

Fig. 4A ENAH

GAPDH

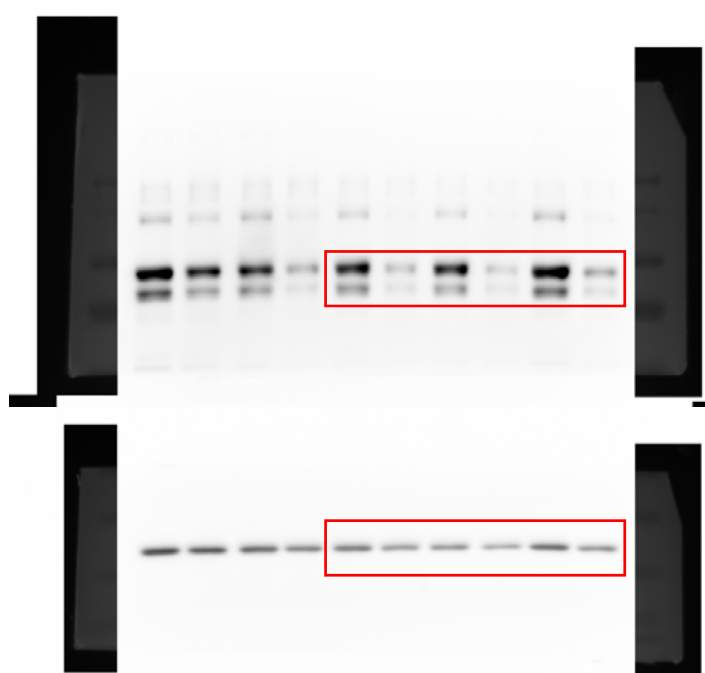


Fig. 4B ENAH

GAPDH

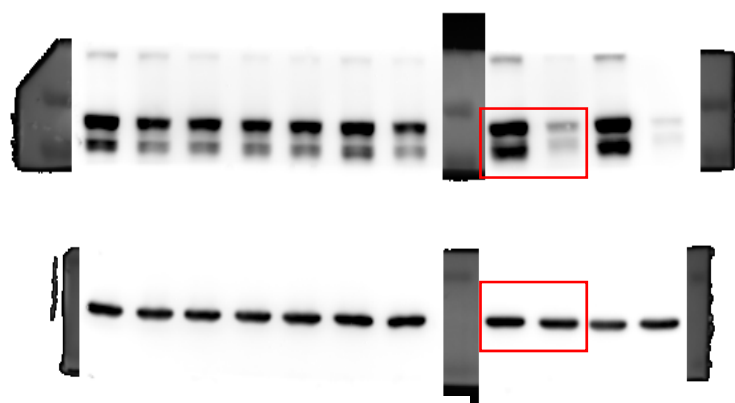


Fig. 4C ENAH

GAPDH

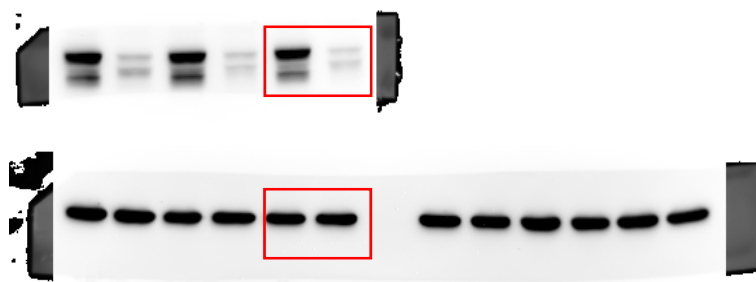


Fig. 4D ENAH



GAPDH

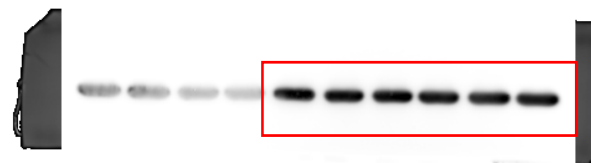


Fig. 4E ENAH



GAPDH

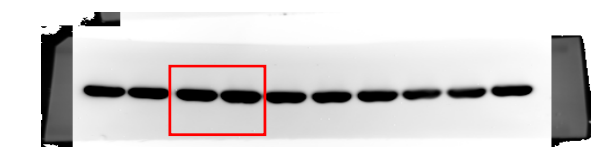


Fig. 4F ENAH



GAPDH

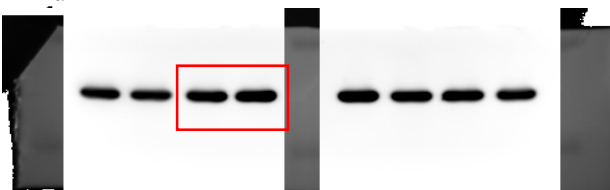


Fig. 5B

ENAH

p-AKT
(Ser473)

AKT

β -catenin

β -catenin
(active)

ITGB5

GAPDH



Fig. 5D ENAH

p-AKT
(Ser473)

