

Figure S1. The figure shows 12 of the 13 intersected genes that didn't reach the statistical significance for survival analysis. Patients were divided into two groups based on the median expression value of each gene respectively. Overall survival analysis for patients with different expression levels of (A) PDZK1IP1, (B) MMP7, (C) CEACAM6, (D) CEACAM7, (E) KRT19, (F) LAMC2, (G) SPP1, (H) RAB25, (I) UBD, (J) DCDC2, (K) EPCAM and (L) MMP11. OS, overall survival; PDZK1IP1, PDZK1 Interacting Protein 1; MMP7, matrix metalloproteinase 7; CEACAM6, carcinoembryonic antigen related adhesion molecules 6; MUC1, mucin 1; CEACAM7, carcinoembryonic antigen related adhesion molecules 7; KRT19, keratin 19; LAMC2, laminin subunit gamma-2; SPP1, secreted phosphoprotein 1; RAB25, Ras Genes from Brain Protein 25; UBD, ubiquitin D; DCDC2, doublecortin domain containing 2; EPCAM, epithelial cell adhesion molecule; MMP11, matrix metalloproteinase 11.

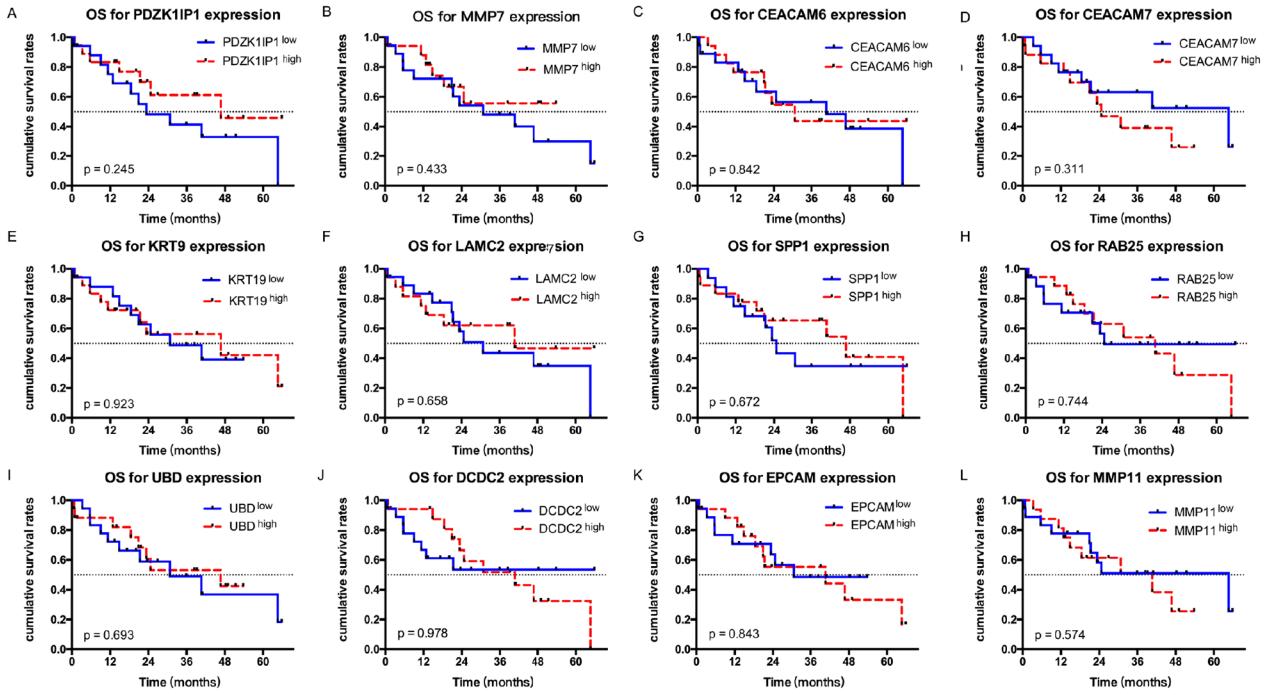


Table SI. PCR primers used in the present study.

Gene symbol	Forward primer	Reverse primer	Amplicon length (bp)
GAPDH	5'-GTCAGCCGCATCTTCTTTG-3'	5'-TTAAAAGCAGCCCTGGTGAC-3'	132
MUC1	5'-GGCATTGGGCTCCTTCTT-3'	5'-TGGAGTGGTAGTCGATGCTAAG-3'	87
MUC1, mucin 1.			

Table SII. List of dysregulated genes associated with cholangiocarcinoma identified by robust rank aggregation.

