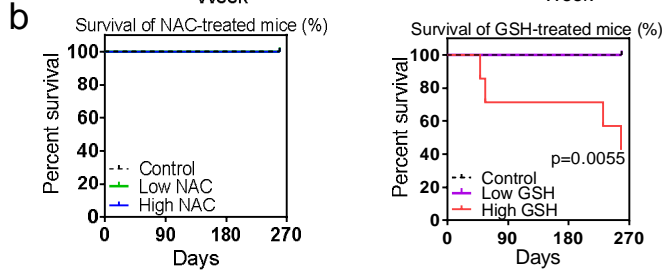
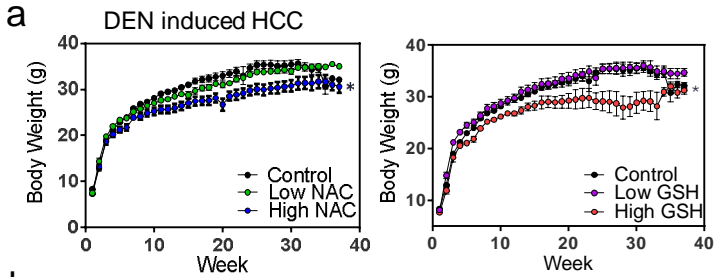
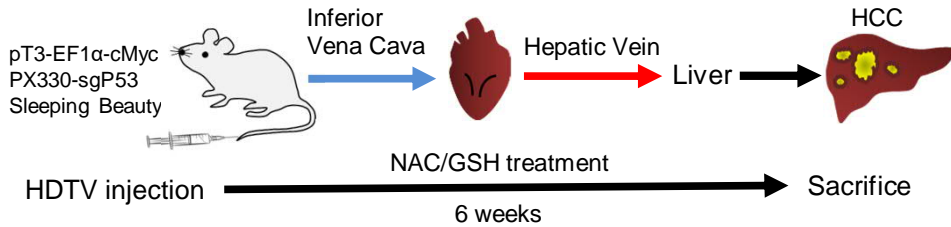


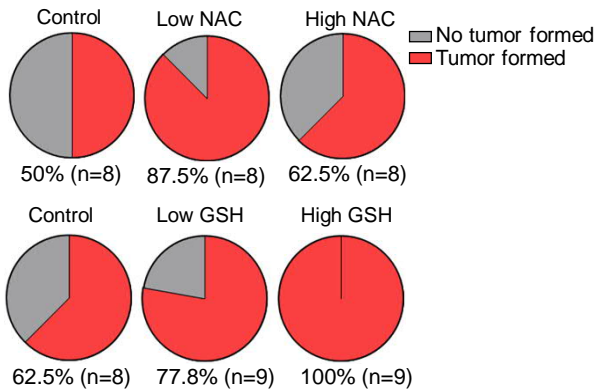
Supplementary Figure 1



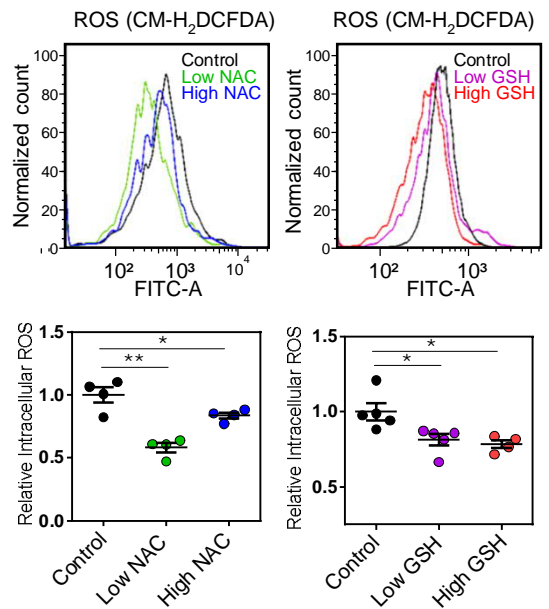
c The hydrodynamic tail vein injection-induced HCC



d Hydrodynamic tail vein injection (p53 KO/c-Myc)
NAC/GSH treatment
6 weeks
Tumor incidence rate

