

Supplementary Data

Material and methods

Generation and culture of canonical and non-canonical Wnt stimulated cholangiocyte organoids from bile and tissue

For the initiation (of culture) of bile cholangiocyte organoids in canonical-Wnt stimulating conditions (cBCOs, n=6), we used a protocol similar to the one previously published by Soroka *et al.*¹. In short, organoids were initiated from a maximum of three mL of bile collected *ex vivo* and from one millilitre of bile collected *in vivo*. Bile was washed twice for five minutes with cold Advanced (Adv)DMEM/F12 (supplemented with 1% penicillin/streptomycin, Life Technologies; 1% Hepes 1 M, Fisher Scientific; 1% Ultra-glutamine 200mM, Fisher Scientific and 0.2% Primocin, Invivogen) at 453g and suspended in 8 mL of cold AdvDMEM/F12. Subsequently, the supernatant was removed and the pellet was suspended in 3 mL of AdvDMEM/F12 and passed through a 100 µm cell strainer to remove debris. After a third wash, the pellet was collected and plated out in a hydrogel (Matrigel, Corning, or diluted Basement Membrane Extract, BME, Cultrex). Medium was added according to the previously described protocol.^{2,3} The first three days the medium was supplemented with 1% anti-anti (Gibco) to prevent bacterial- or fungal infections.

Tissue-derived intrahepatic cholangiocyte organoids in canonical-Wnt stimulating conditions (ciCOs, n=3) and extrahepatic cholangiocyte organoids in canonical-Wnt stimulating conditions (cECOs, n=3) were initiated and cultured as previously published.²⁻⁴ In short, liver or extrahepatic bile duct biopsies (0.5-1.0 cm) were digested using a collagenase solution in (2.5 mg/ml collagenase A1, Roche) EBBS (Hyclone, ThermoFisher) for 30 min at 37 °C. Digestion was stopped by adding cold supplement AdvDMEM/F12 filtered through a 70 µm cell strainer. The cell suspension was spun down for 5 minutes, 4°C at 453g. Pellet was harvested in a hydrogel and specific medium was added according to protocol.^{2,3} Medium was changed twice a week and organoids were passaged in a 1:2 to 1:8 ratio according to growth. Cultures were negative for mycoplasma contamination (data not shown). The ncECO cultures (n=6) were initiated and maintained as previously published.⁵ In short, cells were scraped from the inside of the gallbladders or from the inside of an extrahepatic bile duct (n=1) and collected in cold William's-E Medium (WE). The cells were centrifuged at 453g for 5 minutes at 4°C, subsequently supernatant was removed, and the cell pellet was washed twice with excess cold WE. Next the pellet was collected and plated out in either matrigel or 70% BME diluted with WE and medium was added according to the standard protocol.⁵

Additionally, to investigate the influence of culture conditions on BCOs. Bile cholangiocyte organoids were either created from the same bile samples (n=3) either under cBCO conditions or ncBCO conditions. After three passages a switch from cBCO towards ncBCO conditions and from ncBCO conditions towards cBCO conditions was performed. Two passages later RNA was harvested from these 4 conditions per donor sample and analysed by qRT-PCR.

Ussing Chamber Measurement conditions

The temperature of the chambers was kept 37 °C by warm water bath circulation. Each chamber consisted of 3 mL modified Meyler solution (128 mmol/liter NaCl, 4.7 mmol/liter KCl, 1.3 mmol/liter CaCl₂, 1.0 mmol/liter MgCl₂, 0.3 mmol/liter Na₂HPO₄, 0.4 mmol/liter NaH₂PO₄, 20 mmol/liter NaHCO₃, 10 mmol/liter HEPES, supplemented with glucose (10 mmol/liter) in 95% O₂, 5% CO₂ at pH 7.3). Current was clamped and every second current was recorded. Two voltage spikes (5 millivoltage) were given, to measure resistance along the epithelial layers ($V=I \cdot R$; TEER). CFTR-dependent anion secretion was activated by adding 20 µM Secretin (Sigma) basolateral or 10 µM Forskolin (Sigma) to

both sides of the cells and inhibited by addition of GlyH-101 (20 μM , apical side, Calbiochem) or 100 μM Somatostatin (Sigma) basolateral. Calcium (Ca^{2+}) depended chloride (Cl^-) channels (CaCC) were stimulated by UTP (50 μM , apical side, Sigma) and inhibited by T16Ainh-A01 (50 μM , apical side, Tocris).

