

# 1 **Supplementary Figure Legends**

2 **Supplementary Figure 1.** Histogram of nuclear area of mononuclear hepatocytes for one HCC  
3 patient (A). The shape of the histogram shows a clear trimodal distribution (peaks), which are  
4 positioned proportionally. The first, second and third peaks are representative of diploid, tetraploid  
5 and octoploid nucleus, respectively (nuclear ploidy). Nuclei with area < 200 or > 4000 pixel<sup>2</sup> were  
6 excluded from analysis (non-hepatocytes population and incorrect DAPI segmentation).

7 Immunohistochemistry analysis of positive (left panel) and negative (right panel) PD-L1 expressions on  
8 tumoral cells (B) and immune stroma (C). Upper panel, 200× magnification, scale bar = 200 μm;  
9 lower panel, 400× magnification, scale bar = 50 μm. PD-L1 showed a membrane staining. PD-L1,  
10 programmed death-ligand 1.

11 **Supplementary Figure 2.** Recurrence-free survival and HCC-specific survival stratified by the  
12 Milan criteria (A), AFP model (B) and Metroticket 2.0 criteria (C), and compared using the log-rank  
13 test in the derivation cohort. HCC, hepatocellular carcinoma. HMP–HCC, highly mononuclear  
14 polyploid hepatocellular carcinoma; PMP–HCC, poorly mononuclear polyploid hepatocellular  
15 carcinoma.

16 **Supplementary Figure 3.** Recurrence-free survival and HCC-specific survival according to the  
17 Milan criteria (A), AFP model (B), and Metroticket 2.0 criteria (C) in the validation cohort, and  
18 compared using the log-rank test. HCC, hepatocellular carcinoma. HMP–HCC, highly mononuclear  
19 polyploid hepatocellular carcinoma; PMP–HCC, poorly mononuclear polyploid hepatocellular  
20 carcinoma.

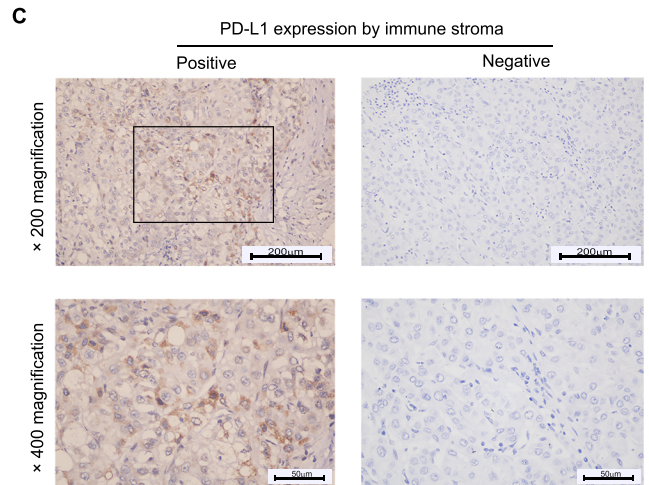
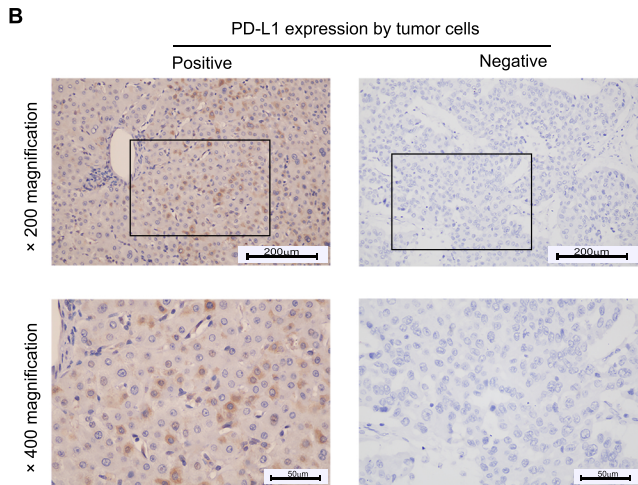
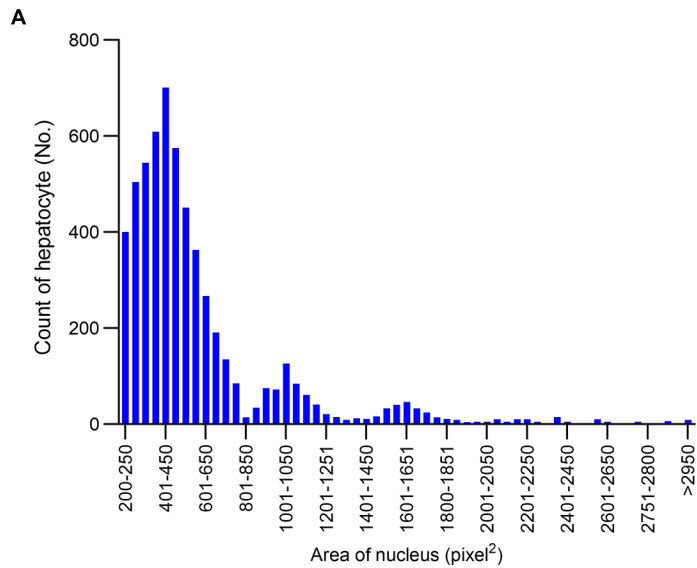
21 **Supplementary Figure 4.** HCC-specific survival curves according to the ploidy distribution (A), and  
22 the Milan criteria (B), AFP model (C), and Metroticket 2.0 criteria (D) in the derivation set. HCC,  
23 hepatocellular carcinoma. HMP–HCC, highly mononuclear polyploid hepatocellular carcinoma;  
24 PMP–HCC, poorly mononuclear polyploid hepatocellular carcinoma.