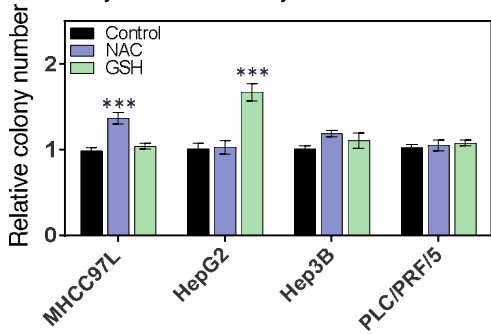


e

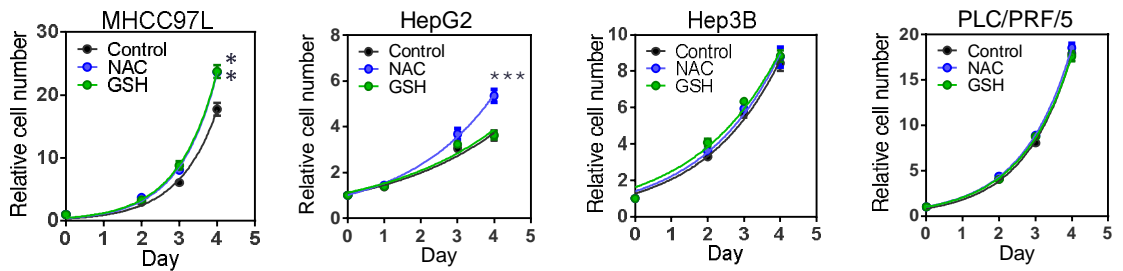


Supplementary Figure 2

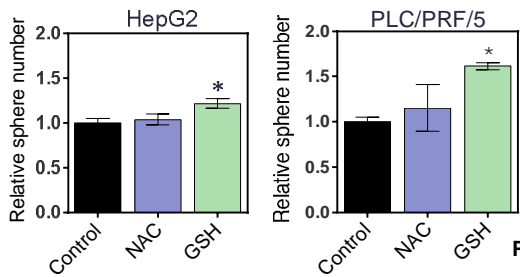
a Colony formation assay



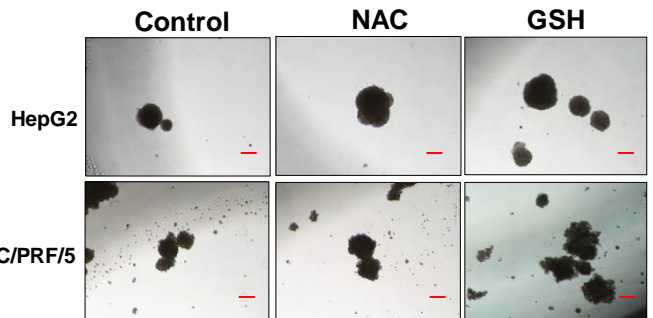
b Cell proliferation assay



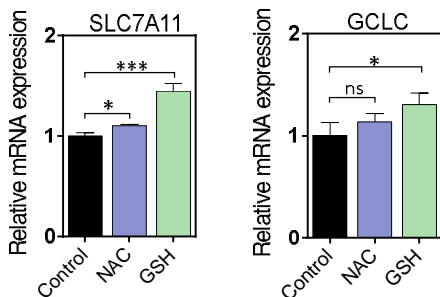
c Sphere formation assay



d Sphere formation assay

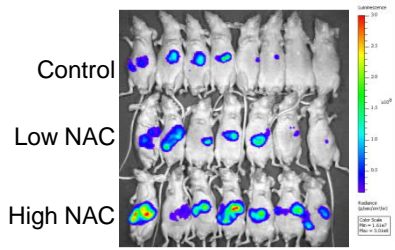


e qPCR MHCC97L cells

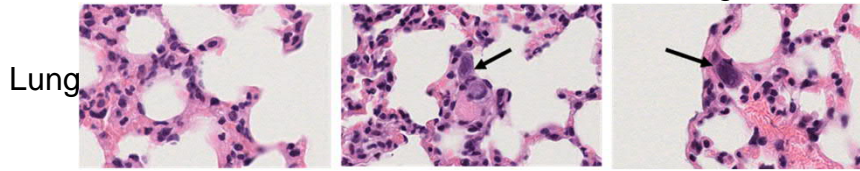


Supplementary Figure 3

a Orthotopic liver injection NAC treatment