AKT

GAPDH

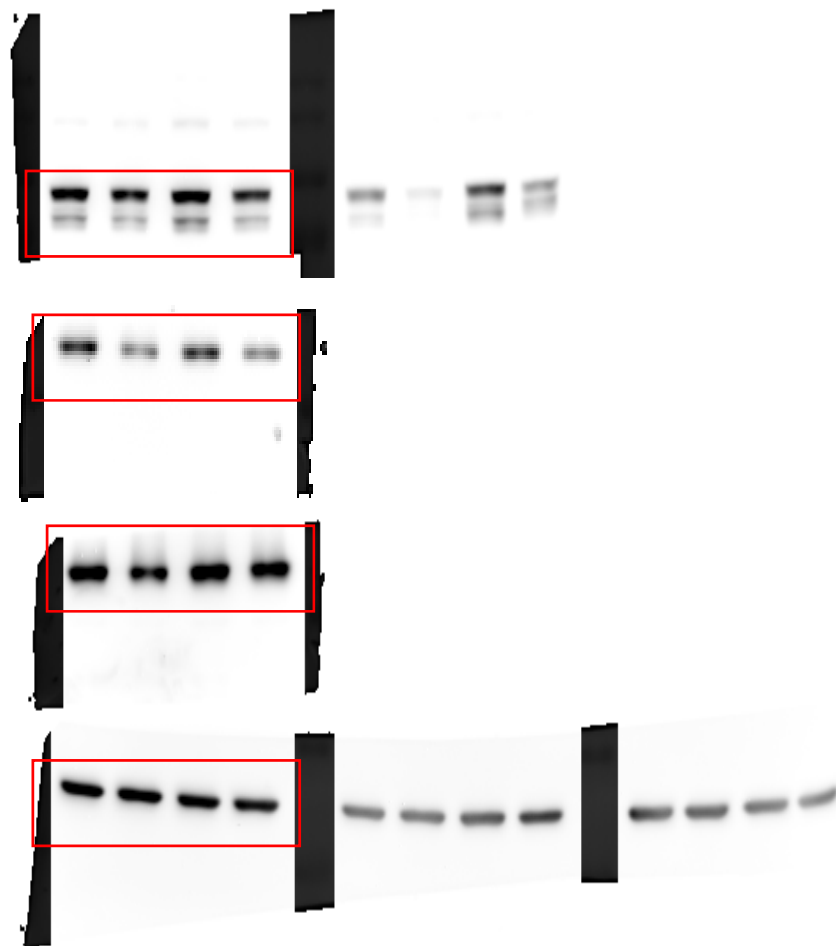
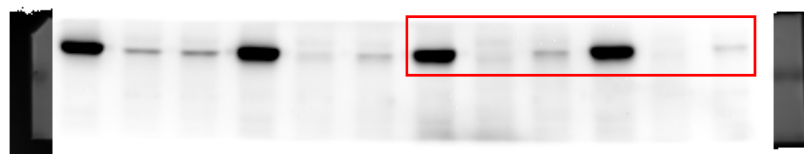


Fig. 5F

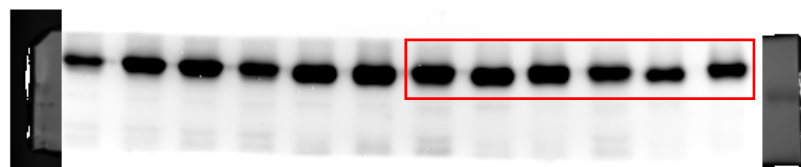
ENAH



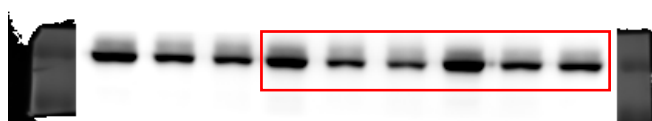
p-AKT
(Ser473)



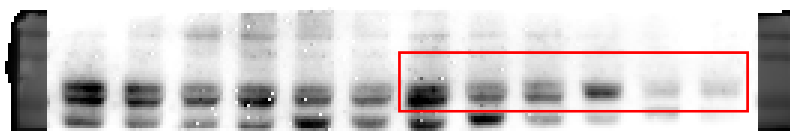
AKT



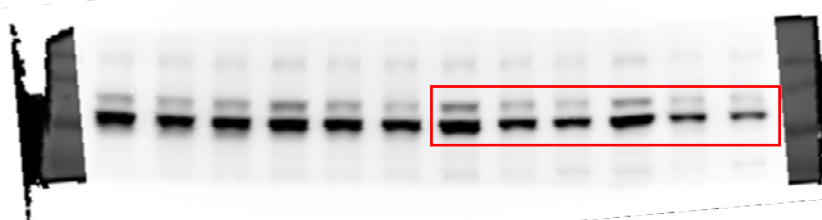
β -catenin



β -catenin
(active)



ITGB5



GAPDH

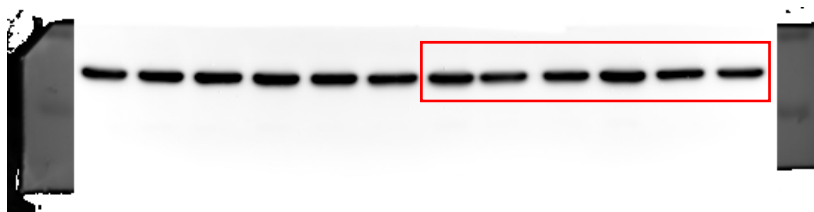


Fig. 5H



β -catenin



GAPDH

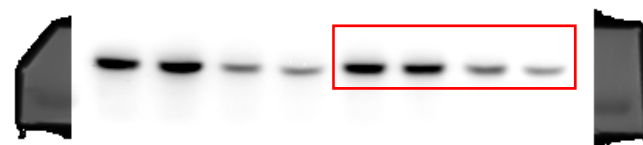


Fig. 5I

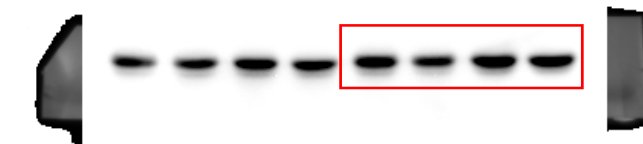
ENAH



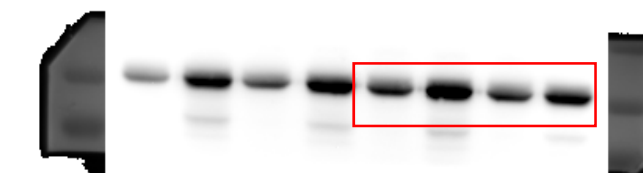
p-AKT
(Ser473)



