

Supplementary Material

Bile Acid Profiles in Primary Sclerosing Cholangitis and their Ability to Predict Hepatic Decompensation

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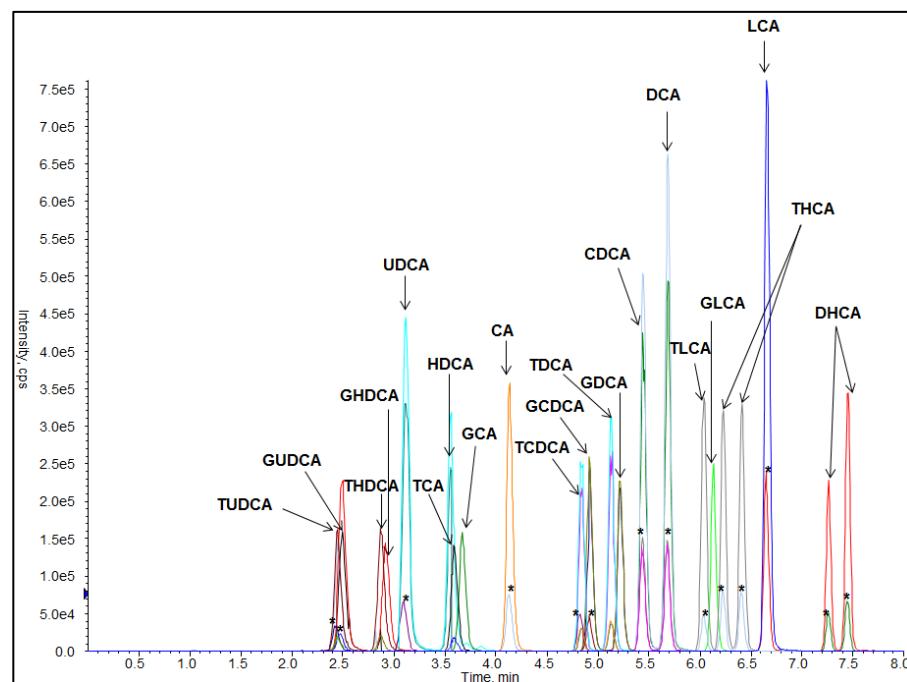
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Supplemental Methods: Bile acid profile method and validation

BA Profiling Method

Samples were prepared by precipitating matrix proteins with an acetonitrile/water (50/50) mixture containing deuterated internal standards at appropriate concentrations. Protein precipitate was removed by filtering through an AcroPrep 0.45 µm GHP filter plate. The filtrate was diluted into water/methanol (80/20) solvent for injection into the LC-MS/MS system (ABSciex API 3200 LC-MS/MS with TurbolonSpray source, LEAP HTC Pal Autosampler, and a Shimadzu LC-20 AB Binary Pump) with an inline Poroshell 120 EC-C18, 2.1x50 mm, 2.7 micron column. Reverse-phase liquid chromatography was initiated with 42% mobile phase A (10 mM Ammonium Acetate in 95% Water/5% Methanol) and 58% mobile phase B (10 mM Ammonium Acetate in 99 % Methanol/ 1% Water) at 0.3 ml/min rate with a binary flow over 13 min. The MS/MS instrument collected 37 scheduled MRMs in negative ion mode with a 60 sec target window and a 0.8 sec scan time. Declustering potential (DP) ranged from -75 volts



to -115 volts, collision energy potential (CEP) ranged from -20 to -30 volts, collision energy (CE) from -20 to -125 volts, and collision cell exit potential (CXP) ranged from -5 to -45 volts across the method. An example chromatogram is shown in **Methods Figure 1**. Analyte concentrations were derived from calculations in Analyst 1.6.2 and peak area ratios with internal labeled standards.

Methods Figure 1. A chromatogram with each measured BA analyte. Cholic Acid (CA), Chenodeoxycholic Acid (CDCA), Deoxycholic Acid (DCA), Ursodeoxycholic Acid (UDCA), Hyodeoxycholic Acid (HDCA), Lithocholic Acid (LCA), Glycocholic Acid (GCA), Glycochenodeoxycholic Acid (GCDCA), Glycodeoxycholic Acid (GDCA), Glycoursodeoxycholic Acid (GUDCA), Glycohyodeoxycholic Acid (GHDCDA), Glycolithocholic Acid (GLCA), Taurocholic Acid (TCA), Taurochenodeoxycholic Acid (TCDCA), Taurodeoxycholic Acid (TDCA), Tauroursodeoxycholic Acid (TUDCA), Taurohyodeoxycholic Acid (THDCA), Taurolithocholic Acid (TLCA), Dihydroxycholestanoic Acid (DHCA), Trihydroxycholestanoic Acid (THCA). The asterisks (*) indicate isotopically labeled internal standards for each analyte with the same retention time. This includes d4-Cholic Acid (CA-d4), d4-Chenodeoxycholic Acid (CDCA-d4), d4-Deoxycholic Acid (DCA-d4), d5-Ursodeoxycholic Acid (UDCA-d5), d4-Lithocholic Acid (LCA-d4), d4-Glycochenodeoxycholic Acid (GCDCA-d4), d5-Glycoursodeoxycholic Acid (GUDCA-d5), d5-Taurochenodeoxycholic Acid (TCDCA-d5),

d5-Tauroursodeoxycholic Acid (TUDCA-d5), d3-Dihydroxycholestanoic Acid (DHCA-d3), and d3-Trihydroxycholestanoic Acid (THCA-d3).

Plasma and Serum Specimen Equivalency

Equivalency between serum and plasma BA analytical performance was shown previously. Reference: Max Scherer, Carsten Gnewuch, Gerd Schmitz, Gerhard Liebisch. Rapid quantification of bile acids and their conjugates in serum by liquid chromatography-tandem mass spectrometry. J Chromatogr B Analyt Technol Biomed Life Sci. 2009 Nov 15;877(3):3920-3925. PMID: 19819765.

Bile Acid Profile Imprecision

Intra-assay imprecision of BA assay was performed with quality control specimens in matrix at low BA concentrations (QC 1) and high concentrations (QC 2) over 20 replicates in a single batch. Inter-assay imprecision was evaluated by preparing and analyzing a low (QC1) and high (QC2) control across 20 separate runs. The calculated coefficient of variation and mean concentrations for each analyte detected by these methods is shown in **Methods Table 1**.

Methods Table 1. BA profile Intra-assay and Inter-assay Imprecision

Analyte	Intra-assay imprecision (N = 20 Specimens)		Inter-assay imprecision (N = 20 Runs)	
	QC1 CV (mean conc., μM)	QC2 CV (mean conc., μM)	QC1 CV (mean conc., μM)	QC2 CV (mean conc., μM)
LCA	4.8% (0.80)	4.2% (7.69)	9.6% (0.81)	4.5% (7.51)
CDCA	5.8% (1.18)	4.2% (7.90)	21.0 % (1.20)	6.4% (7.53)
DCA	5.1% (1.60)	5.4% (8.43)	11.8% (1.59)	8.2% (8.15)
UDCA	5.0% (0.79)	3.6% (7.07)	9.0% (0.79)	8.4% (7.05)
HDCA	18.5% (0.26)	10.1% (2.50)	23.0% (0.29)	25.8% (2.64)
CA	4.9% (1.07)	3.7% (7.41)	7.4% (1.08)	4.5% (7.37)
GLCA	11.1% (0.31)	11.8% (3.15)	22.8% (0.38)	22.6% (3.79)
GCDCA	5.5% (3.16)	4.5% (9.72)	10.3% (3.22)	8.6% (9.56)
GDCA	11.7% (0.88)	9.4% (5.61)	15.6% (0.91)	17.0% (5.76)
GUDCA	7.0% (0.99)	4.6% (7.44)	9.1% (1.00)	6.9% (7.11)
GHDCA	11.8% (0.31)	11.5% (2.76)	29.2% (0.30)	32.7% (2.54)
GCA	6.2% (1.85)	10.5% (9.32)	10.6% (1.78)	14.2% (8.92)
TLCA	18.6% (0.34)	16.8% (3.80)	41.1% (0.45)	52.1% (5.31)
TCDCA	4.2% (1.21)	4.3% (7.64)	7.6% (1.20)	6.6% (7.39)
TDCA	10.2% (0.41)	9.4% (4.68)	29.1% (0.45)	28.0% (4.73)
TUDCA	4.8% (0.72)	4.2% (6.86)	4.2% (0.72)	5.3% (6.65)
THDCA	11.0% (0.56)	8.4% (6.26)	18.2% (0.49)	27.0% (5.51)
TCA	6.2% (0.92)	11.2% (7.68)	14.7% (0.89)	23.5% (7.40)
DHCA	6.9% (0.82)	4.5% (7.66)	22.2% (0.90)	9.3% (7.11)
THCA	9.0% (1.42)	4.6% (8.24)	12.0% (1.47)	4.8% (8.23)

BA Profile Limit of Detection, Biological Limit of Detection and Linearity

The limit of detection (LOD) was determined by the analysis of 12 reagent blanks. The biological limit of detection (BLD) was determined by the analysis (N = 6) of a single serum specimen spiked with BA analyte concentrations near the LOD and measured in triplicate on two separate runs. Linearity was assessed by the analysis of 12-point analyte calibration curves (0.00, 0.05, 0.10, 0.20, 0.50, 1.0, 2.0, 5.0,

10.0, 20.0, 50.0, 100.0 μ M), 5 times, inter-assay. An 11-point calibration curve with each analytical batch was necessary for BA quantitation as a few conjugated bile acids demonstrated correlation coefficients of 0.96-0.98. The LOD and BLD values and linearity results are shown in **Methods Table 2**.

Methods Table 2. Analytical Sensitivity and Linearity

Analyte	LOD (μ M)	BLD (μ M)	Curve 1 Correlation Coefficient	Curve 2 Correlation Coefficient	Curve 3 Correlation Coefficient	Curve 4 Correlation Coefficient	Curve 5 Correlation Coefficient
LCA	0.00	0.01	0.99714	0.99910	0.99944	0.99790	0.99819
CDCA	0.00	0.02	0.99861	0.99961	0.99939	0.99925	0.99866
DCA	0.00	0.02	0.99776	0.99819	0.99788	0.99748	0.99828
UDCA	0.01	0.02	0.99974	0.99977	0.99918	0.99958	0.99954
HDCA	0.00	0.01	0.98951	0.99365	0.99310	0.99533	0.99575
CA	0.00	0.00	0.99793	0.99895	0.99870	0.99770	0.99783
GLCA	0.00	0.02	0.99115	0.99709	0.99449	0.99407	0.99757
GCDCA	0.00	0.01	0.99968	0.99974	0.99948	0.99924	0.99972
GDCA	0.00	0.02	0.98578	0.99065	0.98901	0.98844	0.99669
GUDCA	0.02	0.03	0.99950	0.99927	0.99765	0.99896	0.99715
GHDCA	0.00	0.00	0.98449	0.98087	0.98442	0.98278	0.99136
GCA	0.00	0.02	0.99602	0.99848	0.99804	0.99748	0.99924
TLCA	0.01	0.01	0.98667	0.99129	0.98845	0.98842	0.98399
TCDCA	0.00	0.03	0.99796	0.99775	0.99790	0.99726	0.99822
TDCA	0.00	0.03	0.98190	0.99064	0.96426	0.96745	0.98078
TUDCA	0.01	0.03	0.99955	0.99908	0.99938	0.99847	0.99962
THDCA	0.01	0.02	0.99761	0.99440	0.99238	0.99298	0.99269
TCA	0.00	0.03	0.99771	0.99760	0.99499	0.99735	0.99879
DHCA	0.00	0.01	0.97576	0.98129	0.97869	0.97720	0.97848
THCA	0.00	0.23	0.97297	0.98018	0.98044	0.97546	0.97733

BA Profile Specimen Dilution Limit

As patient specimens with BA concentrations above the analytical measurement range (AMR) of 100 μ M were expected for certain clinical conditions, dilutions limits for each BA analyte were determined by measuring 2 serum specimens spiked to an approximate concentration of 15 μ M for each bile acid. The spiked serum specimens were then diluted 2x, 5x, 10x, 20x, 25x, 40x, and 50x with water, and diluted solutions were carried through the analytical procedure. At a 2-fold dilution, the resulting %difference from expected concentration was near, or less than, 20%. Beyond a 2-fold dilution, various conjugated bile acids resulted in significant overestimation (greater than 25%) of their original concentration. Therefore, it was concluded that only a 2x specimen dilution would yield acceptable BA calculated concentrations, which increased the effective AMR of the assay to 200 μ M for most BAs when completing a specimen dilution.

Use of BA Profile Concentration Values in This Study

As our study is broadly based on comparing BA concentrations between relatively large groups we used the actual reported values for BA concentrations even when they were below the previously determined LOD or above the AMR. Low values approximate a concentration of zero, which could affect results of low-abundance bile acids such as the LCAs and HDCAs. Values above the AMR were only found for GUDCA in 4 of the 400 PSC patients and likely do not significantly impact our findings.

Supplementary Table 1. Bile acids in PSC patients and controls.

	Control	PSC	p-value	Con (n)	PSC (n)
Age in years					
Median (IQR)	56.40 (48.49-65.29)	48.44 (32.99-59.65)	<0.001	302	400
Range	19.28-86.12	10.13-80.48			
Sex (% Male)	58.3	57.8	0.888	302	400
Total BA					
Median (IQR)	1.90 (1.02-3.52)	16.34 (6.04-44.31)	<0.001	302	400
Range	0.19-17.79	0.54-646.47			
CA (total)					
Median (IQR)	0.22 (0.11-0.49)	2.09 (0.73-10.01)	<0.001	302	400
Range	0.01-6.43	0.04-224.36			
CA (unconjugated)					
Median (IQR)	0.04 (0.02-0.12)	0.06 (0.03-0.22)	<0.001	302	400
Range	0.00-5.23	0.00-10.65			
GCA					
Median (IQR)	0.12 (0.05-0.25)	1.60 (0.44-6.12)	<0.001	302	400
Range	0.01-5.84	0.01-165.50			
TCA					
Median (IQR)	0.02 (0.01-0.04)	0.25 (0.06-2.44)	<0.001	302	400
Range	0.00-1.50	0.00-58.61			
CDCA (total)					
Median (IQR)	0.63 (0.33-1.39)	4.09 (1.61-9.81)	<0.001	302	400
Range	0.05-10.77	0.10-268.56			
CDCA (unconjugated)					
Median (IQR)	0.07 (0.03-0.23)	0.15 (0.05-0.46)	<0.001	302	400
Range	0.00-5.28	0.00-14.78			
GCDCA					
Median (IQR)	0.42 (0.21-0.88)	2.87 (1.10-6.90)	<0.001	302	400
Range	0.02-5.38	0.02-214.48			
TCDCA					
Median (IQR)	0.05 (0.03-0.11)	0.47 (0.13-2.12)	<0.001	302	400
Range	0.00-2.49	0.00-52.46			
DCA (total)					
Median (IQR)	0.65 (0.36-1.21)	0.66 (0.03-1.76)	0.328	302	400
Range	0.00-8.26	0.00-24.36			
DCA (unconjugated)					
Median (IQR)	0.33 (0.19-0.58)	0.10 (0.01-0.44)	<0.001	302	400
Range	0.00-4.49	0.00-5.99			

	Control	PSC	p-value	Con (n)	PSC (n)
GDCA					
Median (IQR)	0.21 (0.10-0.46)	0.28 (0.02-0.96)	0.339	302	400
Range	0.00-5.94	0.00-15.49			
TDCA					
Median (IQR)	0.04 (0.02-0.08)	0.06 (0.00-0.27)	0.013	302	400
Range	0.00-1.13	0.00-8.55			
LCA (total)					
Median (IQR)	0.03 (0.02-0.05)	0.04 (0.00-0.15)	0.037	302	400
Range	0.00-0.42	0.00-2.85			
LCA (unconjugated)					
Median (IQR)	0.01 (0.01-0.02)	0.01 (0.00-0.04)	0.512	302	400
Range	0.00-0.21	0.00-0.96			
GLCA					
Median (IQR)	0.01 (0.01-0.02)	0.02 (0.00-0.09)	0.002	302	400
Range	0.00-0.19	0.00-2.14			
TLCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.02)	<0.001	302	400
Range	0.00-0.07	0.00-0.56			
HDCA (total)					
Median (IQR)	0.03 (0.01-0.05)	0.01 (0.00-0.04)	<0.001	302	400
Range	0.00-0.29	0.00-0.73			
HDCA (unconjugated)					
Median (IQR)	0.02 (0.01-0.03)	0.01 (0.00-0.02)	<0.001	302	400
Range	0.00-0.11	0.00-0.24			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.01)	<0.001	302	400
Range	0.00-0.11	0.00-0.73			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.007	302	400
Range	0.00-0.09	0.00-0.07			
UDCA (total)					
Median (IQR)	0.13 (0.07-0.27)	1.95 (0.19-18.27)	<0.001	302	400
Range	0.00-2.15	0.00-537.71			
UDCA (unconjugated)					
Median (IQR)	0.02 (0.01-0.06)	0.17 (0.01-2.02)	<0.001	302	400
Range	0.00-0.92	0.00-52.33			
GUDCA					
Median (IQR)	0.09 (0.05-0.19)	1.53 (0.12-13.52)	<0.001	302	400
Range	0.00-1.78	0.00-472.47			

	Control	PSC	p-value	Con (n)	PSC (n)
TUDCA					
Median (IQR)	0.01 (0.00-0.01)	0.09 (0.01-0.52)	<0.001	302	400
Range	0.00-0.25	0.00-51.06			
Total BA (conj. frac.)					
Median (IQR)	0.64 (0.49-0.79)	0.91 (0.74-0.98)	<0.001	302	400
Range	0.11-0.97	0.13-1.00			
CA (conj. frac.)					
Median (IQR)	0.75 (0.53-0.89)	0.97 (0.85-0.99)	<0.001	302	400
Range	0.21-1.00	0.09-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.87 (0.70-0.94)	0.95 (0.83-0.99)	<0.001	302	400
Range	0.19-1.00	0.12-1.00			
DCA (conj. frac.)					
Median (IQR)	0.44 (0.29-0.60)	0.80 (0.52-0.95)	<0.001	299	345
Range	0.00-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.50 (0.40-0.69)	0.75 (0.56-0.87)	<0.001	274	286
Range	0.00-1.00	0.00-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.20 (0.00-0.50)	1.22 (0.00-0.55)	0.106	261	231
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.83 (0.67-0.94)	0.90 (0.75-0.97)	0.001	301	392
Range	0.00-1.00	0.00-1.00			
Total BA (G:T ratio)					
Median (IQR)	7.66 (4.89-11.17)	8.36 (3.57-17.44)	0.320	300	399
Range	0.91-139.00	0.37-93.11			
CA (G:T ratio)					
Median (IQR)	6.00 (3.50-9.00)	5.56 (2.72-9.23)	0.180	263	393
Range	0.83-31.00	0.27-70.33			
CDCA (G:T ratio)					
Median (IQR)	8.00 (5.33-12.14)	5.76 (2.80-10.30)	<0.001	298	399
Range	1.00-100.00	0.36-92.17			
DCA (G:T ratio)					
Median (IQR)	5.50 (3.74-8.00)	4.90 (2.13-8.26)	0.007	272	292
Range	0.50-61.00	0.22-45.00			
LCA (G:T ratio)					
Median (IQR)	2.00 (1.00-4.25)	3.00 (1.40-6.22)	0.061	79	169
Range	0.50-13.00	0.10-36.00			

	Control	PSC	p-value	Con (n)	PSC (n)
HDCA (G:T ratio)					
Median (IQR)	1.22 (1.00-3.00)	1.17 (1.00-1.88)	0.666	5	12
Range	1.00-4.00	0.71-11.00			
UDCA (G:T ratio)					
Median (IQR)	10.00 (5.00-18.00)	18.40 (7.25-35.75)	<0.001	177	330
Range	1.00-95.00	0.14-145.00			
CA (frac. BA pool)					
Median (IQR)	0.13 (0.08-0.18)	0.18 (0.08-0.36)	<0.001	302	400
Range	0.02-0.55	0.01-0.79			
CDCA (frac. BA pool)					
Median (IQR)	0.35 (0.27-0.45)	0.26 (0.15-0.46)	<0.001	302	400
Range	0.06-0.75	0.04-0.93			
DCA (frac. BA pool)					
Median (IQR)	0.37 (0.24-0.47)	0.04 (0.01-0.14)	<0.001	302	400
Range	0.00-0.78	0.00-0.64			
LCA (frac. BA pool)					
Median (IQR)	0.02 (0.01-0.03)	0.01 (0.00-0.01)	<0.001	302	400
Range	0.00-0.13	0.00-0.12			
HDCA (frac. BA pool)					
Median (IQR)	0.01 (0.01-0.02)	0.00 (0.00-0.01)	<0.001	302	400
Range	0.00-0.14	0.00-0.05			
UDCA (frac. BA pool)					
Median (IQR)	0.07 (0.04-0.12)	0.18 (0.02-0.66)	<0.001	302	400
Range	0.00-0.39	0.00-0.91			
CA:CDCA ratio					
Median (IQR)	0.38 (0.24-0.54)	0.60 (0.35-1.12)	<0.001	302	400
Range	0.06-3.10	0.05-5.26			
CA:DCA ratio					
Median (IQR)	0.36 (0.19-0.73)	2.60 (0.79-26.43)	<0.001	299	345
Range	0.04-103.00	0.09-3796.67			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	3.00 (1.78-5.07)	1.57 (0.22-13.81)	0.001	302	392
Range	0.44-35.00	0.04-1310.00			

Supplementary Table 2. Bile acids in NN PSC patients and controls.

