

Supplementary Materials

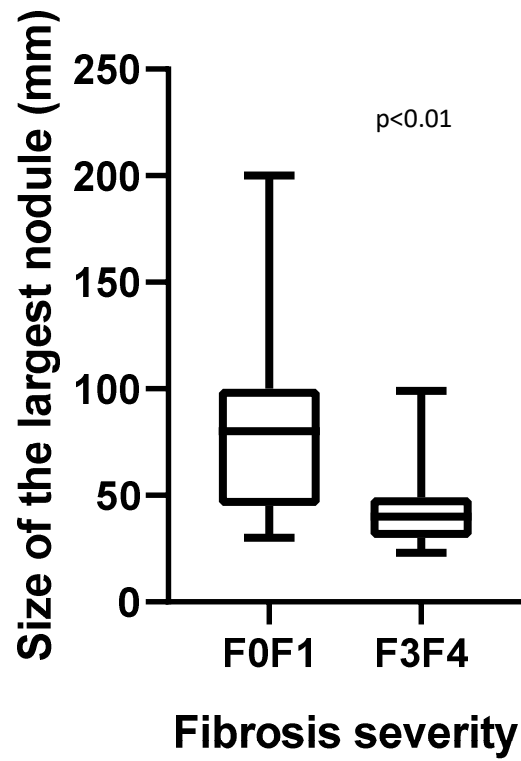


Figure S1. Distribution of the size of the largest nodule according to the severity of fibrosis (n=52).

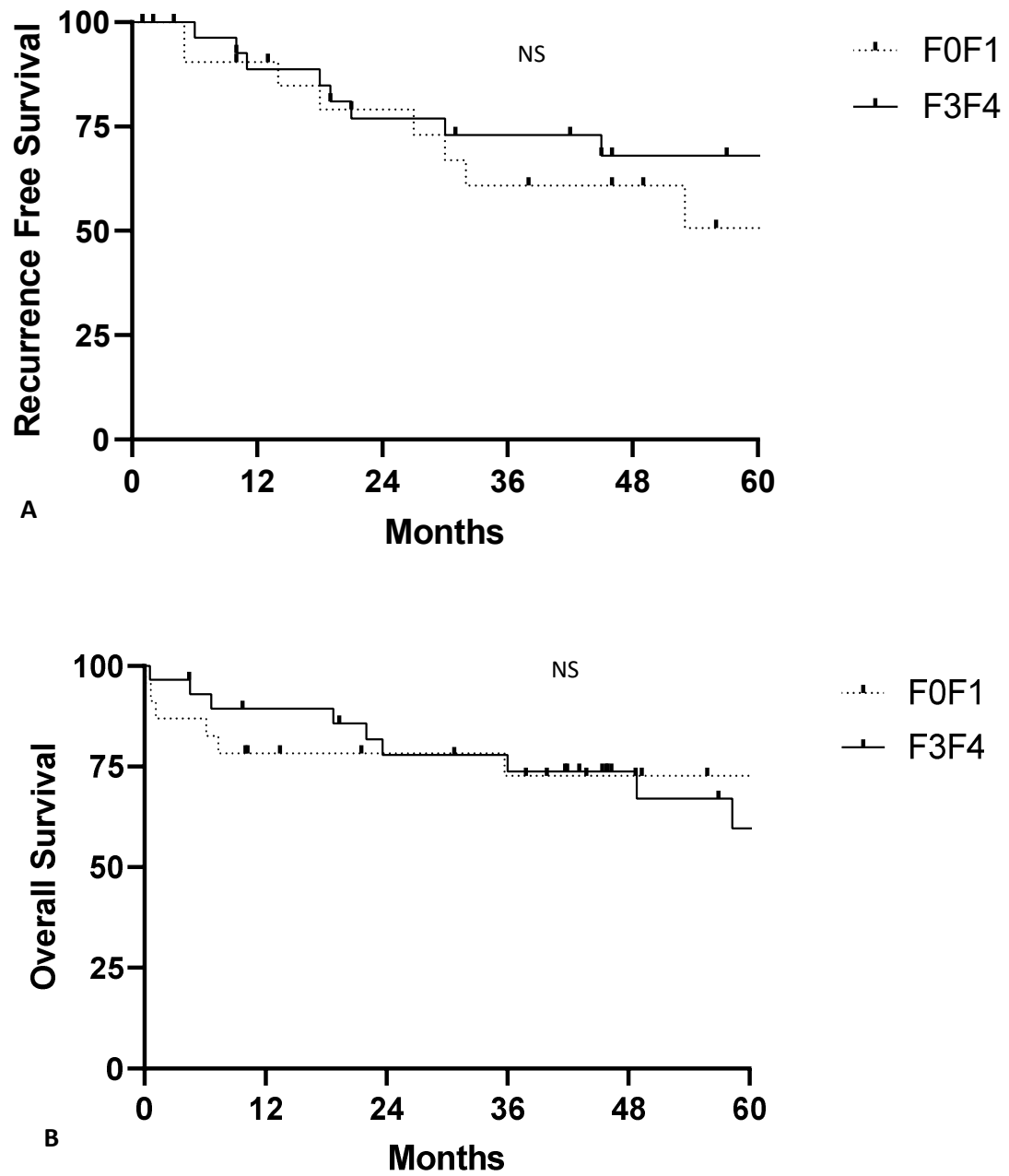
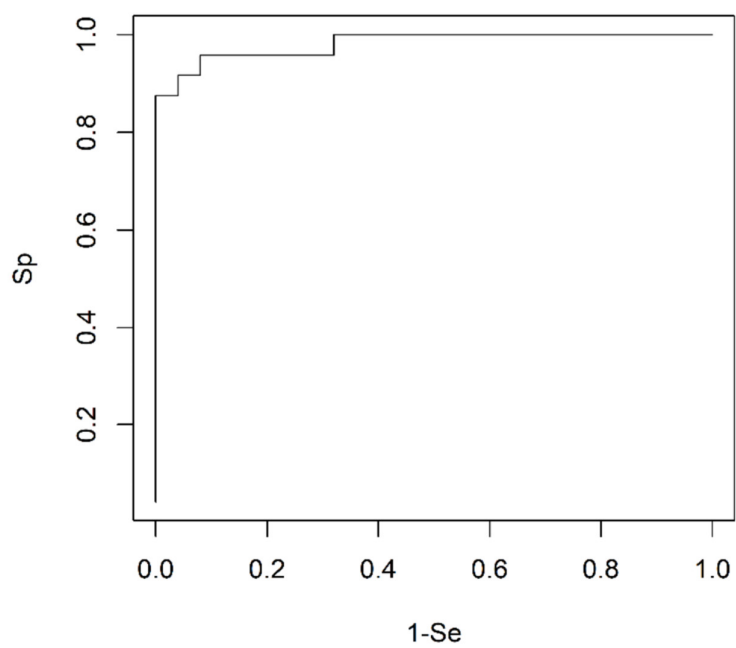


