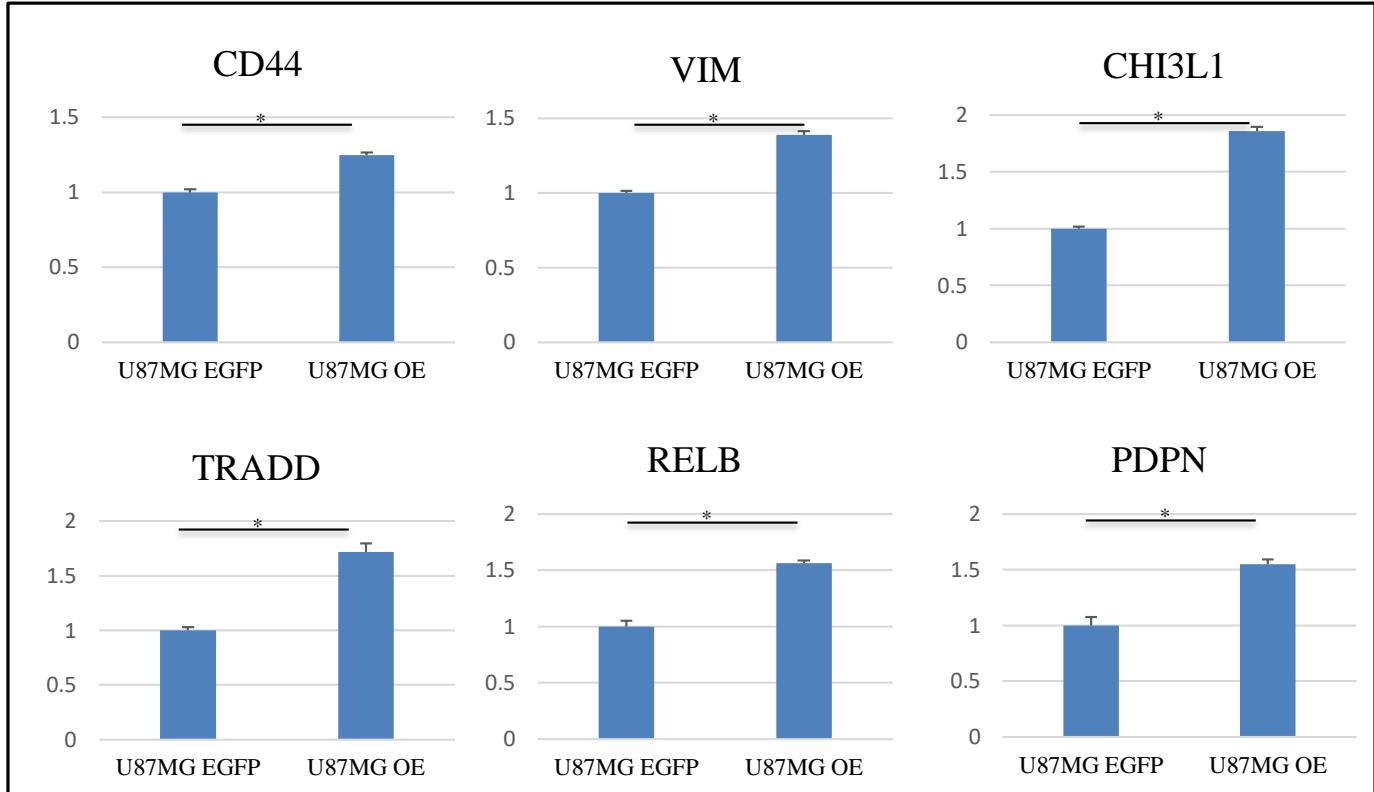


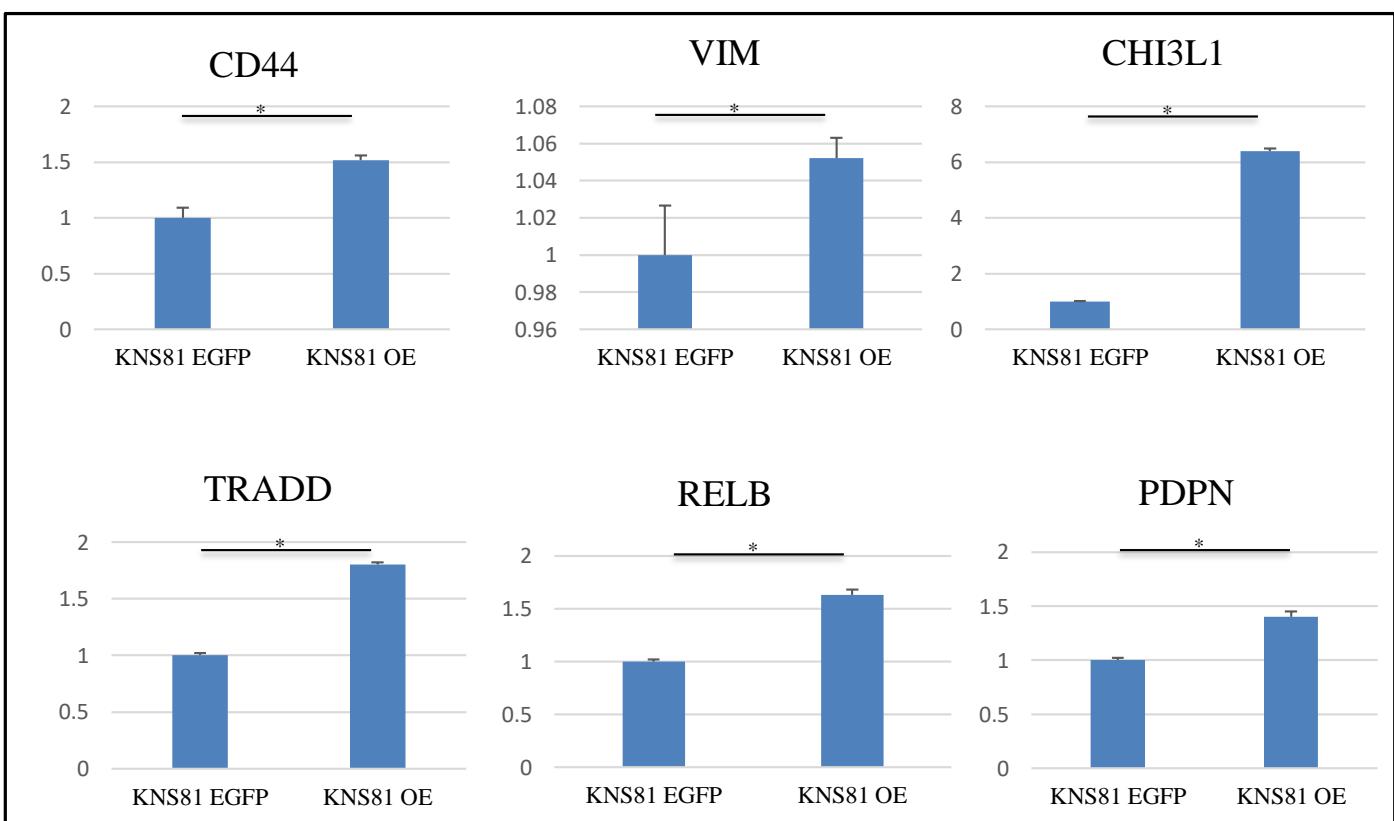
Supplementary Figure S1.

A, The distribution plot of FMNL1 immunohistochemical expression levels. **B**, Kaplan-Meier analysis of overall survival in TCGA GBM patients high or low expressing FMNL1. The former exhibited significantly poorer prognoses than the latter ($P = 0.017$). **C-E**, Representative immunohistochemical staining for FMNL1 in a primary (C) and recurrent tumour (D) are shown, and were compared by paired t-test ($P < 0.01$, E). **F-I**, Enrichment plot of FMNL1 mRNA expression in GBM specimens from TCGA for the gene sets VERHAAK_GLIOMA_CLASSICAL, VERHAAK_GLIOMA_NEURAL, PHILLIPS_GLIOMA_MESENCHYMAL, and PHILLIPS_GLIOMA_PRONEURAL. NES, normalized enrichment score; P, nominal P-value; FDR, False discovery rate.

