

Supplementary Table 2. Baseline characteristics of the included patients

Variables	Buffy coat WTS data (n=175)		Targeted proteome data (n=58)	
	HCC (n=129)	Non-HCC (n=46)	HCC (n=52)	Normal (n=6)
Male	107 (82.9)	34 (73.9)	41 (78.8)	1 (16.7)
Age	59.1±12.0	49.3±13.6	60±10.5	30.0±3.9
Etiology				
Viral (HBV)	95 (73.6)	29 (63)	43 (82.7)	
Non-viral	34 (26.4)	17 (37)	9 (17.3)	
Cirrhosis	109 (84.5)	15 (32.6)	46 (88)	
Platelet	174.8±77.1	202.6±75.7	176.7±78.7	220.7±61.0
AST	62.4±65.0	53.7±35.0	53.4±32.5	17.5±0.7
ALT	42.6±44.9	65.0±61.5	39.0±28.5	12.5±0.7
Albumin	4.2±0.6	4.6±0.5	4.2±0.4	4.4±0.9
Bilirubin	1.3±2.8	0.7±0.4	0.8±0.6	0.5±0.4
AFP	2,898.7±7,630.1	8.2±15.6	3,048.3±8,802	1.4±0.5
PT	13.5±2.2	13.2±1.5	13.1±1.3	

Values are presented as number (%) or mean±standard deviation.

WTS, whole transcriptome sequencing; HCC, hepatocellular carcinoma; HBV, hepatitis B virus; AST, aspartate aminotransferase; ALT, alanine aminotransferase; AFT, alpha-fetoprotein; PT, prothrombin time.