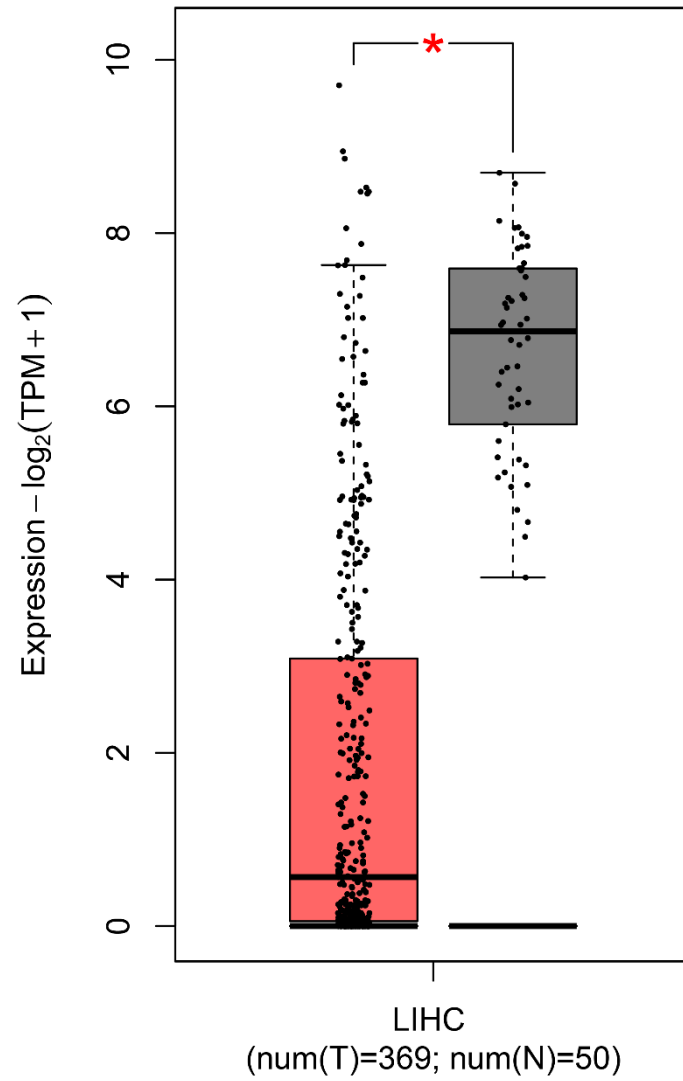
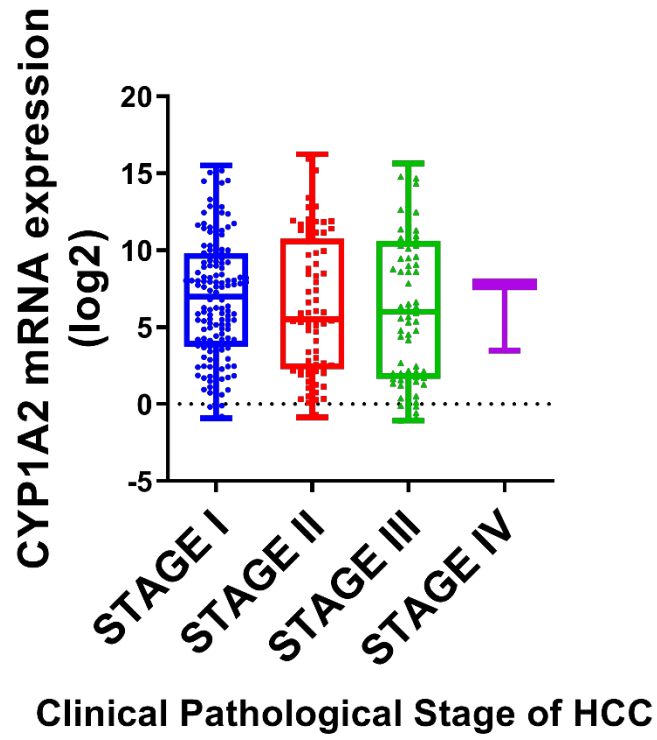