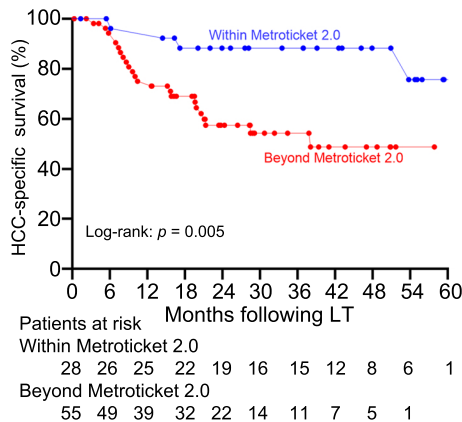
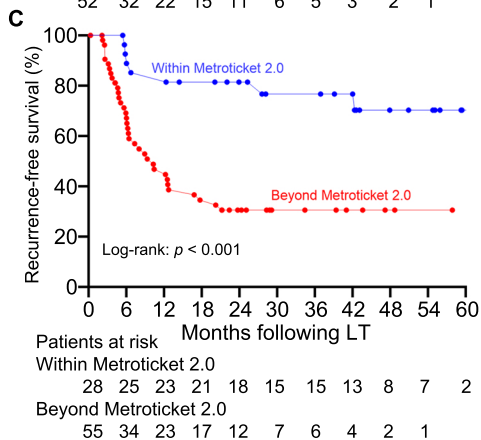
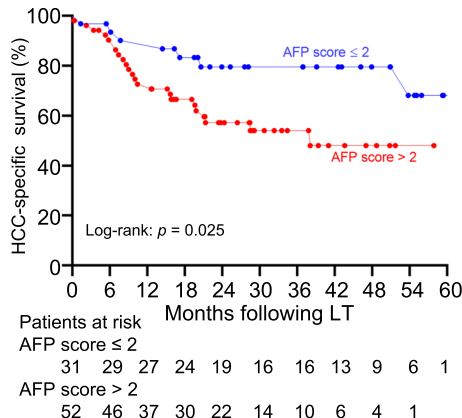
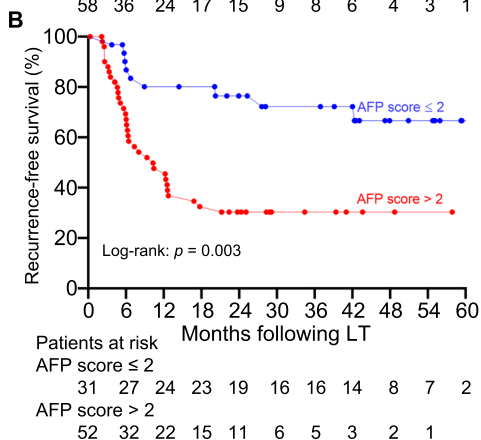
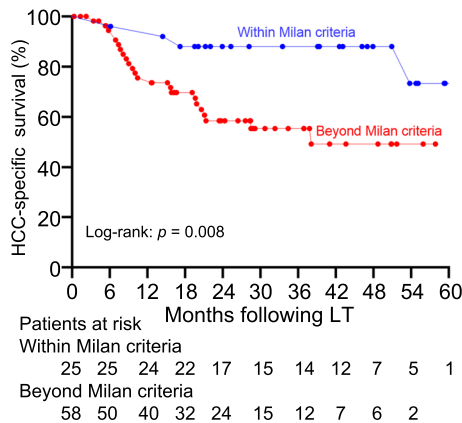
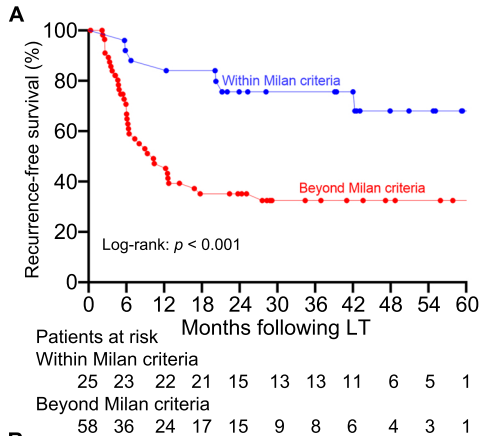
25 **Supplementary Figure 5.** Distributions of density of FoxP3<sup>+</sup> Treg cells (A) and CD8<sup>+</sup> cytotoxic T  
26 cells (B) between HMP–HCC and PMP–HCC. Similar distribution of density of CD4<sup>+</sup> Th cells