Figure S2. Recurrence free survival (A) and overall survival (B) according to the severity of fibrosis (n=52).

**Table 1.** Discriminant metabolites between HCC-F0F1 *versus* NTT-F0F1 according to the number of selections in the aqueous and lipid phases. Upregulated (red) and down-regulated (blue) metabolites in tumors.

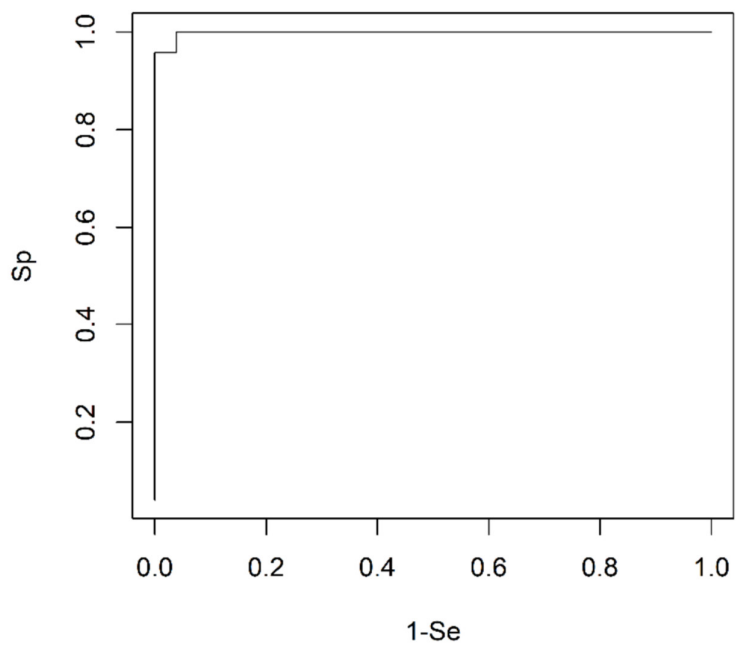
Chemical shift (ppm)	Metabolites	Selections
<b>Aqueous Phase (threshold: &gt;47 selections)</b>		
1.31	Lactate	441
2.14		79
2.43	Glutamine	53
2.56	Glutathione	72
3.22	Phosphocholine	48
3.62		82
3.65		76
3.66	Choline Derivatives	488
3.67		60
3.68		70
4.50	Ascorbic Acid	57
7.11	Histidine	54
7.85		49
3.42	Glucose	50
5.43	Glycogen	104
<b>Lipid Phase (threshold: &gt;39 selections)</b>		
1.52-1.54		81
2.25	Total Cholesterol	111
3.49		67
2.05	MUFA -CH <sub>2</sub> CH=	1339