AKT



β -catenin



GAPDH

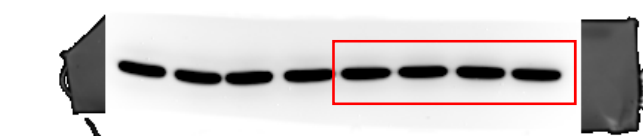


Fig. 5J

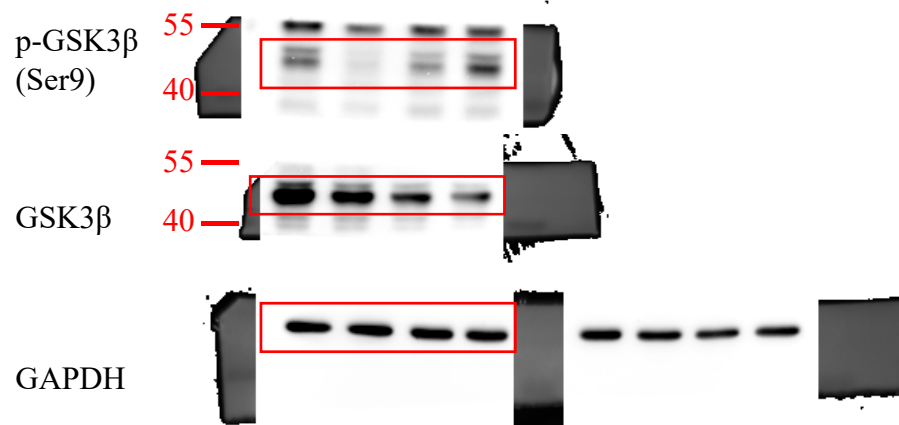


Fig. 6C
(left
Panel)

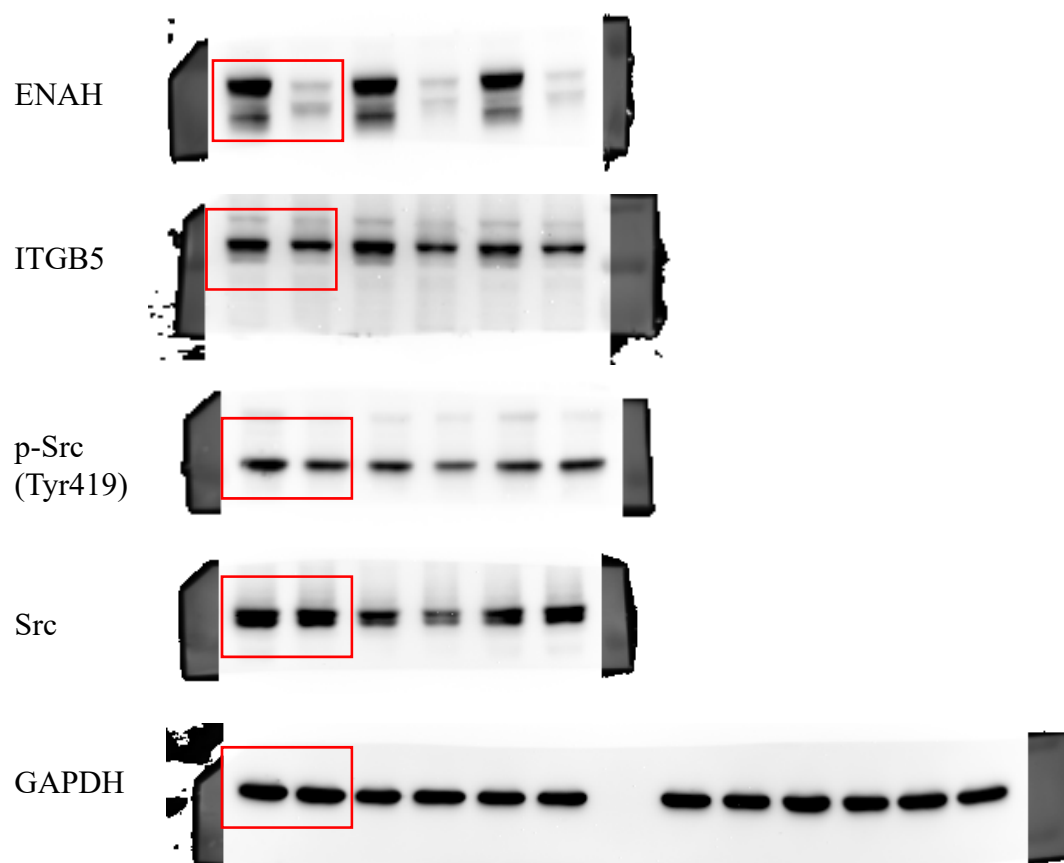
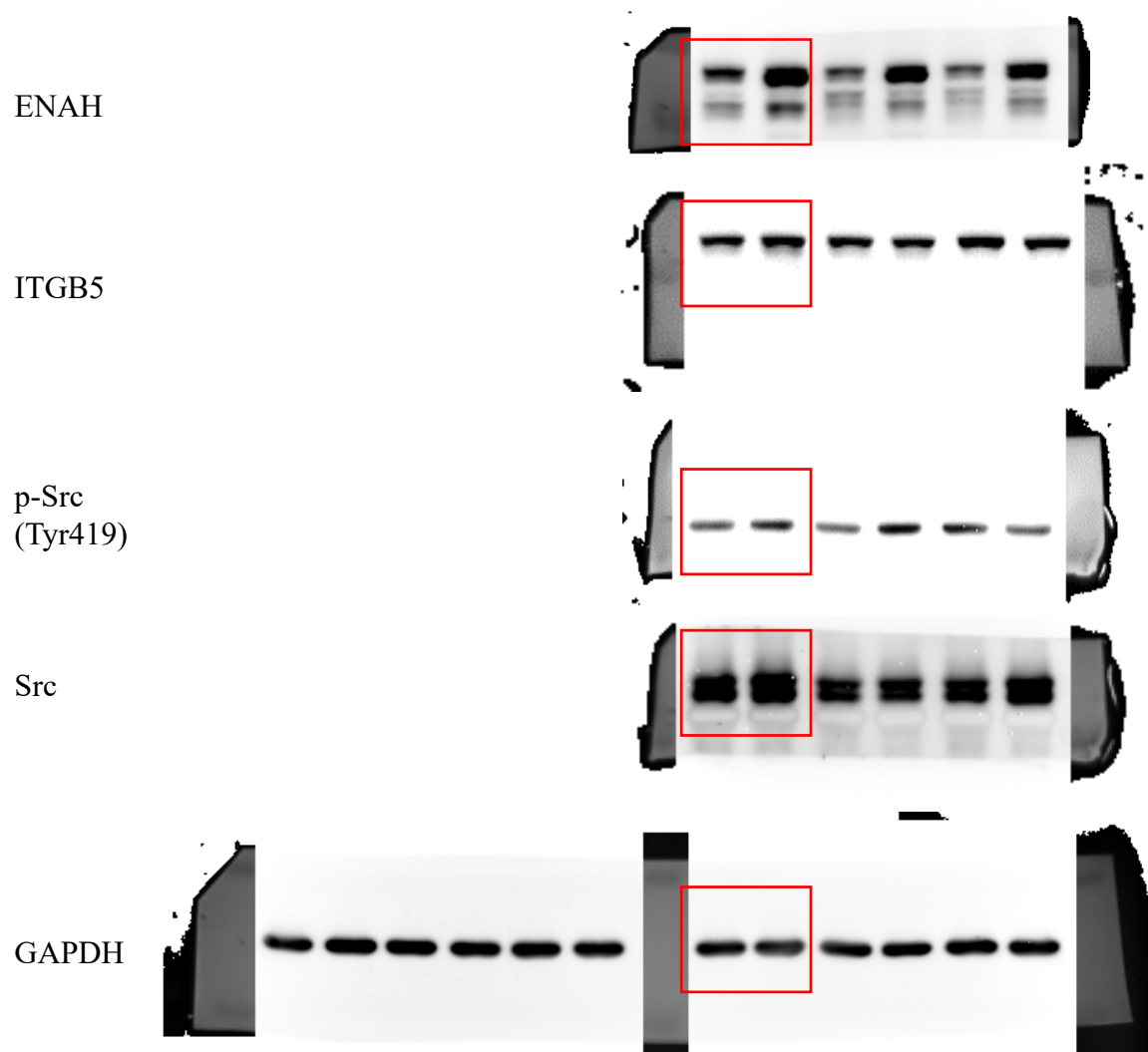


