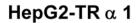
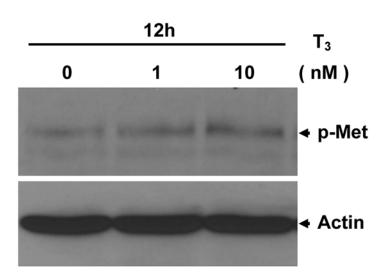
SUPPLEMENTARY FIGURES

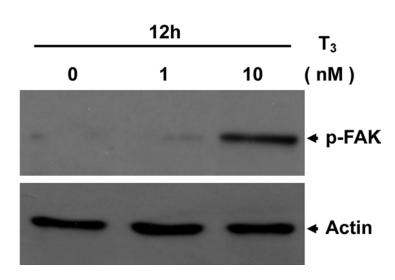
(A)





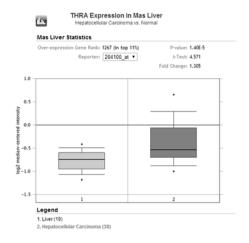
(B)

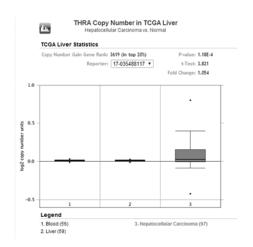
HepG2-TR α 1



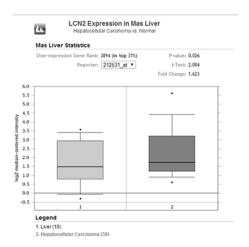
Supplementary Figure S1: T_3 regulates p-Met and p-FAK in HepG2-TR α 1 cells. Expression levels of A. p-Met and B. p-FAK in TR α 1-overexpressing cells were determined via Western blot after 0, 1 or 10 nM T_3 treatment for 12 h.

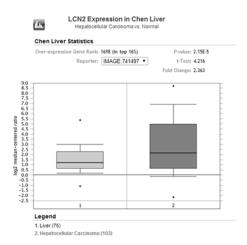






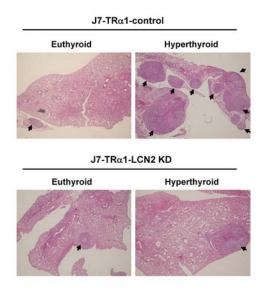


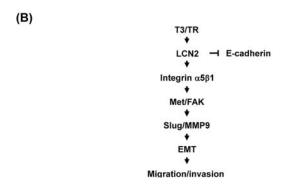




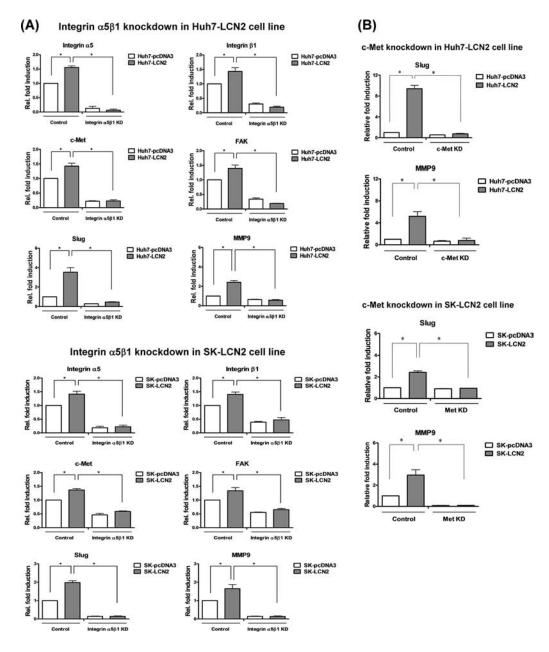
Supplementary Figure S2: Expression of LCN2 and $TR\alpha$ in public Oncomine microarray datasets. In the two Oncomine datasets (Mas and TCGA), $TR\alpha$ was highly expressed A. with a higher copy number in HCC B., compared to normal cells (liver). Similar high LCN2 expression was observed in both Mas and Chen HCC datasets.

(A)

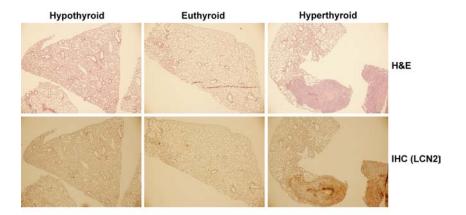


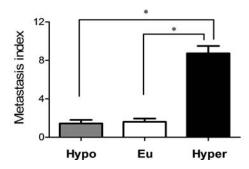


Supplementary Figure S3: *In vivo* model of T3-induced cell invasion under LCN2 depletion. A. The images depict lung tumor foci of mice injected with J7-TR α 1-control (Luc) and J7-TR α 1-LCN2 KD cells, and administered T $_3$ (2 mg/L) in drinking water for hyperthyroid condition (right panel), compared with euthyroid (left panel). The metastatic foci in lung for J7-TR α 1-control and J7-TR α 1-LCN2 KD cell are indicated by arrowhead and staining with H&E. **B.** Schematic diagram showing that T $_3$ /TR-regulated LCN2 promotes hepatoma cell migration through activation of the Integrin α 5 β 1/Met/FAK cascade and modulation of EMT markers, such as slug and MMP-9.



Supplementary Figure S4: Effect of Integrin $\alpha5\beta1$ or c-Met-depletion on LCN2 overexpression cell line. A. The mRNA expression level of Integrin $\alpha5$, Integrin $\beta1$, c-Met, FAK, slug and MMP9 was detected in integrin $\alpha5\beta1$ -depleted Huh7-pcDNA 3.0 and Huh7-LCN2 cells, compared with control cells (upper panel). Similar results were observed in Integrin $\alpha5\beta1$ -depleted SK-pcDNA 3.0 and SK-LCN2 cells (lower panel). B. The mRNA expression level of slug and MMP9 was detected in c-Met-depleted Huh7-pcDNA 3.0 and Huh7-LCN2 cells, compared with control cells (upper panel). Similar results were observed in c-Met depleted SK-pcDNA 3.0 and SK-LCN2 cells (lower panel).





Supplementary Figure S5: Effect of T3 on LCN2-mediated cell migration and invasion. The tumor foci of lung of SCID mice in the hypothyroid, euthyroid and hyperthyroid condition were analyzed with H&E (upper panel) and IHC (lower panel) staining to examine LCN2 expression and tumor foci. The numbers of tumor foci from at least five areas were calculated as a Metastasis index. Differences were analyzed using One-way ANOVA, *P < 0.05.