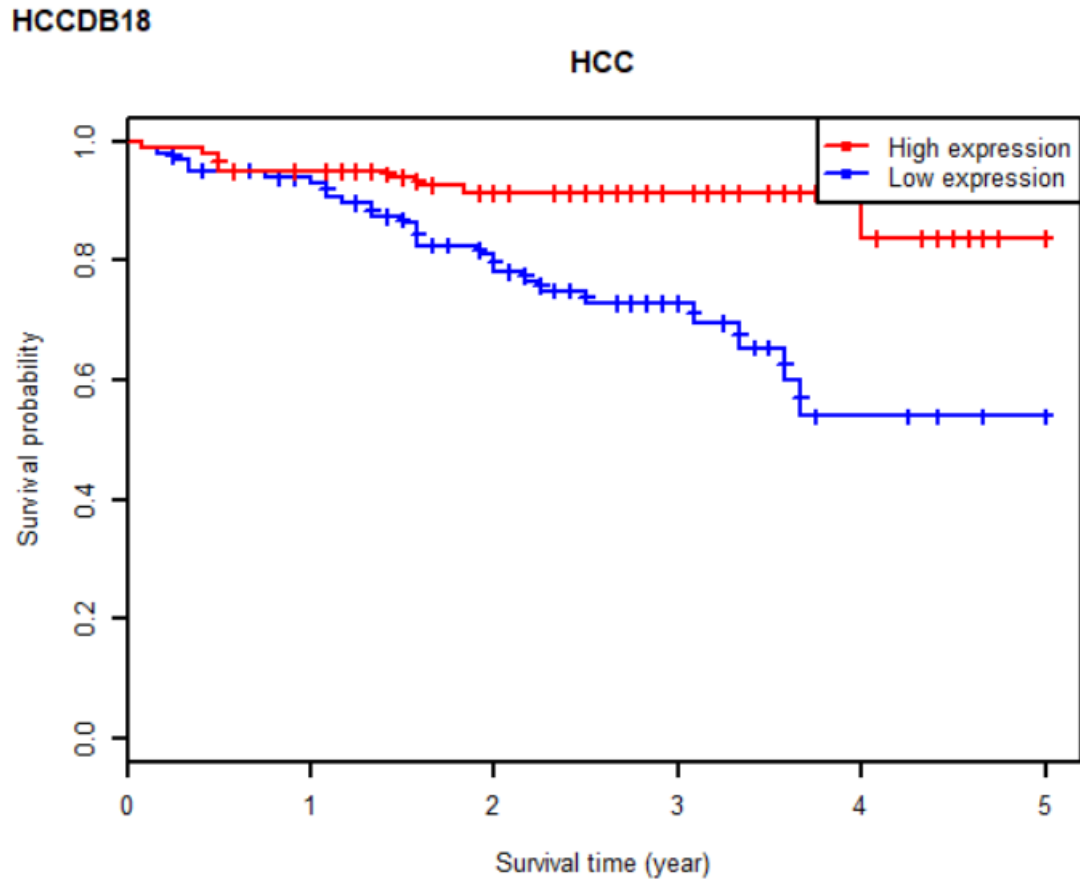
**Figure S1.** The mRNA of CYP1A2 is significantly decreased in TCGA-LIHC dataset from GEPIA ( $*P < 0.05$ ) (N, non-tumor tissues; T, tumor tissues).