	Control	NN PSC	p-value	Con (n)	NN PSC (n)
Age in years					
Median (IQR)	56.40 (48.49-65.29)	51.14 (32.76-61.38)	<0.001	302	137
Range	19.28-86.12	11.82-78.53			
Sex (% Male)	58.3	59.1	0.868	302	137
Total BA					
Median (IQR)	1.90 (1.02-3.52)	4.98 (2.45-8.50)	<0.001	302	137
Range	0.19-17.79	0.59-16.35			
CA (total)					
Median (IQR)	0.22 (0.11-0.49)	1.00 (0.46-2.12)	<0.001	302	137
Range	0.01-6.43	0.04-10.09			
CA (unconjugated)					
Median (IQR)	0.04 (0.02-0.12)	0.07 (0.03-0.23)	<0.001	302	137
Range	0.00-5.23	0.00-2.99			
GCA					
Median (IQR)	0.12 (0.05-0.25)	0.52 (0.27-1.44)	<0.001	302	137
Range	0.01-5.84	0.02-6.74			
TCA					
Median (IQR)	0.02 (0.01-0.04)	0.09 (0.04-0.34)	<0.001	302	137
Range	0.00-1.50	0.00-3.67			
CDCA (total)					
Median (IQR)	0.63 (0.33-1.39)	2.14 (1.22-4.11)	<0.001	302	137
Range	0.05-10.77	0.16-10.18			
CDCA (unconjugated)					
Median (IQR)	0.07 (0.03-0.23)	0.14 (0.05-0.46)	<0.001	302	137
Range	0.00-5.28	0.00-5.23			
GCDCA					
Median (IQR)	0.42 (0.21-0.88)	1.47 (0.69-2.44)	<0.001	302	137
Range	0.02-5.38	0.10-8.29			
TCDCA					
Median (IQR)	0.05 (0.03-0.11)	0.20 (0.08-0.61)	<0.001	302	137
Range	0.00-2.49	0.00-3.25			
DCA (total)					
Median (IQR)	0.65 (0.36-1.21)	0.39 (0.00-1.26)	<0.001	302	137
Range	0.00-8.26	0.00-7.81			
DCA (unconjugated)					
Median (IQR)	0.33 (0.19-0.58)	0.09 (0.00-0.40)	<0.001	302	137
Range	0.00-4.49	0.00-5.99			

	Control	NN PSC	p-value	Con (n)	NN PSC (n)
GDCA					
Median (IQR)	0.21 (0.10-0.46)	0.13 (0.00-0.56)	0.004	302	137
Range	0.00-5.94	0.00-2.41			
TDCA					
Median (IQR)	0.04 (0.02-0.08)	0.03 (0.00-0.12)	0.067	302	137
Range	0.00-1.13	0.00-1.35			
LCA (total)					
Median (IQR)	0.03 (0.02-0.05)	0.02 (0.00-0.06)	<0.001	302	137
Range	0.00-0.42	0.00-1.18			
LCA (unconjugated)					
Median (IQR)	0.01 (0.01-0.02)	0.00 (0.00-0.02)	<0.001	302	137
Range	0.00-0.21	0.00-0.41			
GLCA					
Median (IQR)	0.01 (0.01-0.02)	0.01 (0.00-0.04)	0.022	302	137
Range	0.00-0.19	0.00-0.71			
TLCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.367	302	137
Range	0.00-0.07	0.00-0.06			
HDCA (total)					
Median (IQR)	0.03 (0.01-0.05)	0.01 (0.00-0.04)	<0.001	302	137
Range	0.00-0.29	0.00-0.41			
HDCA (unconjugated)					
Median (IQR)	0.02 (0.01-0.03)	0.01 (0.00-0.02)	<0.001	302	137
Range	0.00-0.11	0.00-0.24			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.01)	0.002	302	137
Range	0.00-0.11	0.00-0.17			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.158	302	137
Range	0.00-0.09	0.00-0.02			
UDCA (total)					
Median (IQR)	0.13 (0.07-0.27)	0.13 (0.03-0.33)	0.280	302	137
Range	0.00-2.15	0.00-1.98			
UDCA (unconjugated)					
Median (IQR)	0.02 (0.01-0.06)	0.01 (0.00-0.06)	0.106	302	137
Range	0.00-0.92	0.00-0.83			
GUDCA					
Median (IQR)	0.09 (0.05-0.19)	0.09 (0.03-0.22)	0.234	302	137
Range	0.00-1.78	0.00-1.71			

	Control	NN PSC	p-value	Con (n)	NN PSC (n)
TUDCA					
Median (IQR)	0.01 (0.00-0.01)	0.01 (0.00-0.02)	0.742	302	137
Range	0.00-0.25	0.00-0.17			
Total BA (conj. frac.)					
Median (IQR)	0.64 (0.49-0.79)	0.86 (0.62-0.94)	<0.001	302	137
Range	0.11-0.97	0.13-1.00			
CA (conj. frac.)					
Median (IQR)	0.75 (0.53-0.89)	0.89 (0.67-0.97)	<0.001	302	137
Range	0.02-1.00	0.09-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.87 (0.70-0.94)	0.93 (0.76-0.98)	0.003	302	137
Range	0.19-1.00	0.12-1.00			
DCA (conj. frac.)					
Median (IQR)	0.44 (0.29-0.60)	0.66 (0.38-0.83)	<0.001	299	102
Range	0.00-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.50 (0.40-0.69)	0.70 (0.50-0.86)	<0.001	274	80
Range	0.00-1.00	0.00-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.20 (0.00-0.50)	0.26 (0.00-0.50)	0.582	261	76
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.83 (0.67-0.94)	0.88 (0.69-0.97)	0.239	301	129
Range	0.00-1.00	0.00-1.00			
Total BA (G:T ratio)					
Median (IQR)	7.66 (4.89-11.17)	6.60 (3.59-10.24)	0.063	300	136
Range	0.91-139.00	0.58-93.11			
CA (G:T ratio)					
Median (IQR)	6.00 (3.50-9.00)	6.00 (3.60-9.38)	0.851	263	135
Range	0.83-31.00	0.50-70.33			
CDCA (G:T ratio)					
Median (IQR)	8.00 (5.33-12.13)	6.53 (3.34-11.17)	0.012	298	136
Range	1.00-100.00	0.38-92.17			
DCA (G:T ratio)					
Median (IQR)	5.50 (3.74-8.00)	5.09 (2.54-8.00)	0.133	272	83
Range	0.50-61.00	0.30-44.50			
LCA (G:T ratio)					
Median (IQR)	2.00 (1.00-4.25)	2.58 (1.08-6.75)	0.363	79	30
Range	0.50-13.00	0.50-17.00			

	Control	NN PSC	p-value	Con (n)	NN PSC (n)
HDCA (G:T ratio)					
Median (IQR)	1.22 (1.00-3.00)	3.50 (3.25-3.75)	0.232	5	2
Range	1.00-4.00	3.00-4.00			
UDCA (G:T ratio)					
Median (IQR)	10.00 (5.00-18.00)	10.50 (5.00-22.00)	0.833	177	73
Range	1.00-95.00	0.71-124.00			
CA (frac. BA pool)					
Median (IQR)	0.13 (0.08-0.17)	0.25 (0.17-0.38)	<0.001	302	137
Range	0.02-0.55	0.04-0.79			
CDCA (frac. BA pool)					
Median (IQR)	0.35 (0.27-0.45)	0.48 (0.38-0.61)	<0.001	302	137
Range	0.06-0.75	0.14-0.93			
DCA (frac. BA pool)					
Median (IQR)	0.37 (0.24-0.47)	0.11 (0.00-0.26)	<0.001	302	137
Range	0.00-0.78	0.00-0.64			
LCA (frac. BA pool)					
Median (IQR)	0.02 (0.01-0.03)	0.00 (0.00-0.01)	<0.001	302	137
Range	0.00-0.13	0.00-0.12			
HDCA (frac. BA pool)					
Median (IQR)	0.01 (0.01-0.02)	0.00 (0.00-0.01)	<0.001	302	137
Range	0.00-0.14	0.00-0.05			
UDCA (frac. BA pool)					
Median (IQR)	0.07 (0.04-0.12)	0.03 (0.01-0.07)	<0.001	302	137
Range	0.00-0.39	0.00-0.29			
CA:CDCA ratio					
Median (IQR)	0.38 (0.24-0.54)	0.50 (0.31-0.80)	<0.001	302	137
Range	0.06-3.10	0.07-3.77			
CA:DCA ratio					
Median (IQR)	0.36 (0.19-0.73)	1.16 (0.45-5.25)	<0.001	299	102
Range	0.04-103.00	0.09-822.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	3.00 (1.78-5.07)	9.33 (4.00-23.00)	<0.001	302	129
Range	0.44-35.00	0.66-422.00			
Total BA*					
Median (IQR)	1.72 (0.94-3.24)	4.54 (2.37-7.83)	<0.001	302	137
Range	0.19-16.79	0.59-16.10			
Total BA (conj. frac.)*					
Median (IQR)	0.62 (0.47-0.78)	0.85 (0.61-0.94)	<0.001	302	137
Range	0.10-0.97	0.12-1.00			

	Control	NN PSC	p-value	Con (n)	NN PSC (n)
Total BA (G:T ratio)*					
Median (IQR)	7.26 (4.62-10.56)	6.43 (3.48-10.00)	0.105	300	136
Range	0.89-119.00	0.58-90.89			
CA (frac. BA pool)*					
Median (IQR)	0.14 (0.09-0.19)	0.25 (0.18-0.39)	<0.001	302	137
Range	0.03-0.56	0.04-0.79			
CDCA (frac. BA pool)*					
Median (IQR)	0.38 (0.29-0.49)	0.51 (0.40-0.65)	<0.001	302	137
Range	0.06-0.91	0.14-0.93			
DCA (frac. BA pool)*					
Median (IQR)	0.41 (0.27-0.53)	0.12 (0.00-0.29)	<0.001	302	137
Range	0.00-0.81	0.00-0.65			
LCA (frac. BA pool)*					
Median (IQR)	0.02 (0.01-0.03)	0.00 (0.00-0.01)	<0.001	302	137
Range	0.00-0.13	0.00-0.12			
HDCA (frac. BA pool)*					
Median (IQR)	0.01 (0.01-0.03)	0.00 (0.00-0.01)	<0.001	302	137
Range	0.00-0.16	0.00-0.06			

*Values with UDCA and its conjugated forms (GUDCA and TUDCA) removed from the analysis;

NN PSC: Normal total bile acids, No UDCA treatment

Supplementary Table 3. Bile acids in NN and NY PSC patients.

	NN PSC	NY PSC	p-value	NN PSC (n)	NY PSC (n)
Age in years					
Median (IQR)	51.14 (32.76-61.38)	43.92 (33.74-54.55)	0.058	137	61
Range	11.82-78.53	10.13-72.71			
Sex (% Male)	59.1	50.8	0.276	137	61
Total BA					
Median (IQR)	4.98 (2.45-8.50)	9.79 (6.73-14.02)	<0.001	137	61
Range	0.59-16.35	3.06-17.51			
CA (total)					
Median (IQR)	1.00 (0.46-2.12)	0.60 (0.31-1.28)	0.002	137	61
Range	0.04-10.09	0.04-5.47			
CA (unconjugated)					
Median (IQR)	0.07 (0.03-0.23)	0.04 (0.02-0.15)	0.018	137	61
Range	0.00-2.99	0.00-0.90			
GCA					
Median (IQR)	0.52 (0.27-1.44)	0.35 (0.18-0.88)	0.032	137	61
Range	0.02-6.74	0.03-4.77			
TCA					
Median (IQR)	0.09 (0.04-0.34)	0.05 (0.02-0.13)	<0.001	137	61
Range	0.00-3.67	0.00-0.66			
CDCA (total)					
Median (IQR)	2.14 (1.22-4.11)	1.19 (0.73-1.92)	<0.001	137	61
Range	0.16-10.18	0.18-4.58			
CDCA (unconjugated)					
Median (IQR)	0.14 (0.05-0.46)	0.17 (0.07-0.44)	0.688	137	61
Range	0.00-5.23	0.01-1.73			
GCDCA					
Median (IQR)	1.47 (0.69-2.44)	0.85 (0.50-1.24)	<0.001	137	61
Range	0.10-8.29	0.10-3.39			
TCDCA					
Median (IQR)	0.20 (0.08-0.61)	0.07 (0.04-0.18)	<0.001	137	61
Range	0.00-3.25	0.01-0.79			
DCA (total)					
Median (IQR)	0.39 (0.00-1.26)	0.62 (0.09-1.22)	0.353	137	61
Range	0.00-7.81	0.00-3.35			
DCA (unconjugated)					
Median (IQR)	0.09 (0.00-0.40)	0.28 (0.04-0.56)	0.031	137	61
Range	0.00-5.99	0.00-1.32			

	NN PSC	NY PSC	p-value	NN PSC (n)	NY PSC (n)
GDCA					
Median (IQR)	0.13 (0.00-0.56)	0.19 (0.04-0.50)	0.526	137	61
Range	0.00-2.41	0.00-2.10			
TDCA					
Median (IQR)	0.03 (0.00-0.12)	0.02 (0.00-0.08)	0.514	137	61
Range	0.00-1.35	0.00-0.50			
LCA (total)					
Median (IQR)	0.02 (0.00-0.06)	0.09 (0.00-0.19)	<0.001	137	61
Range	0.00-1.18	0.00-0.53			
LCA (unconjugated)					
Median (IQR)	0.00 (0.00-0.02)	0.03 (0.00-0.07)	<0.001	137	61
Range	0.00-0.41	0.00-0.23			
GLCA					
Median (IQR)	0.01 (0.00-0.04)	0.04 (0.00-0.10)	<0.001	137	61
Range	0.00-0.71	0.00-0.28			
TLCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.01)	0.010	137	61
Range	0.00-0.06	0.00-0.07			
HDCA (total)					
Median (IQR)	0.01 (0.00-0.04)	0.02 (0.00-0.05)	0.316	137	61
Range	0.00-0.41	0.00-0.14			
HDCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.02)	0.01 (0.00-0.03)	0.501	137	61
Range	0.00-0.24	0.00-0.07			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.01)	0.292	137	61
Range	0.00-0.17	0.00-0.10			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.098	137	61
Range	0.00-0.02	0.00-0.00			
UDCA (total)					
Median (IQR)	0.13 (0.03-0.33)	6.35 (4.51-9.24)	<0.001	137	61
Range	0.00-1.98	2.15-14.82			
UDCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.06)	1.18 (0.60-2.54)	<0.001	137	61
Range	0.00-0.83	0.07-8.58			
GUDCA					
Median (IQR)	0.09 (0.03-0.22)	4.82 (2.79-6.50)	<0.001	137	61
Range	0.00-1.71	1.36-9.59			

	NN PSC	NY PSC	p-value	NN PSC (n)	NY PSC (n)
TUDCA					
Median (IQR)	0.01 (0.00-0.02)	0.09 (0.06-0.18)	<0.001	137	61
Range	0.00-0.17	0.02-0.92			
Total BA (conj. frac.)					
Median (IQR)	0.86 (0.62-0.94)	0.74 (0.63-0.87)	0.015	137	61
Range	0.13-1.00	0.28-0.99			
CA (conj. frac.)					
Median (IQR)	0.89 (0.67-0.97)	0.90 (0.65-0.98)	0.803	137	61
Range	0.09-1.00	0.27-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.93 (0.76-0.98)	0.82 (0.69-0.94)	0.015	137	61
Range	0.12-1.00	0.37-0.99			
DCA (conj. frac.)					
Median (IQR)	0.66 (0.38-0.83)	0.45 (0.29-0.63)	0.006	102	64
Range	0.00-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.70 (0.50-0.86)	0.58 (0.49-0.75)	0.026	80	44
Range	0.00-1.00	0.25-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.26 (0.00-0.50)	0.22 (0.00-0.57)	0.659	76	36
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.88 (0.69-0.97)	0.77 (0.65-0.88)	0.007	129	61
Range	0.00-1.00	0.27-0.99			
Total BA (G:T ratio)					
Median (IQR)	6.60 (3.59-10.24)	21.59 (12.74-33.80)	<0.001	136	61
Range	0.58-93.11	4.30-85.82			
CA (G:T ratio)					
Median (IQR)	6.00 (3.60-9.38)	8.62 (5.64-12.70)	<0.001	135	57
Range	0.50-70.33	2.59-26.00			
CDCA (G:T ratio)					
Median (IQR)	6.53 (3.34-11.17)	9.17 (6.35-13.78)	0.002	136	61
Range	0.38-92.17	1.58-35.11			
DCA (G:T ratio)					
Median (IQR)	5.09 (2.54-8.00)	8.31 (5.00-10.81)	<0.001	83	44
Range	0.30-44.50	2.00-45.00			
LCA (G:T ratio)					
Median (IQR)	2.58 (1.08-6.75)	5.60 (4.18-8.67)	0.035	30	23
Range	0.50-17.00	0.67-22.00			

	NN PSC	NY PSC	p-value	NN PSC (n)	NY PSC (n)
HDCA (G:T ratio)					
Median (IQR)	3.50 (3.25-3.75)	na	na	2	0
Range	3.00-4.00	na			
UDCA (G:T ratio)					
Median (IQR)	10.50 (5.00-22.00)	46.45 (29.94-64.75)	<0.001	73	61
Range	0.71-124.00	7.09-145.00			
CA (frac. BA pool)					
Median (IQR)	0.25 (0.17-0.38)	0.07 (0.03-0.12)	<0.001	137	61
Range	0.04-0.79	0.00-0.39			
CDCA (frac. BA pool)					
Median (IQR)	0.48 (0.38-0.61)	0.12 (0.10-0.17)	<0.001	137	61
Range	0.14-0.93	0.04-0.40			
DCA (frac. BA pool)					
Median (IQR)	0.11 (0.00-0.26)	0.06 (0.01-0.12)	0.111	137	61
Range	0.00-0.64	0.00-0.26			
LCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.01)	0.01 (0.00-0.02)	0.026	137	61
Range	0.00-0.12	0.00-0.08			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.408	137	61
Range	0.00-0.05	0.00-0.01			
UDCA (frac. BA pool)					
Median (IQR)	0.03 (0.01-0.07)	0.70 (0.59-0.77)	<0.001	137	61
Range	0.00-0.29	0.42-0.89			
CA:CDCA ratio					
Median (IQR)	0.50 (0.31-0.80)	0.51 (0.33-0.75)	0.913	137	61
Range	0.07-3.77	0.05-3.09			
CA:DCA ratio					
Median (IQR)	1.16 (0.45-5.25)	0.73 (0.29-3.27)	0.092	102	54
Range	0.09-822.00	0.09-305.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	9.33 (4.00-23.00)	0.20 (0.14-0.27)	<0.001	129	61
Range	0.66-422.00	0.04-0.94			
Total BA*					
Median (IQR)	4.54 (2.37-7.83)	2.93 (1.87-4.28)	0.001	137	61
Range	0.59-16.10	0.61-8.92			
Total BA (conj. frac.)*					
Median (IQR)	0.85 (0.61-0.94)	0.71 (0.54-0.86)	0.005	137	61
Range	0.12-1.00	0.29-0.99			

	Control	NN PSC	p-value	NN PSC (n)	NY PSC (n)
Total BA (G:T ratio)*					
Median (IQR)	6.43 (3.48-10.00)	9.33 (6.65-13.92)	<0.001	136	61
Range	0.58-90.89	2.54-36.50			
CA (frac. BA pool)*					
Median (IQR)	0.25 (0.18-0.39)	0.24 (0.12-0.32)	0.131	136	61
Range	0.04-0.79	0.02-0.73			
CDCA (frac. BA pool)*					
Median (IQR)	0.51 (0.40-0.65)	0.44 (0.30-0.56)	0.010	136	61
Range	0.14-0.93	0.16-0.86			
DCA (frac. BA pool)*					
Median (IQR)	0.12 (0.00-0.29)	0.24 (0.03-0.41)	0.008	136	61
Range	0.00-0.65	0.00-0.58			
LCA (frac. BA pool)*					
Median (IQR)	0.00 (0.00-0.01)	0.03 (0.00-0.06)	<0.001	136	61
Range	0.00-0.12	0.00-0.28			
HDCA (frac. BA pool)*					
Median (IQR)	0.00 (0.00-0.01)	0.01 (0.00-0.01)	0.098	136	61
Range	0.00-0.06	0.00-0.05			

*Values with UDCA and its conjugated forms (GUDCA and TUDCA) removed from the analysis;

NN PSC: Normal total bile acid, No UDCA treatment; NY PSC: Normal total bile acid, Yes UDCA treatment

Supplementary Table 4. Bile acids in HN and HY PSC patients.