Tables

Table S1. Donor and culture characteristics of organoids included.

Age patient (years)	Gender	Cells Source	Donor Type or Indication ERCP/surgery	Culture type	Successful Culture	Characterized at Passage #	Gene-expression by qRT-PCR
13	M	GB, GBB	DBD	ncECO 1, ncBCO 1	+	5-9	+
26	M	GBB	DCD	ncBCO 7**, cBCO 7**	+	5	+
29	M	ERCP	Bile stones	ncBCO 14	+	ND	-
30	M	ERCP	PSC	ncBCO 8 cBCO 8	+	5	+
35	F	GBB, GB	DBD	ncBCO 5, ncECO 5	+	5	+
41	F	EHBD	PSC	ncECO 11	+	5	+
47	M	GBB	DCD	ncBCO 6**, cBCO 6**	+	5	+
50	M	GB	Bile stones	ncECO 12	+	5-8	+
51	F	ERCP	AS	ncBCO 15	+	ND	-
52	M	GBB, GB	DCD	ncBCO 4, ncECO 4	+	5-9	+
54	M	EHBD	DBD	cECO D*	+	5	-
54	F	ERCP	Papiladenoma	ncBCO 10	+	5-8	+
56	F	EHBD	DCD	cECO C*	+	5	-
57	M	Liver, GBB, GBB	DCD	cICO 2, cBCO 2, ncBCO 2	+	5 5 5	+(only cBCO)
58	F	ERCP	AS	ncBCO 16	+	ND	-
59	M	GB, GBB, GBB, Liver	DBD	ncECO 3, cBCO 3, ncBCO 3 cICO 3	+	5-8 5 5 5	+
67	M	Liver	DCD	cICO 13	+	5	+
67	F	EHBD	Cryptogenic Cirrhosis	cECO F*	+	5	-
70	M	ERCP	Bile stones	ncBCO 17	+	ND	-
72	M	ERCP	Bile stones	ncBCO 18	+	ND	-
73	F	ERCP	AS	ncBCO 19	-	ND	-
75	F	GBB	DBD	ncBCO 9**, cBCO 9**	+	5	+

Abbreviations: AS: Anastomotic bile duct Stricture, ncBCO: Bile Cholangiocyte Organoid in non-canonical Wnt stimulating conditions, EHBD: Extrahepatic Bile Duct, cECOs: extrahepatic cholangiocyte organoids in canonical Wnt stimulating conditions, cICOs: intrahepatic cholangiocyte

organoids in canonical Wnt stimulating conditions, cBCO: Bile Cholangiocyte Organoids organoids in canonical Wnt stimulating conditions, DBD: Donation after Brainstem Death, DCD: Donation after Circulatory Death, ncECO: Extrahepatic Cholangiocyte Organoids in non-canonical Wnt stimulating conditions, ERCP: Endoscopic Retrograde Cholangiopancreatography, F: Female, GB: Gallbladder, GBB: Gallbladder Bile, M: Male, PSC: Primary Sclerosing Cholangitis, ND: Not Determined. *These organoid cultures were only analysed for Micro Array analysis. **Used for switching experiment (ncBCOs, are only used for the switching experiment).