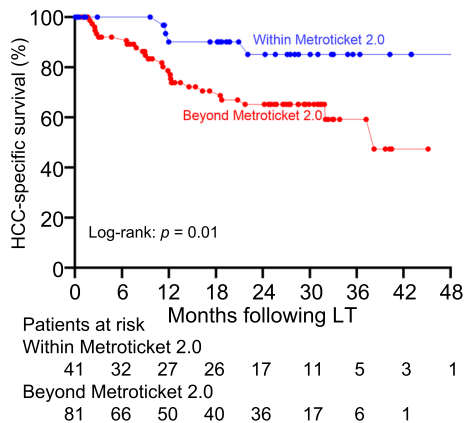
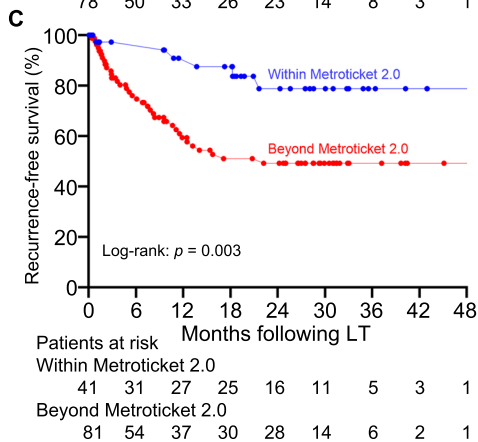
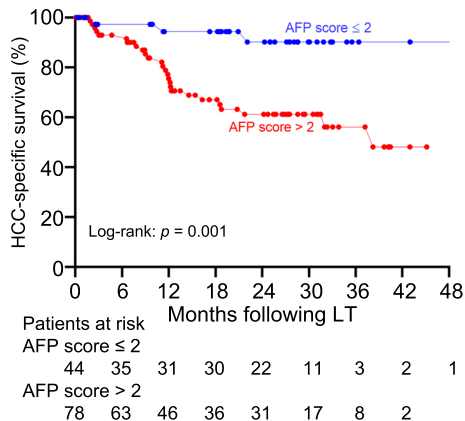
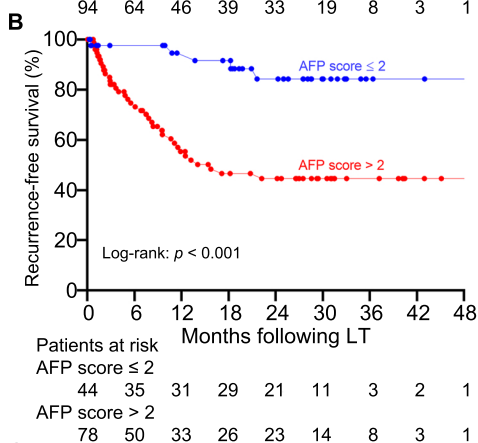
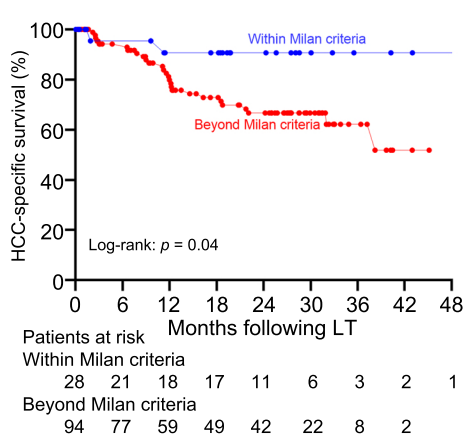
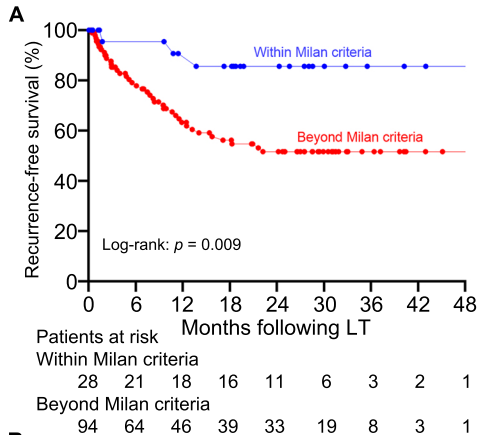
27 between HMP–HCC and PMP–HCC (C). No significant correlation between density of CD4<sup>+</sup> Th  
28 cells and the fraction of mononuclear polyploidy (D). FoxP3, forkhead box P3; HMP–HCC, highly  
29 mononuclear polyploid hepatocellular carcinoma; PMP–HCC, poorly mononuclear polyploid  
30 hepatocellular carcinoma.

