

Table S1 Biochemistry parameters of clinical liver function

index groups	ALT (U/L)	AST (U/L)	ALP (U/L)	D-Bili (μmol/L)	T-Bili (μmol/L)	γ-GT (U/gprot)	TBA (μmol/L)	T-SOD (U/mgprot)	MDA (nmol/mgprot)	GSH-Px (μmol/L)
Control	89.79	164.51	99.26	1.23	1.21	5.64	5.09	20.24	117.40	123.58
	±25.68	±15.39	±8.11	±0.52	±0.50	±2.35	±1.53	±3.90	±30.83	±31.99
YHS	147.1	224.83	203.26	2.41	1.93	7.38	9.26	18.56	133.28	90.90
	±34.49**	±42.66**	±64.73**	±0.99*	±0.47*	±2.40	±2.71**	±3.58	±20.92	±9.74*
GEN	96.42	164.73	108.00	1.24	1.31	5.36	5.80	19.97	113.73	125.79
	±3.86#	±39.10#	±8.71#	±0.52#	±0.30#	±0.62	±0.60#	±1.23	±16.90	±16.54##

Note: *P<0.05, **P<0.01 compared with the control group

#P<0.05, ##P<0.01 compared with the YHS group

Table S2 Identification and trends of change for differential metabolites

NO	Rt(min)	m/z determined	m/z calculated	Error (mDa)	Ion form	Molecular Formula	Metabolite Name	Trend	VIP	T'TEST
1	0.57	146.1653	146.1657	-0.4	[M+H] ⁺	C ₇ H ₁₉ N ₃	Spermidine	↓	1.00	0.0069
2	0.86	126.0222	126.0225	-0.3	[M+H] ⁺	C ₂ H ₇ NO ₃ S	Taurine	↓	3.00	0.0062
3	1.96	314.1202	314.1240	-3.8	[M+H] ⁺	C ₁₄ H ₁₉ NO ₇	Tyramine glucuronide	↓	1.13	0.0297
4	2.10	188.0911	188.0923	-1.2	[M+H] ⁺	C ₈ H ₁₃ NO ₄	2-Keto-6-acetamidocaproate	↓	1.39	0.0096
5	2.70	122.0263	122.0276	-1.3	[M+H] ⁺	C ₃ H ₇ NO ₂ S	L-Cysteine	↓	1.42	0.0001
6	3.17	109.0286	109.0290	-0.4	[M+H] ⁺	C ₆ H ₄ O ₂	1,2-Benzoquinone	↑	1.45	0.0143
7	3.32	168.0662	168.0661	0.1	[M+H] ⁺	C ₈ H ₉ NO ₃	Pyridoxal	↓	1.42	0.0010
8	3.33	315.1193	315.1206	-1.3	[M+H] ⁺	C ₁₄ H ₁₄ N ₆ O ₃	7,8-Dihydropteroic acid	↓	1.61	0.0006
9	3.74	210.0742	210.0766	-2.4	[M+H] ⁺	C ₁₀ H ₁₁ NO ₄	Hydroxyphenylacetyl glycine	↓	4.90	0.0041
10	3.92	184.0972	184.0974	-0.2	[M+H] ⁺	C ₉ H ₁₃ NO ₃	Epinephrine	↑	2.46	0.0442
11	4.67	474.1725	474.1737	-1.2	[M+H] ⁺	C ₂₀ H ₂₃ N ₇ O ₇	10-Formyltetrahydrofolate	↑	1.54	0.0060
12	4.87	235.1081	235.1083	-0.2	[M+H] ⁺	C ₁₂ H ₁₄ N ₂ O ₃	5-Methoxytryptophan	↑	1.38	0.0038
13	4.87	281.1146	281.1137	0.9	[M+H] ⁺	C ₁₃ H ₁₆ N ₂ O ₅	L-beta-aspartyl-L-phenylalanine	↑	1.31	0.0284
14	5.10	190.0504	190.0504	0	[M+H] ⁺	C ₁₀ H ₇ NO ₃	Kynurenic acid	↓	8.04	0.0010
15	5.11	130.0637	130.0657	-2.0	[M+H] ⁺	C ₉ H ₇ N	3-Methylene-indolenine	↑	1.39	0.0011
16	5.64	247.0949	247.0930	1.9	[M+H] ⁺	C ₉ H ₁₄ N ₂ O ₆	5,6-Dihydrouridine	↓	1.20	0.0040
17	6.22	233.1281	233.1290	-0.9	[M+H] ⁺	C ₁₃ H ₁₆ N ₂ O ₂	Melatonin	↑	1.24	0.0013
18	6.38	247.1293	247.1294	-0.1	[M+H] ⁺	C ₁₀ H ₁₈ N ₂ O ₅	L-beta-aspartyl-L-leucine	↑	1.02	0.0075
19	6.75	385.1694	385.1685	0.9	[M+H] ⁺	C ₁₉ H ₂₈ O ₆ S	3b,16a-Dihydroxyandrostene sulfate	↓	1.16	0.0364
20	7.37	300.0913	300.0944	-3.1	[M+H] ⁺	C ₁₀ H ₁₃ N ₅ O ₆	8-Hydroxyguanosine	↓	1.28	0.0024