### NTT-F0F1 vs HCC-F0F1



**A**

### NTT-F0F1 vs HCC-F0F1



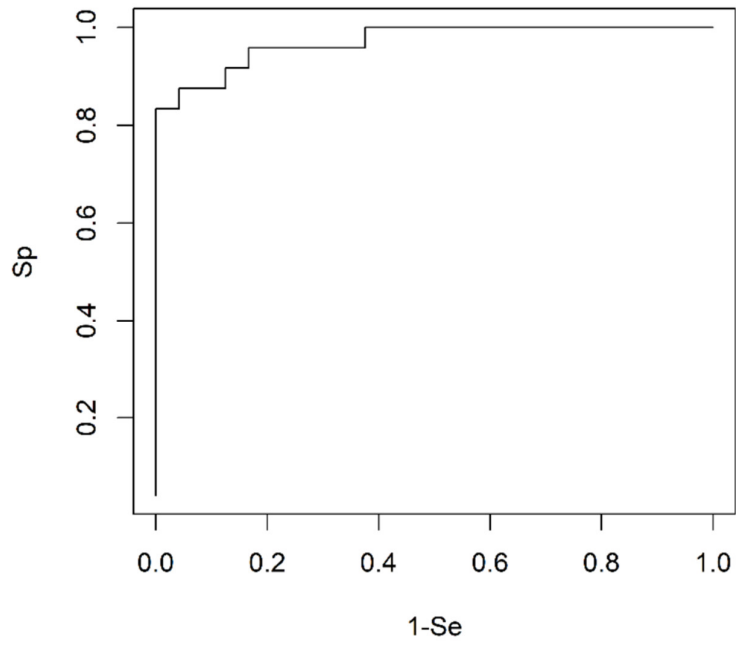
**B**

**Figure S3.** ROC curves of each subset of significant metabolites identified in aqueous (A) and lipid phases (B) for the discrimination of HCC from NTT in F0F1 fibrosis.

**Table S2.** Discriminant metabolites between HCC-F3F4 *versus* NTT-F3F4 according to the number of selections in the aqueous and lipid phases. Upregulated (red) and down-regulated (blue) metabolites in tumors.

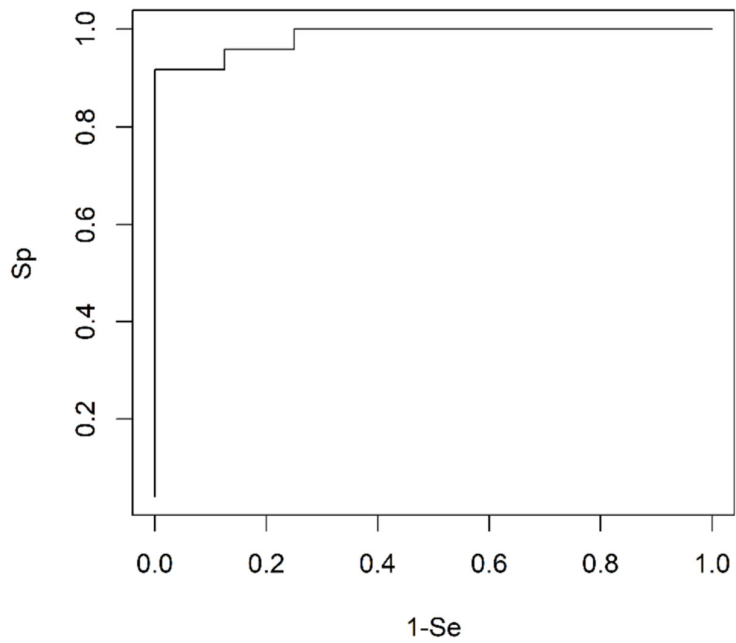
Chemical shift (ppm)	Metabolites	Selections
<b>Aqueous Phase (threshold: &gt;45 selections)</b>		
0.97	BCAA	49
1.02		46
1.31-1.32	Lactate	94
4.10-4.11		71
2.06-2.07	Glutamate	50
2.38		48
2.12-2.14	Glutamine	89
2.42-2.43		48
2.73	Sarcosine	91
3.63		106
8.20	Hypoxanthine	87
9.33	NAD	67
3.67	Choline derivatives	51
3.82		56
4.60	Glucose	49
6.98		64
7.10	Histidine	124
<b>Lipid Phase (threshold: &gt;21 selections)</b>		
1.41-1.43	SFA (CH <sub>2</sub> ) <sub>n</sub>	606
1.83	Total Cholesterol	98
2.22	Free Cholesterol	84
2.06	MUFA -CH <sub>2</sub> CH=	151