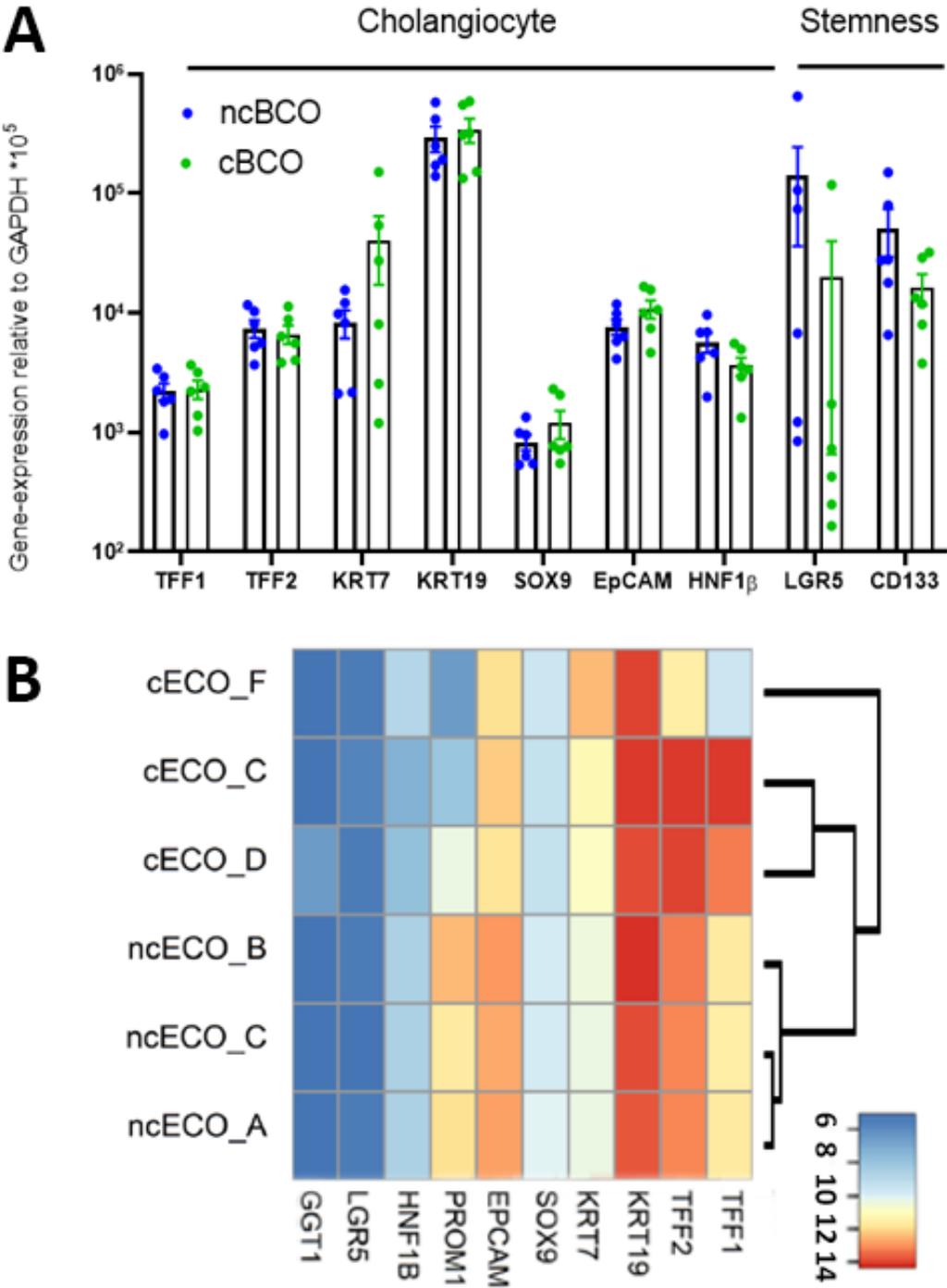
Table S2. List of genes and primers used.