21	0.65	177.0400	177.0399	0.1	[M-H] ⁻	C ₆ H ₁₀ O ₆	L-Gulonolactone	↓	1.79	0.0241
22	0.65	195.0505	195.0505	0	[M-H] ⁻	C ₆ H ₁₂ O ₇	Gluconic acid	↓	7.42	0.0215
23	3.18	347.0977	347.0992	-1.5	[M-H] ⁻	C ₁₅ H ₁₆ N ₄ O ₆	Riboflavin reduced	↑	3.04	0.0077
24	3.49	252.0850	252.0872	-2.2	[M-H] ⁻	C ₁₂ H ₁₅ NO ₅	N-Acetylvanilalanine	↑	1.02	0.0003
25	3.72	176.0375	176.0381	-0.6	[M-H] ⁻	C ₆ H ₁₁ NO ₃ S	N-Formyl-L-methionine	↓	1.23	0.0394
26	3.78	259.1292	259.1292	-0.2	[M-H] ⁻	C ₁₁ H ₂₀ N ₂ O ₅	L-gamma-glutamyl-L-leucine	↑	1.31	0.0083
27	4.52	137.0602	137.0603	-0.1	[M-H] ⁻	C ₈ H ₁₀ O ₂	Tyrosol	↑	2.08	0.0263
28	4.52	167.0341	167.0344	-0.3	[M-H] ⁻	C ₈ H ₈ O ₄	Homogentisic acid	↑	3.44	0.0036
29	5.16	175.0607	175.0607	0.1	[M-H] ⁻	C ₇ H ₁₂ O ₅	2-Isopropylmalic acid	↑	2.37	0.0259
30	5.61	163.0393	163.0395	0.1	[M-H] ⁻	C ₉ H ₈ O ₃	Phenylpyruvic acid	↑	2.45	0.0040
31	6.76	194.0452	194.0453	-0.1	[M-H] ⁻	C ₉ H ₉ NO ₄	Dopaquinone	↑	1.87	0.0024
32	6.96	172.0975	172.0974	1.7	[M-H] ⁻	C ₈ H ₁₅ NO ₃	Isovalerylalanine	↑	11.8	0.0015
33	7.31	211.0600	211.0606	-0.6	[M-H] ⁻	C ₁₀ H ₁₂ O ₅	Vanillactic acid	↓	1.03	0.0104

