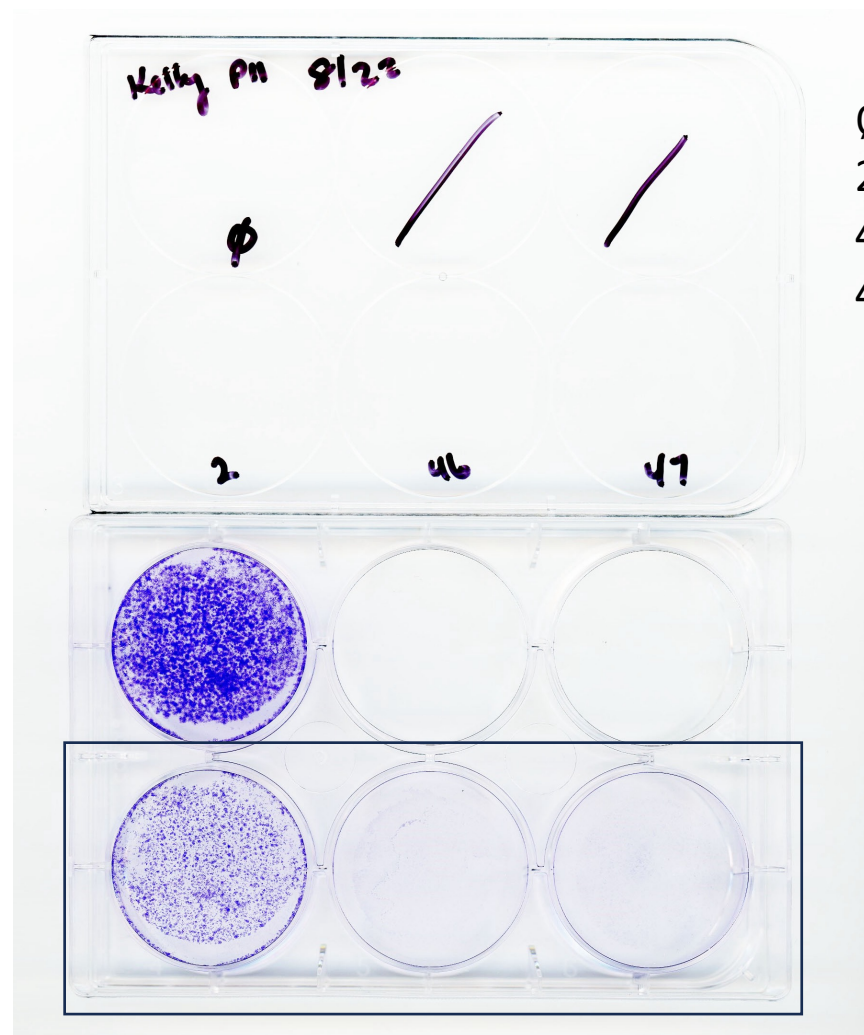
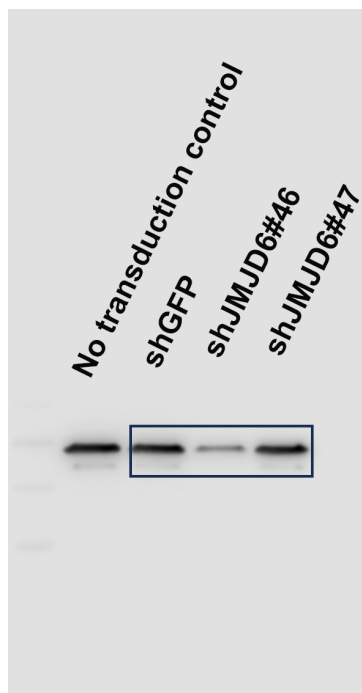
47 = shJMJD6#47