	HN PSC	HY PSC	p-value	HN PSC (n)	HY PSC (n)
Age in years					
Median (IQR)	47.50 (32.21-59.69)	49.13 (34.11-60.88)	0.758	63	115
Range	20.31-79.68	10.74-77.56			
Sex (% Male)	54.0	60.9	0.372	63	115
Total BA					
Median (IQR)	38.73 (23.61-63.57)	48.69 (28.84-116.70)	0.022	63	115
Range	18.53-291.62	17.81-646.47			
CA (total)					
Median (IQR)	18.65 (10.26-32.42)	4.20 (1.89-13.54)	<0.001	63	115
Range	3.65-224.36	0.34-192.53			
CA (unconjugated)					
Median (IQR)	0.07 (0.04-0.17)	0.07 (0.03-0.22)	0.701	63	115
Range	0.01-10.65	0.00-5.72			
GCA					
Median (IQR)	11.67 (5.27-17.57)	3.33 (1.56-10.12)	<0.001	63	115
Range	2.44-165.50	0.19-151.29			
TCA					
Median (IQR)	7.60 (2.74-16.16)	0.53 (0.15-2.94)	<0.001	63	115
Range	0.47-58.61	0.00-41.19			
CDCA (total)					
Median (IQR)	14.21 (9.36-24.92)	6.22 (3.96-16.37)	<0.001	63	115
Range	3.53-71.89	1.05-141.73			
CDCA (unconjugated)					
Median (IQR)	0.07 (0.03-0.21)	0.25 (0.08-0.66)	<0.001	63	115
Range	0.00-14.78	0.01-7.27			
GCDCA					
Median (IQR)	7.80 (5.56-16.44)	4.73 (3.04-13.17)	<0.001	63	115
Range	2.16-39.30	0.85-119.44			
TCDCA					
Median (IQR)	5.22 (3.09-10.93)	0.98 (0.35-3.01)	<0.001	63	115
Range	0.35-42.03	0.04-27.78			
DCA (total)					
Median (IQR)	0.87 (0.18-3.19)	1.15 (0.22-2.71)	0.995	63	115
Range	0.00-24.36	0.00-18.32			
DCA (unconjugated)					
Median (IQR)	0.03 (0.01-0.24)	0.13 (0.01-0.51)	0.006	63	115
Range	0.00-2.63	0.00-2.92			

	HN PSC	HY PSC	p-value	HN PSC (n)	HY PSC (n)
GDCA					
Median (IQR)	0.40 (0.08-1.84)	0.78 (0.14-1.82)	0.458	63	115
Range	0.00-14.96	0.00-15.49			
TDCA					
Median (IQR)	0.34 (0.08-1.08)	0.15 (0.02-0.34)	0.001	63	115
Range	0.00-8.55	0.00-3.41			
LCA (total)					
Median (IQR)	0.04 (0.02-0.12)	0.18 (0.02-0.66)	0.002	63	115
Range	0.00-1.15	0.00-2.85			
LCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.02)	0.05 (0.00-0.15)	<0.001	63	115
Range	0.00-0.67	0.00-0.96			
GLCA					
Median (IQR)	0.02 (0.01-0.06)	0.09 (0.01-0.36)	<0.001	63	115
Range	0.00-0.31	0.00-2.14			
TLCA					
Median (IQR)	0.02 (0.01-0.04)	0.01 (0.00-0.08)	0.951	63	115
Range	0.00-0.17	0.00-0.56			
HDCA (total)					
Median (IQR)	0.01 (0.00-0.03)	0.01 (0.00-0.06)	0.358	63	115
Range	0.00-0.28	0.00-0.73			
HDCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.01)	0.00 (0.00-0.03)	0.328	63	115
Range	0.00-0.11	0.00-0.15			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.01)	0.662	63	115
Range	0.00-0.17	0.00-0.73			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00-0.00	0.100	63	115
Range	0.00-0.07	0.00-0.03			
UDCA (total)					
Median (IQR)	0.45 (0.15-0.91)	31.10 (19.71-63.93)	<0.001	63	115
Range	0.01-2.03	8.87-537.71			
UDCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.03)	3.87 (1.56-6.83)	<0.001	63	115
Range	0.00-0.36	0.24-52.33			
GUDCA					
Median (IQR)	0.29 (0.08-0.73)	27.12 (14.82-52.66)	<0.001	63	115
Range	0.01-1.71	2.72-472.47			

	HN PSC	HY PSC	p-value	HN PSC (n)	HY PSC (n)
TUDCA					
Median (IQR)	0.07 (0.02-0.20)	1.20 (0.49-3.75)	<0.001	63	115
Range	0.00-0.68	0.09-51.06			
Total BA (conj. frac.)					
Median (IQR)	0.99 (0.98-1.00)	0.91 (0.82-0.97)	<0.001	63	115
Range	0.24-1.00	0.37-1.00			
CA (conj. frac.)					
Median (IQR)	1.00 (0.99-1.00)	0.99 (0.91-1.00)	<0.001	63	115
Range	0.24-1.00	0.18-1.00			
CDCA (conj. frac.)					
Median (IQR)	1.00 (0.99-1.00)	0.97 (0.90-0.99)	<0.001	63	115
Range	0.24-1.00	0.39-1.00			
DCA (conj. frac.)					
Median (IQR)	0.97 (0.93-0.99)	0.85 (0.69-0.94)	<0.001	61	107
Range	0.67-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.87 (0.80-1.00)	0.74 (0.63-0.84)	<0.001	55	93
Range	0.33-1.00	0.00-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.41 (0.00-0.67)	0.11 (0.00-0.61)	0.516	37	71
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.98 (0.92-1.00)	0.90 (0.79-0.97)	<0.001	63	115
Range	0.42-1.00	0.28-1.00			
Total BA (G:T ratio)					
Median (IQR)	1.81 (0.99-2.62)	12.52 (7.62-23.15)	<0.001	63	115
Range	0.37-9.61	1.67-67.52			
CA (G:T ratio)					
Median (IQR)	1.69 (0.96-2.80)	6.25 (3.93-10.48)	<0.001	63	114
Range	0.27-9.23	0.62-30.00			
CDCA (G:T ratio)					
Median (IQR)	1.76 (0.99-2.83)	6.28 (4.05-10.21)	<0.001	63	115
Range	0.36-10.69	0.69-22.86			
DCA (G:T ratio)					
Median (IQR)	1.27 (0.71-2.13)	6.02 (3.45-9.35)	<0.001	55	94
Range	0.22-11.33	0.59-22.00			
LCA (G:T ratio)					
Median (IQR)	1.00 (0.68-2.94)	4.25 (2.40-7.80)	<0.001	42	65
Range	0.10-11.00	0.89-36.00			

	HN PSC	HY PSC	p-value	HN PSC (n)	HY PSC (n)
HDCA (G:T ratio)					
Median (IQR)	1.00 (0.81-1.25)	6.25 (3.88-8.62)	0.044	6	2
Range	0.71-1.40	1.50-11.00			
UDCA (G:T ratio)					
Median (IQR)	3.0 (1.76-6.82)	23.01 (13.57-38.67)	<0.001	58	115
Range	0.14-26.67	3.22-102.82			
CA (frac. BA pool)					
Median (IQR)	0.50 (0.35-0.65)	0.10 (0.05-0.17)	<0.001	63	115
Range	0.13-0.79	0.01-0.44			
CDCA (frac. BA pool)					
Median (IQR)	0.40 (0.29-0.49)	0.14 (0.11-0.19)	<0.001	63	115
Range	0.15-0.85	0.04-0.42			
DCA (frac. BA pool)					
Median (IQR)	0.03 (0.00-0.09)	0.02 (0.00-0.06)	0.217	63	115
Range	0.00-0.48	0.00-0.21			
LCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.01)	0.021	63	115
Range	0.00-0.06	0.00-0.04			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.531	63	115
Range	0.00-0.02	0.00-0.01			
UDCA (frac. BA pool)					
Median (IQR)	0.01 (0.00-0.02)	0.68 (0.59-0.77)	<0.001	63	115
Range	0.00-0.10	0.40-0.91			
CA:CDCA ratio					
Median (IQR)	1.17 (0.69-2.10)	0.57 (0.38-1.09)	<0.001	63	115
Range	0.15-5.17	0.06-4.43			
CA:DCA ratio					
Median (IQR)	21.96 (3.34-129.27)	3.37 (1.03-31.84)	<0.001	61	107
Range	0.42-2002.00	0.19-1681.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	31.84 (13.96-109.23)	0.21 (0.15-0.31)	<0.001	63	115
Range	3.62-1310.00	0.004-0.97			
Total BA*					
Median (IQR)	38.02 (22.96-62.69)	14.63 (7.75-36.40)	<0.001	63	115
Range	17.10-290.60	2.41-313.39			
Total BA (conj. frac.)*					
Median (IQR)	0.99 (0.98-1.00)	0.94 (0.86-0.99)	<0.001	63	115
Range	0.24-1.00	0.32-1.00			

	Control	NN PSC	p-value	HN PSC (n)	HY PSC (n)
Total BA (G:T ratio)*					
Median (IQR)	1.79 (0.98-2.59)	6.49 (3.96-10.44)	<0.001	63	115
Range	0.37-9.31	0.67-23.86			
CA (frac. BA pool)*					
Median (IQR)	0.50 (0.35-0.65)	0.31 (0.21-0.45)	<0.001	63	115
Range	0.13-0.81	0.05-0.81			
CDCA (frac. BA pool)*					
Median (IQR)	0.41 (0.30-0.50)	0.47 (0.38-0.64)	0.002	63	115
Range	0.16-0.87	0.18-0.93			
DCA (frac. BA pool)*					
Median (IQR)	0.03 (0.00-0.10)	0.09 (0.01-0.23)	0.026	63	115
Range	0.00-0.50	0.00-0.51			
LCA (frac. BA pool)*					
Median (IQR)	0.00 (0.00-0.00)	0.01 (0.00-0.03)	<0.001	63	115
Range	0.00-0.06	0.00-0.20			
HDCA (frac. BA pool)*					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.017	63	115
Range	0.00-0.02	0.00-0.0			

*Values with UDCA and its conjugated forms (GUDCA and TUDCA) removed from the analysis;

HN PSC: High total bile acid, No UDCA treatment; HY PSC: High total bile acid, Yes UDCA treatment

Supplementary Table 5. Bile acids in NN and HN PSC patients.

	NN PSC	HN PSC	p-value	NN PSC (n)	HN PSC (n)
Age in years					
Median (IQR)	51.14 (32.76-61.38)	47.50 (32.21-59.69)	0.741	137	63
Range	11.82-78.53	20.31-79.68			
Sex (% Male)	59.1	54.0	0.493	137	63
Total BA					
Median (IQR)	4.98 (2.45-8.50)	38.73 (23.61-63.57)	<0.001	137	63
Range	0.59-16.35	18.53-291.62			
CA (total)					
Median (IQR)	1.00 (0.46-2.12)	18.65 (10.26-32.42)	<0.001	137	63
Range	0.04-10.09	3.65-224.36			
CA (unconjugated)					
Median (IQR)	0.07 (0.03-0.23)	0.07 (0.04-0.17)	0.937	137	63
Range	0.00-2.99	0.01-10.65			
GCA					
Median (IQR)	0.52 (0.27-1.44)	11.67 (5.27-17.57)	<0.001	137	63
Range	0.02-6.74	2.44-165.50			
TCA					
Median (IQR)	0.09 (0.04-0.34)	7.60 (2.74-16.16)	<0.001	137	63
Range	0.00-3.67	0.47-58.61			
CDCA (total)					
Median (IQR)	2.14 (1.22-4.11)	14.21 (9.36-24.92)	<0.001	137	63
Range	0.16-10.18	3.53-71.89			
CDCA (unconjugated)					
Median (IQR)	0.14 (0.05-0.46)	0.07 (0.03-0.21)	0.025	137	63
Range	0.00-5.23	0.00-14.78			
GCDCA					
Median (IQR)	1.47 (0.69-2.44)	7.80 (5.56-16.44)	<0.001	137	63
Range	0.10-8.29	2.16-39.30			
TCDCA					
Median (IQR)	0.20 (0.08-0.61)	5.22 (3.09-10.93)	<0.001	137	63
Range	0.00-3.25	0.35-42.03			
DCA (total)					
Median (IQR)	0.39 (0.00-1.26)	0.87 (0.18-3.19)	<0.001	137	63
Range	0.00-7.81	0.00-24.36			
DCA (unconjugated)					
Median (IQR)	0.09 (0.00-0.40)	0.03 (0.01-0.24)	0.237	137	63
Range	0.00-5.99	0.00-2.63			

	NN PSC	HN PSC	p-value	NN PSC (n)	HN PSC (n)
GDCA					
Median (IQR)	0.13 (0.00-0.56)	0.40 (0.08-1.84)	<0.001	137	63
Range	0.00-2.41	0.00-14.96			
TDCA					
Median (IQR)	0.03 (0.00-0.12)	0.34 (0.08-1.08)	<0.001	137	63
Range	0.00-1.35	0.00-8.55			
LCA (total)					
Median (IQR)	0.02 (0.00-0.06)	0.04 (0.02-0.12)	<0.001	137	63
Range	0.00-1.18	0.00-1.15			
LCA (unconjugated)					
Median (IQR)	0.00 (0.00-0.02)	0.01 (0.00-0.02)	0.521	137	63
Range	0.00-0.41	0.00-0.67			
GLCA					
Median (IQR)	0.01 (0.00-0.04)	0.02 (0.01-0.06)	0.005	137	63
Range	0.00-0.71	0.00-0.31			
TLCA					
Median (IQR)	0.00 (0.00-0.00)	0.02 (0.01-0.04)	<0.001	137	63
Range	0.00-0.06	0.00-0.17			
HDCA (total)					
Median (IQR)	0.01 (0.00-0.04)	0.01 (0.00-0.03)	0.996	137	63
Range	0.00-0.41	0.00-0.28			
HDCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.02)	0.01 (0.00-0.01)	0.460	137	63
Range	0.00-0.24	0.00-0.11			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.01)	0.747	137	63
Range	0.00-0.17	0.00-0.17			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.012	137	63
Range	0.00-0.02	0.00-0.07			
UDCA (total)					
Median (IQR)	0.13 (0.03-0.33)	0.45 (0.15-0.91)	<0.001	137	63
Range	0.00-1.98	0.01-2.03			
UDCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.06)	0.01 (0.00-0.03)	0.080	137	63
Range	0.00-0.83	0.00-0.36			
GUDCA					
Median (IQR)	0.09 (0.03-0.22)	0.29 (0.08-0.73)	<0.001	137	63
Range	0.00-1.71	0.01-1.71			

	NN PSC	HN PSC	p-value	NN PSC (n)	HN PSC (n)
TUDCA					
Median (IQR)	0.01 (0.00-0.02)	0.07 (0.02-0.20)	<0.001	137	63
Range	0.00-0.17	0.00-0.68			
Total BA (conj. frac.)					
Median (IQR)	0.86 (0.62-0.94)	0.99 (0.98-1.00)	<0.001	137	63
Range	0.13-1.00	0.24-1.00			
CA (conj. frac.)					
Median (IQR)	0.89 (0.67-0.97)	1.00 (0.99-1.00)	<0.001	137	63
Range	0.09-1.00	0.24-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.93 (0.76-0.98)	1.00 (0.99-1.00)	<0.001	137	63
Range	0.12-1.00	0.24-1.00			
DCA (conj. frac.)					
Median (IQR)	0.66 (0.38-0.83)	0.97 (0.93-0.99)	<0.001	102	61
Range	0.00-1.00	0.67-1.00			
LCA (conj. frac.)					
Median (IQR)	0.70 (0.50-0.86)	0.87 (0.80-1.00)	<0.001	80	55
Range	0.00-1.00	0.33-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.26 (0.00-0.50)	0.41 (0.00-0.67)	0.137	76	37
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.88 (0.69-0.97)	0.98 (0.92-1.00)	<0.001	129	63
Range	0.00-1.00	0.42-1.00			
Total BA (G:T ratio)					
Median (IQR)	6.60 (3.59-10.24)	1.81 (0.99-2.62)	<0.001	136	63
Range	0.58-93.11	0.37-9.61			
CA (G:T ratio)					
Median (IQR)	6.00 (3.60-9.38)	1.69 (0.96-2.80)	<0.001	135	63
Range	0.50-70.33	0.27-9.23			
CDCA (G:T ratio)					
Median (IQR)	6.53 (3.34-11.17)	1.76 (0.99-2.83)	<0.001	136	63
Range	0.38-92.17	0.36-10.69			
DCA (G:T ratio)					
Median (IQR)	5.09 (2.54-8.00)	1.27 (0.71-2.13)	<0.001	83	55
Range	0.30-44.50	0.22-11.33			
LCA (G:T ratio)					
Median (IQR)	2.58 (1.08-6.75)	1.00 (0.68-2.94)	0.004	30	42
Range	0.50-17.00	0.10-11.00			

	NN PSC	HN PSC	p-value	NN PSC (n)	HN PSC (n)
HDCA (G:T ratio)					
Median (IQR)	3.50 (3.25-3.75)	1.00 (0.81-1.25)	0.044	2	6
Range	3.00-4.00	0.71-1.40			
UDCA (G:T ratio)					
Median (IQR)	10.50 (5.00-22.00)	3.00 (1.76-6.82)	<0.001	73	58
Range	0.71-124.00	0.14-26.67			
CA (frac. BA pool)					
Median (IQR)	0.25 (0.17-0.38)	0.50 (0.35-0.64)	<0.001	137	63
Range	0.04-0.79	0.13-0.79			
CDCA (frac. BA pool)					
Median (IQR)	0.48 (0.38-0.61)	0.40 (0.29-0.49)	<0.001	137	63
Range	0.14-0.93	0.15-0.85			
DCA (frac. BA pool)					
Median (IQR)	0.11 (0.00-0.26)	0.03 (0.00-0.09)	0.111	137	63
Range	0.00-0.64	0.00-0.48			
LCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.167	137	63
Range	0.00-0.12	0.00-0.06			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.005	137	63
Range	0.00-0.05	0.00-0.02			
UDCA (frac. BA pool)					
Median (IQR)	0.03 (0.01-0.07)	0.01 (0.00-0.02)	<0.001	137	63
Range	0.00-0.29	0.00-0.10			
CA:CDCA ratio					
Median (IQR)	0.50 (0.31-0.80)	1.17 (0.69-2.10)	<0.001	137	63
Range	0.07-3.77	0.15-5.17			
CA:DCA ratio					
Median (IQR)	1.16 (0.45-5.25)	21.96 (3.34-129.27)	<0.001	102	61
Range	0.09-822.00	0.42-2002.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	9.33 (4.00-23.00)	31.84 (13.96-109.2)	<0.001	129	63
Range	0.66-422.00	3.62-1310.00			

NN PSC: Normal total bile acids, No UDCA treatment; HN PSC: High total bile acids, No UDCA treatment

Supplementary Table 6. Bile acids in NY and HY PSC patients.

	NY PSC	HY PSC	p-value	NY PSC (n)	HY PSC (n)
Age in years					
Median (IQR)	43.92 (33.74-54.55)	49.13 (34.11-60.88)	0.088	61	115
Range	10.13-72.71	10.74-77.56			
Sex (% Male)	50.8	60.9	0.199	61	115
Total BA					
Median (IQR)	9.79 (6.73-14.02)	48.69 (28.84-116.7)	<0.001	61	115
Range	3.06-17.51	17.81-646.47			
CA (total)					
Median (IQR)	0.60 (0.31-1.28)	4.20 (1.89-13.54)	<0.001	61	115
Range	0.04-5.47	0.34-192.53			
CA (unconjugated)					
Median (IQR)	0.04 (0.02-0.15)	0.07 (0.03-0.22)	0.061	61	115
Range	0.00-0.90	0.00-5.72			
GCA					
Median (IQR)	0.35 (0.18-0.88)	3.33 (1.56-10.12)	<0.001	61	115
Range	0.03-4.77	0.19-151.29			
TCA					
Median (IQR)	0.05 (0.02-0.13)	0.53 (0.15-2.94)	<0.001	61	115
Range	0.00-0.66	0.00-41.19			
CDCA (total)					
Median (IQR)	1.19 (0.73-1.92)	6.22 (3.96-16.37)	<0.001	61	115
Range	0.18-4.58	1.05-141.73			
CDCA (unconjugated)					
Median (IQR)	0.17 (0.07-0.44)	0.25 (0.08-0.66)	0.109	61	115
Range	0.01-1.73	0.01-7.27			
GCDCA					
Median (IQR)	0.85 (0.50-1.24)	4.73 (3.04-13.17)	<0.001	61	115
Range	0.10-3.39	0.85-119.44			
TCDCA					
Median (IQR)	0.07 (0.04-0.18)	0.98 (0.35-3.01)	<0.001	61	115
Range	0.01-0.79	0.04-27.78			
DCA (total)					
Median (IQR)	0.62 (0.09-1.22)	1.15 (0.22-2.71)	<0.001	61	115
Range	0.00-3.35	0.00-18.32			
DCA (unconjugated)					
Median (IQR)	0.28 (0.04-0.56)	0.13 (0.01-0.51)	0.331	61	115
Range	0.00-1.32	0.00-2.92			

	NY PSC	HY PSC	p-value	NY PSC (n)	HY PSC (n)
GDCA					
Median (IQR)	0.19 (0.04-0.50)	0.78 (0.14-1.82)	<0.001	61	115
Range	0.00-2.10	0.00-15.49			
TDCA					
Median (IQR)	0.02 (0.00-0.08)	0.15 (0.02-0.34)	<0.001	61	115
Range	0.00-0.50	0.00-3.41			
LCA (total)					
Median (IQR)	0.09 (0.00-0.19)	0.18 (0.02-0.66)	0.005	61	115
Range	0.00-0.53	0.00-2.85			
LCA (unconjugated)					
Median (IQR)	0.03 (0.00-0.07)	0.05 (0.00-0.15)	0.060	61	115
Range	0.00-0.23	0.00-0.96			
GLCA					
Median (IQR)	0.04 (0.00-0.10)	0.09 (0.01-0.36)	0.004	61	115
Range	0.00-0.28	0.00-2.14			
TLCA					
Median (IQR)	0.00 (0.00-0.01)	0.01 (0.00-0.08)	<0.001	61	115
Range	0.00-0.07	0.00-0.56			
HDCA (total)					
Median (IQR)	0.02 (0.00-0.05)	0.01 (0.00-0.06)	0.841	61	115
Range	0.00-0.14	0.00-0.73			
HDCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.03)	0.00 (0.00-0.03)	0.978	61	115
Range	0.00-0.07	0.00-0.15			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.01)	0.237	61	115
Range	0.00-0.10	0.00-0.73			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.036	61	115
Range	0.00-0.00	0.00-0.03			
UDCA (total)					
Median (IQR)	6.35 (4.51-9.24)	31.10 (19.71-63.93)	<0.001	61	115
Range	2.15-14.82	8.87-537.71			
UDCA (unconjugated)					
Median (IQR)	1.18 (0.60-2.54)	3.87 (1.56-6.83)	<0.001	61	115
Range	0.07-8.58	0.24-52.33			
GUDCA					
Median (IQR)	4.84 (2.79-6.50)	27.12 (14.82-52.66)	<0.001	61	115
Range	1.36-9.59	2.72-472.47			