U87MG



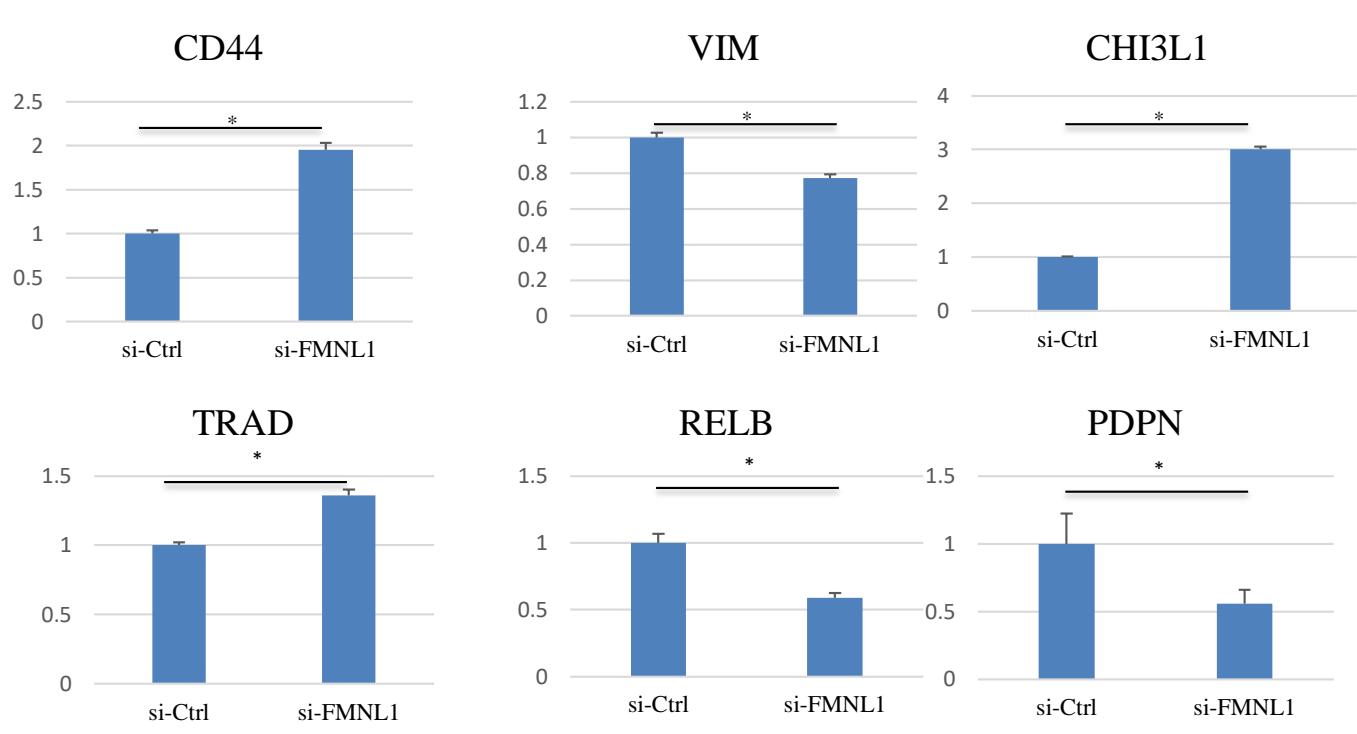
KNS81



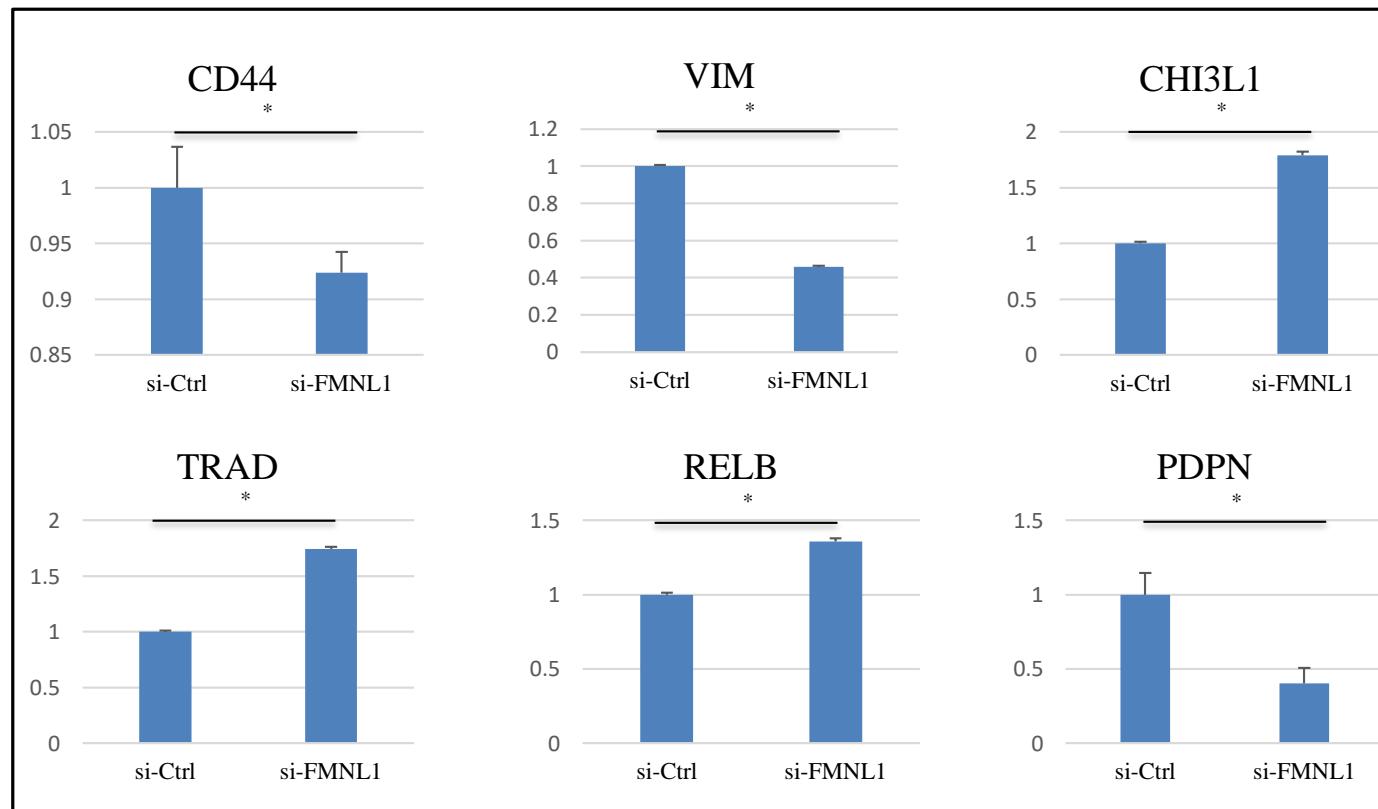
Supplementary Figure S2.

FMNL1 induces mesenchymal markers expression. Mesenchymal markers, measured by RT-PCR, were significantly upregulated in U87MG and KNS81 cells overexpressing FMNL1, compared with cells overexpressing EGFP. * $P < 0.01$