Fig. 6C
(right
Panel)



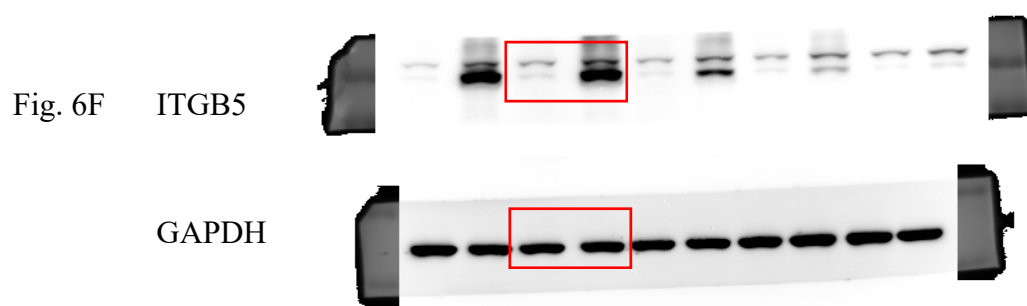
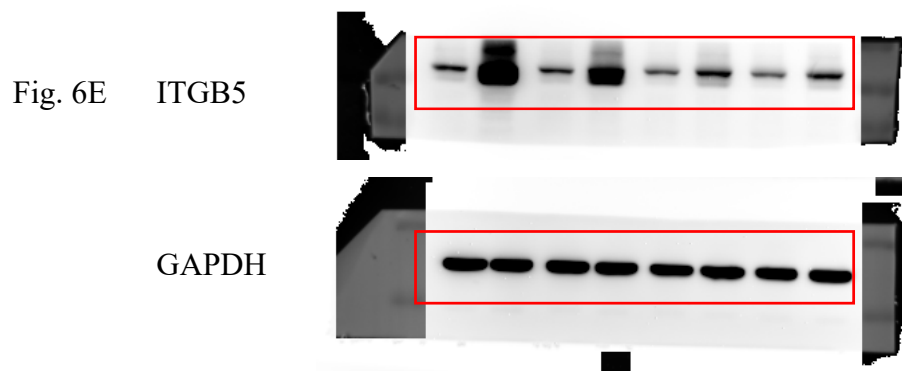
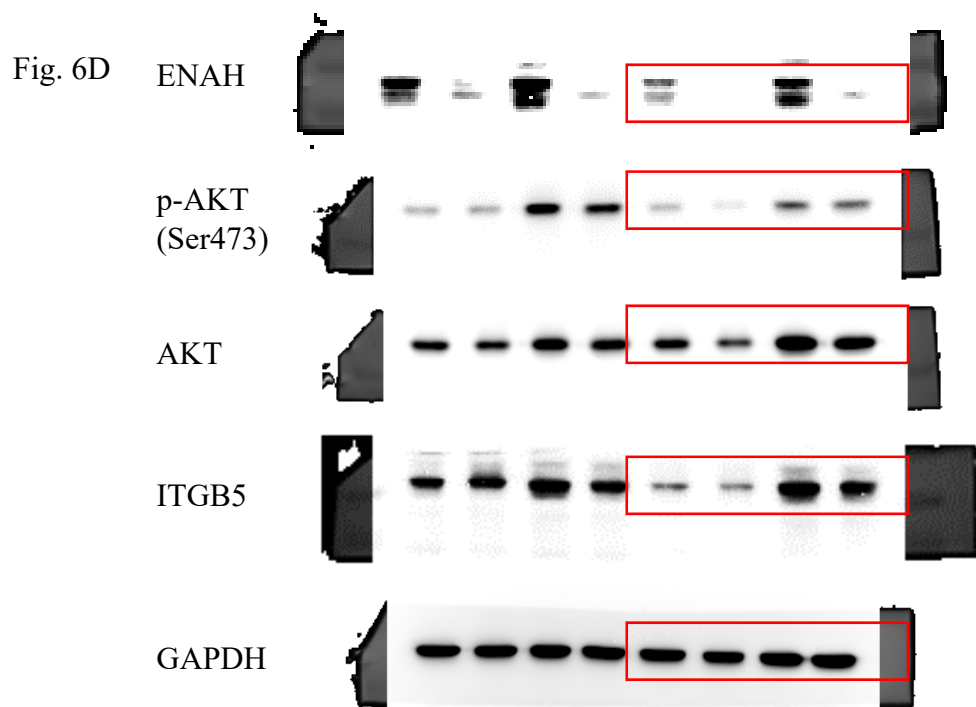