	NY PSC	HY PSC	p-value	NY PSC (n)	HY PSC (n)
TUDCA					
Median (IQR)	0.09 (0.06-0.18)	1.20 (0.49-3.75)	<0.001	61	115
Range	0.02-0.92	0.09-51.06			
Total BA (conj. frac.)					
Median (IQR)	0.74 (0.63-0.87)	0.91 (0.82-0.97)	<0.001	61	115
Range	0.28-0.99	0.37-1.00			
CA (conj. frac.)					
Median (IQR)	0.90 (0.65-0.98)	0.99 (0.91-1.00)	<0.001	61	115
Range	0.27-1.00	0.18-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.82 (0.69-0.94)	0.97 (0.90-0.99)	<0.001	61	115
Range	0.37-0.99	0.39-1.00			
DCA (conj. frac.)					
Median (IQR)	0.45 (0.29-0.63)	0.85 (0.69-0.94)	<0.001	54	107
Range	0.00-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.58 (0.49-0.75)	0.74 (0.63-0.84)	<0.001	44	93
Range	0.25-1.00	0.00-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.22 (0.00-0.57)	0.11 (0.00-0.61)	0.986	36	71
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.77 (0.65-0.88)	0.90 (0.79-0.97)	<0.001	61	115
Range	0.27-0.99	0.28-1.00			
Total BA (G:T ratio)					
Median (IQR)	21.59 (12.74-33.80)	12.52 (7.62-23.15)	<0.001	61	115
Range	4.30-85.82	1.67-67.52			
CA (G:T ratio)					
Median (IQR)	8.62 (5.64-12.70)	6.25 (3.93-10.48)	0.016	57	114
Range	2.59-26.00	0.62-30.00			
CDCA (G:T ratio)					
Median (IQR)	9.17 (6.35-13.78)	6.28 (4.05-10.21)	<0.001	61	115
Range	1.58-35.11	0.69-22.86			
DCA (G:T ratio)					
Median (IQR)	8.31 (5.00-10.81)	6.02 (3.45-9.35)	0.011	44	94
Range	2.00-45.00	0.59-22.00			
LCA (G:T ratio)					
Median (IQR)	5.60 (4.18-8.67)	4.25 (2.40-7.80)	0.277	23	65
Range	0.67-22.00	0.89-36.00			

	NY PSC	HY PSC	p-value	NY PSC (n)	HY PSC (n)
HDCA (G:T ratio)					
Median (IQR)	na	6.25 (3.88-8.62)	na	0	2
Range	na	1.50-11.00			
UDCA (G:T ratio)					
Median (IQR)	46.45 (29.94-64.75)	23.01 (13.57-38.67)	<0.001	61	115
Range	7.09-145.00	3.22-102.82			
CA (frac. BA pool)					
Median (IQR)	0.07 (0.03-0.12)	0.10 (0.05-0.17)	0.034	61	115
Range	0.00-0.39	0.01-0.44			
CDCA (frac. BA pool)					
Median (IQR)	0.12 (0.10-0.17)	0.14 (0.11-0.19)	0.186	61	115
Range	0.04-0.40	0.04-0.42			
DCA (frac. BA pool)					
Median (IQR)	0.06 (0.01-0.12)	0.02 (0.00-0.06)	0.003	61	115
Range	0.00-0.26	0.00-0.21			
LCA (frac. BA pool)					
Median (IQR)	0.01 (0.00-0.02)	0.00 (0.00-0.01)	0.034	61	115
Range	0.00-0.08	0.00-0.04			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.002	61	115
Range	0.00-0.01	0.00-0.01			
UDCA (frac. BA pool)					
Median (IQR)	0.70 (0.59-0.77)	0.68 (0.59-0.77)	0.873	61	115
Range	0.42-0.89	0.40-0.91			
CA:CDCA ratio					
Median (IQR)	0.51 (0.33-0.75)	0.57 (0.38-1.09)	0.068	61	115
Range	0.05-3.09	0.06-4.43			
CA:DCA ratio					
Median (IQR)	0.73 (0.29-3.27)	3.37 (1.03-31.84)	<0.001	54	107
Range	0.09-305.00	0.19-1681.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	0.20 (0.14-0.27)	0.21 (0.15-0.31)	0.260	61	115
Range	0.04-0.94	0.04-0.97			

NY PSC: Normal total bile acids, Yes UDCA treatment; HY PSC: High total bile acids, Yes UDCA treatment

Supplementary Table 7. Bile acids in Control and No IBD PSC patients.

	Control	No IBD PSC	p-value	Con (n)	No IBD PSC (n)
Age in years					
Median (IQR)	56.40 (48.49-65.29)	55.82 (43.32-61.82)	0.032	302	65
Range	19.28-86.12	14.25-79.68			
Sex (% Male)	58.3	49.2	0.182	302	65
Tot BA / UDCA Group					
NN - # (%)	na	19 (29.2)	na	na	65
NY - # (%)	na	10 (15.4)			
HN - # (%)	na	13 (20.0)			
HY - # (%)	na	23 (35.4)			
Total BA					
Median (IQR)	1.90 (1.02-3.52)	23.57 (7.71-45.45)	<0.001	302	65
Range	0.19-17.79	0.88-646.47			
CA (total)					
Median (IQR)	0.22 (0.11-0.49)	1.84 (0.86-8.72)	<0.001	302	65
Range	0.01-6.43	0.13-68.19			
CA (unconjugated)					
Median (IQR)	0.04 (0.02-0.12)	0.05 (0.03-0.12)	0.169	302	65
Range	0.00-5.23	0.00-4.02			
GCA					
Median (IQR)	0.12 (0.05-0.25)	1.11 (0.45-5.13)	<0.001	302	65
Range	0.01-5.84	0.06-53.99			
TCA					
Median (IQR)	0.02 (0.01-0.04)	0.30 (0.07-2.09)	<0.001	302	65
Range	0.00-1.50	0.00-30.29			
CDCA (total)					
Median (IQR)	0.63 (0.33-1.39)	3.85 (2.14-9.80)	<0.001	302	65
Range	0.05-10.77	0.30-141.73			
CDCA (unconjugated)					
Median (IQR)	0.07 (0.03-0.23)	0.16 (0.05-0.47)	<0.001	302	65
Range	0.00-5.28	0.01-7.27			
GCDCA					
Median (IQR)	0.42 (0.21-0.88)	2.99 (1.32-6.33)	<0.001	302	65
Range	0.02-5.38	0.18-119.44			
TCDCA					
Median (IQR)	0.05 (0.03-0.11)	0.56 (0.16-2.77)	<0.001	302	65
Range	0.00-2.49	0.02-42.03			

	Control	No IBD PSC	p-value	Con (n)	No IBD PSC (n)
DCA (total)					
Median (IQR)	0.65 (0.36-1.21)	1.43 (0.76-2.23)	<0.001	302	65
Range	0.00-8.26	0.00-24.36			
DCA (unconjugated)					
Median (IQR)	0.33 (0.19-0.58)	0.29 (0.09-0.65)	0.263	302	65
Range	0.00-4.49	0.00-3.63			
GDCA					
Median (IQR)	0.21 (0.10-0.46)	0.64 (0.35-1.07)	<0.001	302	65
Range	0.00-5.94	0.00-15.49			
TDCA					
Median (IQR)	0.04 (0.02-0.08)	0.15 (0.05-0.31)	<0.001	302	65
Range	0.00-1.13	0.00-8.55			
LCA (total)					
Median (IQR)	0.03 (0.02-0.05)	0.10 (0.06-0.21)	<0.001	302	65
Range	0.00-0.42	0.00-2.85			
LCA (unconjugated)					
Median (IQR)	0.01 (0.01-0.02)	0.03 (0.01-0.07)	<0.001	302	65
Range	0.00-0.21	0.00-0.29			
GLCA					
Median (IQR)	0.01 (0.01-0.02)	0.05 (0.02-0.13)	<0.001	302	65
Range	0.00-0.19	0.00-2.14			
TLCA					
Median (IQR)	0.00 (0.00-0.01)	0.01 (0.00-0.03)	<0.001	302	65
Range	0.00-0.07	0.00-0.42			
HDCA (total)					
Median (IQR)	0.03 (0.01-0.05)	0.02 (0.00-0.05)	0.322	302	65
Range	0.00-0.29	0.00-0.17			
HDCA (unconjugated)					
Median (IQR)	0.02 (0.01-0.03)	0.02 (0.00-0.04)	0.448	302	65
Range	0.00-0.11	0.00-0.13			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.003	302	65
Range	0.00-0.11	0.00-0.06			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	<0.001	302	65
Range	0.00-0.09	0.00-0.07			
UDCA (total)					
Median (IQR)	0.13 (0.07-0.27)	3.33 (0.29-18.37)	<0.001	302	65
Range	0.00-2.15	0.02-537.71			

	Control	No IBD PSC	p-value	Con (n)	No IBD PSC (n)
UDCA (unconjugated)					
Median (IQR)	0.02 (0.01-0.06)	0.12 (0.02-2.29)	<0.001	302	65
Range	0.00-0.92	0.00-52.33			
GUDCA					
Median (IQR)	0.09 (0.05-0.19)	1.73 (0.24-15.78)	<0.001	302	65
Range	0.00-1.78	0.01-472.47)			
TUDCA					
Median (IQR)	0.01 (0.00-0.01)	0.14 (0.03-0.52)	<0.001	302	65
Range	0.00-0.25	0.00-35.47			
Total BA (conj. frac.)					
Median (IQR)	0.64 (0.49-0.79)	0.92 (0.74-0.98)	<0.001	302	65
Range	0.11-0.97	0.20-1.00			
CA (conj. frac.)					
Median (IQR)	0.75 (0.53-0.89)	0.97 (0.83-1.00)	<0.001	302	65
Range	0.02-1.00	0.09-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.87 (0.70-0.94)	0.96 (0.79-0.99)	<0.001	302	65
Range	0.19-1.00	0.29-1.00			
DCA (conj. frac.)					
Median (IQR)	0.44 (0.29-0.60)	0.78 (0.52-0.93)	<0.001	299	63
Range	0.00-1.00	0.18-1.00			
LCA (conj. frac.)					
Median (IQR)	0.50 (0.40-0.69)	0.71 (0.54-0.80)	<0.001	274	62
Range	0.00-1.00	0.00-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.20 (0.00-0.50)	0.00 (0.00-0.41)	0.275	261	48
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.83 (0.67-0.94)	0.90 (0.75-0.97)	0.068	301	65
Range	0.00-1.00	0.28-1.00			
Total BA (G:T ratio)					
Median (IQR)	7.66 (4.89-11.17)	8.09 (2.80-18.65)	0.959	300	65
Range	0.91-139.00	0.62-69.38			
CA (G:T ratio)					
Median (IQR)	6.00 (3.50-9.00)	5.12 (2.33-9.54)	0.171	263	64
Range	0.83-31.00	0.50-27.00			
CDCA (G:T ratio)					
Median (IQR)	8.00 (5.33-12.13)	5.43 (2.37-9.74)	<0.001	298	65
Range	1.00-100.00	0.67-35.50			

	Control	No IBD PSC	p-value	No IBD Con (n)	No IBD PSC (n)
DCA (G:T ratio)					
Median (IQR)	5.50 (3.74-8.00)	5.09 (2.17-8.60)	0.263	272	61
Range	0.50-61.00	0.50-45.00			
LCA (G:T ratio)					
Median (IQR)	2.00 (1.00-4.25)	3.00 (1.50-7.00)	0.062	79	37
Range	0.50-13.00	0.50-36.00			
HDCA (G:T ratio)					
Median (IQR)	1.22 (1.00-3.00)	0.71 (0.71-0.71)	0.137	5	1
Range	1.00-4.00	0.71-0.71			
UDCA (G:T ratio)					
Median (IQR)	10.00 (5.00-18.00)	13.81 (6.47-31.96)	0.073	177	59
Range	1.00-95.00	0.14-101.14			
CA (frac. BA pool)					
Median (IQR)	0.13 (0.08-0.17)	0.12 (0.05-0.28)	0.691	302	65
Range	0.02-0.55	0.01-0.79			
CDCA (frac. BA pool)					
Median (IQR)	0.35 (0.27-0.45)	0.23 (0.13-0.38)	<0.001	302	65
Range	0.06-0.75	0.04-0.79			
DCA (frac. BA pool)					
Median (IQR)	0.37 (0.24-0.47)	0.08 (0.03-0.19)	<0.001	302	65
Range	0.00-0.78	0.00-0.64			
LCA (frac. BA pool)					
Median (IQR)	0.02 (0.01-0.03)	0.01 (0.00-0.01)	<0.001	302	65
Range	0.00-0.13	0.00-0.04			
HDCA (frac. BA pool)					
Median (IQR)	0.01 (0.01-0.02)	0.00 (0.00-0.00)	<0.001	302	65
Range	0.00-0.14	0.00-0.03			
UDCA (frac. BA pool)					
Median (IQR)	0.07 (0.04-0.12)	0.43 (0.02-0.70)	0.024	302	65
Range	0.00-0.39	0.00-0.89			
CA:CDCA ratio					
Median (IQR)	0.38 (0.24-0.54)	0.51 (0.30-0.91)	0.002	302	65
Range	0.06-3.10	0.06-5.17			
CA:DCA ratio					
Median (IQR)	0.36 (0.19-0.73)	1.18 (0.44-5.94)	<0.001	299	63
Range	0.04-103.00	0.12-698.25			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	3.00 (1.78-5.07)	0.48 (0.19-11.13)	0.031	302	65
Range	0.44-35.00	0.05-414.18			

Supplementary Table 8. Bile acids in No IBD and UC PSC patients.

	No IBD PSC	UC PSC	p-value	No IBD PSC (n)	UC PSC (n)
Age in years					
Median (IQR)	55.82 (43.32-61.82)	47.61 (31.69-61.33)	0.044	65	231
Range	14.25-79.68	10.74-77.56			
Sex (% Male)	49.2	61.9	0.066	65	231
Tot BA / UDCA Group					
NN - # (%)	19 (29.2)	83 (35.9)	0.706	65	231
NY - # (%)	10 (15.4)	38 (16.5)			
HN - # (%)	13 (20.0)	37 (16.0)			
HY - # (%)	23 (35.4)	73 (31.6)			
Total BA					
Median (IQR)	23.57 (7.71-45.45)	15.93 (5.90-39.39)	0.206	65	231
Range	0.88-646.47	1.00-530.95			
CA (total)					
Median (IQR)	1.84 (0.86-8.72)	2.21 (0.81-8.82)	0.691	65	231
Range	0.13-68.19	0.04-224.36			
CA (unconjugated)					
Median (IQR)	0.05 (0.03-0.12)	0.07 (0.03-0.24)	0.244	65	231
Range	0.00-4.02	0.00-10.65			
GCA					
Median (IQR)	1.11 (0.45-5.13)	1.74 (0.51-5.36)	0.682	65	231
Range	0.06-53.99	0.03-165.50			
TCA					
Median (IQR)	0.30 (0.07-2.09)	0.25 (0.07-1.67)	0.957	65	231
Range	0.00-30.29	0.00-58.61			
CDCA (total)					
Median (IQR)	3.85 (2.14-9.80)	3.91 (1.52-9.02)	0.426	65	231
Range	0.30-141.73	0.29-120.63			
CDCA (unconjugated)					
Median (IQR)	0.16 (0.05-0.47)	0.14 (0.06-0.38)	0.539	65	231
Range	0.01-7.27	0.00-14.78			
GCDCA					
Median (IQR)	2.99 (1.32-6.33)	2.71 (1.14-6.53)	0.648	65	231
Range	0.18-119.44	0.21-92.80			
TCDCA					
Median (IQR)	0.56 (0.16-2.77)	0.45 (0.13-1.71)	0.360	65	231
Range	0.02-42.03	0.01-35.85			

	No IBD PSC	UC PSC	p-value	No IBD PSC (n)	UC PSC (n)
DCA (total)					
Median (IQR)	1.43 (0.76-2.23)	0.48 (0.01-1.54)	<0.001	65	231
Range	0.00-24.36	0.00-16.23			
DCA (unconjugated)					
Median (IQR)	0.29 (0.09-0.65)	0.06 (0.00-0.41)	<0.001	65	231
Range	0.00-3.63	0.00-5.99			
GDCA					
Median (IQR)	0.64 (0.35-1.07)	0.19 (0.01-0.84)	<0.001	65	231
Range	0.00-15.49	0.00-10.84			
TDCA					
Median (IQR)	0.15 (0.05-0.31)	0.04 (0.00-0.18)	<0.001	65	231
Range	0.00-8.55	0.00-4.02			
LCA (total)					
Median (IQR)	0.10 (0.06-0.21)	0.03 (0.00-0.14)	<0.001	65	231
Range	0.00-2.85	0.00-2.18			
LCA (unconjugated)					
Median (IQR)	0.03 (0.01-0.07)	0.01 (0.00-0.04)	<0.001	65	231
Range	0.00-0.29	0.00-0.88			
GLCA					
Median (IQR)	0.05 (0.02-0.13)	0.01 (0.00-0.08)	<0.001	65	231
Range	0.00-2.14	0.00-1.63			
TLCA					
Median (IQR)	0.01 (0.00-0.03)	0.00 (0.00-0.01)	0.009	65	231
Range	0.00-0.42	0.00-0.43			
HDCA (total)					
Median (IQR)	0.02 (0.00-0.05)	0.01 (0.00-0.04)	0.006	65	231
Range	0.00-0.17	0.00-0.28			
HDCA (unconjugated)					
Median (IQR)	0.02 (0.00-0.04)	0.00 (0.00-0.02)	0.001	65	231
Range	0.00-0.13	0.00-0.14			
GHDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.01)	0.372	65	231
Range	0.00-0.06	0.00-0.17			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.051	65	231
Range	0.00-0.07	0.00-0.05			
UDCA (total)					
Median (IQR)	3.33 (0.29-18.37)	1.82 (0.13-18.98)	0.163	65	231
Range	0.02-537.71	0.00-355.15			

	No IBD PSC	UC PSC	p-value	No IBD PSC (n)	UC PSC (n)
UDCA (unconjugated)					
Median (IQR)	0.12 (0.02-2.29)	0.17 (0.01-2.25)	0.328	65	231
Range	0.00-52.33	0.00-19.26			
GUDCA					
Median (IQR)	1.73 (0.24-15.78)	1.31 (0.09-13.86)	0.200	65	231
Range	0.01-472.47	0.00-297.47			
TUDCA					
Median (IQR)	0.14 (0.03-0.52)	0.08 (0.01-0.53)	0.072	65	231
Range	0.00-35.47	0.00-51.06			
Total BA (conj. frac.)					
Median (IQR)	0.92 (0.74-0.98)	0.90 (0.76-0.98)	0.855	65	231
Range	0.20-1.00	0.24-1.00			
CA (conj. frac.)					
Median (IQR)	0.97 (0.83-1.00)	0.97 (0.85-0.99)	0.652	65	231
Range	0.09-1.00	0.18-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.96 (0.79-0.99)	0.95 (0.85-0.99)	0.961	65	231
Range	0.29-1.00	0.24-1.00			
DCA (conj. frac.)					
Median (IQR)	0.78 (0.52-0.93)	0.81 (0.55-0.96)	0.307	63	187
Range	0.18-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.71 (0.54-0.80)	0.75 (0.57-0.87)	0.089	62	146
Range	0.00-1.00	0.00-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.00 (0.00-0.41)	0.33 (0.00-0.61)	0.020	48	121
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.90 (0.75-0.97)	0.90 (0.77-0.98)	0.790	65	225
Range	0.28-1.00	0.00-1.00			
Total BA (G:T ratio)					
Median (IQR)	8.09 (2.80-18.65)	9.01 (4.08-18.02)	0.578	65	231
Range	0.62-69.38	0.37-67.52			
CA (G:T ratio)					
Median (IQR)	5.12 (2.33-9.54)	6.00 (3.08-9.56)	0.311	64	230
Range	0.50-27.00	0.27-30.00			
CDCA (G:T ratio)					
Median (IQR)	5.43 (2.37-9.74)	6.28 (2.93-10.32)	0.247	65	231
Range	0.67-35.50	0.36-51.00			

	No IBD PSC	UC PSC	p-value	No IBD PSC (n)	UC PSC (n)
DCA (G:T ratio)					
Median (IQR)	5.09 (2.17-8.60)	4.69 (1.96-9.00)	0.648	61	153
Range	0.50-45.00	0.29-44.50			
LCA (G:T ratio)					
Median (IQR)	3.00 (1.50-7.00)	3.00 (1.80-7.00)	0.881	37	91
Range	0.50-36.00	0.10-25.00			
HDCA (G:T ratio)					
Median (IQR)	0.71 (0.71-0.71)	1.40 (1.00-4.00)	0.143	1	5
Range	0.71-0.71	0.75-11.00			
UDCA (G:T ratio)					
Median (IQR)	13.81 (6.47-31.96)	20.23 (7.58-37.57)	0.324	59	185
Range	0.14-101.14	0.33-124.00			
CA (frac. BA pool)					
Median (IQR)	0.12 (0.05-0.28)	0.19 (0.10-0.38)	0.022	65	231
Range	0.01-0.79	0.00-0.79			
CDCA (frac. BA pool)					
Median (IQR)	0.23 (0.13-0.38)	0.28 (0.14-0.47)	0.211	65	231
Range	0.04-0.79	0.04-0.83			
DCA (frac. BA pool)					
Median (IQR)	0.08 (0.03-0.19)	0.02 (0.00-0.12)	<0.001	65	231
Range	0.00-0.64	0.00-0.56			
LCA (frac. BA pool)					
Median (IQR)	0.01 (0.00-0.01)	0.00 (0.00-0.01)	<0.001	65	231
Range	0.00-0.04	0.00-0.08			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.005	65	231
Range	0.00-0.03	0.00-0.04			
UDCA (frac. BA pool)					
Median (IQR)	0.43 (0.02-0.70)	0.18 (0.01-0.67)	0.226	65	231
Range	0.00-0.89	0.00-0.91			
CA:CDCA ratio					
Median (IQR)	0.51 (0.30-0.91)	0.67 (0.40-1.16)	0.012	65	231
Range	0.06-5.17	0.05-4.43			
CA:DCA ratio					
Median (IQR)	1.18 (0.44-5.94)	3.12 (0.90-60.92)	<0.001	63	187
Range	0.12-698.25	0.09-2002.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	0.48 (0.19-11.13)	1.44 (0.21-18.25)	0.317	65	225
Range	0.05-414.18	0.04-1310.00			

Supplementary Table 9. Bile acids in No IBD and CD PSC patients.