U251MG



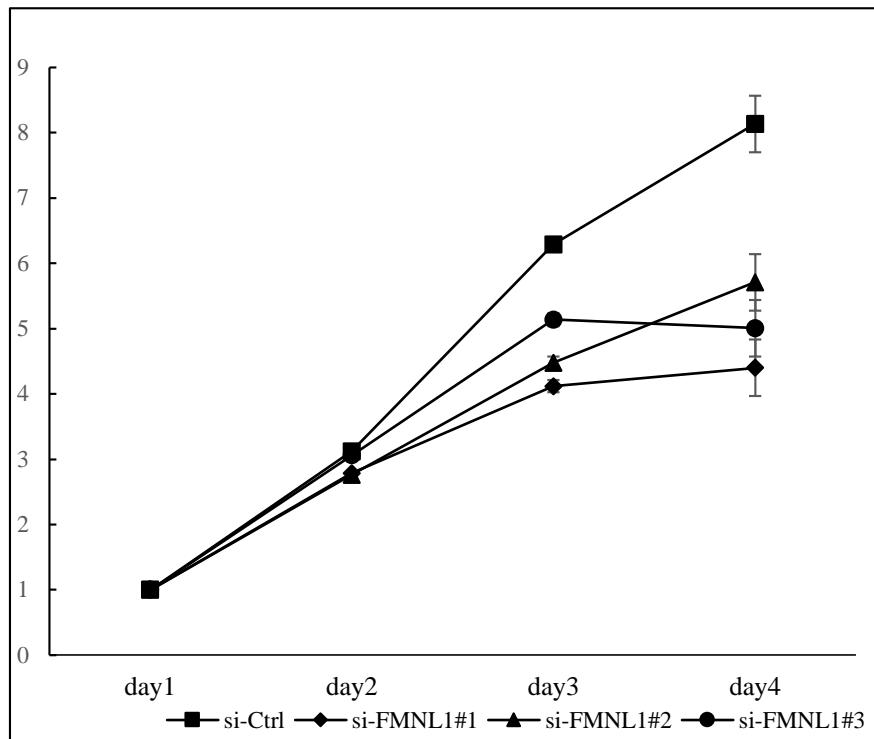
DBTR-05MG



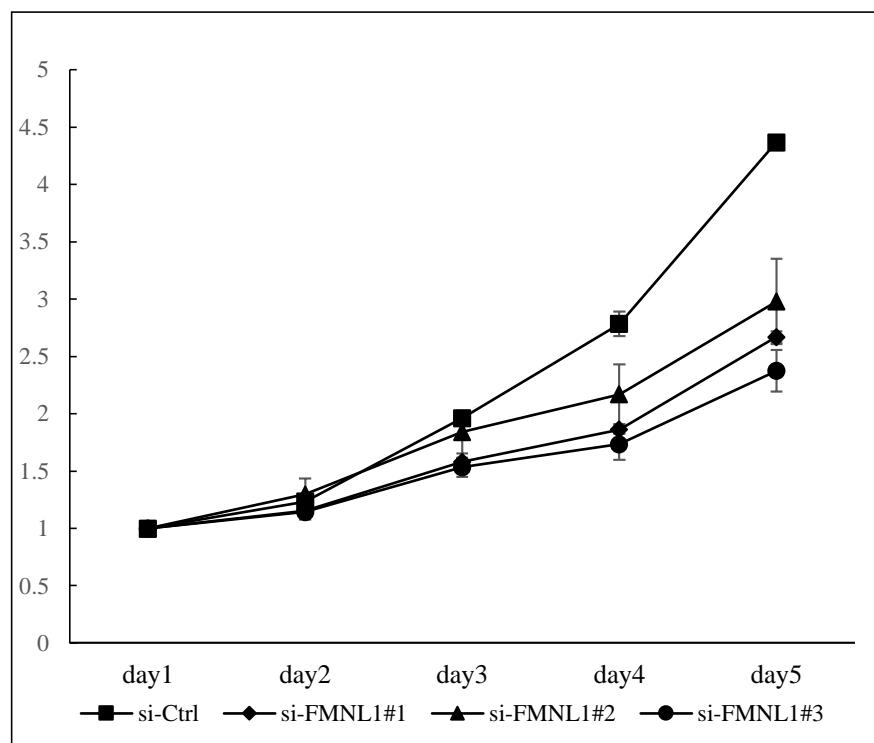
Supplementary Figure S3.

FMNL1 knockdown effect on mesenchymal markers. * $P < 0.01$

U251MG

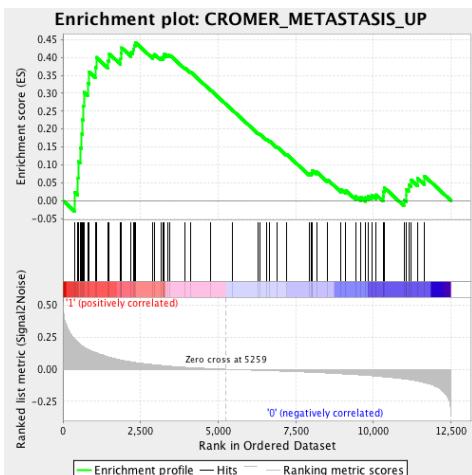
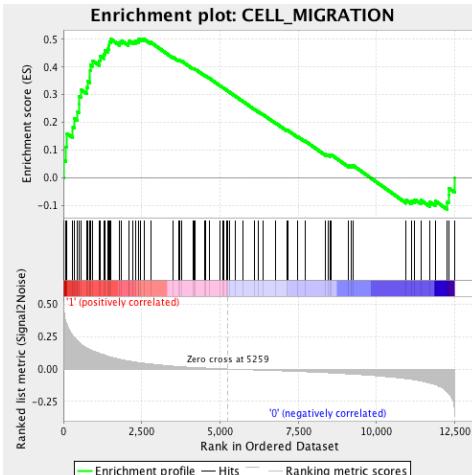
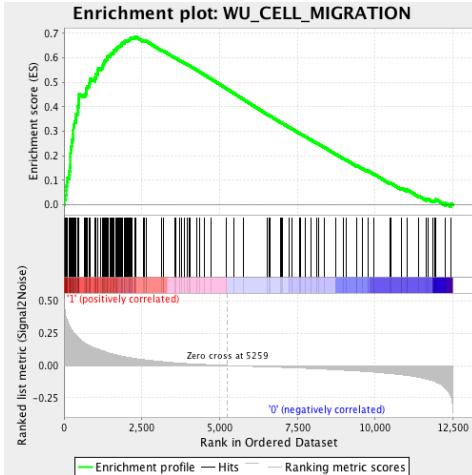


DBTRG-05MG



Supplementary Figure S4.