Fig. 6H ITGB5

p-Src
(Tyr419)

Src

GAPDH

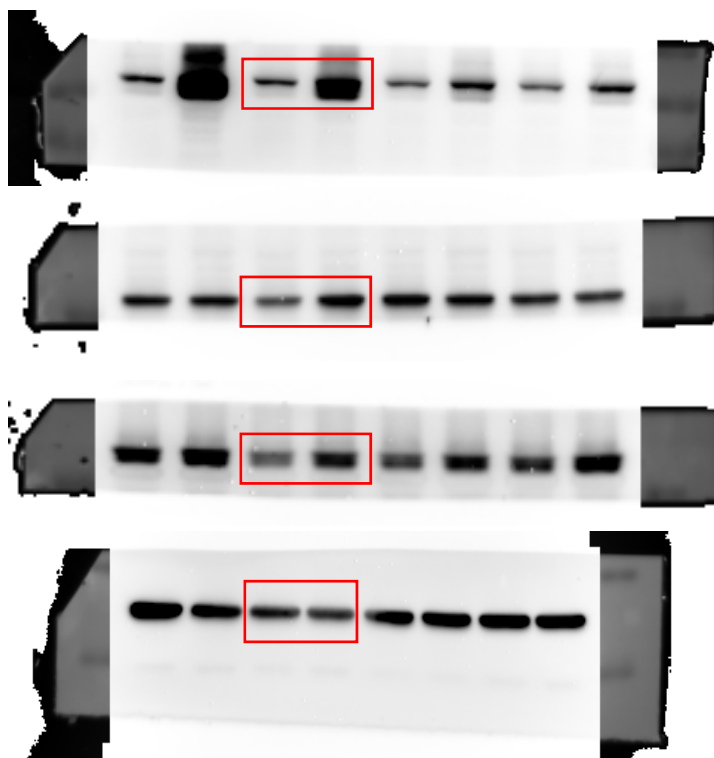


Fig. 7A ENAH

p-AKT
(Ser473)

AKT

GAPDH

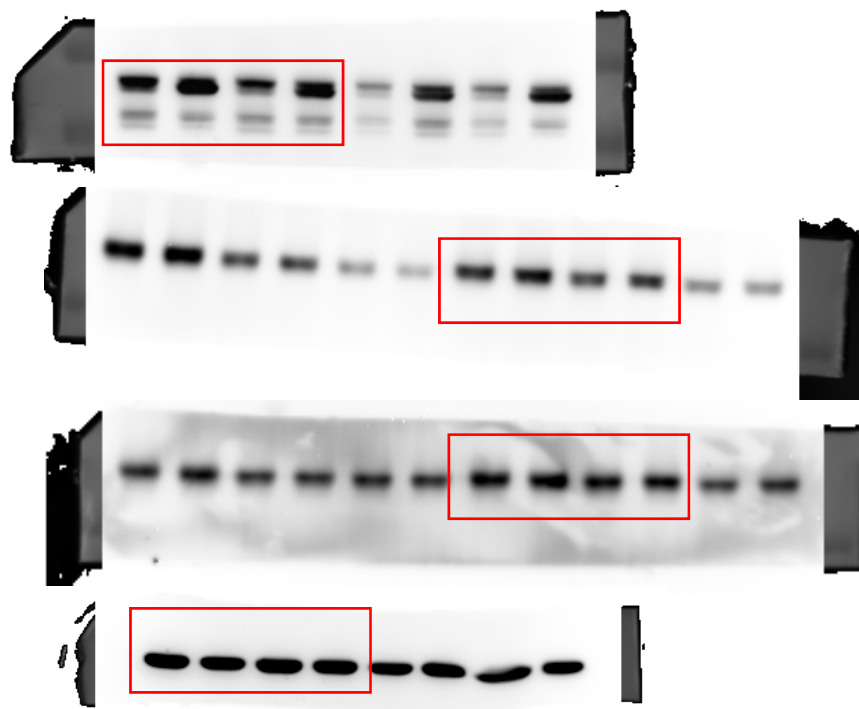


Fig. 7B ENAH

p-AKT
(Ser473)