	No IBD PSC	CD PSC	p-value	No IBD PSC (n)	CD PSC (n)
Age in years					
Median (IQR)	55.82 (43.32-61.82)	48.17 (38.38-55.93)	0.150	65	26
Range	14.25-79.68	10.13-71.10			
Sex (% Male)	49.2	38.5	0.352	65	26
Tot BA / UDCA Group					
NN - # (%)	19 (29.2)	17 (65.4)	0.003	65	26
NY - # (%)	10 (15.4)	5 (19.2)			
HN - # (%)	13 (20.0)	3 (11.5)			
HY - # (%)	23 (35.4)	1 (3.8)			
Total BA					
Median (IQR)	23.57 (7.71-45.45)	6.79 (3.08-12.15)	<0.001	65	26
Range	0.88-646.47	0.59-124.14			
CA (total)					
Median (IQR)	1.84 (0.86-8.72)	0.98 (0.41-1.54)	0.015	65	26
Range	0.13-68.19	0.04-31.62			
CA (unconjugated)					
Median (IQR)	0.05 (0.03-0.12)	0.12 (0.03-0.28)	0.127	65	26
Range	0.00-4.02	0.01-0.76			
GCA					
Median (IQR)	1.11 (0.45-5.13)	0.54 (0.16-1.24)	0.009	65	26
Range	0.06-53.99	0.02-19.29			
TCA					
Median (IQR)	0.30 (0.07-2.09)	0.07 (0.02-0.22)	0.010	65	26
Range	0.00-30.29	0.00-20.10			
CDCA (total)					
Median (IQR)	3.85 (2.14-9.80)	2.21 (0.91-4.99)	0.021	65	26
Range	0.30-141.73	0.16-24.66			
CDCA (unconjugated)					
Median (IQR)	0.16 (0.05-0.47)	0.31 (0.08-0.63)	0.318	65	26
Range	0.01-7.27	0.00-5.23			
GCDCA					
Median (IQR)	2.99 (1.32-6.33)	1.35 (0.37-3.32)	0.007	65	26
Range	0.18-119.44	0.10-17.57			
TCDCA					
Median (IQR)	0.56 (0.16-2.77)	0.12 (0.04-0.69)	0.008	65	26
Range	0.02-42.03	0.00-15.94			

	No IBD PSC	CD PSC	p-value	No IBD PSC (n)	CD PSC (n)
DCA (total)					
Median (IQR)	1.43 (0.76-2.23)	0.81 (0.15-2.20)	0.063	65	26
Range	0.00-24.36	0.00-11.70			
DCA (unconjugated)					
Median (IQR)	0.29 (0.09-0.65)	0.37 (0.03-0.66)	0.598	65	26
Range	0.00-3.63	0.00-3.67			
GDCA					
Median (IQR)	0.64 (0.35-1.07)	0.17 (0.05-0.85)	0.009	65	26
Range	0.00-15.49	0.00-7.51			
TDCA					
Median (IQR)	0.15 (0.05-0.31)	0.02 (0.00-0.25)	0.014	65	26
Range	0.00-8.55	0.00-3.41			
LCA (total)					
Median (IQR)	0.10 (0.06-0.21)	0.03 (0.01-0.09)	0.001	65	26
Range	0.00-2.85	0.00-1.44			
LCA (unconjugated)					
Median (IQR)	0.03 (0.01-0.07)	0.01 (0.00-0.02)	0.001	65	26
Range	0.00-0.29	0.00-0.41			
GLCA					
Median (IQR)	0.05 (0.02-0.13)	0.01 (0.01-0.06)	0.005	65	26
Range	0.00-2.14	0.00-0.71			
TLCA					
Median (IQR)	0.01 (0.00-0.03)	0.00 (0.00-0.01)	0.017	65	26
Range	0.00-0.42	0.00-0.56			
HDCA (total)					
Median (IQR)	0.02 (0.00-0.05)	0.01 (0.00-0.05)	0.701	65	26
Range	0.00-0.17	0.00-0.41			
HDCA (unconjugated)					
Median (IQR)	0.02 (0.00-0.04)	0.01 (0.00-0.03)	0.719	65	26
Range	0.00-0.13	0.00-0.24			
GHDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.01)	0.311	65	26
Range	0.00-0.06	0.00-0.17			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.291	65	26
Range	0.00-0.07	0.00-0.01			
UDCA (total)					
Median (IQR)	3.33 (0.29-18.37)	0.49 (0.12-1.15)	0.005	65	26
Range	0.02-537.71	0.00-60.88			

	No IBD PSC	CD PSC	p-value	No IBD PSC (n)	CD PSC (n)
UDCA (unconjugated)					
Median (IQR)	0.12 (0.02-2.29)	0.06 (0.01-0.40)	0.135	65	26
Range	0.00-52.33	0.00-6.51			
GUDCA					
Median (IQR)	1.73 (0.24-15.78)	0.24 (0.06-0.73)	0.003	65	26
Range	0.01-472.47	0.00-54.41			
TUDCA					
Median (IQR)	0.14 (0.03-0.52)	0.02 (0.00-0.05)	<0.001	65	26
Range	0.00-35.47	0.00-4.77			
Total BA (conj. frac.)					
Median (IQR)	0.92 (0.74-0.98)	0.75 (0.47-0.91)	0.014	65	26
Range	0.20-1.00	0.28-1.00			
CA (conj. frac.)					
Median (IQR)	0.97 (0.83-1.00)	0.85 (0.55-0.96)	0.006	65	26
Range	0.09-1.00	0.25-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.96 (0.79-0.99)	0.87 (0.59-0.93)	0.015	65	26
Range	0.29-1.00	0.24-1.00			
DCA (conj. frac.)					
Median (IQR)	0.78 (0.52-0.93)	0.58 (0.23-0.80)	0.042	63	24
Range	0.18-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.71 (0.54-0.80)	0.75 (0.50-0.89)	0.386	62	21
Range	0.00-1.00	0.33-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.00 (0.00-0.41)	0.00 (0.00-0.41)	0.737	48	17
Range	0.00-1.00	0.00-0.67			
UDCA (conj. frac.)					
Median (IQR)	0.90 (0.75-0.97)	0.81 (0.60-0.92)	0.032	65	24
Range	0.28-1.00	0.27-1.00			
Total BA (G:T ratio)					
Median (IQR)	8.09 (2.80-18.65)	8.19 (4.70-20.57)	0.818	65	25
Range	0.62-69.38	0.46-85.82			
CA (G:T ratio)					
Median (IQR)	5.12 (2.33-9.54)	6.29 (3.07-10.25)	0.453	64	23
Range	0.50-27.00	0.57-22.80			
CDCA (G:T ratio)					
Median (IQR)	5.43 (2.37-9.74)	7.85 (3.81-13.20)	0.126	65	25
Range	0.67-35.50	0.36-67.00			

	No IBD PSC	CD PSC	p-value	No IBD PSC (n)	CD PSC (n)
DCA (G:T ratio)					
Median (IQR)	5.09 (2.17-8.60)	6.86 (5.09-8.60)	0.456	61	19
Range	0.50-45.00	0.22-23.50			
LCA (G:T ratio)					
Median (IQR)	3.00 (1.50-7.00)	4.25 (0.79-8.21)	0.733	37	8
Range	0.50-36.00	0.20-13.00			
HDCA (G:T ratio)					
Median (IQR)	0.71 (0.71-0.71)	3.00 (3.00-3.00)	0.317	1	1
Range	0.71-0.71	3.00-3.00			
UDCA (G:T ratio)					
Median (IQR)	13.81 (6.47-31.96)	21.00 (11.41-51.25)	0.310	59	17
Range	0.14-101.14	0.42-145.00			
CA (frac. BA pool)					
Median (IQR)	0.12 (0.05-0.28)	0.17 (0.12-0.27)	0.404	65	26
Range	0.01-0.79	0.01-0.56			
CDCA (frac. BA pool)					
Median (IQR)	0.23 (0.13-0.38)	0.39 (0.25-0.56)	0.004	65	26
Range	0.04-0.79	0.08-0.85			
DCA (frac. BA pool)					
Median (IQR)	0.08 (0.03-0.19)	0.12 (0.04-0.22)	0.404	65	26
Range	0.00-0.64	0.00-0.61			
LCA (frac. BA pool)					
Median (IQR)	0.01 (0.00-0.01)	0.01 (0.00-0.01)	0.626	65	26
Range	0.00-0.04	0.00-0.12			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.01)	0.463	65	26
Range	0.00-0.03	0.00-0.05			
UDCA (frac. BA pool)					
Median (IQR)	0.43 (0.02-0.70)	0.05 (0.04-0.20)	0.150	65	26
Range	0.00-0.89	0.00-0.83			
CA:CDCA ratio					
Median (IQR)	0.51 (0.30-0.91)	0.38 (0.26-0.71)	0.366	65	26
Range	0.06-5.17	0.15-1.45			
CA:DCA ratio					
Median (IQR)	1.18 (0.44-5.94)	1.03 (0.40-5.40)	0.812	63	24
Range	0.12-698.25	0.09-182.50			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	0.48 (0.19-11.13)	4.27 (0.87-11.00)	0.189	65	24
Range	0.05-414.18	0.09-50.65			

Supplementary Table 10. Bile acids in No IBD and Indeterminate IBD PSC patients.

	No IBD PSC	Ind IBD PSC	p-value	No IBD PSC (n)	Ind IBD PSC (n)
Age in years					
Median (IQR)	55.82 (43.32-61.82)	47.03 (29.80-59.11)	0.187	65	25
Range	14.25-79.68	18.90-78.53			
Sex (% Male)	49.2	52.0	0.814	65	25
Tot BA / UDCA Group					
NN - # (%)	19 (29.2)	11 (44.0)	0.465	65	25
NY - # (%)	10 (15.4)	4 (16.0)			
HN - # (%)	13 (20.0)	5 (20.0)			
HY - # (%)	23 (35.4)	5 (20.0)			
Total BA					
Median (IQR)	23.57 (7.71-45.45)	12.06 (6.73-38.40)	0.184	65	25
Range	0.88-646.47	0.96-53.68			
CA (total)					
Median (IQR)	1.84 (0.86-8.72)	1.85 (0.60-7.10)	0.564	65	25
Range	0.13-68.19	0.16-25.90			
CA (unconjugated)					
Median (IQR)	0.05 (0.03-0.12)	0.04 (0.03-0.14)	0.910	65	25
Range	0.00-4.02	0.01-2.65			
GCA					
Median (IQR)	1.11 (0.45-5.13)	1.26 (0.33-4.17)	0.525	65	25
Range	0.06-53.99	0.07-12.90			
TCA					
Median (IQR)	0.30 (0.07-2.09)	0.15 (0.03-2.10)	0.452	65	25
Range	0.00-30.29	0.01-12.74			
CDCA (total)					
Median (IQR)	3.85 (2.14-9.80)	4.68 (1.48-8.18)	0.675	65	25
Range	0.30-141.73	0.70-24.40			
CDCA (unconjugated)					
Median (IQR)	0.16 (0.05-0.47)	0.14 (0.04-0.56)	0.708	65	25
Range	0.01-7.27	0.01-3.68			
GCDCA					
Median (IQR)	2.99 (1.32-6.33)	3.14 (1.00-5.53)	0.656	65	25
Range	0.18-119.44	0.36-16.22			
TCDCA					
Median (IQR)	0.56 (0.16-2.77)	0.42 (0.13-2.06)	0.441	65	25
Range	0.02-42.03	0.03-8.40			

	No IBD PSC	Ind IBD PSC	p-value	No IBD PSC (n)	Ind IBD PSC (n)
DCA (total)					
Median (IQR)	1.43 (0.76-2.23)	0.61 (0.24-2.34)	0.073	65	25
Range	0.00-24.36	0.00-6.13			
DCA (unconjugated)					
Median (IQR)	0.29 (0.09-0.65)	0.12 (0.02-0.39)	0.054	65	25
Range	0.00-3.63	0.00-3.06			
GDCA					
Median (IQR)	0.64 (0.35-1.07)	0.33 (0.08-1.27)	0.091	65	25
Range	0.00-15.49	0.00 – 4.71			
TDCA					
Median (IQR)	0.15 (0.05-0.31)	0.08 (0.01-0.32)	0.239	65	25
Range	0.00-8.55	0.00-1.35			
LCA (total)					
Median (IQR)	0.10 (0.06-0.21)	0.04 (0.01-0.11)	0.005	65	25
Range	0.00-2.85	0.00-1.13			
LCA (unconjugated)					
Median (IQR)	0.03 (0.01-0.07)	0.01 (0.00-0.03)	0.002	65	25
Range	0.00-0.29	0.00-0.96			
GLCA					
Median (IQR)	0.05 (0.02-0.13)	0.01 (0.00-0.05)	0.008	65	25
Range	0.00-2.14	0.00-0.30			
TLCA					
Median (IQR)	0.01 (0.00-0.03)	0.00 (0.00-0.03)	0.237	65	25
Range	0.00-0.42	0.00-0.11			
HDCA (total)					
Median (IQR)	0.02 (0.00-0.05)	0.01 (0.00-0.04)	0.224	65	25
Range	0.00-0.17	0.00-0.15			
HDCA (unconjugated)					
Median (IQR)	0.02 (0.00-0.04)	0.01 (0.00-0.03)	0.178	65	25
Range	0.00-0.13	0.00-0.15			
GHDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.850	65	25
Range	0.00-0.06	0.00-0.06			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.723	65	25
Range	0.00-0.07	0.00-0.02			
UDCA (total)					
Median (IQR)	3.33 (0.29-18.37)	0.67 (0.21-7.33)	0.254	65	25
Range	0.02-537.71	0.01-44.31			

	No IBD PSC	Ind IBD PSC	p-value	No IBD PSC (n)	Ind IBD PSC (n)
UDCA (unconjugated)					
Median (IQR)	0.12 (0.02-2.29)	0.06 (0.01-0.76)	0.238	65	25
Range	0.00-52.33	0.00-5.65			
GUDCA					
Median (IQR)	1.73 (0.24-15.78)	0.49 (0.20-4.89)	0.266	65	25
Range	0.01-472.47	0.01-39.65			
TUDCA					
Median (IQR)	0.14 (0.03-0.52)	0.07 (0.02-0.17)	0.161	65	25
Range	0.00-35.47	0.00-2.63			
Total BA (conj. frac.)					
Median (IQR)	0.92 (0.74-0.98)	0.94 (0.78-0.97)	0.811	65	25
Range	0.20-1.00	0.21-1.00			
CA (conj. frac.)					
Median (IQR)	0.97 (0.83-1.00)	0.95 (0.88-0.99)	0.488	65	25
Range	0.09-1.00	0.15-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.96 (0.79-0.99)	0.97 (0.84-0.99)	0.853	65	25
Range	0.29-1.00	0.24-1.00			
DCA (conj. frac.)					
Median (IQR)	0.78 (0.52-0.93)	0.83 (0.59-0.94)	0.565	63	23
Range	0.18-1.00	0.20-1.00			
LCA (conj. frac.)					
Median (IQR)	0.71 (0.54-0.80)	0.75 (0.58-0.96)	0.151	62	19
Range	0.00-1.00	0.15-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.00 (0.00-0.41)	0.00 (0.00-0.53)	0.558	48	15
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.90 (0.75-0.97)	0.92 (0.78-0.98)	0.766	65	25
Range	0.28-1.00	0.44-1.00			
Total BA (G:T ratio)					
Median (IQR)	8.09 (2.80-18.65)	8.47 (2.41-22.54)	0.989	65	25
Range	0.62-69.38	0.62-93.11			
CA (G:T ratio)					
Median (IQR)	5.12 (2.33-9.54)	5.19 (2.87-9.27)	0.671	64	25
Range	0.50-27.00	0.66-70.33			
CDCA (G:T ratio)					
Median (IQR)	5.43 (2.37-9.74)	4.54 (2.33-12.08)	0.715	65	25
Range	0.67-35.50	0.56-92.17			

	No IBD PSC	Ind IBD PSC	p-value	No IBD PSC (n)	Ind IBD PSC (n)
DCA (G:T ratio)					
Median (IQR)	5.09 (2.17-8.60)	4.05 (2.00-7.74)	0.462	61	18
Range	0.50-45.00	0.59-14.00			
LCA (G:T ratio)					
Median (IQR)	3.00 (1.50-7.00)	1.45 (1.00-5.00)	0.131	37	9
Range	0.50-36.00	0.33-8.00			
HDCA (G:T ratio)					
Median (IQR)	0.71 (0.71-0.71)	1.50 (1.50-1.50)	0.317	1	1
Range	0.71-0.71	1.50-1.50			
UDCA (G:T ratio)					
Median (IQR)	13.81 (6.47-31.96)	15.37 (4.33-29.00)	0.840	59	21
Range	0.14-101.14	2.41-87.17			
CA (frac. BA pool)					
Median (IQR)	0.12 (0.05-0.28)	0.23 (0.10-0.33)	0.240	65	25
Range	0.01-0.79	0.02-0.67			
CDCA (frac. BA pool)					
Median (IQR)	0.23 (0.13-0.38)	0.43 (0.18-0.57)	0.023	65	25
Range	0.04-0.79	0.08-0.74			
DCA (frac. BA pool)					
Median (IQR)	0.08 (0.03-0.19)	0.07 (0.01-0.12)	0.240	65	25
Range	0.00-0.64	0.00-0.55			
LCA (frac. BA pool)					
Median (IQR)	0.01 (0.00-0.01)	0.00 (0.00-0.01)	0.010	65	25
Range	0.00-0.04	0.00-0.04			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.317	65	25
Range	0.00-0.03	0.00-0.01			
UDCA (frac. BA pool)					
Median (IQR)	0.43 (0.02-0.70)	0.06 (0.02-0.59)	0.479	65	25
Range	0.00-0.89	0.01-0.89			
CA:CDCA ratio					
Median (IQR)	0.51 (0.30-0.91)	0.53 (0.34-0.72)	0.960	65	25
Range	0.06-5.17	0.11-2.14			
CA:DCA ratio					
Median (IQR)	1.18 (0.44-5.94)	2.71 (0.89-16.38)	0.139	63	23
Range	0.12-698.25	0.18-1295.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	0.48 (0.19-11.13)	7.78 (0.27-18.80)	0.075	65	25
Range	0.05-414.18	0.09-75.00			

Supplementary Table 11. Bile acids in cirrhotic and non-cirrhotic PSC patients.

	Non-cirrhotic	Cirrhotic	p-value	NC (n)	Cir (n)
Age in years					
Median (IQR)	47.99 (32.95-59.36)	55.37 (33.51-61.41)	0.143	340	60
Range	10.13-80.48	17.27-79.68			
Sex (% Male)	58.5	53.3	0.453	340	60
Total BA					
Median (IQR)	13.50 (5.38-32.27)	48.84 (22.67-119.16)	<0.001	340	60
Range	0.54-291.62	1.86-646.47			
CA (total)					
Median (IQR)	1.80 (0.64-6.38)	13.34 (3.04-32.12)	<0.001	340	60
Range	0.04-224.36	0.46-192.53			
CA (unconjugated)					
Median (IQR)	0.06 (0.03-0.20)	0.08 (0.03-0.26)	0.766	340	60
Range	0.00-10.65	0.01-9.85			
GCA					
Median (IQR)	1.33 (0.41-4.53)	8.07 (1.96-18.62)	<0.001	340	60
Range	0.01-165.50	0.18-151.29			
TCA					
Median (IQR)	0.18 (0.05-1.23)	4.19 (0.49-11.65)	<0.001	340	60
Range	0.00-58.61	0.01-41.19			
CDCA (total)					
Median (IQR)	3.53 (1.37-80.3)	14.80 (4.69-42.63)	<0.001	340	60
Range	0.10-71.89	0.73-268.56			
CDCA (unconjugated)					
Median (IQR)	0.16 (0.05-0.46)	0.11 (0.05-0.46)	0.823	340	60
Range	0.00-14.78	0.01-9.35			
GCDCA					
Median (IQR)	2.39 (0.97-5.39)	7.88 (2.95-23.59)	<0.001	340	60
Range	0.02-47.74	0.51-214.48			
TCDCDA					
Median (IQR)	0.36 (0.11-1.32)	5.07 (1.11-11.79)	<0.001	340	60
Range	0.00-35.85	0.01-52.46			
DCA (total)					
Median (IQR)	0.66 (0.02-1.78)	0.55 (0.17-1.57)	0.540	340	60
Range	0.00-24.36	0.00-9.94			
DCA (unconjugated)					
Median (IQR)	0.13 (0.01-0.46)	0.03 (0.01-0.19)	0.032	340	60
Range	0.00-5.99	0.00-3.63			

	Non-cirrhotic	Cirrhotic	p-value	NC (n)	Cir (n)
GDCA					
Median (IQR)	0.27 (0.01-1.03)	0.31 (0.08-0.83)	0.483	340	60
Range	0.00-15.49	0.00-8.20			
TDCA					
Median (IQR)	0.05 (0.00-0.25)	0.17 (0.04-0.32)	0.007	340	60
Range	0.00-8.55	0.00-3.44			
LCA (total)					
Median (IQR)	0.04 (0.00-0.13)	0.06 (0.02-0.26)	0.006	340	60
Range	0.00-2.18	0.00-2.85			
LCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.04)	0.01 (0.00-0.07)	0.151	340	60
Range	0.00-0.96	0.00-0.70			
GLCA					
Median (IQR)	0.02 (0.00-0.08)	0.02 (0.01-0.14)	0.070	340	60
Range	0.00-1.63	0.00-2.14			
TLCA					
Median (IQR)	0.00 (0.00-0.02)	0.01 (0.00-0.06)	<0.001	340	60
Range	0.00-0.56	0.00-0.42			
HDCA (total)					
Median (IQR)	0.01 (0.00-0.04)	0.01 (0.00-0.03)	0.310	340	60
Range	0.00-0.41	0.00-0.73			
HDCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.03)	0.00 (0.00-0.01)	0.125	340	60
Range	0.00-0.24	0.00-0.13			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.204	340	60
Range	0.00-0.17	0.00-0.73			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.018	340	60
Range	0.00-0.07	0.00-0.01			
UDCA (total)					
Median (IQR)	2.00 (0.19-14.29)	1.09 (0.22-33.79)	0.207	340	60
Range	0.00-194.12	0.00-537.71			
UDCA (unconjugated)					
Median (IQR)	0.17 (0.01-2.00)	0.14 (0.01-2.33)	0.784	340	60
Range	0.00-26.13	0.00-52.33			
GUDCA					
Median (IQR)	1.62 (0.12-12.06)	0.88 (0.09-30.72)	0.239	340	60
Range	0.00-175.83	0.00-472.47			