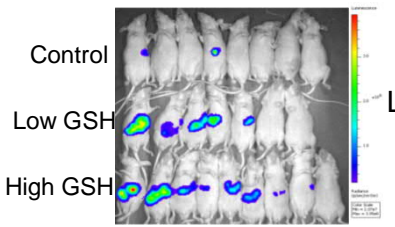
b Control Low NAC High NAC



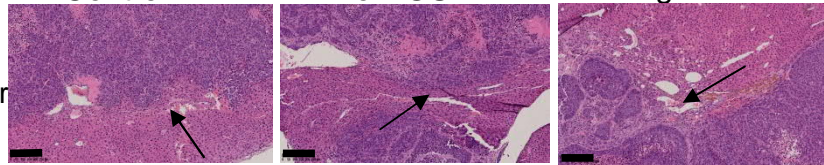
The metastatic foci

Control	Low NAC	High NAC
0.23 (0-1)	0.37 (0-1.8)	0.22 (0-1.1)

c Orthotopic liver injection GSH treatment



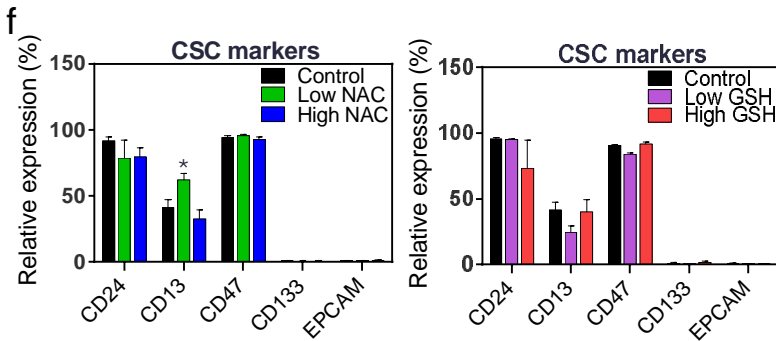
d Control Low GSH High GSH



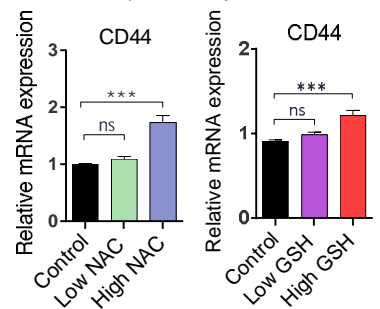
e

Tumor incidence rate		Control	Low NAC	High NAC
MHCC97L cells	Number of cells injected			
	2,000	4/8	6/8	4/8
Frequency	20,000	4/8	6/8	7/8
	200,000	7/8	8/8	8/8