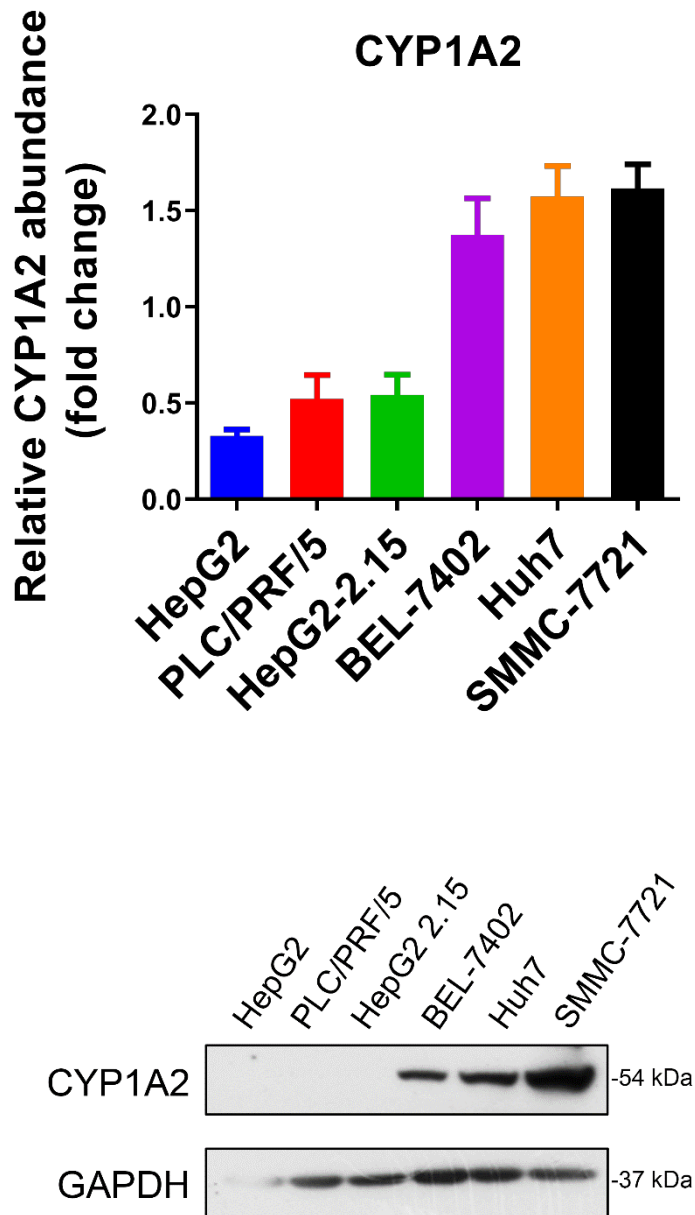
**Figure S2.** No significant difference was observed between the expression of CYP1A2 mRNA and the clinical pathological stage of HCC in TCGA-LIHC dataset. Stage I: n = 163; Stage II: n = 77; Stage III: n = 76; Stage IV: n = 3.



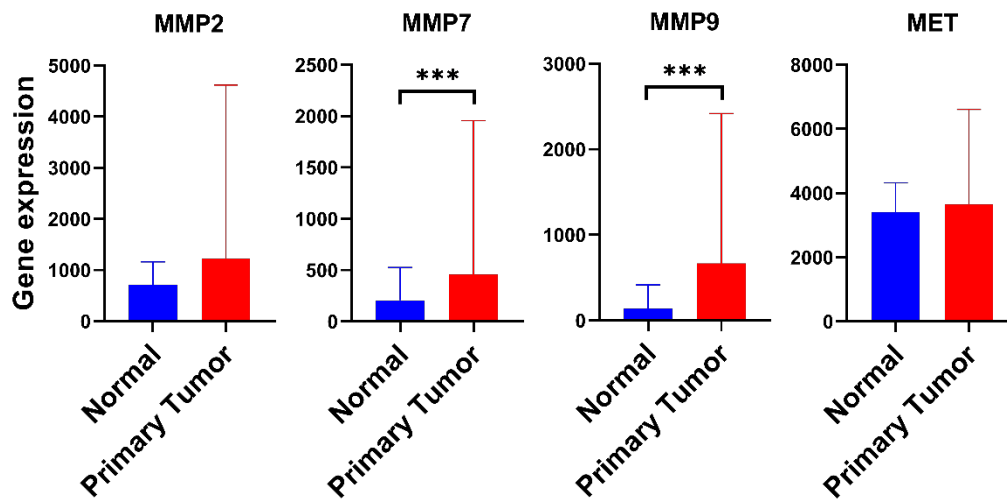
**Figure S3.** The HCC patients with higher expression of CYP1A2 displayed better survival in HCCDB online database. The data was acquired from <http://lifeome.net/database/hccdb/home.html>