AKT

GAPDH

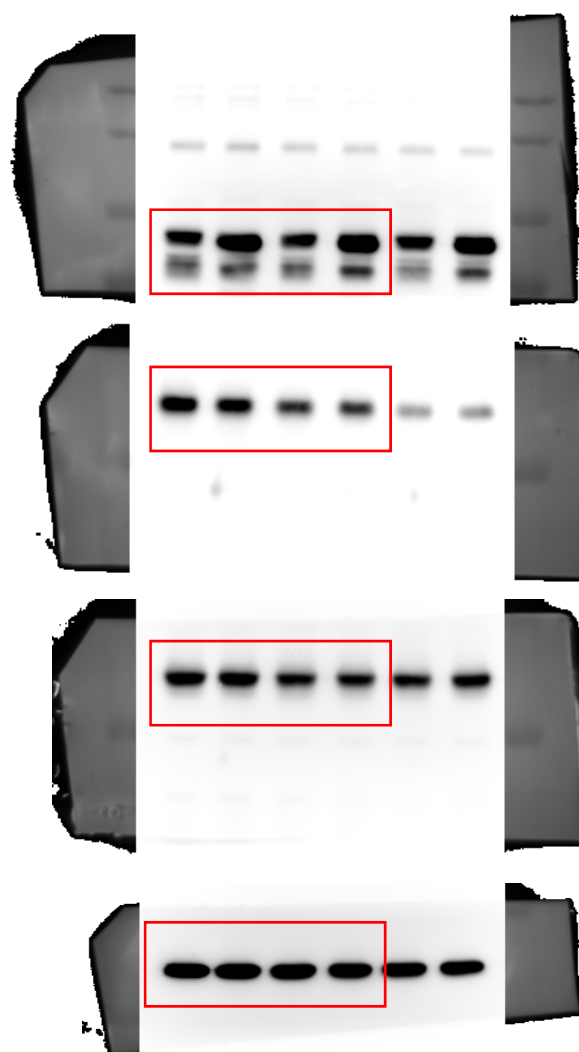


Fig. 7C

ENAH

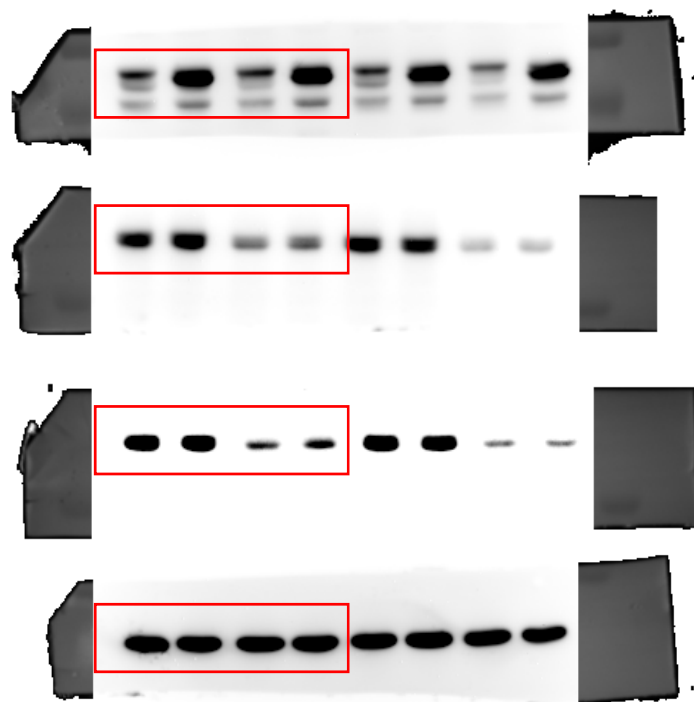


Fig. 7D

ENAH

ITGB5

p-Src
(Tyr419)

Src

GAPDH

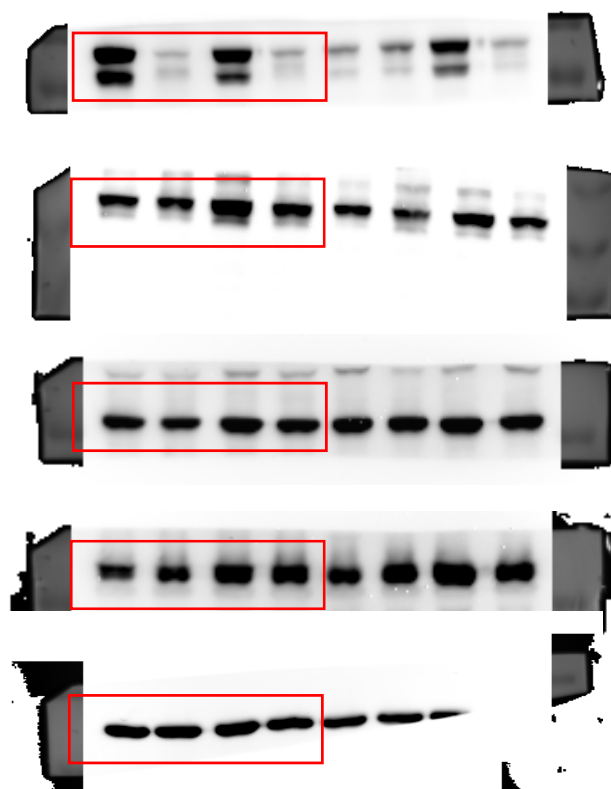
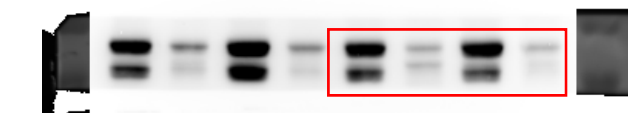
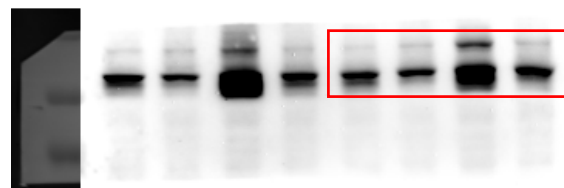


Fig. 7E

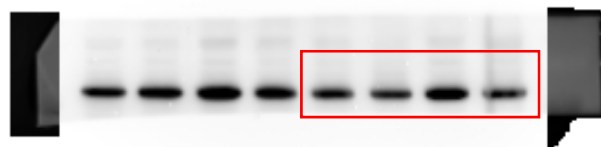
ENAH



ITGB5



p-Src
(Tyr419)



Src

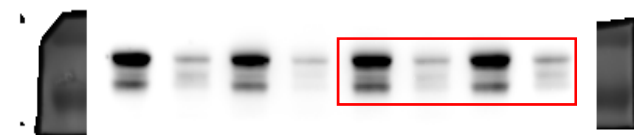


GAPDH

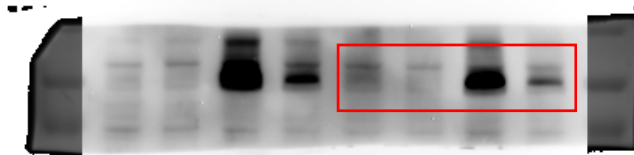


Fig. 7F

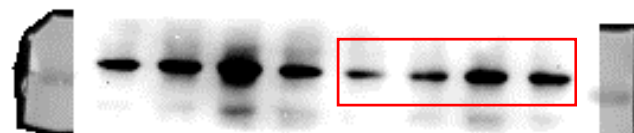
ENAH



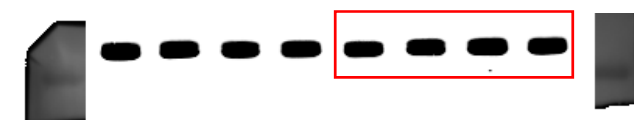
ITGB5



p-Src
(Tyr419)



