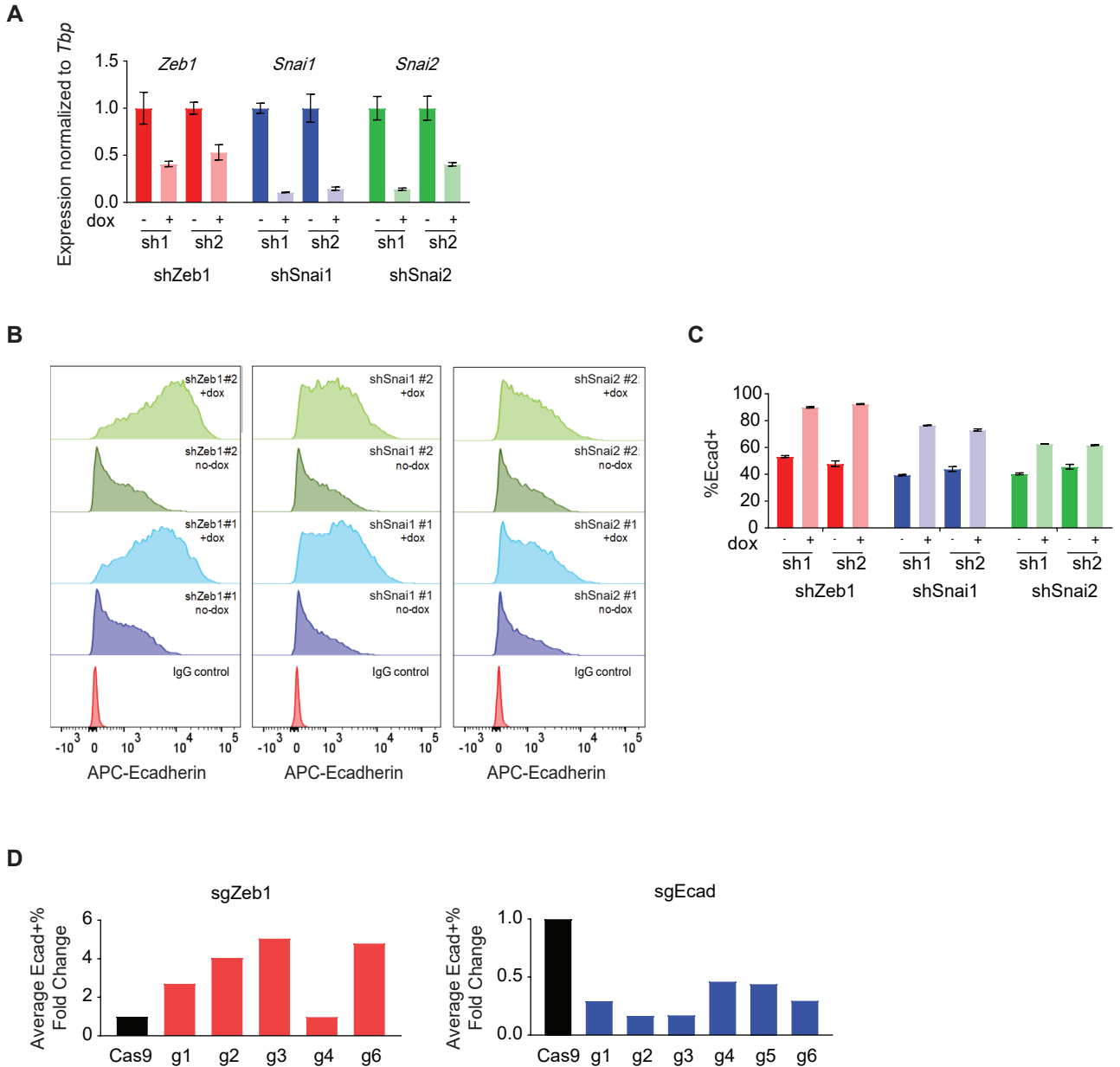


Supplementary Figure S2



Supplementary Figure S2. Identifying EMT-TFs as controls for screen.

(A) Knockdown efficiencies of dox-inducible shRNAs targeting *Zeb1*, *Snai1*, and *Snai2* in cell line 3077 c4 were determined by qPCR analysis ($n = 3$, mean \pm SEM). Two shRNAs were tested per gene.

(B-C) Representative flow histograms (B) and quantification (C) of Ecad⁺ cells in 3077 c4 expressing the indicated dox-inducible shRNAs ($n = 3$, mean \pm SEM). IgG was used to determine Ecad⁻ and Ecad⁺ gates.

(D) Quantification of Ecad⁺% fold change of sgRNAs targeting *Zeb1* and *Ecad*.