Supplementary materials for

Proteomic dissection of cell type-specific H2AX-interacting protein complex associated with hepatocellular carcinoma

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Table 1S. Ratio of the ectopic to endogenous H2AX $/\gamma$ -H2AX expression in IB assay by densitometry.

Protein name ecto	Ratio pic/endogenous and SD
H2AX	0.81±0.05
γ-H2AX	1.15±0.04

(Standard deviation (SD) was obtained from 3 replicated measurements. Empirically, we selected the ratio> 1.2 for showing the up-regulated protein expression, and ratio<0.8 for showing the down-regulated protein expression.)

Table 2S. Ratio of protein expression in BLM(+) cells to that in BLM(-) cells in IB assay after IP by densitometry.

Protein name	e Ratio BLM(+)/BLM(-) and SD
PARP-1	1.34±0.10
PEP1	1.27±0.04
14-3-3ζ	1.51±0.06
CFL1	1.40±0.02
γ-H2AX	1.65 ± 0.09
FLAG	1.05±0.03

(Standard deviation (SD) was obtained from 3 replicated measurements. Empirically, we selected the ratio> 1.2 for showing the up-regulated protein expression, and ratio<0.8 for showing the down-regulated protein expression.)

Table 3S. Ratio of protein expression in 7703 cells to that in 7701 cells in IB assay after IP by densitometry.

Protein name Ratio

	7703/7701 and SD	
PARP-1	1.44 ± 0.07	
PEP1	1.03 ± 0.04	
CFL1	1.32 ± 0.07	
14-3-3ζ	1.41 ± 0.09	
γ-H2AX	0.92 ± 0.03	
FLAG	0.96 ± 0.02	

(Standard deviation (SD) was obtained from 3 replicated measurements. Empirically, we selected the ratio> 1.2 for distinguishing the up-regulated protein expression.)