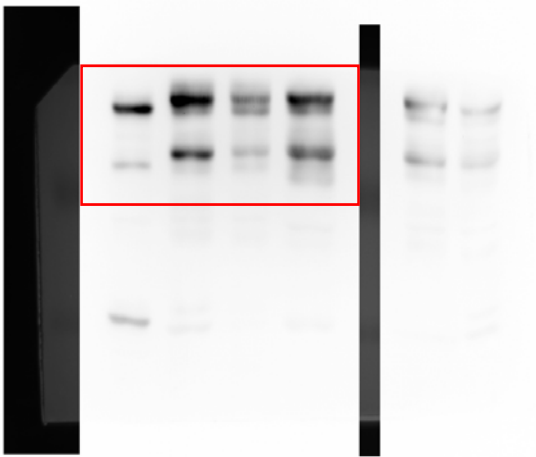
Src



GAPDH



Fig. S3B ENAH



VCP

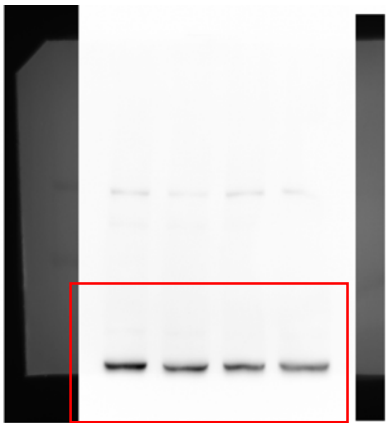


Fig. S3C ENAH
(left panel)



GAPDH

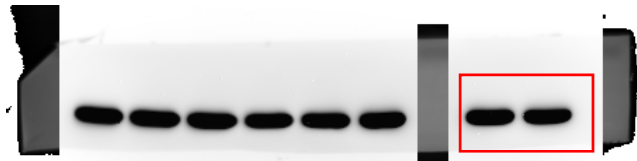
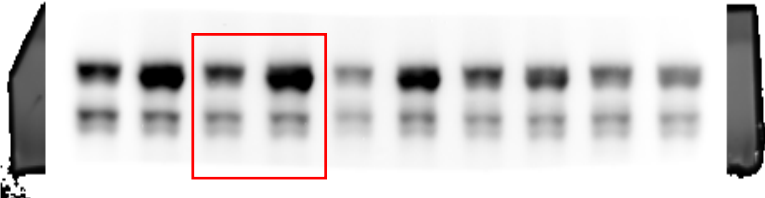


Fig. S3C ENAH
(right panel)



GAPDH

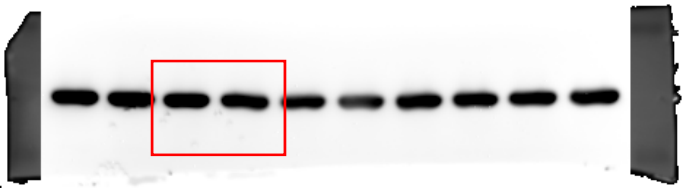
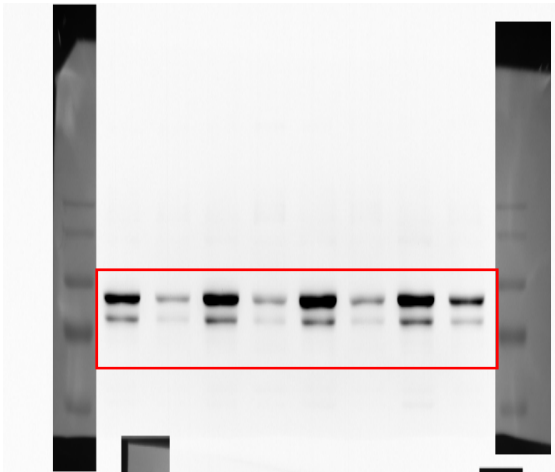


Fig. S4A ENAH



GAPDH



Fig. S4B ENAH



GAPDH

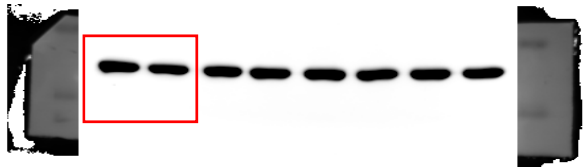


Fig. S4C ENAH



GAPDH



Fig. S4D ENAH



GAPDH

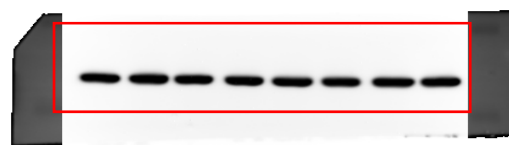
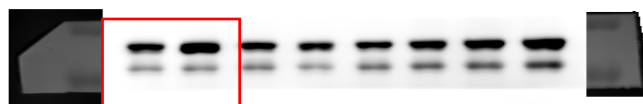


Fig. S4E ENAH



GAPDH



Fig. S4F ENAH



GAPDH



Western blot analysis showing protein levels of ENAH, p-AKT (Ser473), AKT, and GAPDH. The blot displays four rows of bands. The first three rows (ENAH, p-AKT, AKT) have red boxes highlighting the first two lanes. The fourth row (GAPDH) has a red box highlighting the first two lanes. The bands in the first two lanes are significantly more intense than in the other lanes.

Western blot analysis of ENAH, p-AKT (Ser473), AKT, and GAPDH protein levels in H1299 cells. The blot shows four rows of bands. The first three rows are labeled ENAH, p-AKT (Ser473), and AKT. The fourth row is labeled GAPDH. Each row has four lanes. Red boxes highlight the bands in the first two lanes of each row. The ENAH row shows a strong band in the second lane. The p-AKT (Ser473) row shows a strong band in the second lane. The AKT row shows a strong band in the second lane. The GAPDH row shows a strong band in the second lane.

Western blot analysis showing protein levels of ENAH, p-AKT (Ser473), AKT, and GAPDH across 12 lanes. The lanes are grouped into four sets of three, each corresponding to a different treatment condition. Red boxes highlight the bands for ENAH, p-AKT (Ser473), AKT, and GAPDH in lanes 5-8.