**NTT-F3F4 vs HCC-F3F4**



**A**

**NTT-F3F4 vs HCC-F3F4**



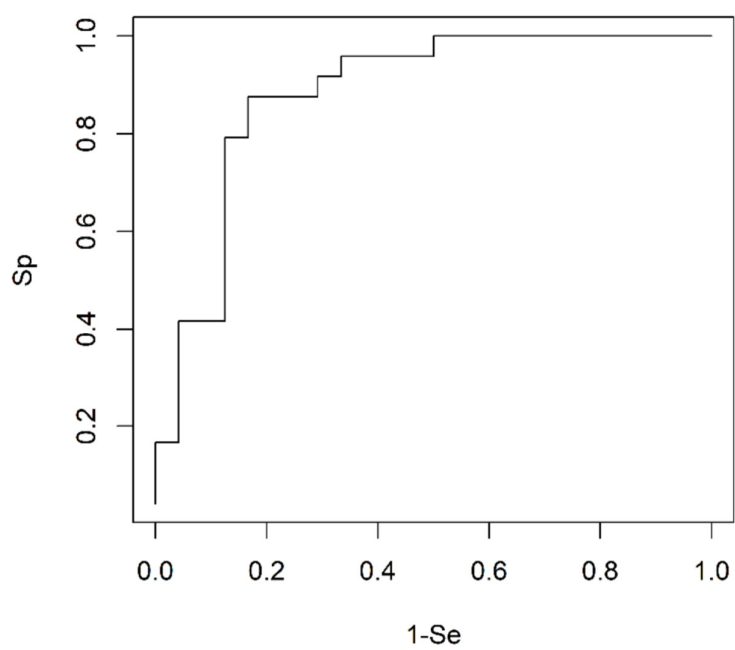
**B**

**Figure S4:** ROC curves of each subset of significant metabolites identified in aqueous (A) and lipid phases (B) for the discrimination of HCC from NTT in F3F4 fibrosis.

**Table S3.** Discriminant metabolites between HCC-F0F1 *versus* HCC-F3F4 according to the number of selections in the aqueous and lipid phases. Upregulated (red) and down-regulated (blue) metabolites in HCC-F0F1.

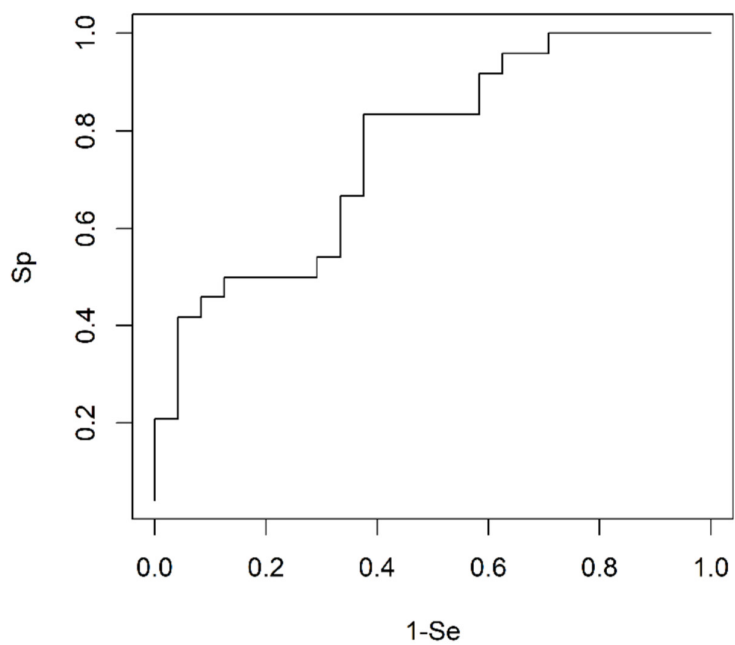
Chemical Shift (ppm)	Metabolites	Selections
<b>Aqueous Phase (threshold: &gt;45 selections)</b>		
3.50		2756
3.74	Glucose	55
5.23		61
3.62	Choline derivatives	2086
<b>Lipid Phase (threshold: &gt;50 selections)</b>		
2.11		61
5.46	MUFA	64
3.05-3.07	Phosphoethanolamine	374
4.12-4.22	TAG	531

### HCC-F0F1 vs HCC-F3F4



A

### HCC-F0F1 vs HCC-F3F4



B

**Figure S5:** ROC curves of each subset of significant metabolites identified in aqueous (A) and lipid phases (B) for the discrimination of HCC-F3F4 from HCC-F0F1.