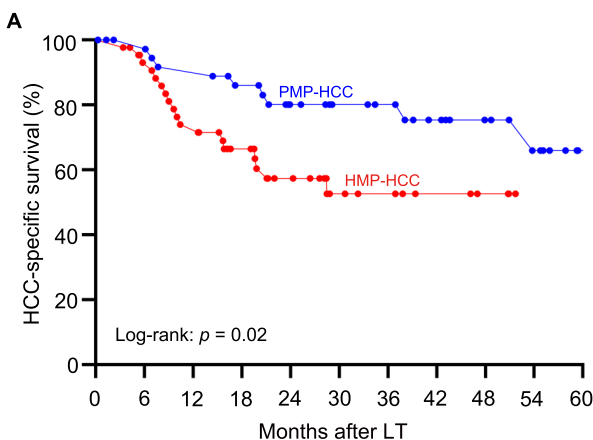
31 **Supplementary Figure 6.** Recurrence-free survival stratified according to the ploidy distribution (A),  
32 and with the combinations of the Milan criteria (B), AFP model (C), and Metroticket 2.0 criteria (D) in  
33 the validation set. HCC, hepatocellular carcinoma. HMP–HCC, highly mononuclear polyploid  
34 hepatocellular carcinoma; PMP–HCC, poorly mononuclear polyploid hepatocellular carcinoma.