Figure 2C- KELLY

JMJD6



GAPDH



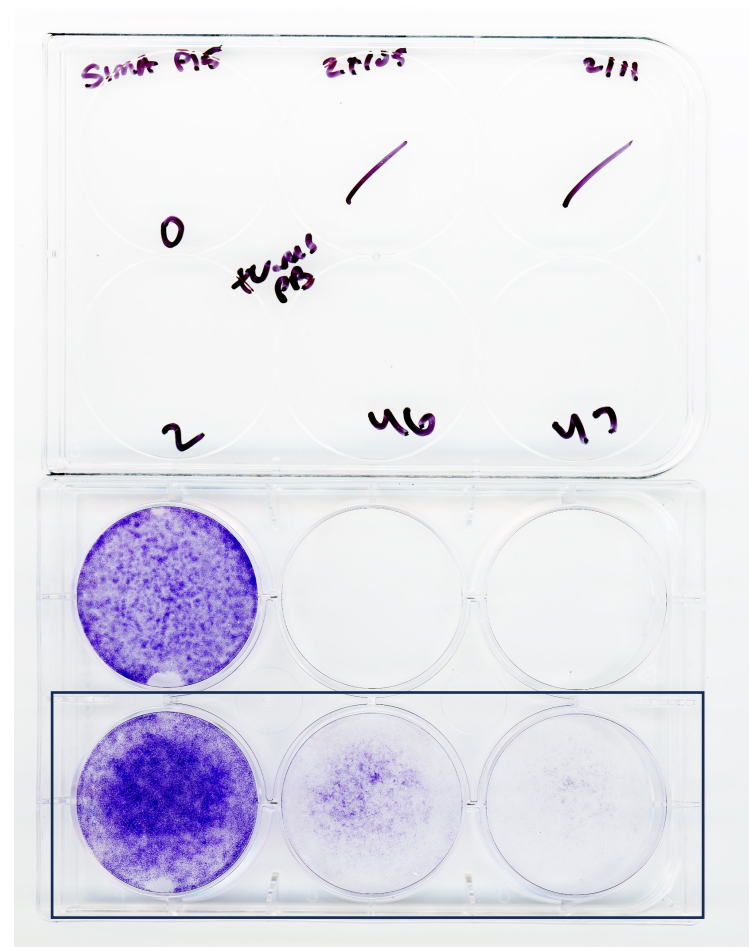
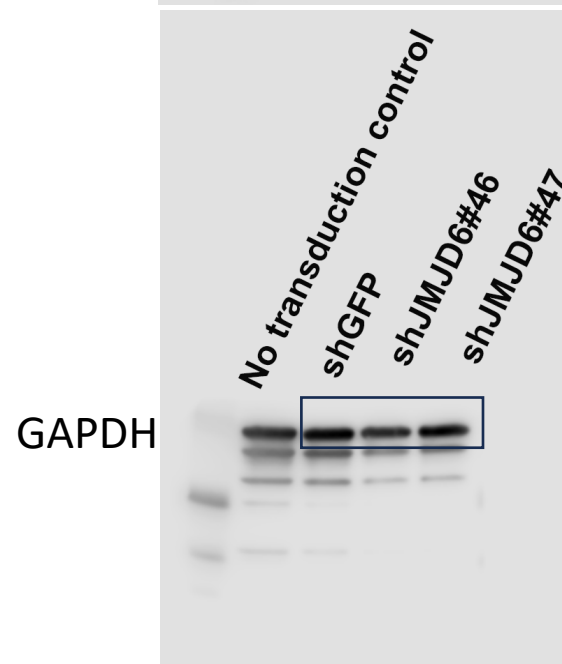
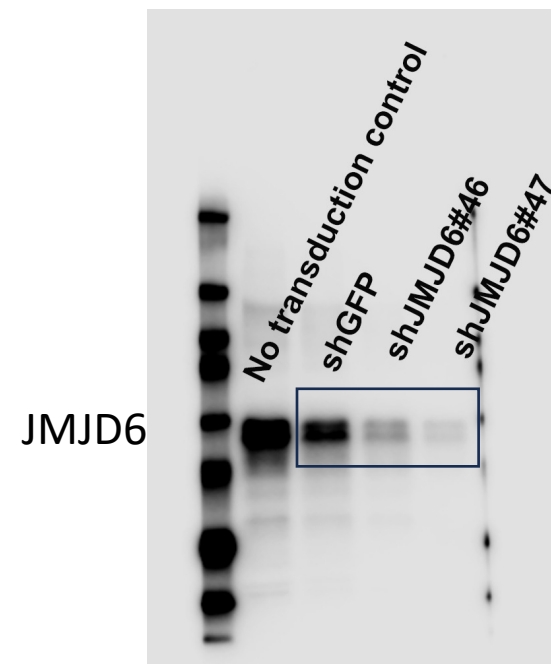
Ø = non transduction control