Note: ↑, ↓ compared with the control group

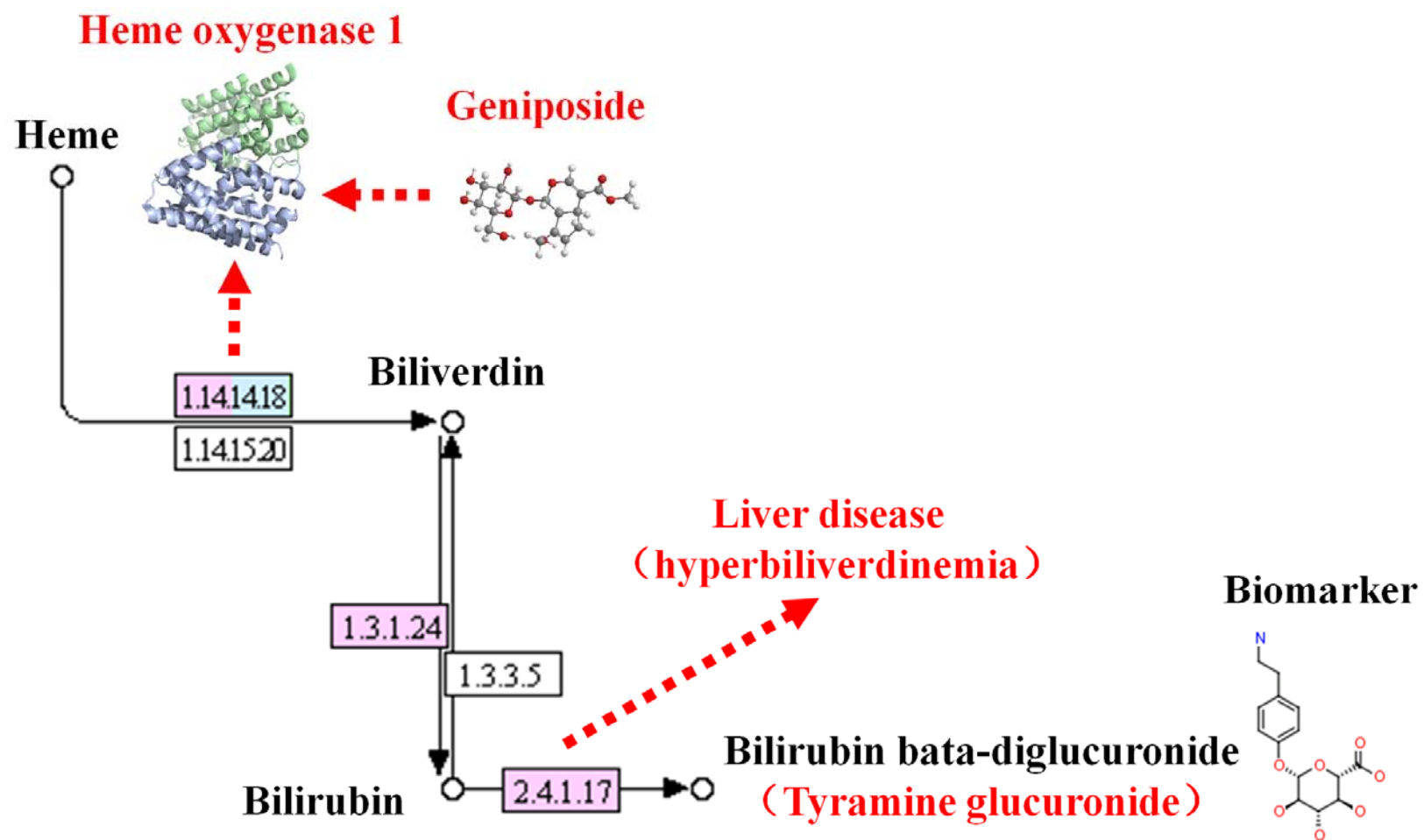


Fig. S1 The integrated analysis of metabolomics and network pharmacology