35 **Supplementary Figure 7.** HCC-specific survival according to the ploidy distribution (A), and with  
36 the combinations of the Milan criteria (B), AFP model (C), and Metroticket 2.0 criteria (D) in the  
37 validation set. HCC, hepatocellular carcinoma. HMP–HCC, highly mononuclear polyploid  
38 hepatocellular carcinoma; PMP–HCC, poorly mononuclear polyploid hepatocellular carcinoma.



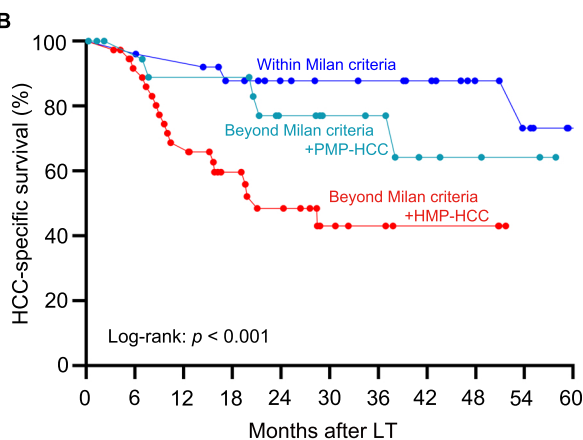






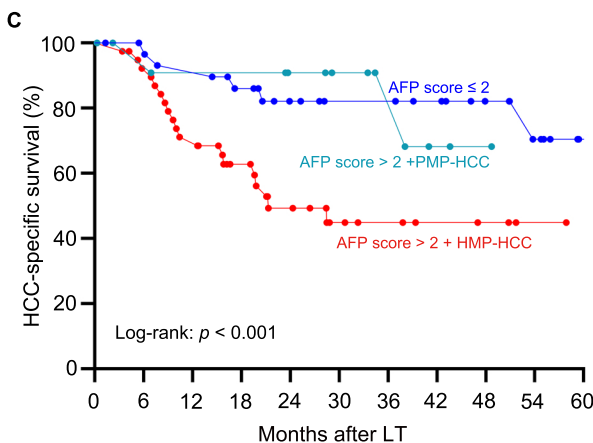
Patients at risk

PMP-HCC	39	36	33	30	24	20	18	14	10	7	1
HMP-HCC	44	39	31	24	17	10	8	5	3		



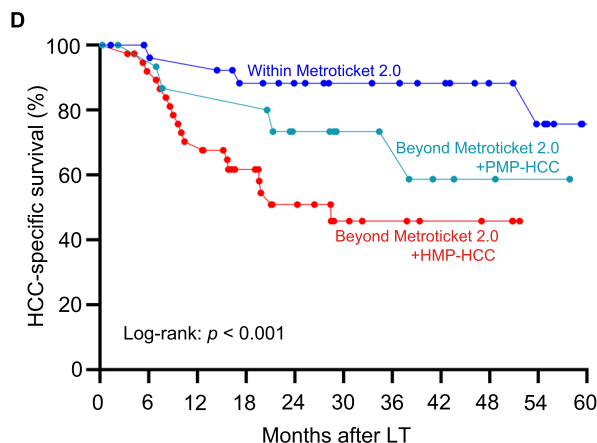
Patients at risk

Within Milan criteria	25	25	24	21	17	15	14	12	7	5	1
Beyond Milan criteria + PMP-HCC	21	18	16	16	11	8	7	4	3	2	
Beyond Milan criteria + HMP-HCC	37	32	24	17	13	7	5	3	3		