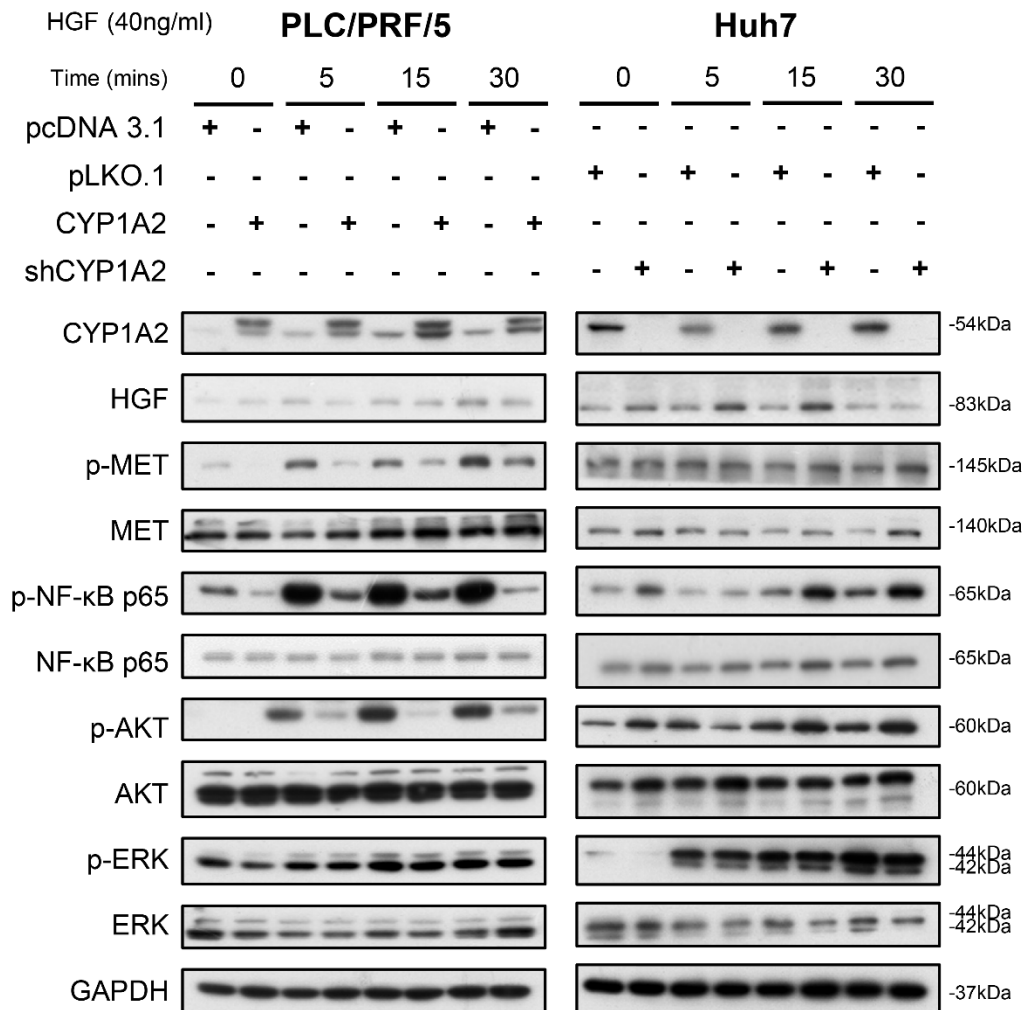
**Figure S4.** The endogenous expression of CYP1A2 in different HCC cell lines.