A, Upregulated genes

COL1A1, PDZK1IP1, MMP7, SPP1, CEACAM6, UBD, RAB25, MUC1, CEACAM7, SULF1, KRT7, KRT19, MMP11, FAP, DCDC2, EPCAM, COL1A2, VCAN, LAMC2

B, Downregulated genes

TAT, MT1M, MT1G, FBP1, BHMT, METTL7A, ADH1C, MT1H, ALDOB, MTTP, MT1X, APOC3, SEPP1, MT1E, OTC, ADH4, APOF, FOLH1, GSTA2, APOA1, CPS1, GBA3, SERPINC1, SLC22A1, HPD, HMGCS2, KMO, HSD17B13, SULT2A1, MBL2, MT1F, XDH, SLC27A5, PCK1, AHSG, CIDEB, SELENBP1, HSD17B6, DNASE1L3, F9, CYP4A11, GC, MT1A, CYP4F2, C8B, GLYAT, CREB3L3, HPR, INSIG1, GSTA1, SLC10A1, CYP3A4, HGFAC, AFM, RBP4, EPHX2, HAMP, PON3, HAO1, FMO5, CYP2A6, F12, SARDH, ADH1A, DCXR, UGT2B10, FTCD, HBB, SLC27A2, SLC7A2, HPX, CYP2C9, ARG1, DAK, HRG, F2, SERPIND1, ITIH4, RBP5, ABCG8, GCGR, ORM1, MAT1A, ALAS1, OIT3, BCHE, MLXIPL, CYB5A, ADH1B, APOH, KHK, C4BPA, SPP2, SLC2A2, UGT2B11, HP, ACOX2, SLCO1B1, CPB2, ANG, KNG1, AQP9, TM6SF2, SERPINF2, SLC39A5, CRYL1, ACSL1, AASS, CYP2C8, FETUB, ADH6, TTR, PLA2G12B, C6, GNMT, CTH, HRSP12, ITIH1, HAO2, PLG, CYP3A43, ABCB4, FST, PRG4, CYP8B1, CYP27A1, G6PC, ORM2, C8A, GADD45G

Table SIII. Continued.

B, Downregulated genes (n=705)

PRAMEF33, AC007406.1, CYP3A43, AC090796.1, BX547991.1, AC244394.1, AC007298.1, MIR4290HG, AC092071.1, C2, AKR1C8P, NTF3, ABCB4, SORD, AMDHD1, FBP1, ADH4, OIT3, LINC01018, LINC02153, AC025423.3, GLYAT, AC004160.2, ADH1A, LINC01780, PITPNM2-AS1, ANGPTL6, AP006216.2, MT1CP, SLC17A2, AC063919.1, GSTA12P, SAA1, KNG1, ABCG5, LINC02037, GPAM, KRT16P3, CHRM2, SEC14L2, ASS1P2, AC120042.1, UGT2B24P, MAT1A, XAGE3, CYP17A1, LECT2, AL663023.1, CFHR2, ITIH1, PLG, UBTFL10, AOC4P, AC026461.2, AC024559.2, SLC13A5, CYP2C8, TSLP, TMEM220, SPP2, AP001065.3, AC112206.2, LINC01818, HAO2, PLGLA, PLGLB1, AC090227.2, SAA4, U91324.1, AC080129.1, AZGP1P2, DCXR, HAAO, ADH1B, CYP2A7P1, MIR146B, AC007298.2, AC005495.1, KCNN2, SARDH, HFE2, AC008549.1, LINC01714, SLC6A12, TPRG1-AS1, MYRIP, EVA1A, AC006329.2, COX6A2, CPS1, CNDP1, SLC2A2, PRAMEF30P, FCN3, GBA3, AL592182.2, CES5A, RD3L, TCTEX1D1, AC137723.2, CLEC4G, AKR1C6P, RNU1-70P, OTC, PPP1R1A, ABCB11, CYP2B6, ORM2, AHSG, APOC4-APOC2, RTL4, FXYD1, HPX, PRAMEF10, KMO, CPS1-IT1, MASP1, AC083902.1, AC097063.1, AC080128.1, CFHR3, G6PC, MARCO, ADH6, AGXT, MT1H, LCAT, APOA4, AC138430.1, INHBC, ADORA2BP1, F12, TDO2, HSD11B1, MT1A, LDLRAD4-AS1, AC005304.2, UGT2B27P, C4BPA, ALB, SAA2, ALDOB, HAMP, SERPINC1, AL391095.2, AC007423.1, NR1I2, AC104809.1, GMNC, AC004160.1, APOA1, CYP8B1, LRRC37A7P, ASCL1, TTPA, FTCD, HULC, AC244100.1, ARL4AP3, SLC2A9, AC011591.1, MTND4P20, GOLGA6A, AL355877.1, JAKMIP2-AS1, ACOX2, TM6SF2, CPN2, AP001781.2, STAB2, AC068631.1, AGMAT, HGFAC, AC121758.1, MT2A, PHYHD1, SYT9, AR, MTND6P4, HAO2-IT1, AC107396.1, LINC01847, AC006254.1, SLC22A10, PZP, AC087392.1, AC106822.1, MEP1B, AP000851.2, AOX1, CP, KCNJ8, HSD17B6, UGT1A3, EHHADH, HEPACAM, CES1, PIPOX, MT2P1, SERPINA10, INSIG1, AC022784.6, FAM99A, CYP3A4, BAAT, APOC1, TPP2, SLC27A5, ACSM2B, PRODH2, AC244100.4, AC083841.1, AP000355.1, GLS2, C6, PAH, C9, AC027688.1, LINC02348, AC021744.1, RFPL1, AC008250.2, PDE3B