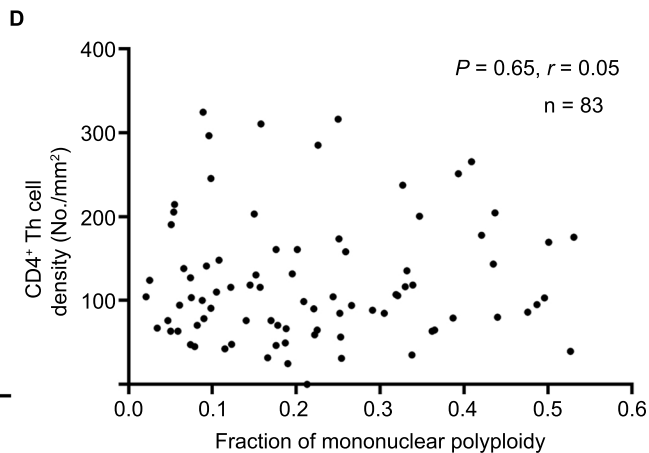
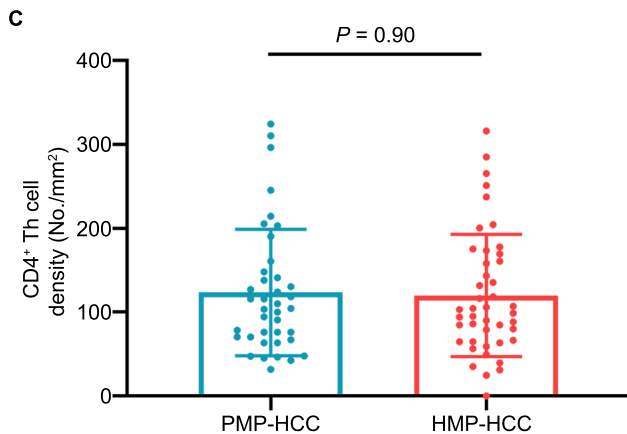
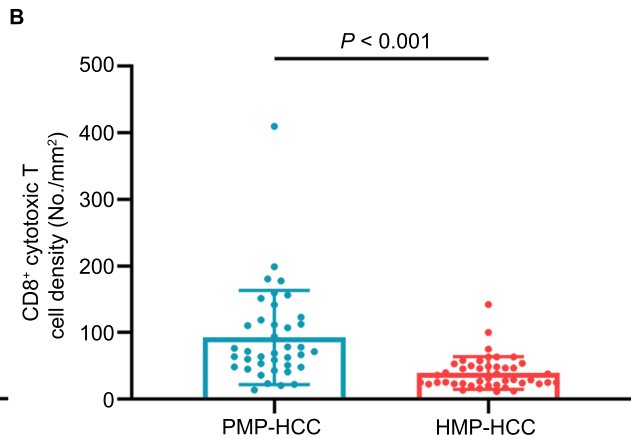
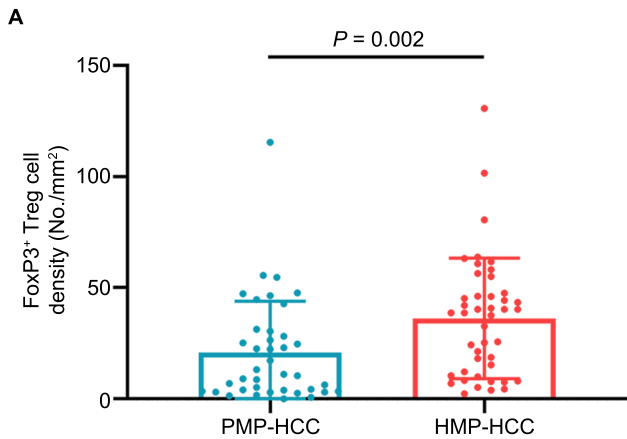
Patients at risk

