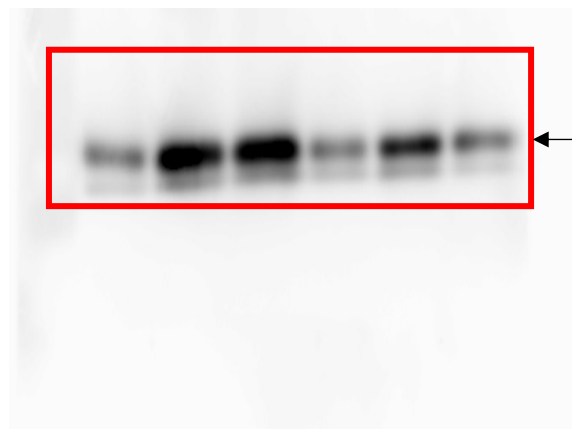
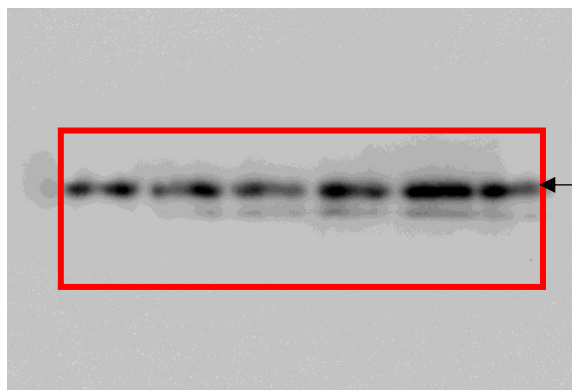
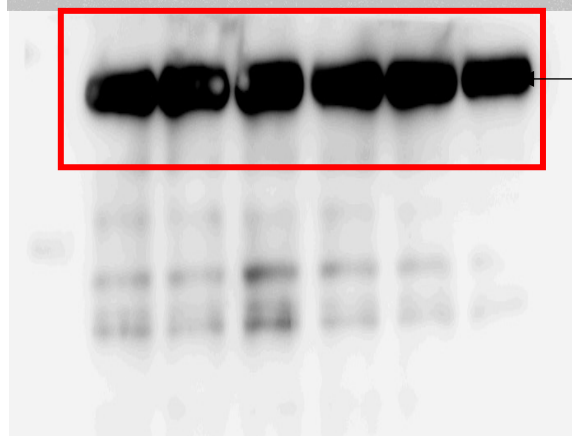