FMNL1 knockdown effect on GBM cell proliferation.
Cell viability was measured using the MTT assay.



NES: 1.616

P-value: 0.0

FDR: 0.054

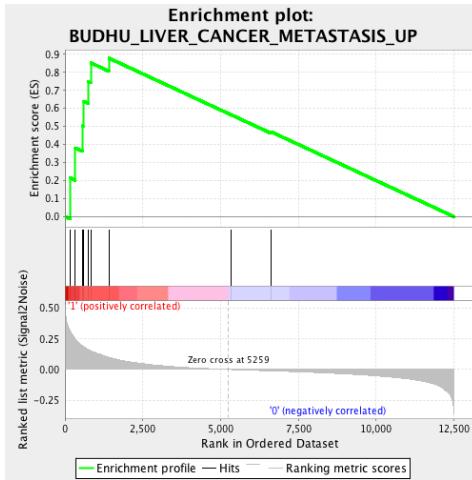
NES: 1.522

P-value: 0.014

FDR: 0.053

NES: 1.623

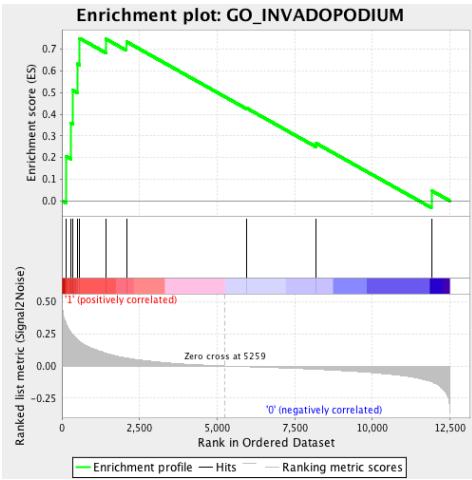
P-value: 0.014



NES: 1.553

P-value: 0.004

FDR: 0.052

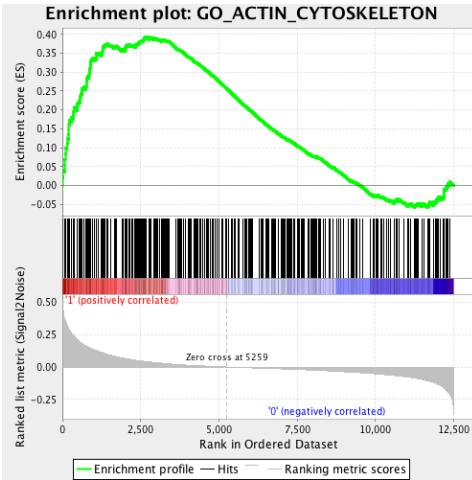


NES: 1.388

P-value: 0.068

FDR: 0.108

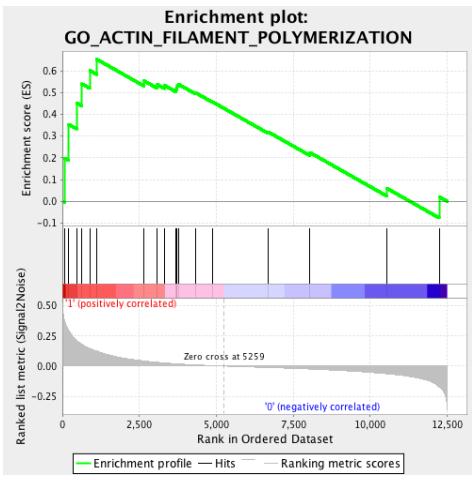
NES: 1.600
P-value: 0.011
FDR: 0.053



NES: 1.556

P-value: 0.002

FDR: 0.055



NES: 1.682

P-value: 0.015

FDR: 0.015

Supplementary Figure S5.

Enrichment plot of FMNL1 mRNA expression in TCGA GBM patients for the gene sets GO_LAMELLIPODUM, KEGG_FOCAL_ADHESION, GO_INVADOPodium, BUDHU_LIVER_CANCER_METASTASIS_UP, CROMER_METASTASIS_UP, CELL_MIGRATION, WU_CELL_MIGRATION, GO_ACTIN_CYTOSKELETON, and GO_ACTIN_FILAMENT_Polymerization. NES, normalized enrichment score; P, nominal *P*-value; FDR, false discovery rate.