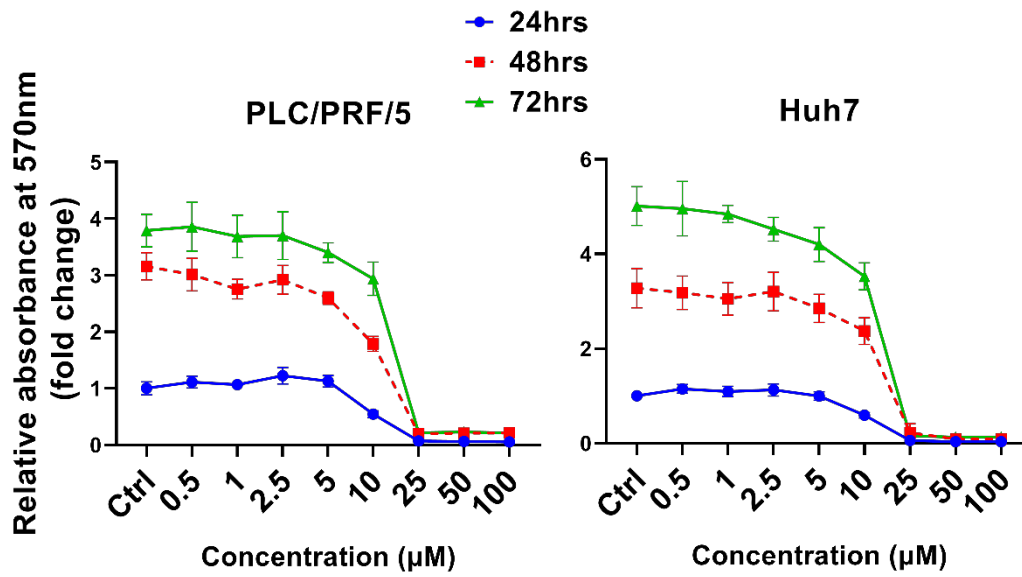
**Figure S5.** The expression of MET and MMPs in the TCGA-LIHC dataset ( $***P < 0.001$ ).



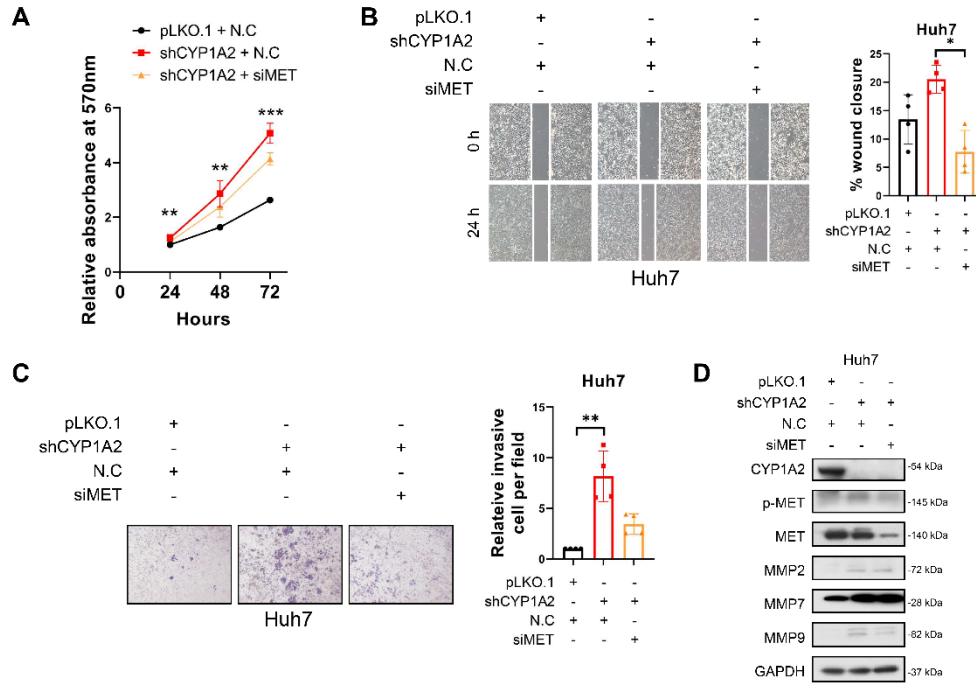
**Figure S6. The effects of CYP1A2 on HGF/MET signaling after administration of HGF (40ng/ml).**



**Figure S7.** MTT assays showed the effect of PHA-665752 on PLC/PRF/5 and Huh7 cells at different time points. Ctrl, DMSO control.