D

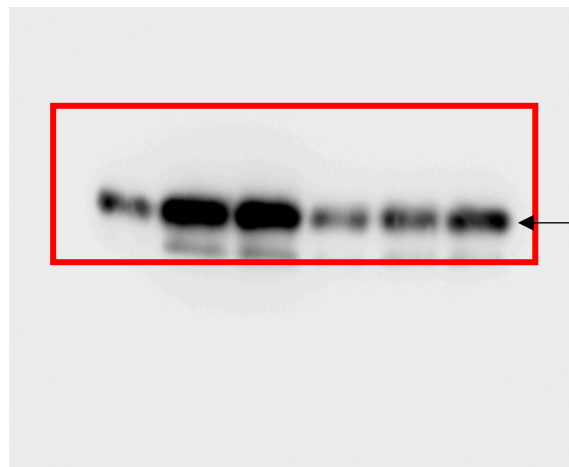
pFAK,
125 kDa



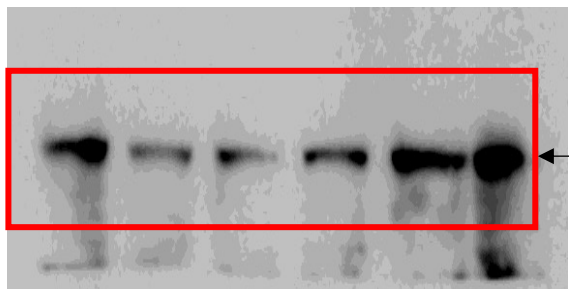
tFAK,
125 kDa



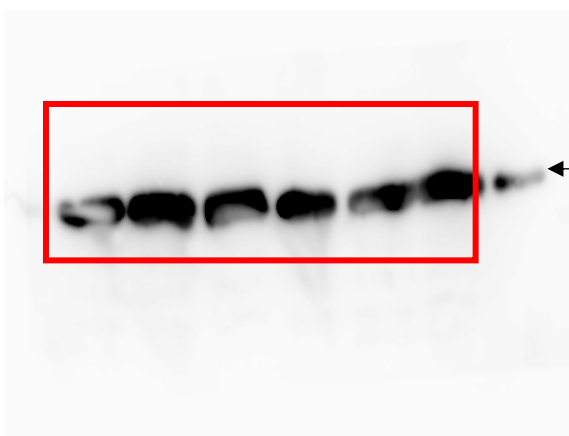
GAPDH,
36 kDa

E

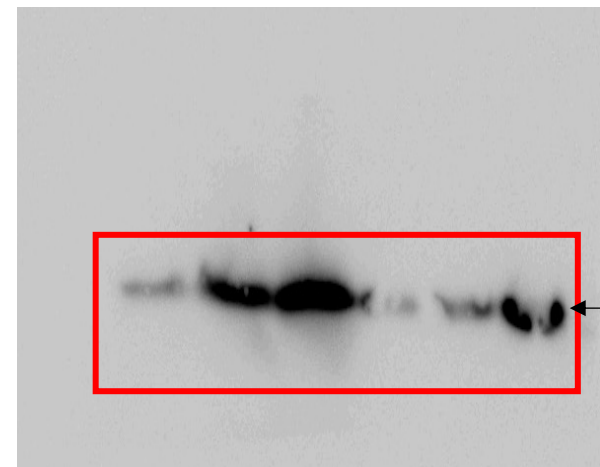
pFAK,
125 kDa



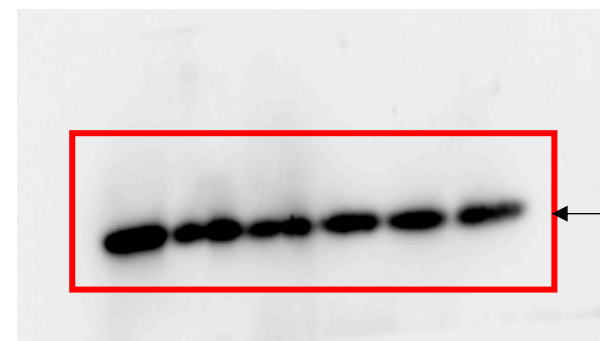
tFAK,
125 kDa



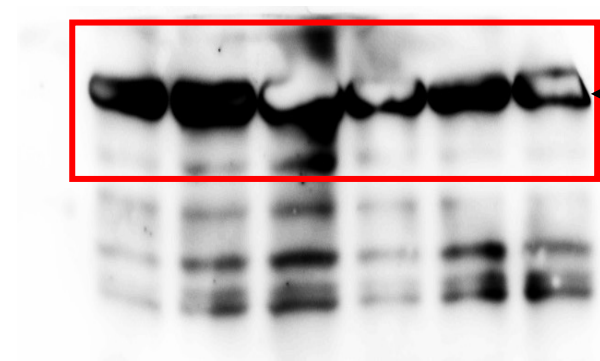
GAPDH,
36 kDa

F

pFAK,
125 kDa

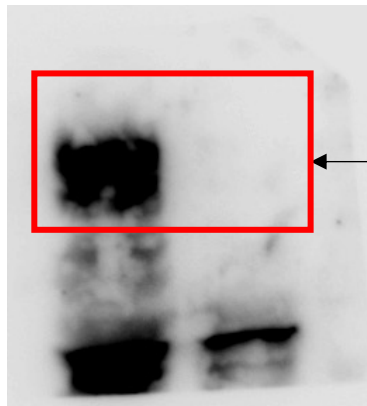


tFAK,
125 kDa



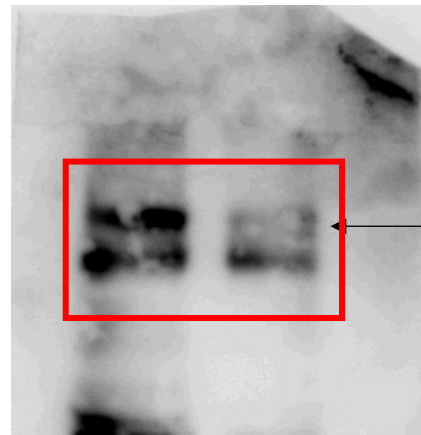
GAPDH,
36 kDa

G



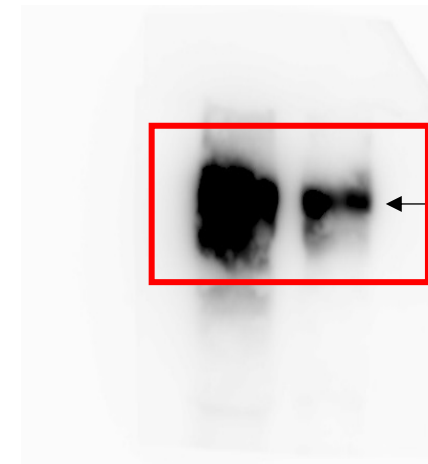
← **ITGB1, 115 kDa**

H

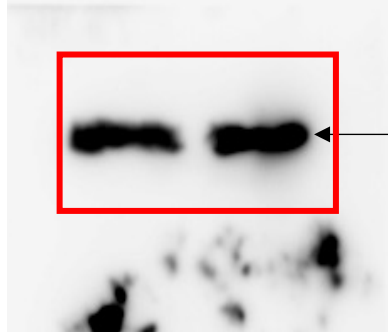


← **ITGAV, 135 kDa**

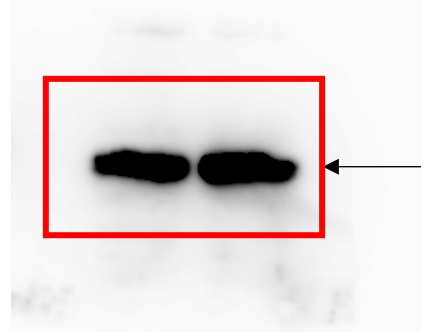
I



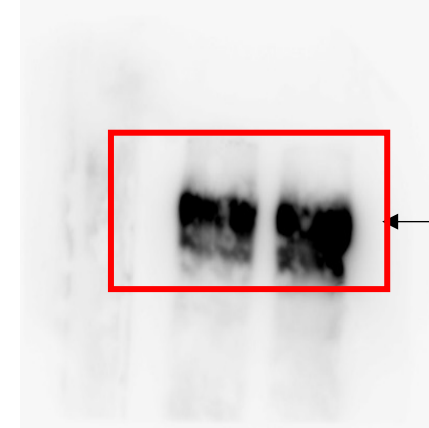
← **ITGA6, 127 kDa**



← **GAPDH,
36 kDa**



← **GAPDH,
36 kDa**



← **GAPDH,
36 kDa**