Supplementary information to the manuscript:

Deviations of the immune cell landscape between healthy liver and hepatocellular carcinoma

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Cell type	Correlation	p-value
B cells	0.52	<2e-16
Macrophages	0.22	1.60e-05
NK cells	0.23	3.70e-06
$CD8^+$ T cells	0.21	3.60e-05

Supplementary Table 1. Pearson's correlation of CIBERSORT and EPIC deconvolution results.

Supplementary Table 2. Comparison of immune cell abundance in HCC calculated by xCell and CIBERSORT 3algorithms.

Cell type	xCell	score, mean+- SD				
/parameter			Pearson	Qualitat		
			betwee	n xCell and	ive	
			CIBE	RSORT	consiste	
						ncy
	HCC tissue	Tumor adjacent	p-value	correlatio	p-value	xCell vs
		tissue (TaT)	(HCC vs	n		CIBERS
			TaT)	coefficient		ORT
Immunoscore	0.42+0.10	0.54+0.09	4 10 ⁻²⁰	n.d.	n.d.	(yes)
Plasma cells	0.025+0.026	0.0047+0.011	10 ⁻¹¹	0.42	<10 -16	yes
T cells CD8+	0.091+0.048	0.11+0.06	5 · 10 ⁻⁵	0.53	<10 -16	yes
Tregs	0.16+ 0.06	0.14+ 0.06	8 · 10 ⁻³	-0.19	0.00013	no
Total	0.038+0.028	0.012+0.015	2 10 ⁻¹⁴	n.d.	n.d.	yes
Macrophages						
M1	0.0023+0.0049	0.00034+0.00111	4 10-4	0.02	0.7	(yes)
macrophages						
M2	0.11+0.03	0.096+0.039	10-3	0.29	4 [·] 10 ⁻⁹	yes
macrophages						

	Γ		Viral s	tatus			ANOVA p-
CIBERSORT results	negativ	e (n=3)	HCV (n=57)	HBV (n=17)	values
	mean	stdev	mean	stdev	mean	stdev	(between the
							groups)
B_cells_naive	0,026	0,028	0,048	0,039	0,040	0,041	0,539
B_cells_memory	0,011	0,018	0,009	0,020	0,012	0,019	0,859
Plasma_cells	0,006	0,010	0,006	0,025	0,008	0,025	0,980
T_cells_CD8	0,099	0,022	0,117	0,081	0,091	0,056	0,442
T_cells_CD4_memory_resting	0,247	0,047	0,222	0,106	0,237	0,052	0,793
T_cells_CD4_memory_activated	0,028	0,016	0,023	0,030	0,012	0,020	0,315
T_cells_follicular_helper	0,015	0,013	0,047	0,046	0,039	0,051	0,472
Tregs	0,027	0,046	0,020	0,029	0,017	0,028	0,850
T_cells_gamma_delta	0,000	0,000	0,001	0,005	0,000	0,000	0,843
NK_cells_resting	0,035	0,037	0,018	0,035	0,026	0,038	0,570
NK_cells_activated	0,040	0,042	0,044	0,043	0,029	0,033	0,443
Monocytes	0,000	0,000	0,005	0,010	0,003	0,008	0,609
Macrophages_M0	0,048	0,035	0,029	0,039	0,022	0,030	0,514
Macrophages_M1	0,072	0,034	0,110	0,040	0,105	0,033	0,244
Macrophages_M2	0,126	0,043	0,145	0,084	0,157	0,068	0,787
Dendritic_cells_resting	0,022	0,027	0,022	0,030	0,034	0,035	0,422
Dendritic_cells_activated	0,000	0,000	0,000	0,001	0,000	0,000	0,843
Mast_cells_resting	0,095	0,082	0,080	0,059	0,107	0,071	0,280
Mast_cells_activated	0,051 ^{*#}	0,089	0,002*	0,016	0,005 [#]	0,016	0,001
Eosinophils	0,000	0,000	0,001	0,004	0,000	0,000	0,734
Neutrophils	0,052	0,029	0,052	0,035	0,057	0,037	0,882
P_value	0,173	0,146	0,122	0,174	0,142	0,228	0,844
total_Tcells	0,514	0,051	0,472	0,078	0,450	0,085	0,363
total_Bcells	0,037	0,025	0,056	0,036	0,052	0,039	0,617
total_NK	0,075	0,009	0,062	0,042	0,055	0,034	0,695
total_Mast_cells	0,146	0,008	0,082	0,058	0,112	0,065	0,051
total_MF	0,247	0,038	0,284	0,084	0,283	0,058	0,718
total_DC	0,022	0,027	0,023	0,030	0,034	0,035	0,429
M1_M2_ratio	0,716	0,633	1,033	0,946	0,840	0,491	0,625

Supplementary Table 3. Impact of the viral status in HCC patients on CIBERSORT immune cell frequencies.

Viral status was available for a subset of 78 patients. Of these, n= 57 were HCV positive, n=17 were HBV positive, n=3 were HBV/HCV negative and n=1 was positive for both HBV/HCV. Double positive patient was excluded from analysis. Means and standard deviations were calculated for each group. * p=0.001 vs HCV after Bonferroni correction; # p=0.003 vs HBV after Bonferroni correction

Dataset	Sex	Age,	Condition	Reference
		years		
GSM372247	female	27	Healthy adult, sudden death	33
GSM372248	female	29	Healthy adult, sudden death	33
GSM372249	male	29	Healthy adult, sudden death	33
GSM372599	female	48	Liver transplant	33
GSM372600	male	34	Liver transplant	33
GSM373314	male	53	Liver transplant	33
GSM373315	female	48	Liver transplant	33
GSM373324	female	47	Liver transplant	33
GSM35982	NA	NA	Healthy liver	34
GSM155926	NA	NA	Healthy liver	35,36
GSM155988	NA	NA	Normal liver tissue	35,36
GSM176332	male	NA	Normal liver tissue	35
GSM80730	female	39	Healthy adult, sudden death	35,37
GSM319287	NA	NA	Liver transplant	35,39
E-AFMX-11HL4	male	29	Normal liver tissue	35,38
E-AFMX-11HL5	Male	NA	Normal liver tissue	35,38

Supplementary Table 4. Sample characteristics of healthy liver datasets.