2 = shGFP

46 = shJMJD6#46

47 = shJMJD6#47

Figure 2C- SIMA



\emptyset = non transduction control
 2 = shGFP
 46 = shJMJD6#46
 47 = shJMJD6#47

Figure 2C- IMR32 and CHLA20

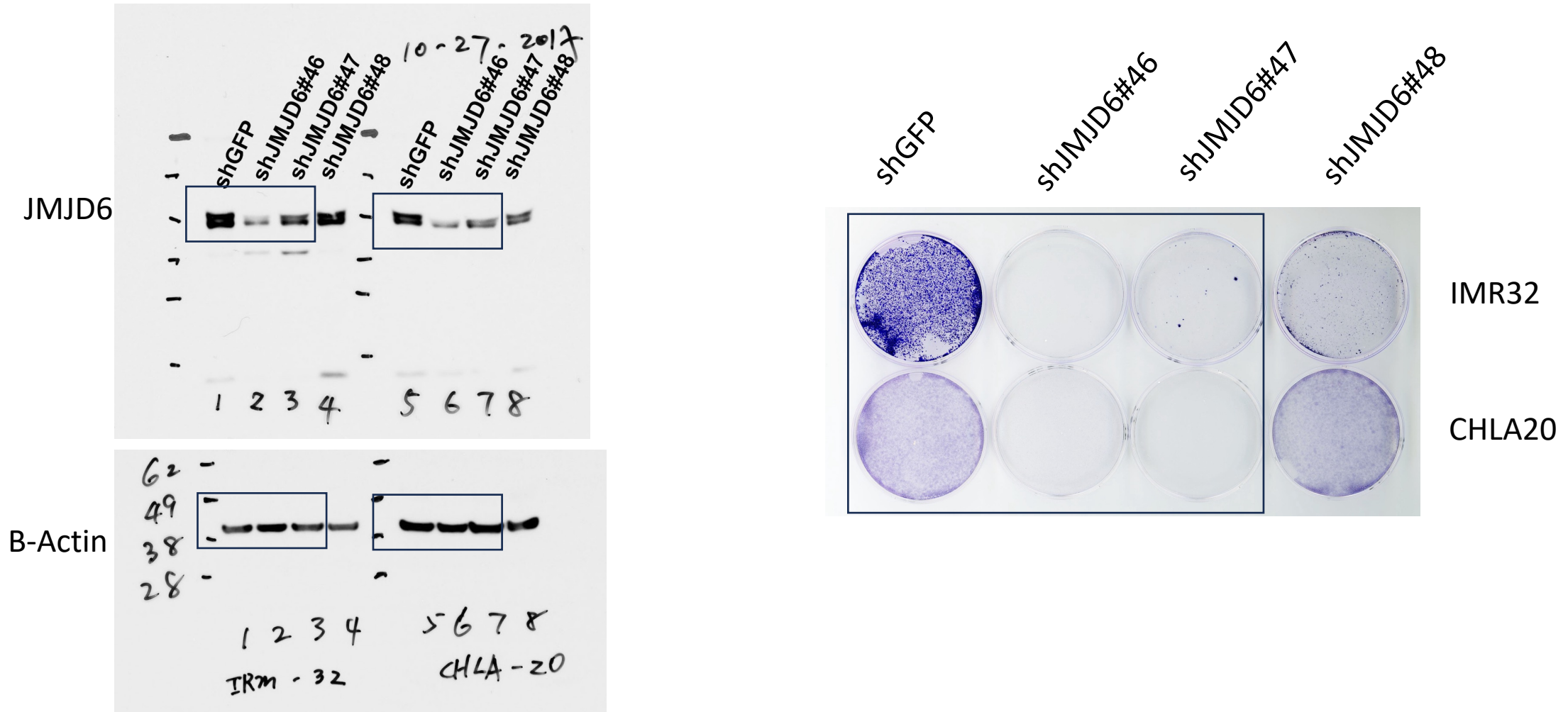


Figure 2F