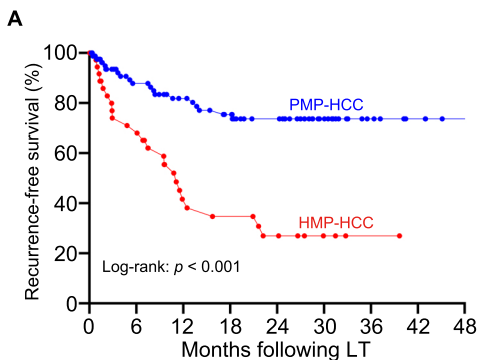
AFP score $\leq 2$	31	29	27	24	19	16	16	13	9	6	1
AFP score $> 2$ + PMP-HCC	13	11	10	10	8	6	4	2	1		
Beyond Milan criteria + HMP-HCC	39	35	27	20	14	8	6	4	3	1	



Patients at risk

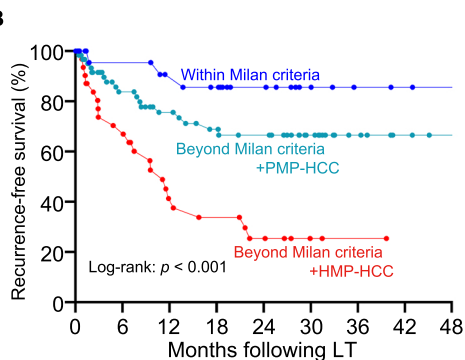
Within Metroticket 2.0	28	26	25	22	19	16	15	12	8	6	1
Beyond Metroticket 2.0 + PMP-HCC	17	15	13	13	9	6	5	3	2	1	
Beyond Metroticket 2.0 + HMP-HCC	38	34	26	19	13	8	6	4	3		





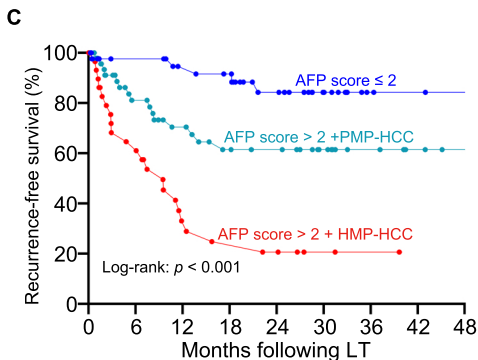
Patients at risk

PMP-HCC	85	61	52	45	37	22	10	5	2
HMP-HCC	37	24	12	10	7	3	1		



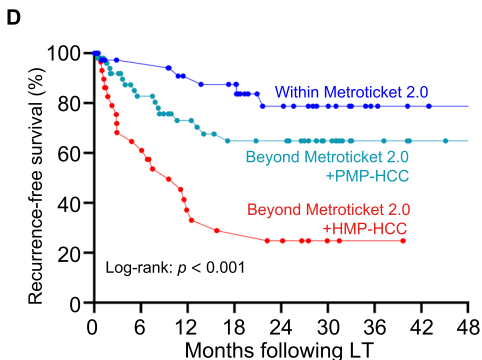
Patients at risk

Within Milan criteria	28	21	18	16	11	6	3	2	1
Beyond Milan criteria + PMP-HCC	63	43	35	30	27	17	7	3	1
Beyond Milan criteria + HMP-HCC	31	21	11	9	6	2	1		