	Non-cirrhotic	Cirrhotic	p-value	NC (n)	Cir (n)
TUDCA					
Median (IQR)	0.08 (0.01-0.47)	0.17 (0.04-2.63)	<0.001	340	60
Range	0.00-18.60	0.00-51.06			
Total BA (conj. frac.)					
Median (IQR)	0.89 (0.72-0.97)	0.98 (0.88-0.99)	<0.001	340	60
Range	0.13-1.00	0.20-1.00			
CA (conj. frac.)					
Median (IQR)	0.96 (0.82-0.99)	0.99 (0.96-1.00)	<0.001	340	60
Range	0.09-1.00	0.09-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.94 (0.82-0.99)	0.99 (0.93-1.00)	<0.001	340	60
Range	0.09-1.00	0.24-1.00			
DCA (conj. frac.)					
Median (IQR)	0.77 (0.50-0.93)	0.95 (0.80-0.99)	<0.001	286	59
Range	0.00-1.00	0.18-1.00			
LCA (conj. frac.)					
Median (IQR)	0.75 (0.55-0.86)	0.79 (0.57-1.00)	0.081	231	55
Range	0.00-1.00	0.00-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.20 (0.00-0.52)	0.33 (0.00-0.67)	0.314	199	32
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.88 (0.74-0.97)	0.97 (0.86-0.99)	0.002	333	59
Range	0.00-1.00	0.46-1.00			
Total BA (G:T ratio)					
Median (IQR)	9.04 (4.44-18.55)	3.80 (1.77-9.92)	<0.001	339	60
Range	0.47-93.11	0.37-55.50			
CA (G:T ratio)					
Median (IQR)	6.00 (3.38-9.67)	2.44 (1.42-5.80)	<0.001	333	60
Range	0.47-70.33	0.27-22.96			
CDCA (G:T ratio)					
Median (IQR)	6.46 (3.27-10.71)	2.52 (1.74-5.60)	<0.001	339	60
Range	0.38-92.17	0.36-51.00			
DCA (G:T ratio)					
Median (IQR)	5.47 (2.61-8.75)	2.17 (1.29-3.98)	<0.001	241	51
Range	0.29-45.00	0.22-38.50			
LCA (G:T ratio)					
Median (IQR)	3.82 (2.00-7.00)	1.45 (1.00-2.85)	<0.001	132	37
Range	0.50-36.00	0.10-12.00			

	Non-cirrhotic	Cirrhotic	p-value	NC (n)	Cir (n)
HDCA (G:T ratio)					
Median (IQR)	1.37 (1.00-2.62)	1.00 (1.00-1.00)	0.382	10	2
Range	0.71-11.00	1.00-1.00			
UDCA (G:T ratio)					
Median (IQR)	21.00 (8.88-38.21)	7.55 (4.25-18.05)	<0.001	276	54
Range	0.14-145.00	0.33-89.15			
CA (frac. BA pool)					
Median (IQR)	0.17 (0.08-0.34)	0.26 (0.12-0.43)	0.008	340	60
Range	0.00-0.79	0.01-0.76			
CDCA (frac. BA pool)					
Median (IQR)	0.26 (0.14-0.45)	0.31 (0.20-0.50)	0.136	340	60
Range	0.04-0.93	0.06-0.88			
DCA (frac. BA pool)					
Median (IQR)	0.05 (0.00-0.14)	0.01 (0.00-0.04)	0.026	340	60
Range	0.00-0.64	0.00-0.46			
LCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.01)	0.992	340	60
Range	0.00-0.12	0.00-0.06			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.039	340	60
Range	0.00-0.05	0.00-0.02			
UDCA (frac. BA pool)					
Median (IQR)	0.21 (0.03-0.68)	0.08 (0.01-0.60)	0.029	340	60
Range	0.00-0.97	0.00-0.88			
CA:CDCA ratio					
Median (IQR)	0.59 (0.34-1.07)	0.66 (0.44-1.45)	0.106	340	60
Range	0.05-5.26	0.06-4.43			
CA:DCA ratio					
Median (IQR)	1.91 (0.59-17.09)	19.69 (2.81-162.95)	<0.001	286	59
Range	0.09-3796.67	0.36-2002.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	1.36 (0.21-11.07)	5.27 (0.30-36.10)	0.017	333	59
Range	0.04-1310.00	0.06-422.00			
Bile acid / UDCA group					
NN	127 (39.8%)	10 (17.5%)			
NY	60 (18.8%)	1 (1.8%)			
HN	41 (12.9%)	22 (38.6%)			
HY	91 (28.5%)	24 (42.1%)			

Supplementary Table 12. Bile acids in autoimmune hepatitis (AIH) overlap and non-overlap large duct PSC patients.

	non-overlap "Large duct" PSC	AIH overlap	p-value	No (n)	AIH (n)
Age in years					
Median (IQR)	48.83 (33.42-60.14)	45.84 (25.23-64.63)	0.716	362	14
Range	10.13-80.48	19.44-67.75			
Sex (% Male)	57.2	57.1	0.998	362	14
Total BA					
Median (IQR)	16.34 (5.89-45.87)	25.92 (12.76-55.13)	0.234	362	14
Range	0.54-646.47	2.56-184.74			
CA (total)					
Median (IQR)	2.12 (0.76-10.54)	2.26 (1.12-5.25)	0.894	362	14
Range	0.04-224.36	0.18-110.08			
CA (unconjugated)					
Median (IQR)	0.07 (0.03-0.23)	0.04 (0.02-0.11)	0.347	362	14
Range	0.00-10.65	0.01-0.77			
GCA					
Median (IQR)	1.70 (0.46-6.41)	1.96 (0.69-3.87)	0.975	362	14
Range	0.01-165.50	0.16-51.45			
TCA					
Median (IQR)	0.27 (0.07-2.57)	0.27 (0.08-1.27)	0.911	362	14
Range	0.00-58.57	0.01-58.61			
CDCA (total)					
Median (IQR)	4.09 (1.69-10.26)	4.50 (3.06-8.57)	0.660	362	14
Range	0.10-268.56	0.73-71.89			
CDCA (unconjugated)					
Median (IQR)	0.15 (0.05-0.46)	0.29 (0.14-0.63)	0.170	362	14
Range	0.00-14.78	0.01-1.55			
GCDCA					
Median (IQR)	2.87 (1.10-7.18)	3.59 (1.81-5.57)	0.651	362	14
Range	0.02-214.48	0.40-36.01			
TCDCA					
Median (IQR)	0.49 (0.13-2.25)	0.62 (0.20-2.17)	0.678	362	14
Range	0.00-52.46	0.06-35.85			
DCA (total)					
Median (IQR)	0.65 (0.03-1.73)	1.43 (0.32-2.02)	0.279	362	14
Range	0.00-24.36	0.00-6.45			
DCA (unconjugated)					
Median (IQR)	0.09 (0.01-0.41)	0.38 (0.07-0.54)	0.160	362	14
Range	0.00-5.99	0.00-0.85			

	Non-overlap	AIH overlap	p-value	No (n)	AIH (n)
GDCA					
Median (IQR)	0.28 (0.02-0.93)	0.65 (0.10-1.36)	0.386	362	14
Range	0.00-15.49	0.00-4.16			
TDCA					
Median (IQR)	0.06 (0.00-0.29)	0.11 (0.01-0.39)	0.350	362	14
Range	0.00-8.55	0.00-1.98			
LCA (total)					
Median (IQR)	0.04 (0.00-0.15)	0.10 (0.01-0.24)	0.379	362	14
Range	0.00-2.85	0.00-1.71			
LCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.04)	0.04 (0.00-0.06)	0.160	362	14
Range	0.00-0.96	0.00-0.55			
GLCA					
Median (IQR)	0.02 (0.00-0.09)	0.03 (0.00-0.13)	0.672	362	14
Range	0.00-2.14	0.00-1.03			
TLCA					
Median (IQR)	0.00 (0.00-0.02)	0.01 (0.00-0.02)	0.440	362	14
Range	0.00-0.56	0.00-0.13			
HDCA (total)					
Median (IQR)	0.01 (0.00-0.04)	0.01 (0.00-0.07)	0.341	362	14
Range	0.00-0.73	0.00-0.09			
HDCA (unconjugated)					
Median (IQR)	0.00 (0.00-0.02)	0.01 (0.00-0.03)	0.872	362	14
Range	0.00-0.24	0.00-0.09			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.01 (0.00-0.01)	0.101	362	14
Range	0.00-0.73	0.00-0.05			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.894	362	14
Range	0.00-0.07	0.00-0.04			
UDCA (total)					
Median (IQR)	1.80 (0.17-18.69)	8.73 (1.47-24.76)	0.106	362	14
Range	0.00-537.71	0.03-78.99			
UDCA (unconjugated)					
Median (IQR)	0.15 (0.01-2.03)	2.79 (0.14-5.47)	0.060	362	14
Range	0.00-52.33	0.00-16.90			
GUDCA					
Median (IQR)	1.41 (0.11-13.86)	5.80 (1.08-15.36)	0.124	362	14
Range	0.00-472.47	0.02-66.92			

	Non-overlap	AIH overlap	p-value	No (n)	AIH (n)
TUDCA					
Median (IQR)	0.08 (0.01-0.53)	0.28 (0.10-0.70)	0.114	362	14
Range	0.00-51.06	0.01-8.13			
Total BA (conj. frac.)					
Median (IQR)	0.91 (0.74-0.98)	0.84 (0.68-0.93)	0.371	362	14
Range	0.13-1.00	0.63-1.00			
CA (conj. frac.)					
Median (IQR)	0.97 (0.84-0.99)	0.98 (0.89-0.99)	0.591	362	14
Range	0.09-1.00	0.44-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.96 (0.83-0.99)	0.91 (0.77-0.98)	0.449	362	14
Range	0.12-1.00	0.57-1.00			
DCA (conj. frac.)					
Median (IQR)	0.81 (0.54-0.96)	0.71 (0.51-0.81)	0.359	313	13
Range	0.00-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.75 (0.57-0.88)	0.68 (0.33-0.79)	0.099	258	11
Range	0.00-1.00	0.00-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.25 (0.00-0.55)	0.59 (0.06-0.97)	0.111	205	10
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.90 (0.75-0.97)	0.80 (0.74-0.96)	0.394	354	14
Range	0.00-1.00	0.58-1.00			
Total BA (G:T ratio)					
Median (IQR)	8.26 (3.39-17.30)	11.42 (5.03-20.53)	0.612	361	14
Range	0.37-93.11	0.92-47.57			
CA (G:T ratio)					
Median (IQR)	5.53 (2.64-9.00)	5.82 (2.98-11.86)	0.663	356	14
Range	0.27-70.33	0.88-21.30			
CDCA (G:T ratio)					
Median (IQR)	5.68 (2.77-10.33)	5.75 (2.67-9.27)	0.684	361	14
Range	0.36-92.17	0.97-16.33			
DCA (G:T ratio)					
Median (IQR)	4.69 (2.08-8.20)	4.82 (3.07-6.57)	0.900	265	11
Range	0.22-44.50	0.63-18.38			
LCA (G:T ratio)					
Median (IQR)	3.00 (1.45-6.00)	5.05 (1.66-6.25)	0.657	153	8
Range	0.10-36.00	1.00-7.92			

	Non-overlap	AIH overlap	p-value	No (n)	AIH (n)
HDCA (G:T ratio)					
Median (IQR)	1.33 (1.00-2.25)	0.75 (0.75-0.75)	0.185	11	1
Range	0.71-11.00	0.75-0.75			
UDCA (G:T ratio)					
Median (IQR)	17.44 (7.00-34.58)	27.54 (8.80-35.44)	0.552	299	14
Range	0.14-145.00	1.11-62.84			
CA (frac. BA pool)					
Median (IQR)	0.19 (0.09-0.37)	0.08 (0.04-0.18)	0.066	362	14
Range	0.00-0.79	0.03-0.60			
CDCA (frac. BA pool)					
Median (IQR)	0.27 (0.15-0.45)	0.17 (0.12-0.36)	0.141	362	14
Range	0.04-0.88	0.06-0.55			
DCA (frac. BA pool)					
Median (IQR)	0.03 (0.00-0.14)	0.06 (0.01-0.10)	0.743	362	14
Range	0.00-0.64	0.00-0.23			
LCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.01)	0.538	362	14
Range	0.00-0.12	0.00-0.06			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.385	362	14
Range	0.00-0.05	0.00-0.01			
UDCA (frac. BA pool)					
Median (IQR)	0.17 (0.02-0.66)	0.65 (0.18-0.72)	0.096	362	14
Range	0.00-0.91	0.00-0.88			
CA:CDCA ratio					
Median (IQR)	0.63 (0.35-1.15)	0.51 (0.34-0.86)	0.374	362	14
Range	0.05-5.26	0.15-1.53			
CA:DCA ratio					
Median (IQR)	2.72 (0.81-27.51)	1.08 (0.76-3.56)	0.448	313	14
Range	0.09-3796.67	0.28-1650.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	1.88 (0.22-14.22)	0.27 (0.17-3.66)	0.159	354	14
Range	0.04-1310.00	0.06-462.67			
Bile acid / UDCA group					
NN	128 (37.4%)	1 (7.7%)	0.185	342	13
NY	52 (15.2%)	3 (23.1%)			
HN	58 (17.0%)	3 (23.1%)			
HY	104 (30.4%)	6 (46.2%)			

Supplementary Table 13. Bile acids in small duct (SD) and large duct (LD) PSC patients.

	large duct PSC	small duct PSC	p-value	LD (n)	SD (n)
Age in years					
Median (IQR)	48.83 (33.42-60.14)	38.20 (29.06-50.11)	0.013	362	26
Range	10.13-80.48	11.82-68.15			
Sex (% Male)	57.2	65.4	0.413	362	26
Total BA					
Median (IQR)	16.34 (5.89-45.87)	11.36 (5.95-28.85)	0.184	362	26
Range	0.54-646.47	0.88-60.71			
CA (total)					
Median (IQR)	2.12 (0.76-10.54)	1.18 (0.37-5.82)	0.020	362	26
Range	0.04-224.36	0.30-24.40			
CA (unconjugated)					
Median (IQR)	0.07 (0.03-0.23)	0.04 (0.02-0.08)	0.034	362	26
Range	0.00-10.65	0.01-0.50			
GCA					
Median (IQR)	1.70 (0.46-6.41)	0.92 (0.26-2.66)	0.064	362	26
Range	0.01-165.50	0.06-13.69			
TCA					
Median (IQR)	0.27 (0.07-2.57)	0.12 (0.03-0.44)	0.040	362	26
Range	0.00-58.57	0.00-4.92			
CDCA (total)					
Median (IQR)	4.09 (1.69-10.26)	3.35 (1.19-5.82)	0.114	362	26
Range	0.10-268.56	0.30-24.40			
CDCA (unconjugated)					
Median (IQR)	0.15 (0.05-0.46)	0.13 (0.07-0.38)	0.773	362	26
Range	0.00-14.78	0.02-1.67			
GCDCA					
Median (IQR)	2.87 (1.10-7.18)	2.06 (0.86-4.46)	0.183	362	26
Range	0.02-214.48	0.18-19.54			
TCDCA					
Median (IQR)	0.49 (0.13-2.25)	0.33 (0.07-0.94)	0.117	362	26
Range	0.00-52.46	0.02-8.04			
DCA (total)					
Median (IQR)	0.65 (0.03-1.73)	0.64 (0.01-1.71)	0.565	362	26
Range	0.00-24.36	0.00-5.98			
DCA (unconjugated)					
Median (IQR)	0.09 (0.01-0.41)	0.30 (0.00-0.44)	0.545	362	26
Range	0.00-5.99	0.00-1.23			

	large duct PSC	small duct PSC	p-value	LD (n)	SD (n)
GDCA					
Median (IQR)	0.28 (0.02-0.93)	0.23 (0.01-1.02)	0.605	362	26
Range	0.00-15.49	0.00-4.76			
TDCA					
Median (IQR)	0.06 (0.00-0.29)	0.04 (0.00-0.12)	0.160	362	26
Range	0.00-8.55	0.00-0.59			
LCA (total)					
Median (IQR)	0.04 (0.00-0.15)	0.04 (0.00-0.13)	0.752	362	26
Range	0.00-2.85	0.00-0.60			
LCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.04)	0.01 (0.00-0.06)	0.666	362	26
Range	0.00-0.96	0.00-0.23			
GLCA					
Median (IQR)	0.02 (0.00-0.09)	0.01 (0.00-0.07)	0.686	362	26
Range	0.00-2.14	0.00-0.33			
TLCA					
Median (IQR)	0.00 (0.00-0.02)	0.00 (0.00-0.01)	0.300	362	26
Range	0.00-0.56	0.00-0.15			
HDCA (total)					
Median (IQR)	0.01 (0.00-0.04)	0.01 (0.00-0.04)	0.668	362	26
Range	0.00-0.73	0.00-0.12			
HDCA (unconjugated)					
Median (IQR)	0.00 (0.00-0.02)	0.01 (0.00-0.03)	0.310	362	26
Range	0.00-0.24	0.00-0.08			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.555	362	26
Range	0.00-0.73	0.00-0.04			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.176	362	26
Range	0.00-0.07	0.00-0.00			
UDCA (total)					
Median (IQR)	1.80 (0.17-18.69)	3.56 (0.23-11.21)	0.954	362	26
Range	0.00-537.71	0.02-52.22			
UDCA (unconjugated)					
Median (IQR)	0.15 (0.01-2.03)	0.51 (0.01-1.49)	0.792	362	26
Range	0.00-52.33	0.00-5.84			
GUDCA					
Median (IQR)	1.41 (0.11-13.86)	2.62 (0.19-9.51)	0.869	362	26
Range	0.00-472.47	0.01-44.53			

	large duct PSC	small duct PSC	p-value	LD (n)	SD (n)
TUDCA					
Median (IQR)	0.08 (0.01-0.53)	0.10 (0.00-0.29)	0.498	362	26
Range	0.00-51.06	0.00-2.61			
Total BA (conj. frac.)					
Median (IQR)	0.91 (0.74-0.98)	0.87 (0.74-0.93)	0.317	362	26
Range	0.13-1.00	0.51-1.00			
CA (conj. frac.)					
Median (IQR)	0.97 (0.84-0.99)	0.95 (0.88-0.99)	0.811	362	26
Range	0.09-1.00	0.37-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.96 (0.83-0.99)	0.91 (0.81-0.98)	0.164	362	26
Range	0.12-1.00	0.45-1.00			
DCA (conj. frac.)					
Median (IQR)	0.81 (0.54-0.96)	0.63 (0.42-0.83)	0.049	313	20
Range	0.00-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.75 (0.57-0.88)	0.62 (0.47-0.79)	0.041	258	18
Range	0.00-1.00	0.25-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.25 (0.00-0.55)	0.00 (0.00-0.31)	0.072	205	18
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.90 (0.75-0.97)	0.88 (0.80-0.96)	0.864	354	26
Range	0.00-1.00	0.50-1.00			
Total BA (G:T ratio)					
Median (IQR)	8.26 (3.39-17.30)	10.64 (5.99-17.69)	0.133	361	26
Range	0.37-93.11	2.07-69.38			
CA (G:T ratio)					
Median (IQR)	5.53 (2.64-9.00)	6.00 (4.17-10.56)	0.105	356	25
Range	0.27-70.33	2.08-14.00			
CDCA (G:T ratio)					
Median (IQR)	5.68 (2.77-10.33)	6.79 (4.58-10.42)	0.301	361	26
Range	0.36-92.17	2.02-31.00			
DCA (G:T ratio)					
Median (IQR)	4.69 (2.08-8.20)	8.06 (4.13-10.67)	0.036	265	17
Range	0.22-44.50	1.83-45.00			
LCA (G:T ratio)					
Median (IQR)	3.00 (1.45-6.00)	3.86 (1.40-7.00)	0.583	153	9
Range	0.10-36.00	1.00-14.00			

	large duct PSC	small duct PSC	p-value	LD (n)	SD (n)
HDCA (G:T ratio)					
Median (IQR)	1.33 (1.00-2.25)	NA	NA	11	0
Range	0.71-11.00	NA			
UDCA (G:T ratio)					
Median (IQR)	17.44 (7.00-34.58)	29.78 (19.47-44.73)	0.019	299	19
Range	0.14-145.00	5.91-101.14			
CA (frac. BA pool)					
Median (IQR)	0.19 (0.09-0.37)	0.12 (0.06-0.28)	0.054	362	26
Range	0.00-0.79	0.02-0.62			
CDCA (frac. BA pool)					
Median (IQR)	0.27 (0.15-0.45)	0.26 (0.14-0.47)	0.780	362	26
Range	0.04-0.88	0.05-0.93			
DCA (frac. BA pool)					
Median (IQR)	0.03 (0.00-0.14)	0.08 (0.00-0.13)	0.828	362	26
Range	0.00-0.64	0.00-0.39			
LCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.01)	0.01 (0.00-0.01)	0.421	362	26
Range	0.00-0.12	0.00-0.04			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.232	362	26
Range	0.00-0.05	0.00-0.03			
UDCA (frac. BA pool)					
Median (IQR)	0.17 (0.02-0.66)	0.53 (0.04-0.66)	0.417	362	26
Range	0.00-0.91	0.00-0.86			
CA:CDCA ratio					
Median (IQR)	0.63 (0.35-1.15)	0.48 (0.30-0.64)	0.044	362	26
Range	0.05-5.26	0.07-2.31			
CA:DCA ratio					
Median (IQR)	2.72 (0.81-27.51)	0.80 (0.38-4.20)	0.030	313	20
Range	0.09-3796.67	0.16-520.00			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	1.88 (0.22-14.22)	0.41 (0.22-9.28)	0.504	354	26
Range	0.04-1310.00	0.07-375.67			
Bile acid / UDCA group					
NN	128 (37.4%)	8 (34.8%)	0.245	342	23
NY	52 (15.2%)	7 (30.4%)			
HN	58 (17.0%)	2 (8.7%)			
HY	104 (30.4%)	6 (26.1%)			

Supplementary Table 14. Univariate time to event (hepatic decompensation) analyses.

