

**Supplementary table S3.** Demographic and clinical characteristics of the cirrhotic patients of the study populations A and B, subgrouped according to the respective median value of Δ-MELDNA during the waitlist time

	Cirrhotics of the study population A with low Δ-MELDNA	Cirrhotics of the study population A with high Δ-MELDNA	P value	Cirrhotics of the study population B with low Δ-MELDNA	Cirrhotics of the study population B with high Δ-MELDNA	P value
<b>Age, median (IQR)</b>	51.0 (45.0-59.0)	54.0 (49.0-60.8)	0.239	56.5 (52.3-60.3)	59.1 (52.1-63.7)	
<b>Male sex, n (%)</b>	10 (76.9)	12 (85.7)	0.648	12 (80.0)	15 (93.8)	
<b>BMI (Kg/m<sup>2</sup>), median (IQR)</b>	24.1 (22.6-28.4)	24.3 (22.9-26.7)	0.943	25.9 (24.0-29.7)	25.7 (24.5-28.7)	
<b>Cirrhosis aetiology, n (%):</b>						
MAFLD	7 (53.8)	9 (64.3)	0.704	9 (60.0)	8 (50.0)	0.576
Alcohol	5 (38.5)	5 (35.7)	1	10 (66.7)	6 (37.5)	0.156
Viral	9 (69.2)	10 (71.4)	1	4 (26.7)	8 (50.0)	0.273
<b>HCC, n (%)</b>	8 (61.5)	6 (42.9)	0.449	9 (60.0)	8 (50.0)	0.576
<b>Diuretic treatment, n (%):</b>						
No	4 (30.8)	4 (28.6)		5 (33.3)	6 (37.5)	
K sparing diuretics only	0 (0)	3 (21.4)	0.331	1 (6.7)	3 (18.8)	0.621
Loop diuretics plus K sparing diuretics	9 (69.2)	7 (50.0)		9 (60.0)	7 (43.8)	
<b>MELDNA at transplant, median (IQR)</b>	16.0 (12.2-23.0)	19.0 (14.8-21.8)	0.402	14.0 (10.0-23.0)	22.0 (14.0-28.8)	<0.001
<b>MELDNA at listing, median (IQR)</b>	15.9 (12.2-25.7)	14.6 (11.4-17.8)	0.280	15.9 (12.8-22.5)	17.6 (9.8-28.8)	0.830
<b>Δ-MELDNA, median (IQR)</b>	0.0 (-1.31 – 0.0)	1.09 (0.24-3.51)	0.059	-0.31 (-1.84-0.08)	3.23 (1.34-10.83)	<0.001
<b>Serum AST (IU/L), median (IQR)</b>	41.0 (35.5-70.0)	88.0 (58.5-115.3)	0.009	43.0 (28.0-59.0)	93.0 (61.0-139.5)	0.001
<b>Serum ALT (IU/L), median, (IQR)</b>	32.0 (24.5-59.0)	51.0 (43.5-95.3)	0.089	29.0 (20.0-48.0)	59.5 (36.3-84.8)	0.002
<b>Serum Mg, mg/dL, median (IQR)</b>	NAV	NAV	-	1.90 (1.70-2.11)	1.75 (1.63-1.94)	0.188
<b>Tissue Mg, (µg/g), median, (IQR) *</b>	118.7 (113.3-140.6)	114.7 (102.8-125.8)	0.280	NAV	NAV	

Percentage of weak positive hepatocytes to TRPM7 stain, median (IQR)	NAV	NAV	-	42.8 (24.6-56.2)	50.2 (40.7-56.3)	0.299
Percentage of intense positive hepatocytes to TRPM7 stain, median, (IQR)	NAV	NAV	-	0.6 (0.0-4.0)	7.0 (1.8-17.3)	<b>0.001</b>
Percentage of total positive hepatocytes to TRPM7 stain, median (IQR)	NAV	NAV	-	46.8 (24.8-56.2)	55.4 (50.9-73.6)	0.045