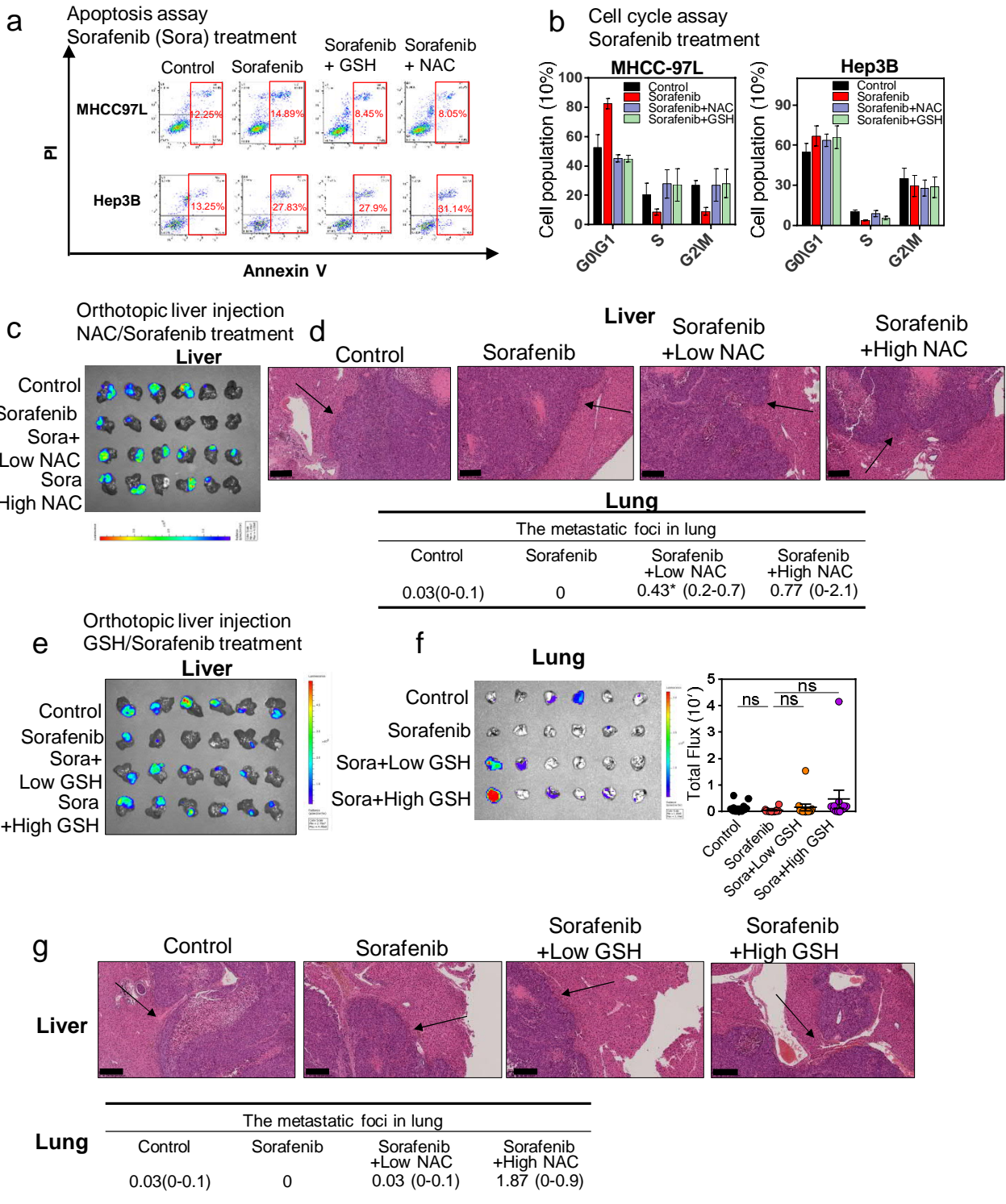
Tumor incidence rate		Control	Low GSH	High GSH
MHCC97L cells	Number of cells injected			
	2,000	4/8	5/8	7/8
Frequency	20,000	6/8	7/8	8/8
	200,000	7/8	7/8	8/8



g CD44 mRNA expression in mice orthotopic liver injection tumors



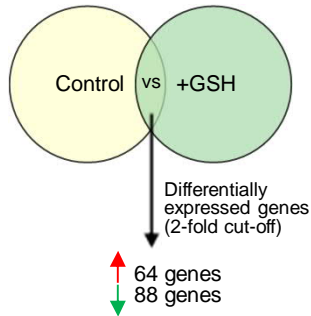
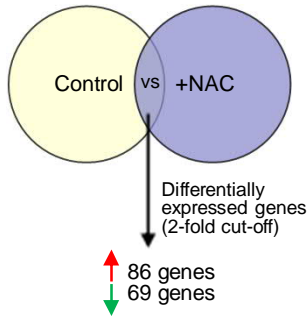
Supplementary Figure 4