Supplementary Table 5. Estimated absolute mast cell infiltration in HCC and TaT using "absolute score" feature of CIBERSORT.

	HCC (n=183)	TaT (n=65)	p-value for means
			comparison by ANOVA
Mast cells resting	0,029+0,031*	0,046+0,039 [*]	0,004
Mast cell activated	0,0063+0,014	0,0031+0,0077	0,522
Mast cell total	0,035+0,028 [#]	0,049+0,035 [#]	0,028

Only the datasets with p<0.05 of CIBERSORT calculations in "absolute score" mode were selected for analysis.

#, * the corresponding TaT and HCC values differ significantly at p=0.05.

	sex	Age	AFP	tumor size (cm)	tumor- grading	underlying disease	Mq	Nd	vascular invasion
Patient1	male	65	n.a.	10	3	HCV	0	NO	1
Patient2	male	49	2,9	1,1	1	HCV	0	NO	0
Patient3	male	55	n.a.	6	2	HCV	n.a.	n.a.	0
Patient4	female	68	7,8	6	2	HCV	0	NO	0
Patient5	male	55	4,4	1,1	2	HCV	0	NO	0
Patient6	male	61	641	0,5	2	HCV	0	NO	0
Patient7	male	50	85	0,5	2	HBV+HCV	0	NO	0
Patient8	male	58	38,9	0,6	1	HCV	0	NO	0
Patient9	male	58	22	2,2	2	HCV	0	NO	0
Patient10	male	65	8,2	2,4	2	HCV	0	NO	0
mean		58	101	3,0					
stdev		6	220	3,2					

Supplementary Table 6. Clinical characteristics of HCC patients used for immunohistochemical quantitation of mast cells in tumor and tumor adjacent tissues shown in Fig. 6C.

Supplementary Figure 1



Supplementary Fig.1 . Natural Killer (NK) cells and dendritic cells (DC) in human HCC tumor tissue (HCC), adjacent tissue (TaT) and healthy liver (HL). CIBERSORT immune cell fractions were determined for each patient; each dot represents one patient. Mean values and standard deviations for each cell subset including resting NK cells (A), activated NK cells (B), resting dendritic cells (C) and activated dendritic cells (D) were calculated for each patient group and compared using one-way ANOVA. * p<0.05; ** p<0.01

Supplementary Fig. 2

	Healthy liver (n=17)			HCC tumor tissue (n=258)			Tumor a tissue	idjacent (n=46)
	MC rest.	MC activ.		MC rest.	MC activ.		MC rest.	MC activ.
B_cells_naive	ns	ns		-0,11	ns		-0,275	<u>0,352</u>
B_cells_memory	ns	ns		-0,13	0,13		ns	ns
Plasma_cells	ns	-0,51		<u>-0,30</u>	ns		-0,322	ns
T_cells_CD8	ns	ns		ns	-0,15		-0,285	ns
T_cells_CD4_naive	ns	ns		ns	ns		ns	ns
T_cells_CD4_memory_resting	ns	ns		0,14	ns		ns	<u>-0,384</u>
T_cells_CD4_memory_activated	ns	ns		<u>-0,26</u>	ns		ns	ns
T_cells_follicular_helper	ns	-0,47		<u>-0,41</u>	<u>0,15</u>		<u>-0,553</u>	ns
Tregs	ns	-0,50		ns	ns		-0,257	ns
T_cells_gamma_delta	ns	ns		<u>-0,19</u>	<u>0,16</u>		ns	0,323
NK_cells_resting	<u>0,56</u>	-0,48		ns	ns		ns	ns
NK_cells_activated	ns	ns		<u>0,21</u>	ns		ns	ns
Monocytes	ns	0,50		<u>0,28</u>	<u>-0,19</u>		<u>0,343</u>	ns
Macrophages_M0	ns	0,52		<u>0,42</u>	<u>-0,15</u>		0,329	-0,263
Macrophages_M1	ns	ns		<u>0,22</u>	<u>-0,164</u>		ns	<u>0,431</u>
Macrophages_M2	ns	-0,51		<u>-0,288</u>	<u>0,17</u>		ns	ns
Dendritic_cells_resting	ns	ns			<u>-0,159</u>		-0,29	ns
Dendritic_cells_activated	ns	<u>0,59</u>		<u>-0,232</u>	<u>0,197</u>		ns	<u>0,386</u>
Eosinophils	ns	<u>0,60</u>		<u>-0,347</u>	<u>0,291</u>		-0,311	<u>0,438</u>
Neutrophils	ns	ns		<u>-0,21</u>	<u>0,167</u>		ns	ns
bold	p<0.05							
bold and underscored	p<0.01							
ns	no signific	ant correla	atio	ו				
red	positive correlation (r ² >0)							
blue	negative correlation (r ² <0)							

suppl. Fig.2. Pearson correlation coefficients between resting /activated mast cells and other immune cell populations in healthy livers, HCC tumor tissues and tumor adjacent tissues. Cell populations for each patient were calculated by CIBERSORT as decribed in Materials and Methods.