Parameter	HR (95% CI)	p-value	c-stat	n
Age (per 10 years)	1.50 (1.24-1.83)	<0.001	0.69	400
Male sex	0.75 (0.43-1.30)	0.302	0.52	400
IBD type (vs. No IBD)				347
Indeterminate IBD	0.75 (0.21-2.70)	0.66	0.53	
Crohn's Disease (CD)	0.25 (0.03-1.96)	0.19	0.53	
Ulcerative colitis (UC)	0.78 (0.39-1.57)	0.49	0.53	
PSC subtypes (vs. non-subtype)				400
Cirrhotic	5.29 (3.02-9.25)	<0.001	0.66	
Small duct disease	0.24 (0.03-1.75)	0.16	0.53	
AIH overlap	2.94 (1.17-7.40)	0.02	0.53	
Total bile acids				
Total	1.15 (1.11-1.18)	<0.001	0.80	400
Total (excluding UDCA)	1.15 (1.11-1.19)	<0.001	0.80	400
Conjugated fraction	10.90 (3.47-34.24)	<0.001	0.72	400
G:T conjugation ratio	0.47 (0.30-0.72)	<0.001	0.69	399
CA				
Total	1.06 (1.04-1.08)	<0.001	0.80	400
Unconjugated	0.98 (0.90-1.06)	0.60	0.56	400
GCA	1.05 (1.03-1.07)	<0.001	0.79	400
TCA	1.03 (1.02-1.03)	<0.001	0.82	400
Conjugated fraction	42.27 (5.09-351.22)	<0.001	0.77	400
G:T conjugation ratio	0.30 (0.17-0.53)	<0.001	0.72	393
Fraction of bile acid pool	1.31 (1.04-1.65)	0.02	0.55	400
CDCA				
Total	1.16 (1.11-1.20)	<0.001	0.80	400
Unconjugated	0.98 (0.87-1.11)	0.75	0.53	400
GCDCA	1.13 (1.09-1.16)	<0.001	0.78	400
TCDCA	1.09 (1.07-1.11)	<0.001	0.83	400
Conjugated fraction	12.68 (3.30-48.76)	<0.001	0.75	400
G:T conjugation ratio	0.20 (0.10-0.39)	<0.001	0.74	399
Fraction of bile acid pool	0.74 (0.50-1.11)	0.14	0.54	400
DCA				
Total	1.02 (0.91-1.15)	0.71	0.56	400
Unconjugated	0.71 (0.48-1.06)	0.09	0.56	400
GDCA	1.02 (0.94-1.11)	0.58	0.56	400
TDCA	1.03 (0.99-1.07)	0.18	0.70	400
Conjugated fraction	7.35 (3.04-17.73)	<0.001	0.71	345
G:T conjugation ratio	0.24 (0.12-0.45)	<0.001	0.74	292
Fraction of bile acid pool	0.08 (0.02-0.39)	<0.001	0.61	400

Parameter	HR (95% CI)	p-value	c-stat	n
LCA				
Total	1.12 (1.07-1.17)	<0.001	0.66	400
Unconjugated	1.12 (1.08-1.15)	<0.001	0.62	400
GLCA	1.06 (1.02-1.10)	0.01	0.62	400
TLCA	1.05 (1.03-1.08)	<0.001	0.75	400
Conjugated fraction	1.08 (0.71-1.65)	0.70	0.53	286
G:T conjugation ratio	0.55 (0.33-0.91)	0.02	0.65	169
Fraction of bile acid pool	1.00 (0.65-1.53)	0.99	0.49	400
HDCA				
Total	0.98 (0.75-1.29)	0.90	0.52	400
Unconjugated	1.06 (0.83-1.36)	0.65	0.49	400
GHDCA	0.94 (0.80-1.11)	0.47	0.52	400
THDCA	na (0.00-na)	Na	na	400
Conjugated fraction	0.81 (0.46-1.42)	0.46	0.52	231
G:T conjugation ratio	na (0.00-na)	Na	na	12
Fraction of bile acid pool	0.17 (0.04-0.76)	0.02	0.59	400
UDCA				
Total	1.04 (1.03-1.06)	<0.001	0.66	400
Unconjugated	1.05 (1.02-1.08)	<0.001	0.57	400
GUDCA	1.04 (1.02-1.05)	<0.001	0.65	400
TUDCA	1.02 (1.01-1.02)	<0.001	0.75	400
Conjugated fraction	4.18 (1.97-8.87)	<0.001	0.66	392
G:T conjugation ratio	0.30 (0.17-0.54)	<0.001	0.72	330
Fraction of bile acid pool	1.22 (0.81-1.83)	0.34	0.50	400
High total BA (>17.79 µmol/L)	8.00 (3.60-17.77)	<0.001	0.72	400
UDCA treatment (predicted)	1.65 (0.94-2.90)	0.08	0.57	376
CA:CDCA Ratio	1.19 (1.05-1.35)	0.01	0.62	400
CA:DCA Ratio	1.00 (1.00-1.00)	0.67	0.71	345
Bilirubin (total)	1.15 (1.09-1.21)	<0.001	0.77	370
Alkaline phosphatase (AP)				
AP x ULN (upper limit of normal)	1.21 (1.10-1.33)	<0.001	0.70	375
AP > 1.0 x ULN	7.93 (2.47-25.51)	<0.001	0.65	375
AP > 1.5 x ULN	2.81 (1.51-5.23)	0.001	0.64	375

Supplementary Table 15. Univariate event rate (hepatic decompensation) by quartiles analyses.

Parameter	Quartile of parameter value			
	Q1	Q2	Q3	Q4
Total bile acids				
Total	3.18	4.42	14.56	45.51
Total (excluding UDCA)	1.07	9.56	12.35	45.75
Conjugated fraction	3.20	11.79	15.32	31.71
G:T conjugation ratio	26.13	15.83	14.48	4.29
CA				
Total	1.05	3.27	21.90	40.81
Unconjugated	17.71	17.58	13.66	10.89
GCA	1.04	5.59	20.26	38.52
TCA	0.00	4.43	18.42	44.10
Conjugated fraction	2.05	8.18	15.19	41.70
G:T conjugation ratio	37.62	13.21	8.24	3.35
Fraction of bile acid pool	10.26	17.94	11.83	19.66
CDCA				
Total	4.21	7.55	9.99	47.97
Unconjugated	20.91	9.83	17.19	12.34
GCDCA	3.12	9.83	14.88	38.55
TCDCA	0.00	7.52	13.81	48.07
Conjugated fraction	0.00	14.13	14.85	34.74
G:T conjugation ratio	34.84	20.37	4.31	3.30
Fraction of bile acid pool	16.40	17.64	16.50	9.23
DCA				
Total	4.44	18.72	23.18	14.15
Unconjugated	11.47	30.51	8.68	11.40
GDCA	5.61	18.67	21.63	14.22
TDCA	2.21	5.38	25.99	29.62
Conjugated fraction	3.38	14.82	35.28	9.36
G:T conjugation ratio	36.66	17.59	4.43	3.42
Fraction of bile acid pool	8.11	37.57	14.33	2.27
LCA				
Total	5.53	13.34	13.41	28.74
Unconjugated	10.80	12.59	11.28	25.80
GLCA	8.04	11.84	15.13	25.52
TLCA	1.06	8.21	10.36	44.74
Conjugated fraction	19.95	15.60	19.56	5.43
G:T conjugation ratio	38.75	18.80	3.35	2.22
Fraction of bile acid pool	4.39	30.18	15.04	12.42
HDCA				
Total	11.03	20.91	11.53	16.72
Unconjugated	11.09	16.61	19.70	12.55
GHDCA	14.84	5.29	28.01	12.79

Parameter	Quartile of parameter value			
	Q1	Q2	Q3	Q4
HDCA (cont.)				
THDCA	13.26	18.92	6.87	21.08
Conjugated fraction	15.01	16.07	5.83	23.05
G:T conjugation ratio	29.81	25.55	5.58	3.08
Fraction of bile acid pool	6.72	31.92	15.32	7.79
UDCA				
Total	11.91	10.76	6.51	33.66
Unconjugated	19.19	4.68	11.36	24.87
GUDCA	15.53	8.07	3.25	36.78
TUDCA	4.16	10.69	11.48	37.41
Conjugated fraction	3.11	12.29	24.77	21.43
G:T conjugation ratio	34.27	21.17	5.64	3.07
Fraction of bile acid pool	21.46	4.50	19.11	15.12
CA:CDCA Ratio	6.85	9.69	16.79	26.52
CA:DCA Ratio	2.24	14.33	36.40	9.37

Event rate calculated as Number of Events / Person Years

Supplementary Table 16. Variables and relative influence on GBM Model

Variable	Relative Influence	Influence direction [#]
Age (per decade)	14.684	Increase
TUDCA	14.475	Increase
TCDCA	12.421	Increase
TLCA	8.865	Increase
TCA	8.603	Increase
TBA	4.234	Increase
DCA (fraction of BA pool)	3.606	Decrease
GDCA	3.023	Decrease
CDCA (total)	2.890	Increase
LCA (total)	2.462	Increase
LCA (conjugated fraction)	2.198	Decrease
CA (fraction of BA pool)	1.831	Increase
GCDCA	1.814	Increase
CA (conjugated fraction)	1.775	Increase
GUDCA	1.739	Increase
TDCA	1.421	Decrease
TBA (conjugated fraction)	1.386	Increase
CA (total)	1.356	Increase
DCA (conjugated fraction)	1.207	Increase
UDCA (fraction of BA pool)	1.191	Decrease
UDCA (total)	1.123	Increase
CDCA (conjugated fraction)	1.086	Decrease
GLCA	0.947	Decrease
GCA	0.873	Decrease
UDCA (conjugated fraction)	0.849	Increase
DCA (total)	0.787	Decrease
LCA (fraction of BA pool)	0.659	Increase
CDCA (unconjugated)	0.576	Increase
UDCA (unconjugated)	0.451	Decrease
HDCA (unconjugated)	0.426	Increase
CDCA (fraction of BA pool)	0.361	Decrease
CA (unconjugated)	0.144	Decrease
HDCA (fraction of BA pool)	0.121	Decrease
HDCA (total)	0.102	Increase
Male sex	0.100	Decrease
DCA (unconjugated)	0.070	Decrease
GHDCA	0.038	Increase
HDCA (conjugated fraction)	0.000	Decrease
THDCA	0.000	Decrease
UDCA use (calculated)	0.000	Increase

Increase or decreased risk of hepatic decompensation with increase in value of variable based on visualization of functional form plots. For instance, increased TUDCA concentration increased risk in the model whereas increased GDCA concentration lowered the risk.

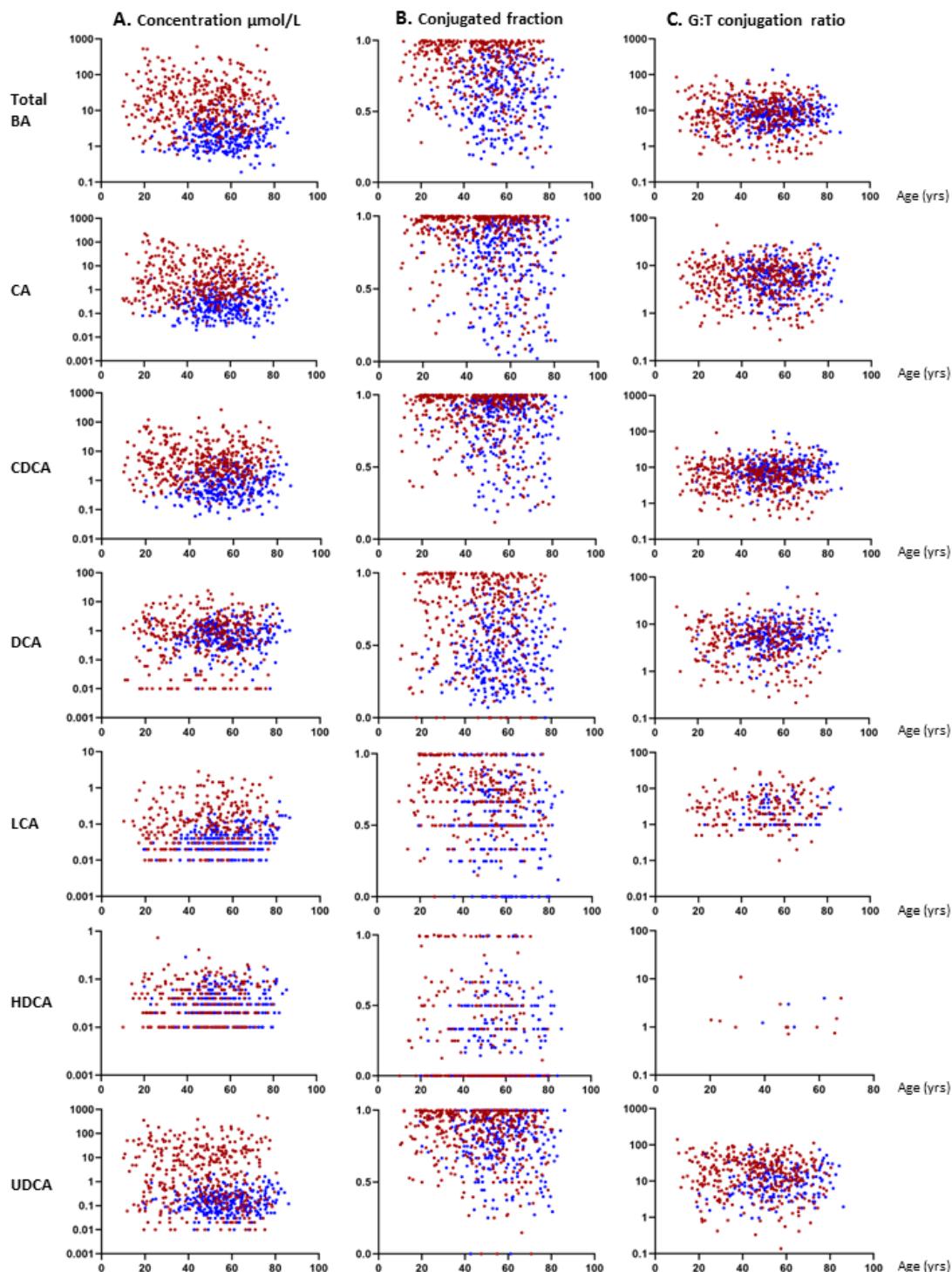
Supplementary Table 17. Bile acids in PSC patients: Derivation and Validation cohorts.

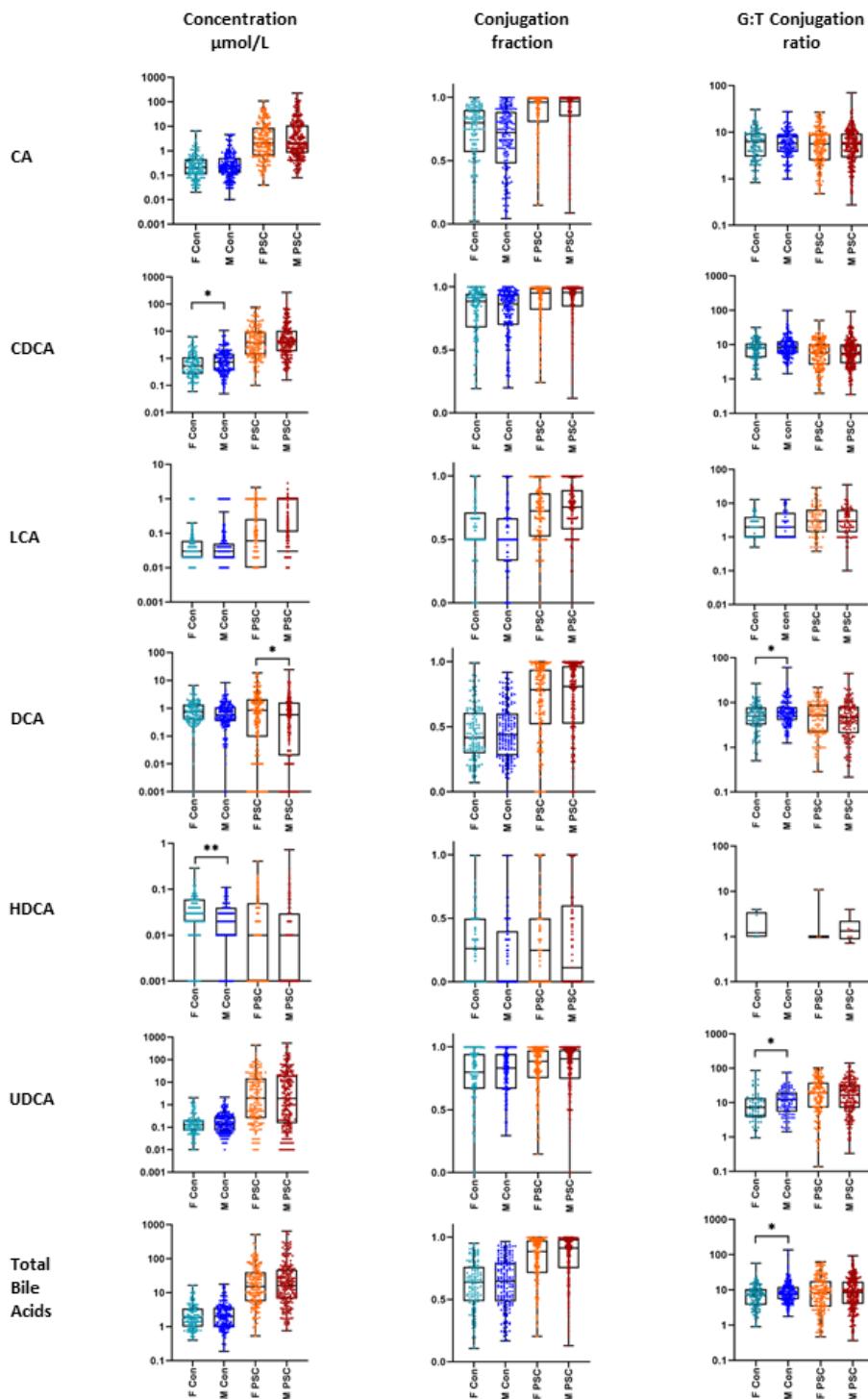
	PSC (derivation)	PSC (validation)	p-value	PSC Der (n)	PSC Val (n)
Total BA					
Median (IQR)	16.34 (6.04-44.31)	14.31 (5.12-59.30)	0.895	400	108
Range	0.54-646.47	0.74-457.45			
CA (total)					
Median (IQR)	2.09 (0.73-10.01)	3.01 (1.26-20.19)	0.019	400	108
Range	0.04-224.36	0.15-251.95			
CA (unconjugated)					
Median (IQR)	0.06 (0.03-0.22)	0.04 (0.02-0.09)	0.003	400	108
Range	0.00-10.65	0.00-3.92			
GCA					
Median (IQR)	1.60 (0.44-6.12)	2.03 (0.98-15.33)	0.010	400	108
Range	0.01-165.50	0.11-192.78			
TCA					
Median (IQR)	0.25 (0.06-2.44)	0.42 (0.14-6.64)	0.003	400	108
Range	0.00-58.61	0.01-149.57			
CDCA (total)					
Median (IQR)	4.09 (1.61-9.81)	4.30 (1.86-12.04)	0.482	400	108
Range	0.10-268.56	0.31-125.26			
CDCA (unconjugated)					
Median (IQR)	0.15 (0.05-0.46)	0.07 (0.02-0.20)	<0.001	400	108
Range	0.00-14.78	0.01-5.13			
GCDCA					
Median (IQR)	2.87 (1.10-6.90)	2.77 (1.30-7.63)	0.672	400	108
Range	0.02-214.48	0.19-87.38			
TCDCA					
Median (IQR)	0.47 (0.13-2.12)	0.92 (0.30-3.98)	0.003	400	108
Range	0.00-52.46	0.02-67.72			
DCA (total)					
Median (IQR)	0.66 (0.03-1.76)	0.61 (0.16-1.79)	0.702	400	108
Range	0.00-24.36	0.00-20.69			
DCA (unconjugated)					
Median (IQR)	0.10 (0.01-0.44)	0.05 (0.00-0.16)	0.025	400	108
Range	0.00-5.99	0.00-2.96			
GDCA					
Median (IQR)	0.28 (0.02-0.96)	0.38 (0.10-1.13)	0.158	400	108
Range	0.00-15.49	0.00-16.97			