Gene	Primer sequence (5' à 3')	Gene	Primer sequence (5' à 3')
KRT7	F GGGGACGACCTCCGGAATAC R CTTGGCAGCTGGTTCTTGA	IGFBP3	F CAAGGGGAAGGAGGACGTGCA R GCTGCTGGTCATGTCCTTGCC
KRT19	F GCACTACAGCCACTACTACACGA R CTCATGCGCAGAGCCTGTT	KLK7	F ACCACCTGTACTGTCTCCGGCT R AACCTTCGTGCACTCCTGGGG
HNF1β	F TCACAGATAACCAGCAGCATCAGT R GGGCATCACCAGGCTTGTA	IGFBP1	F AGGCACAGGAGACATCAGGAGA R CACCAGCAGAGTCCCGCCTC
HPRT	F GCTATAAATTCTTTGCTGACCTGCG R CTTGCTGGGGTCCTTTTCACC	SLC2a3	F ATTGCTCTTCCCTCCGCTGC R CCTCAAAGTCTGCCACGGGT
ALB	F CTGCCTGCCTGTTGCCAAAGC R GGCAAGGTCCGCCCTGTCATC	TFF1	F ACAAGCTGCTGTACACGGACA R AAGTTTCCAGGGCCGGGAAT
GAPDH	F CTTTTGCGTCGCCAGCCGAG R CCAGGCGCCAATACGACCA	TFF2	F TCTGTCCTGCCTCCCTGATCCA R CTCTGGCAGCTGAATCCCGGT
HNF4α	F GTACTCCTGCAGATTGACCC R CTGTCCTCATAGCTTGACCT	GC	F ATGGCCAAAGAGCTGCCTGA R TGGGCAGCTGGCATGAAGTA
AQP1	F GGCCAGCGAGTTCAAGAAGAA R TCACACCATCAGCCAGGTCAT	A1AT	F TGAGGAGAGCAGGAAAGGACA R CTCAGCCAGGGAGACAGG
NKCC1	F ACCAAGGATGTGGTAGTAAGTGTGG R GGATTCTTTTTCAACAGTGGTTGA	ALDH1A	F TGAAGTCATCGGTCCGGCTTGA R ACCCTACGATGACACTTGTGCC
GGT	F TGGTGGACATCATAGGTGGGGA R ATGACGGCAGCACCTCACTT	LYZ	F GGCAAAACCCAGGAGCAGTT R TGCCACCCATGCTCTAATGCCT
ASBT	F GGTGGCCTTTGACATCCTCCC R GCATCATTCCGAGGGCAAGC	AGR2	F TCCTAGCCGCCGACTCACACA R GGTTTGACTGTGGTATCTCTGGCCA
SOX9	F ACCAGTACCCGCACTTGAC R GCGCCTGAAGATGGCGTTG	S100A6	F GGCAGGGAGGGTGACAAGCA R CCTCCTGGTCCTTGTCCGGT
EpCAM	F GACTTTTGCCGCAGCTCAGGA R AGCAGTTTACGGCCAGCTTGT	TACSTD2	F CGAGCTTGTAGGTACCCGGCG R TGCGCCGAGGAATCAGGAAGC
CFTR	F TGGCGGTCACTCGGCAATTT R TCCAGCAACCGCCAACAAT	LCN2	F ACCACATCGTCTTCCCTGTCCC R TGGCACTGTGGGGAGGGTCTC
LGR5	F GTCAGCTGCTCCCGAATCCC R TGAAACAGCTTGGGGGCACA	KRT18	F TCCAAAAGACCACCACCCGCC R TGCTCCCCAAAGTGTACCTGCT
Cyp3A4	F AGCAAAGAGCAACACAGAGCTGAA R CAGAGGTGTGGGCCCTGGAAT	CES1	F GCCCCGAACCACAGAGATGCT R AGCTCATCCCCGTGGTCTCCT
LOLX4	F CTGGGCCGGACTGACTTTTCGT R CTTGTGCCCTCAGCCACCTT	CD133	F CCTGGGGCTGCTGTTTATTA R ATCACCAACAGGGAGATTGC
MMP1	F GGCCCAAAACCCAAAAGCG R CGGGTAGAAGGGATTTGTGCGC	LGR5	F GTCAGCTGCTCCCGAATCCC R TGAAACAGCTTGGGGGCACA
AGR2	F TCCTAGCCGCCGACTCACACA R GGTTTGACTGTGGTATCTCTGGCCA		
CTSE	F ATTGGCTCCCCACCACAGAA R CTGGACTGGGAAGGCTGGAA		

Table S3. Overview of cholangiocyte organoids cultured in this manuscript.