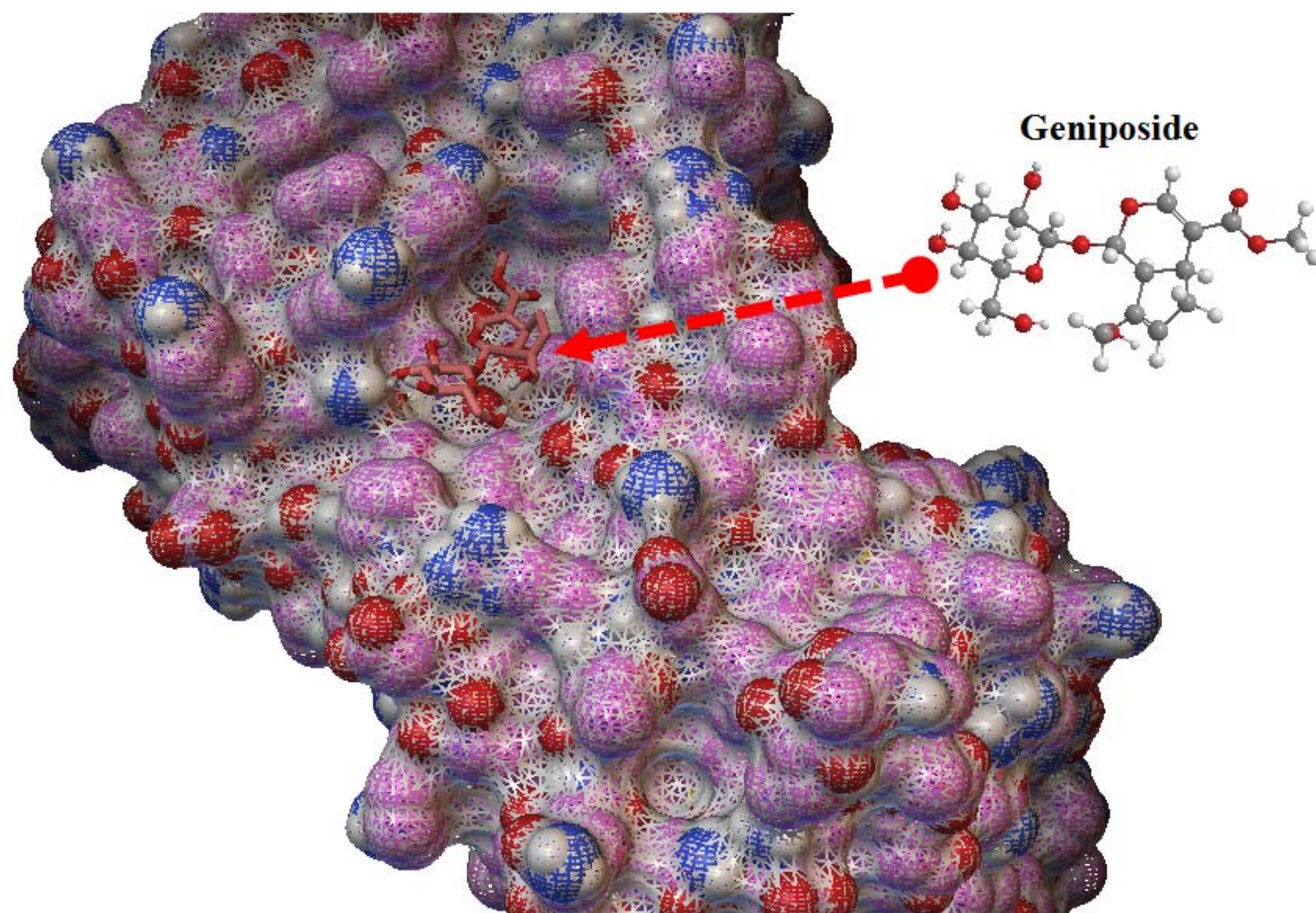


Fig. S2 The molecular docking result between heme oxygenase 1 and geniposide