Supplementary Figure 5

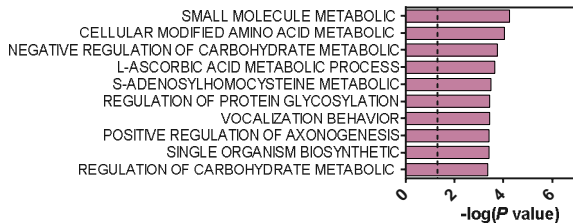
a

RNA-seq
MHCC97L cells

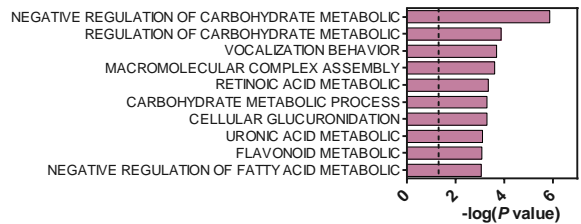


b RNA-seq
MHCC97L cells

GO biological process: upregulated genes by NAC

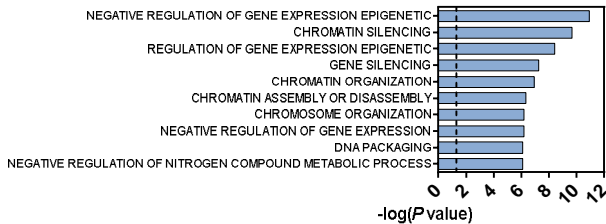


GO biological process: upregulated genes by GSH

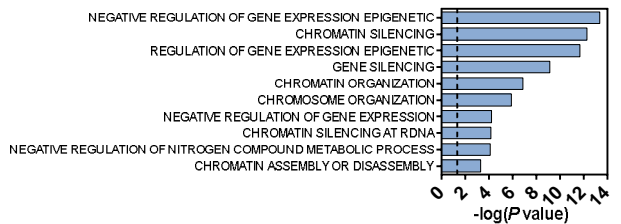


c

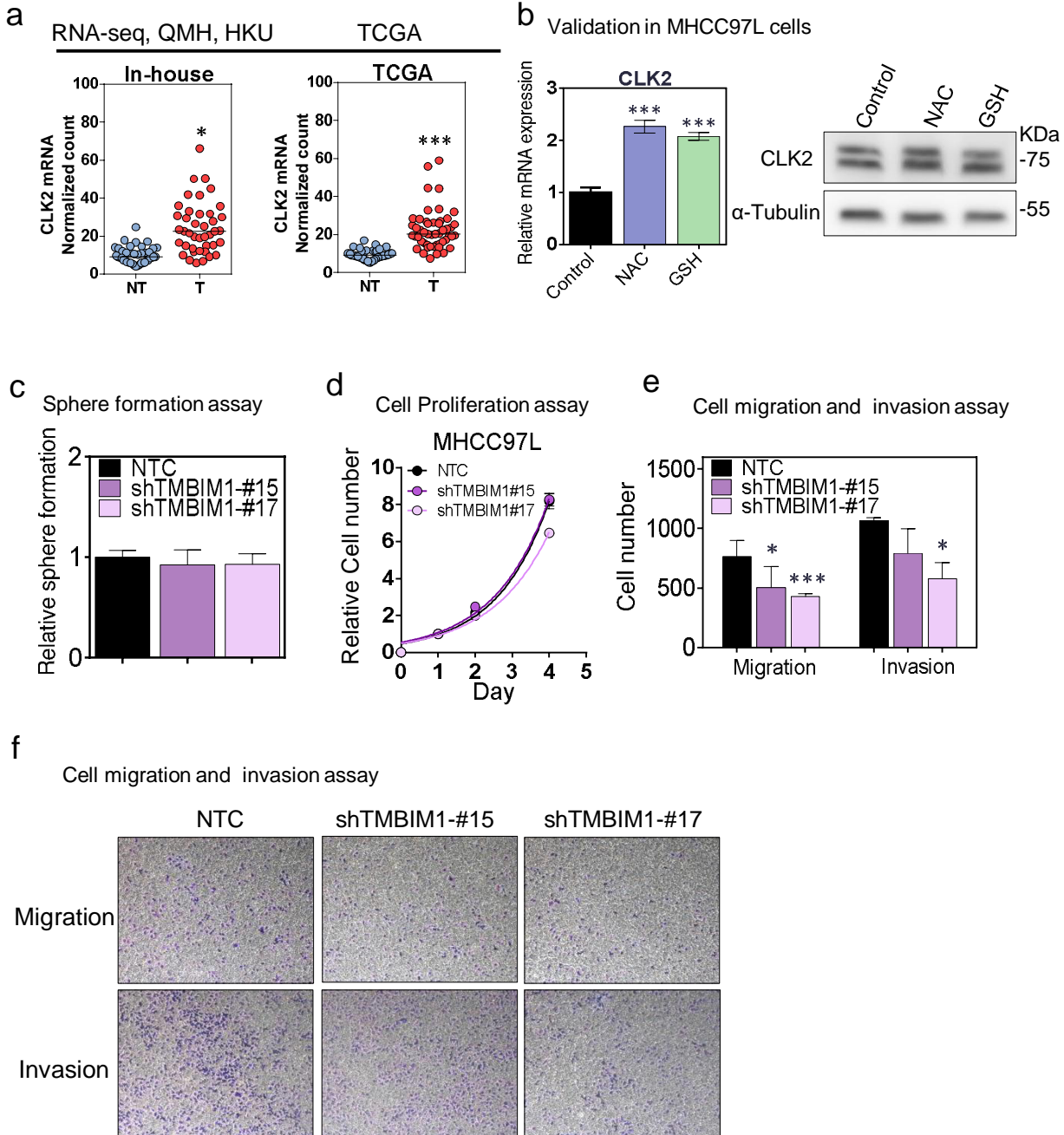
GO biological process: down-regulated genes by NAC