Fig. S7C
(left
panel)

ENAH

ITGB5

p-Src
(Tyr419)

Src

GAPDH

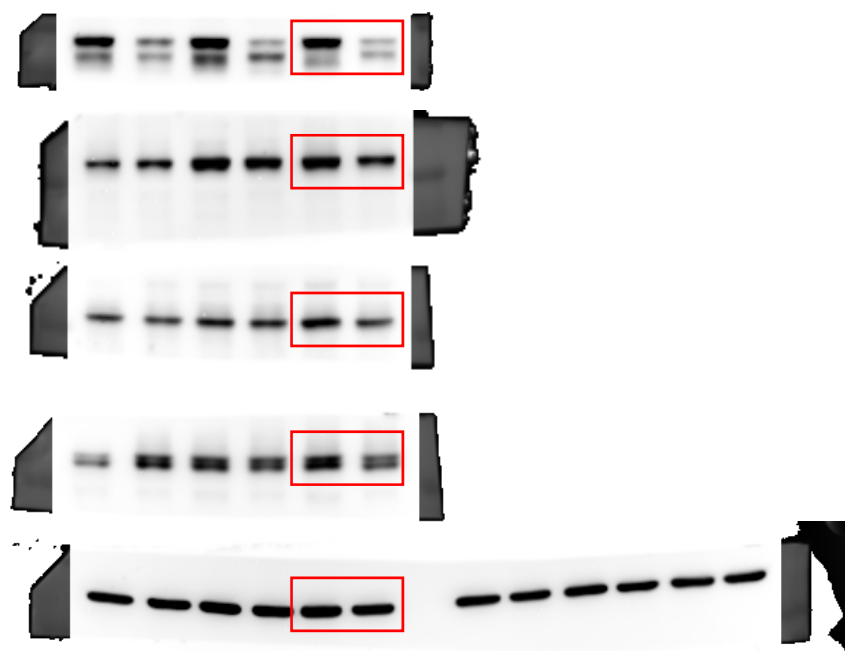


Fig. S7C
(right
panel)

ENAH

ITGB5

p-Src
(Tyr419)

Src

GAPDH

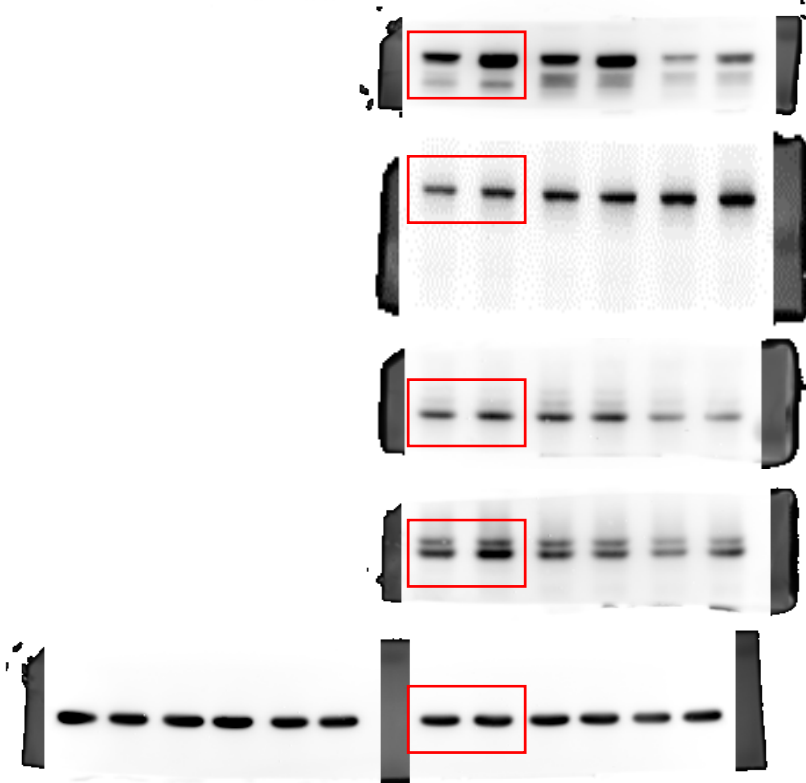
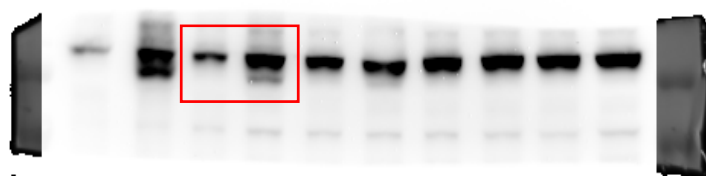


Fig. S7D ITGB5

GAPDH



Fig. S7E ITGB5



GAPDH

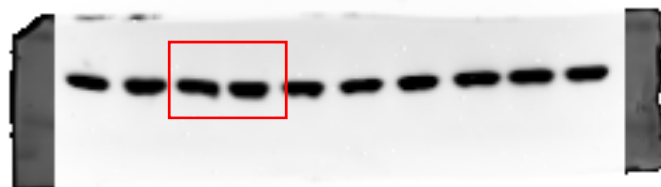
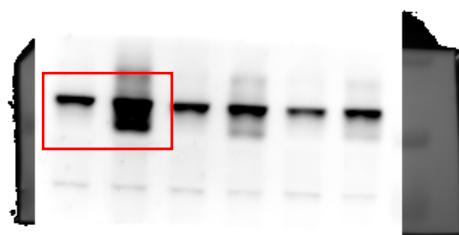


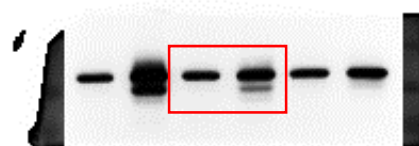
Fig. S7F ITGB5



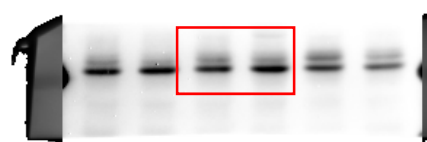
GAPDH



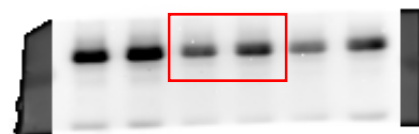
Fig. S7G ITGB5



p-Src
(Tyr419)



Src



GAPDH