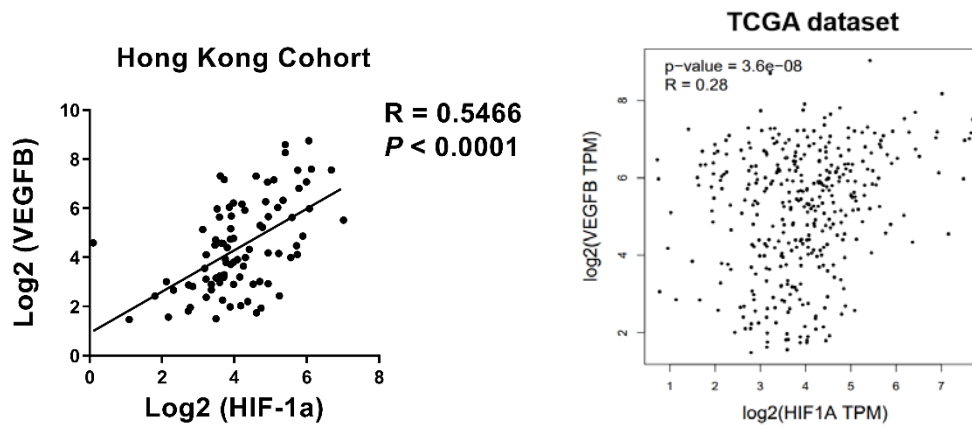


**Figure S8.** MET silencing inhibited the MMP2 and MMP9 expression and reversed the increased proliferation, migration, and invasion abilities in the CYP1A2-knockdown Huh7 cells. In experiment A, \*\*\* represents “pLKO.1 + N.C” verses “shCYP1A2 + N.C”.

