

ORIGINAL RESEARCH ARTICLE

Quantitative proteomic analysis reveals regulatory networks of extracellular matrix receptor interaction pathways in endothelial cells after myocardial infarction

Raw Western blot image

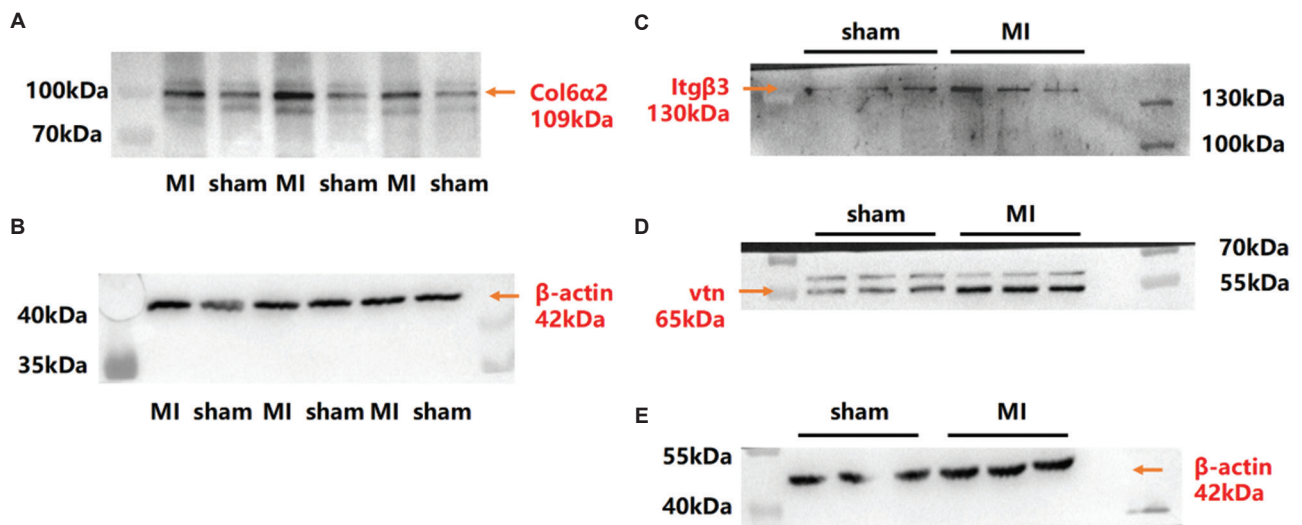


Figure 1. Raw Western blot images of Figure 8A. (A) Western blotting for Col6 α 2 (109 kDa), (B) Western blotting for β -actin (42 kDa), (C) Western blotting for Itg β 3 (130 kDa), (D) Western blotting for vtn (65 kDa), and (E) Western blotting for β -actin (42 kDa).