Patients at risk

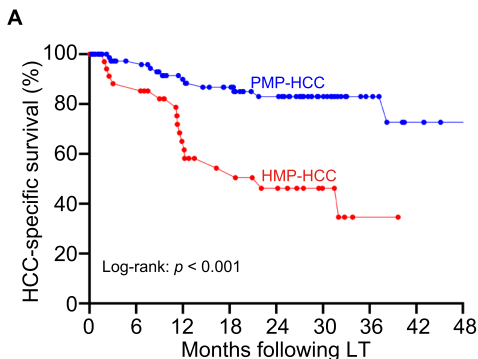
AFP score $\leq 2$	44	35	31	29	21	11	3	2	1
AFP score $> 2$ + PMP-HCC	48	32	25	20	18	12	7	3	1
AFP score $> 2$ + HMP-HCC	30	18	8	6	5	2	1		



Patients at risk

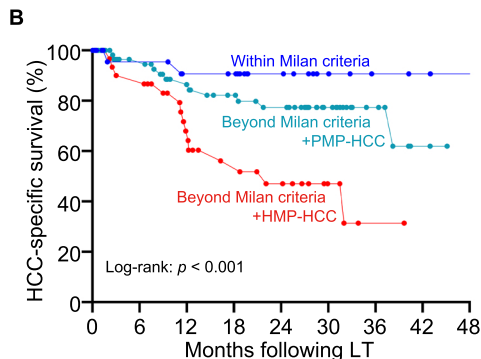
Within Metroticket 2.0	41	31	27	25	16	11	5	3	1
Beyond Metroticket 2.0 + PMP-HCC	52	36	28	23	22	12	5	2	1
Beyond Metroticket 2.0 + HMP-HCC	29	18	9	7	6	2	1		





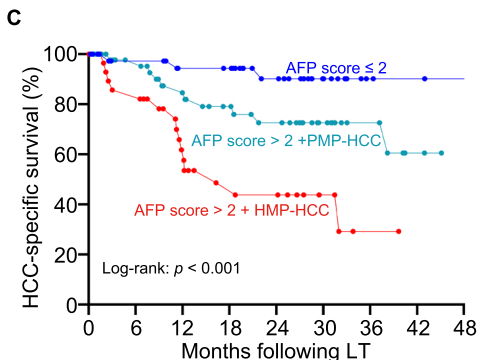
Patients at risk

PMP-HCC	85	68	58	52	42	23	10	4	1
HMP-HCC	37	30	19	14	11	5	1		



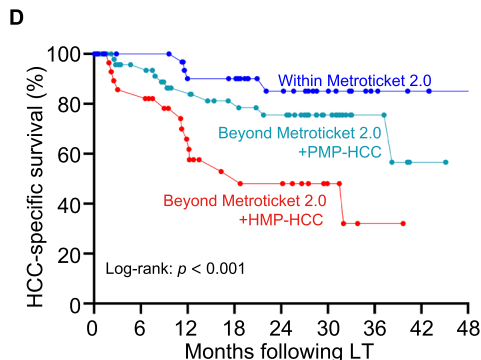
Patients at risk

Within Milan criteria	28	21	18	17	11	6	3	2	1
Beyond Milan criteria + PMP-HCC	63	50	41	36	32	18	7	2	
Beyond Milan criteria + HMP-HCC	31	27	18	13	10	4	1		



Patients at risk

AFP score $\leq 2$	44	35	31	30	22	11	3	2	1
AFP score $> 2$ + PMP-HCC	48	39	31	26	22	13	7	2	
AFP score $> 2$ + HMP-HCC	30	24	15	10	9	4	1		



Patients at risk

Within Metroticket 2.0	41	32	27	26	17	11	5	3	1
Beyond Metroticket 2.0 + PMP-HCC	52	42	34	29	26	13	5	1	
Beyond Metroticket 2.0 + HMP-HCC	29	24	16	11	10	4	1		