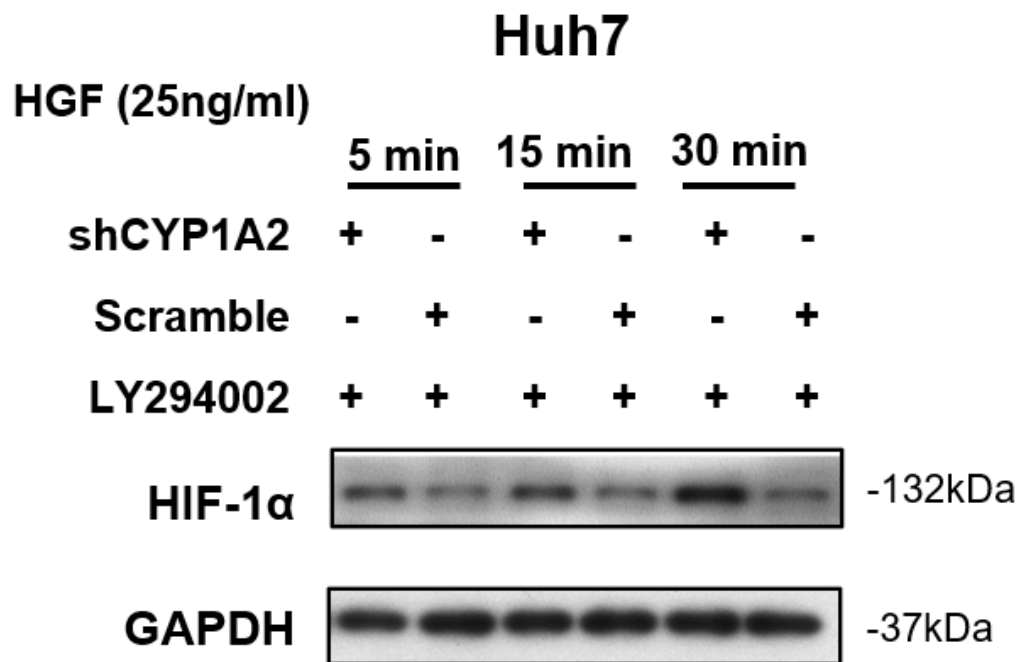




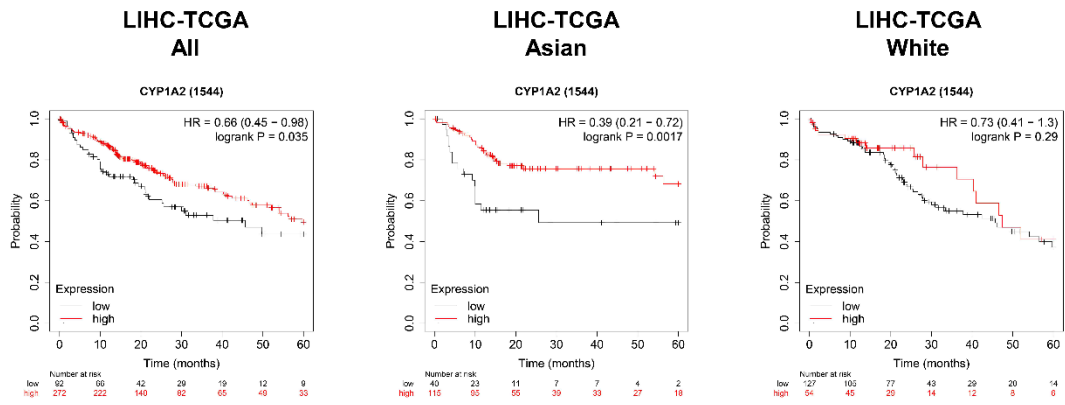
**Figure S9.** The positive correlation between HIF-1 $\alpha$ -targeted gene VEGFB in Hong Kong HCC cohort and in TCGA-LIHC dataset.



**Figure S10.** Western blot assay showed that LY294002 had no impact on HIF-1 $\alpha$  expression.



**Figure S11.** The five-year OS from the LIHC-TCGA dataset in accordance with CYP1A2 expression. Data were collected from <https://kmplot.com/analysis/> and the patients was automatically split by the website using the Auto Select Best Cutoff.



**Table S1.** Sequences of primers used

<b>Primer names</b>	<b>Sequences (5' -&gt; 3')</b>
<b>qPCR</b>	
CYP1A2-F	GGACACAACGCTGAATG
CYP1A2-R	CATCATCTTCTCACTCAAGG
MMP2-F	GTCTGAAGAGCGTGAAG
MMP2-R	AGGTAGGAGTGAGAATGC
MMP7-F	GCCTACCTATAACTGGAATG
MMP7-R	AAGCCTTTGACACTAATCG
MMP9-F	CGTGACCTATGACATCC
MMP9-R	CCTCCAGAACAGAATACC
HIF-1 $\alpha$ -F	TTCCAGTTACGTTCTTCGATCA
HIF-1 $\alpha$ -R	TTTGAGGACTTGCGCTTTCA
MET-F	GCACCCTAAAGCCGAAATG
MET-R	GATGACAACAGAGAAGGATACG
HGF-F	TGATACCACACGAACAC
HGF-R	AACTTCTGAACACTGAGG
GAPDH-F	CACTGGCGTCTTCACC
GAPDH-R	GAGGCTGTTGTCATACTTC
<b>Oligo design for shRNA cloning</b>	
shCYP1A2-F	CCGGCAAGGGACACAACGCTGAAT GCTCGAGCATTTCAGCGTTGTGTCCC TTGTTTTTG
shCYP1A2-R	AATTCAAAAACAAGGGACACAACGC TGAATGCTCGAGCATTTCAGCGTTGT GTCCCTTG

Abbreviations: qPCR, real-time polymerase chain reaction

**Table S2.** Antibodies used in Western blot assay and Immunohistochemistry

	<b>Company</b>	<b>Catalog No.</b>
CYP1A2	Santa Cruz Biotechnology Dallas, TX	sc-53241
phospho-NF- $\kappa$ B p65 ser536	Cell signaling Danvers, MA	#3033
NF- $\kappa$ B p65	Cell signaling	#8242
N-cadherin	Cell signaling	#13116
HGF	Abcam	ab83760
GAPDH	Santa Cruz Biotechnology	sc-32233
phospho-Akt (Ser473)	Cell signaling	#4060
Akt	ImmunoWay Biotechnology Plano, TX	YT0178
phospho-Met (Tyr1234/1235)	Cell signaling	#3077
Met	Cell signaling	#8198
Phospho-p44/42 MAPK (Erk1/2)	Cell signaling	#4370
p44/42 MAPK (Erk1/2)	Cell signaling	#4695
Phospho-p38 MAPK (Thr180/Tyr182)	Cell signaling	#4511
p38 MAPK	Cell signaling	#9212
MMP2	Research And Diagnostic Systems, Minneapolis, MN	AF902
MMP7	Research And Diagnostic Systems, Minneapolis, MN	MAB9071
MMP9	Abcam	ab137867
E-cadherin	Cell signaling	#14472
N-cadherin	Cell signaling	#13116