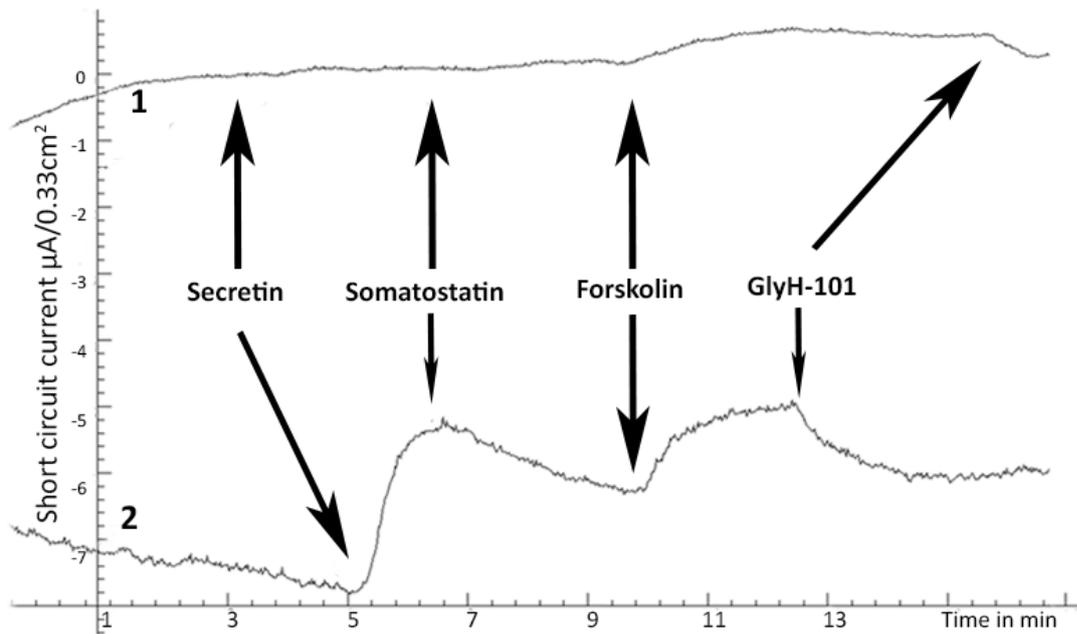
Name	Abbreviation	Source	Culturing conditions	Protocol/reference
Bile Cholangiocyte Organoid in canonical Wnt stimulating conditions	cBCO	Bile	Canonical Wnt	Soroka <i>et al.</i> ¹
Intrahepatic Cholangiocyte Organoid in canonical Wnt stimulating conditions	cICOs	Intrahepatic cholangiocytes obtained via a liver biopsy	Canonical Wnt	Huch <i>et al.</i> ² ; Broutier <i>et al.</i> ³
Extrahepatic Cholangiocyte Organoid in canonical Wnt stimulating conditions	cECOs	Extrahepatic cholangiocytes obtained from an extrahepatic bile duct biopsy	Canonical Wnt	Rimland <i>et al.</i> ⁴
Extrahepatic Cholangiocyte Organoid in non-canonical Wnt stimulating conditions	ncECO	Extrahepatic cholangiocytes	Non-canonical Wnt	Sampaziotis <i>et al.</i> ⁵ ; Tysoe <i>et al.</i> ⁶
Bile Cholangiocyte Organoid in non-canonical Wnt stimulating conditions	ncBCO	Bile	Non-canonical Wnt	This protocol

Figures



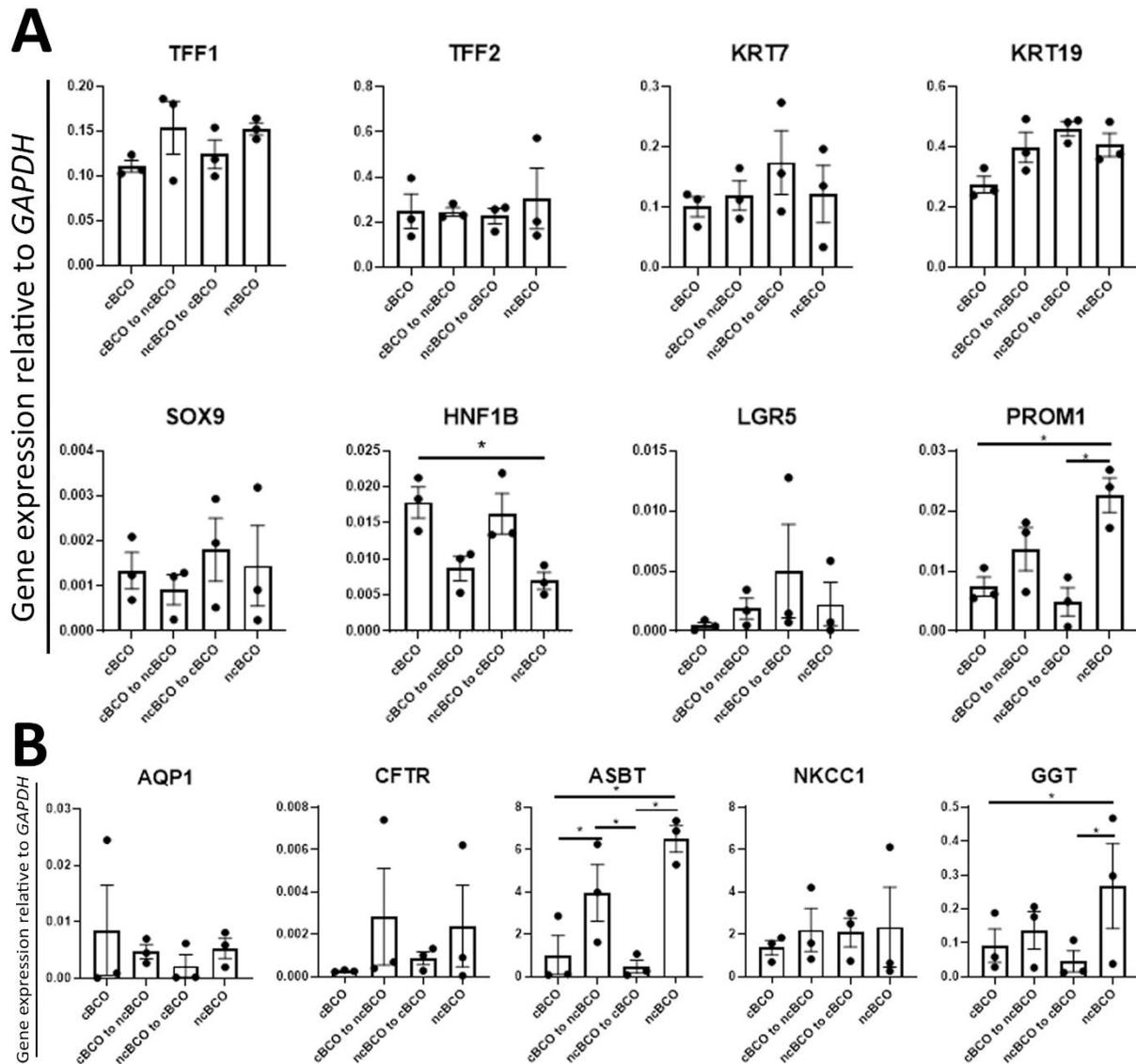
Supplementary Figure 1. Classical cholangiocyte markers are similar expressed between non-canonical Wnt and canonical-Wnt stimulating conditions.