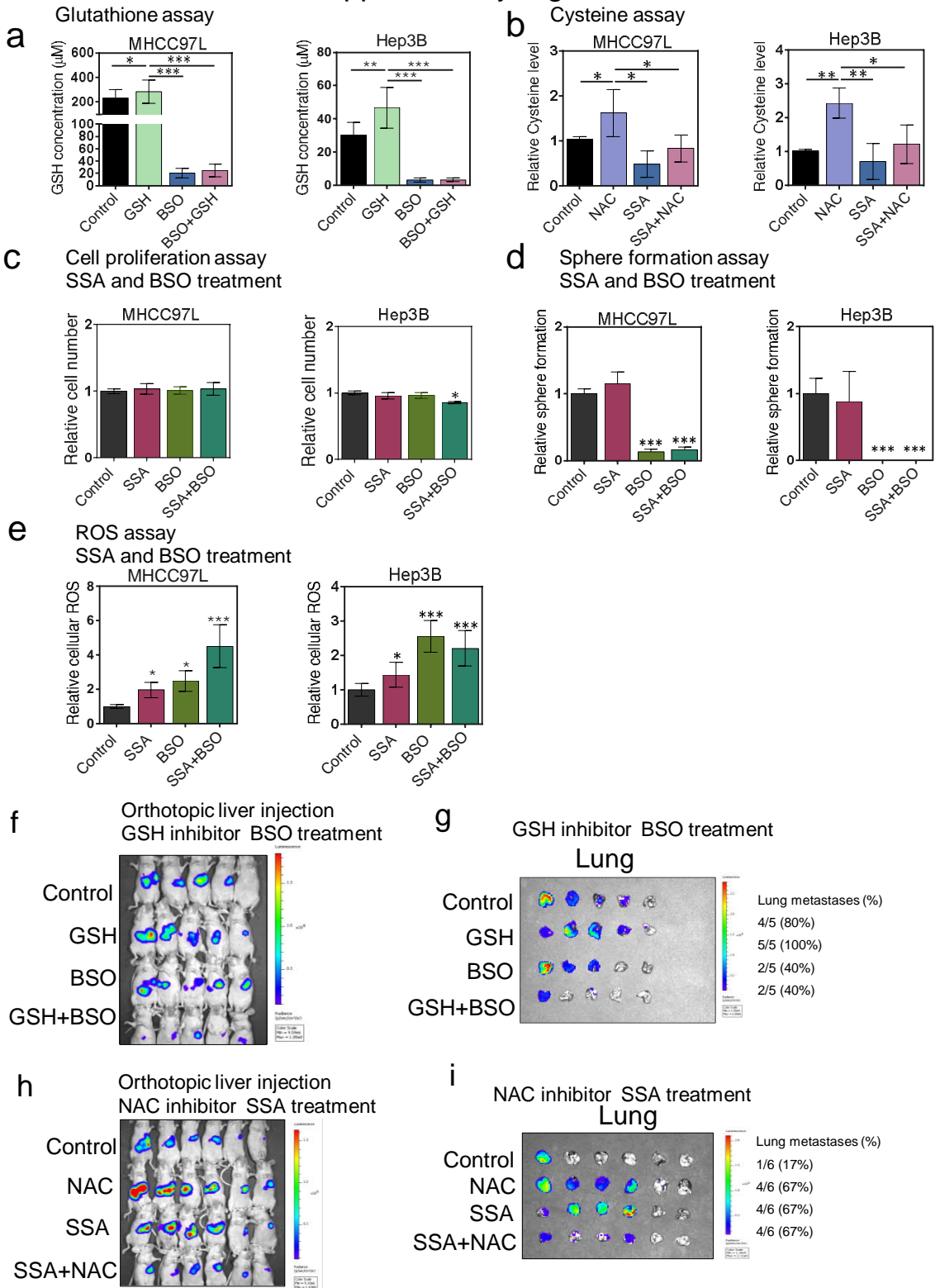
GO biological process: down-regulated genes by GSH