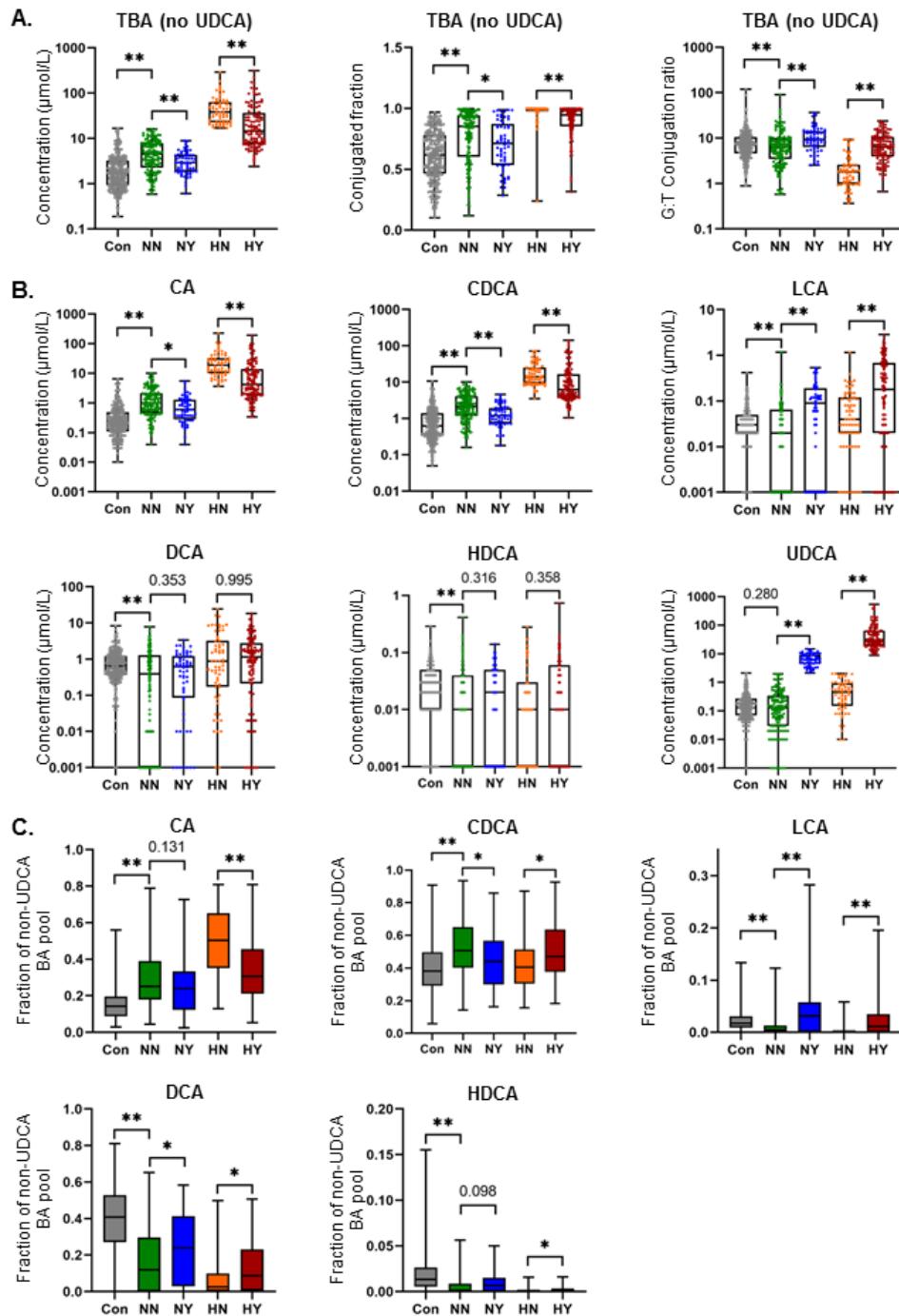
	PSC (derivation)	PSC (validation)	p-value	PSC Der (n)	PSC Val (n)
TDCA					
Median (IQR)	0.06 (0.00-0.27)	0.07 (0.01-0.25)	0.266	400	108
Range	0.00-8.55	0.00-3.71			
LCA (total)					
Median (IQR)	0.04 (0.00-0.15)	0.02 (0.00-0.05)	0.010	400	108
Range	0.00-2.85	0.00-1.82			
LCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.04)	0.01 (0.00-0.02)	0.043	400	108
Range	0.00-0.96	0.00-0.23)			
GLCA					
Median (IQR)	0.02 (0.00-0.09)	0.00 (0.00-0.00)	<0.001	400	108
Range	0.00-2.14	0.00-1.38			
TLCA					
Median (IQR)	0.00 (0.00-0.02)	0.01 (0.00-0.02)	0.037	400	108
Range	0.00-0.56)	0.00-0.26			
HDCA (total)					
Median (IQR)	0.01 (0.00-0.04)	0.01 (0.00-0.03)	0.117	400	108
Range	0.00-0.73	0.00-0.18			
HDCA (unconjugated)					
Median (IQR)	0.01 (0.00-0.02)	0.00 (0.00-0.02)	0.063	400	108
Range	0.00-0.24	0.00-0.16			
GHDCA					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.004	400	108
Range	0.00-0.73	0.00-0.09			
THDCA					
Median (IQR)	0.00 (0.00-0.00)	0.00 (0.00-0.00)	0.065	400	108
Range	0.00-0.07	0.00-0.10			
UDCA (total)					
Median (IQR)	1.95 (0.19-18.27)	0.36 (0.09-10.56)	0.039	400	108
Range	0.00-537.71	0.00-308.01			
UDCA (unconjugated)					
Median (IQR)	0.17 (0.01-2.02)	0.05 (0.00-0.53)	<0.001	400	108
Range	0.00-52.33	0.00-17.16			
GUDCA					
Median (IQR)	1.53 (0.12-13.52)	0.27 (0.06-9.42)	0.063	400	108
Range	0.00-472.47	0.00-278.90			
TUDCA					
Median (IQR)	0.09 (0.01-0.52)	0.06 (0.01-0.60)	0.428	400	108
Range	0.00-51.06	0.00-44.90			

	PSC (derivation)	PSC (validation)	p-value	PSC Der (n)	PSC Val (n)
Total BA (conj. frac.)					
Median (IQR)	0.91 (0.74-0.98)	0.95 (0.88-0.99)	<0.001	400	108
Range	0.13-1.00	0.23-1.00			
CA (conj. frac.)					
Median (IQR)	0.97 (0.85-0.99)	0.98 (0.94-1.00)	<0.001	400	108
Range	0.09-1.00	0.185-1.00			
CDCA (conj. frac.)					
Median (IQR)	0.95 (0.83-0.99)	0.98 (0.92-1.00)	<0.001	400	108
Range	0.12-1.00	0.23-1.00			
DCA (conj. frac.)					
Median (IQR)	0.80 (0.52-0.95)	0.92 (0.74-0.98)	<0.001	345	95
Range	0.00-1.00	0.00-1.00			
LCA (conj. frac.)					
Median (IQR)	0.75 (0.56-0.87)	0.71 (0.40-0.88)	0.069	286	77
Range	0.00-1.00	0.00-1.00			
HDCA (conj. frac.)					
Median (IQR)	0.22 (0.00-0.55)	0.00 (0.00-0.48)	0.134	231	55
Range	0.00-1.00	0.00-1.00			
UDCA (conj. frac.)					
Median (IQR)	0.90 (0.75-0.97)	0.95 (0.86-1.00)	<0.001	392	105
Range	0.00-1.00	0.23-1.00			
Total BA (G:T ratio)					
Median (IQR)	8.36 (3.57-17.44)	4.70 (2.55-8.21)	<0.001	399	108
Range	0.37-93.11	0.45-57.86			
CA (G:T ratio)					
Median (IQR)	5.56 (2.72-9.23)	4.71 (2.25-8.07)	0.038	393	108
Range	0.27-70.33	0.49-19.00			
CDCA (G:T ratio)					
Median (IQR)	5.76 (2.80-10.30)	3.24 (1.83-5.53)	<0.001	399	108
Range	0.36-92.17	0.37-21.00			
DCA (G:T ratio)					
Median (IQR)	4.90 (2.13-8.26)	4.97 (2.82-9.25)	0.290	292	89
Range	0.22-45.00	0.68-30.29			
LCA (G:T ratio)					
Median (IQR)	3.00 (1.40-6.22)	2.43 (1.00-5.00)	0.174	169	23
Range	0.10-36.00	0.50-11.57			
HDCA (G:T ratio)					
Median (IQR)	1.17 (1.00-1.88)	1.50 (1.50-1.80)	0.336	12	5
Range	0.71-11.00	1.00-2.67			

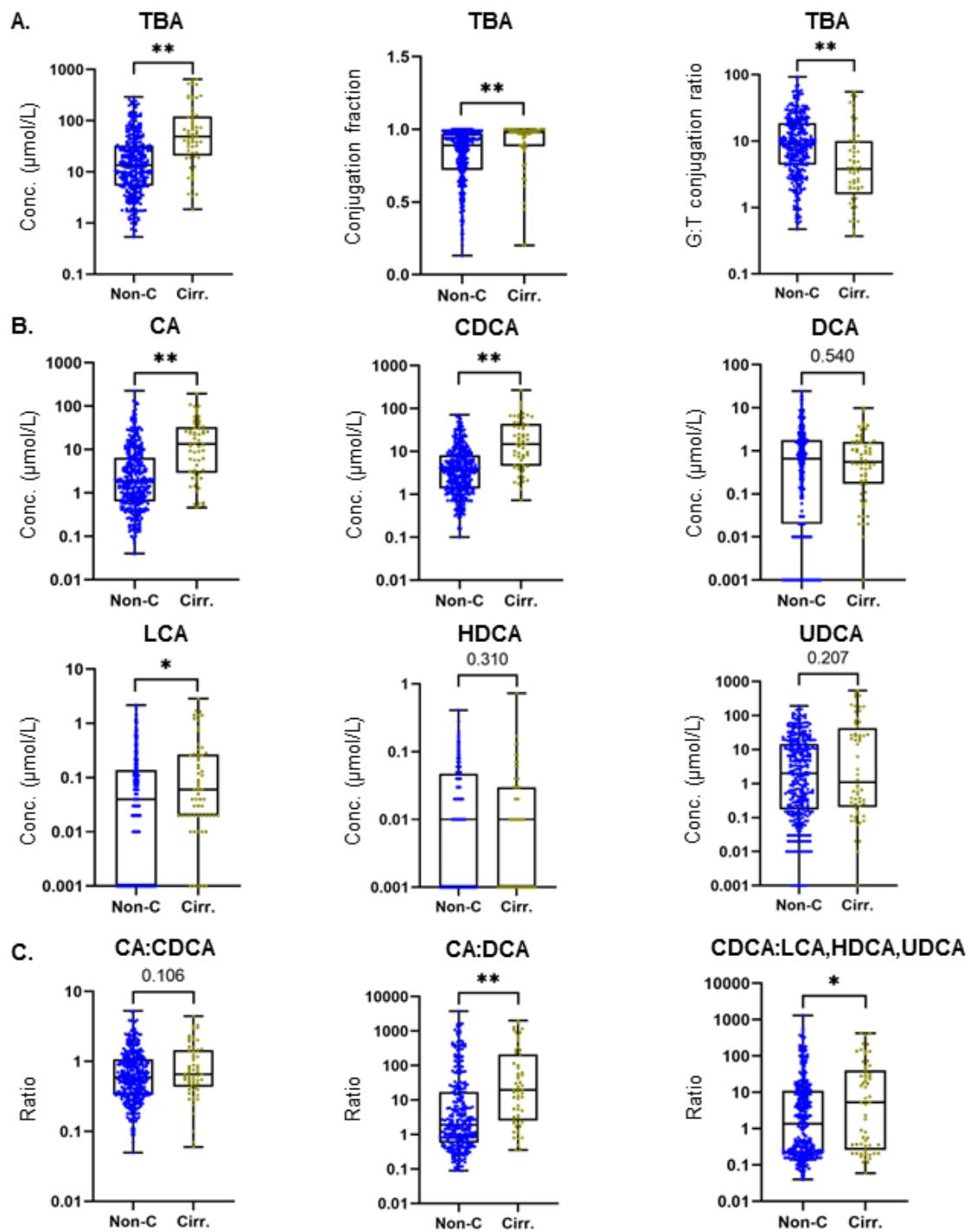
	PSC (derivation)	PSC (validation)	p-value	PSC Der (n)	PSC Val (n)
UDCA (G:T ratio)					
Median (IQR)	18.40 (7.25-35.75)	9.26 (4.45-18.00)	<0.001	330	83
Range	0.14-145.00	0.27-107.33			
CA (frac. BA pool)					
Median (IQR)	0.18 (0.08-0.36)	0.29 (0.18-0.44)	<0.001	400	108
Range	0.01-0.79	0.02-0.87			
CDCA (frac. BA pool)					
Median (IQR)	0.26 (0.15-0.46)	0.32 (0.19-0.46)	0.153	400	108
Range	0.04-0.93	0.09-0.79			
DCA (frac. BA pool)					
Median (IQR)	0.04 (0.01-0.14)	0.03 (0.01-0.15)	0.532	400	108
Range	0.00-0.64	0.00-0.54			
LCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.004	400	108
Range	0.00-0.12	0.00-0.05			
HDCA (frac. BA pool)					
Median (IQR)	0.00 (0.00-0.01)	0.00 (0.00-0.00)	0.064	400	108
Range	0.00-0.05	0.00-0.04			
UDCA (frac. BA pool)					
Median (IQR)	0.18 (0.02-0.66)	0.05 (0.01-0.47)	<0.001	400	108
Range	0.00-0.91	0.00-0.84			
CA:CDCA ratio					
Median (IQR)	0.60 (0.35-1.12)	0.89 (0.49-1.60)	<0.001	400	108
Range	0.05-5.26	0.13-7.15			
CA:DCA ratio					
Median (IQR)	2.60 (0.79-26.43)	4.47 (1.38-32.18)	0.101	345	95
Range	0.09-3796.67	0.09-1683.67			
CDCA:LCA+HDCA+UDCA					
Median (IQR)	1.57 (0.22-13.81)	5.93 (0.38-22.31)	0.002	392	105
Range	0.04-1310.00	0.12-673.13			

Supplementary Figure 1. Bile acids by age in PSC and controls.**Supplementary Figure 1.** Plots showing bile acid (A) concentration, (B) conjugated fraction and (C) G:T conjugation ratio (Y axis) by age in years (X axis). Red dots indicate PSC patients and blue dots indicate controls.

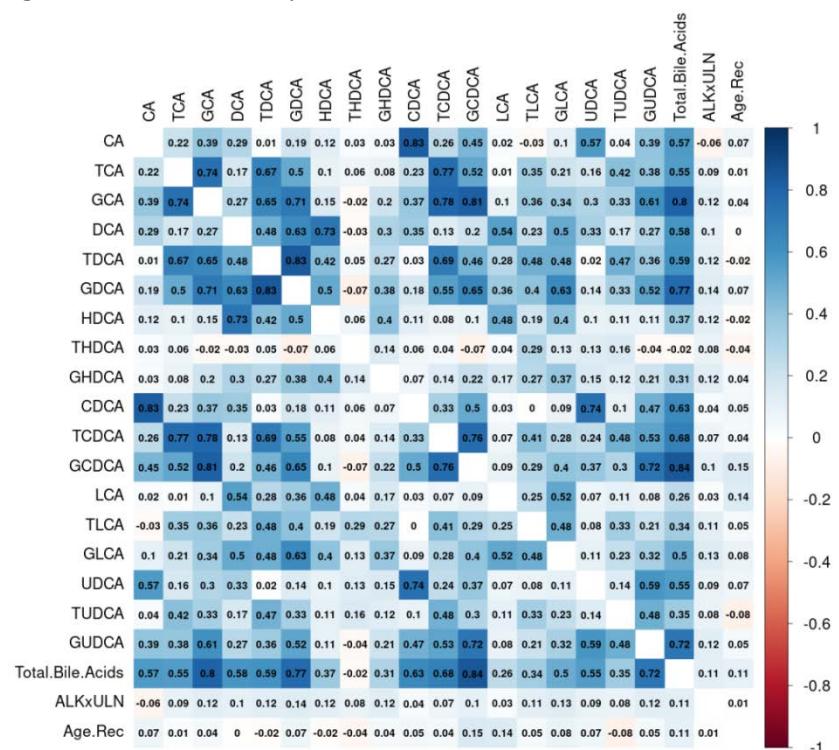
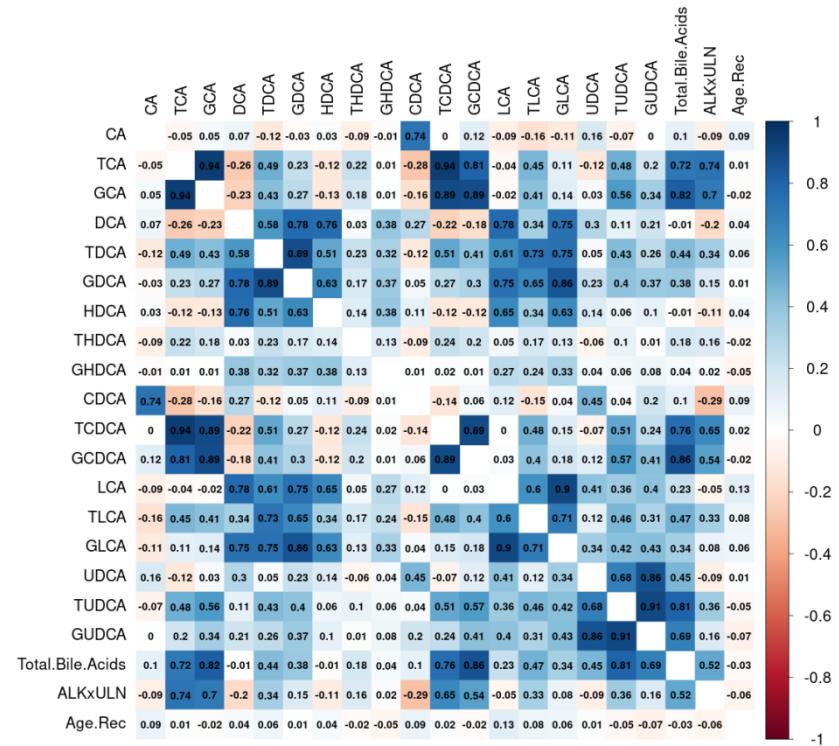
Supplementary Figure 2. Bile acids by sex in PSC and controls.**Supplementary Figure 2.** Plots showing bile acid (A) concentration, (B) conjugated fraction and (C) G:T conjugation ratio by comparing values in females to males within PSC and control groups. * $p<0.05$, ** $p<0.01$.

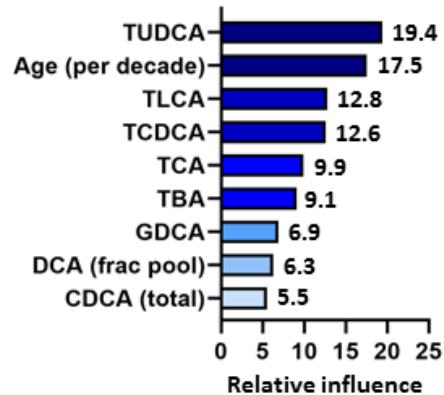
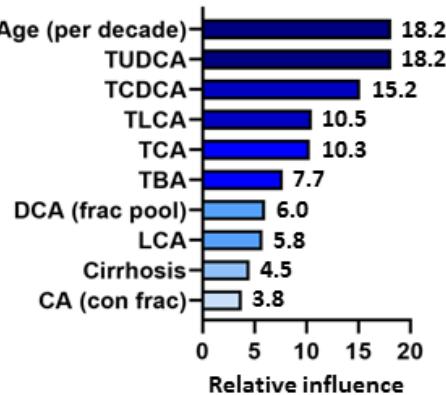
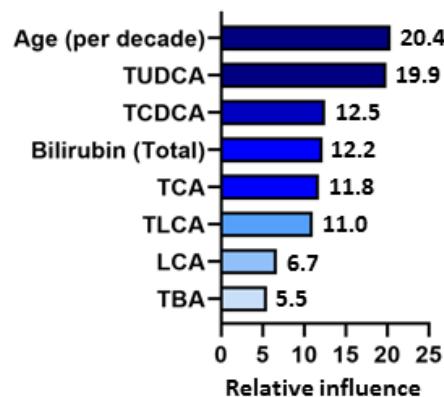
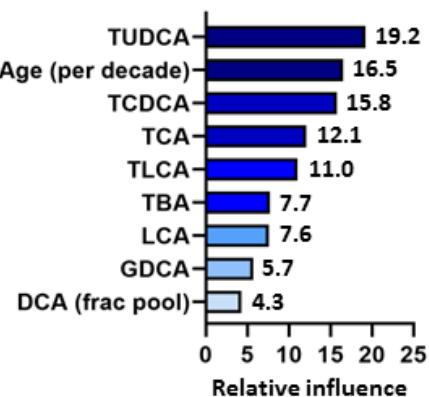
Supplementary Figure 3. Bile acids in controls and PSC patients by bile acid group.

Supplementary Figure 3. Plots showing (A) concentration, conjugation fraction, and G:T conjugation ratio of total bile acids (TBA) with UDCA, GUDCA and TUDCA removed; (B) concentration of bile acid “families” (i.e., the sum of conjugated and unconjugated forms); and (C) bile acid “families” as a fraction of the non-UDCA BA pool in controls (Con) and PSC patients separated by BA group. NN: normal TBA, no UDCA; NY: normal TBA, yes UDCA; HN: high TBA, no UDCA; HY: high TBA, yes UDCA. * $p < 0.05$, ** $p < 0.001$

Supplementary Figure 4. Bile acids in cirrhotic and non-cirrhotic PSC patients

Supplementary Figure 4. Plots showing (A) concentration, conjugation fraction and G:T conjugation ratio of total bile acid (TBA); (B) concentration of bile acid “families” (i.e., total of conjugated and unconjugated forms); and (C) bile acid ratios in PSC patients who were non-cirrhotic (Non-C) or cirrhotic (Cirr.) at time of sample collection. * $p < 0.05$, ** $p < 0.001$

Supplementary Figure 5. BA correlation plots in PSC and controls.**A. Controls****B. PSC****Supplementary Figure 5.** Spearman correlation plots of BA concentrations in (A) controls and (B) PSC patients. Dark blue values indicate high correlation whereas red values indicate low correlation.

Supplementary Figure 6. Alternate GBM models using BA profiles**A. Remove SD and AIH patients from GBM model – Concordance = 0.95****B. Add Cirrhosis phenotype to GBM model – Concordance = 0.95****C. Add Bilirubin (Total) to GBM model Concordance = 0.95****D. Add ALP (times upper limit of normal) to GBM model – Concordance = 0.96**

Supplementary Figure 6. Alternate GBM models using BA profiles with (A) PSC patients with small duct (SD) disease and/or having overlapping autoimmune hepatitis (AIH) removed, (B) cirrhosis phenotype (yes/no) added to the model, (C) Bilirubin (total) added to the model, and (D) alkaline phosphatase (ALP) added to the model.