(a) qRT-PCR showing comparison between ncBCOs (n=6) and cBCOs (n=6) for different cholangiocyte- and stemness/Wnt-target related genes, relative to the housekeeping gene *GAPDH*, error bars are presented with standard error of the mean. (b) Gene-expression as analysed by Micro Array for ncECOs (n=3) and cECOs (n=3) for the similar genes as displayed in panel a. Color key represents the log₂ transformed signal intensities after variance stabilizing normalization.



Supplementary Figure 2. Cholangiocyte related and ion-channel and basolateral receptor activity in ncECOs and cICOs.

Representative ion-channel functionality of 2D-grown tissue derived canonical-Wnt stimulated organoids (cICOs, line 1) and tissue derived non-canonical Wnt stimulated organoids (ncECOs, line 2) in an Ussing-chamber. Stimulation with cAMP-activator(forskolin), resulted in an increase in short circuit current, however secretin stimulation (to the basolateral side) only gave a response in the ncECOs. In similar fashion, somatostatin (basolateral addition) only give a response in ncECOs and not in cICOs, while Cystic Fibrosis Transmembrane conductance Regulator (CFTR)-inhibitor, GlyH-101 (luminal addition), resulted in an inhibition of the channel in both organoid-types. Indicating the presence of functional CFTR channels in both organoids, but only somatostatin and secretin-receptors are functional in ncECOs. Moreover, the CFTR responses in ncECOs seems more pronounced compared to cICOs.



Supplementary Figure 3. Gene-expression profiles of bile cholangiocyte organoids under different culture conditions.

(a) Gene-expression profiles as analysed by qRT-PCR for BCOs started up in either canonical Wnt (cBCOs) or non-canonical Wnt culture (ncBCOs) conditions for the classical cholangiocyte and stem cell/Wnt-target genes. After three passages pallet was split in half and organoids were either continued in their original conditions or switch to the other ($n=3$, all conditions). Error bars indicate standard error of the mean. **(b)** Gene-expression profiles of the cholangiocyte markers related to function as analysed by qRT-PCR for the same samples as displayed in panel a. Error bars indicate standard error of the mean. *is considered a significant difference ($p<0.05$).

Supplementary references

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2. Huch M, Gehart H, van Boxtel R, *et al.* Long-Term Culture of Genome-Stable Bipotent Stem Cells from Adult Human Liver. *Cell*. 2015 Jan 15; 160(1-2): 299–312.
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