Supplementary Figure 6

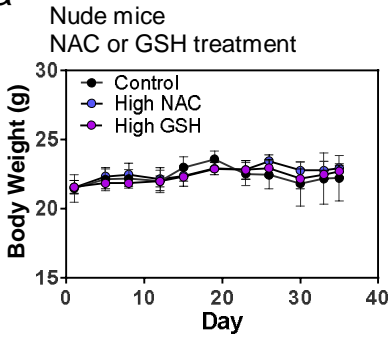


Supplementary Figure 7

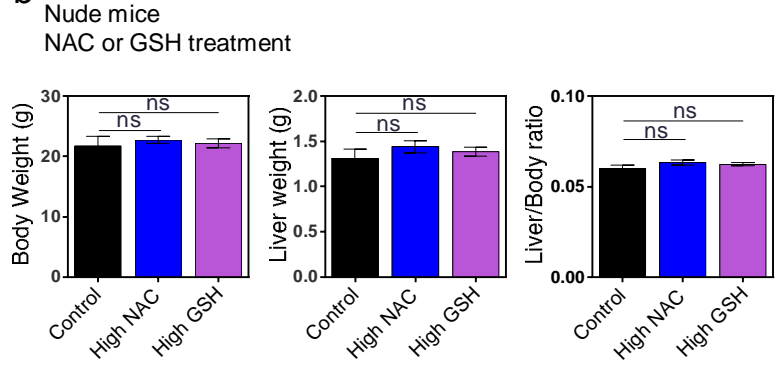


Supplementary Figure 8

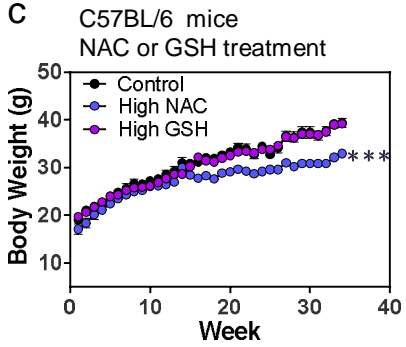
a



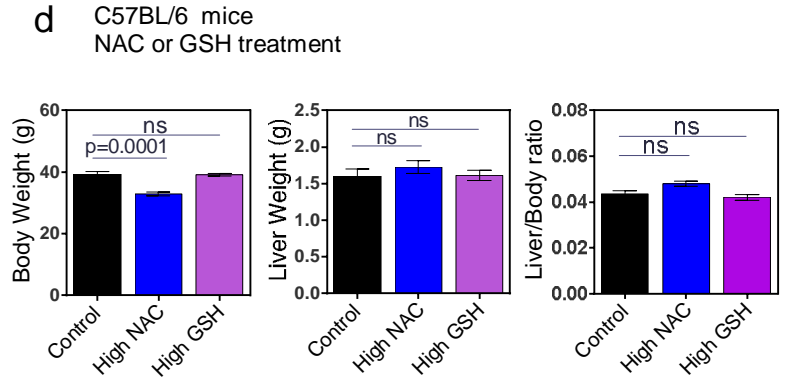
b



c



d



Supplementary Figure 9

Test Results:	DNA Marker	MHCC97L ⁴	MHCC97L (L-171218744P)
		AMEL	X, Y
	CSFIPO	11, 13	11, 13
	D13S317	8	8
	D16S539	12	12
	D5S818	12,13	12,13
	D7S820	10	10
	TH01	9	9
	TPOX	8	8
	vWA	14	14
	D18S51	--	13, 22
	D21S11	--	31,2
	D3S1358	--	15, 16
	D8S1179	--	12, 13
	FGA	--	21, 24
	Penta D	--	8, 9
	Penta E	--	11, 17
	Number of shared alleles		12
	Total number of alleles in the reference profile		12
	Percent match		100%

Comment: The tested cell line has 100% match with MHCC97L. Therefore, they are considered to be related from a common ancestry.

Figure S9. STR authentication of MHCC-97L cells.

Supplementary Figure 10

a Cell proliferation assay
Sorafenib, SSA and BSO treatment

