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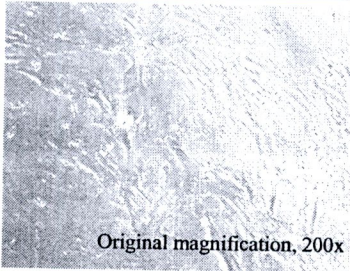
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APPENDICES

APPENDIX A
PRIMARY CULTURE FIBROBLASTS AND HUMAN BILIARY
EPITHELIAL CELL LINES

A-1 Primary culture fibroblasts

Cell Name	Cf1
Species	<i>Homo Sapiens</i>
Sex	Male
Age	56-year-old
Diagnosis	Cholangiocarcinoma
Histological type	Papillary
Establisher	Chuaysri, C. et al. 2009
Medium	10% FBS containing DMEM with 20% epidermal growth factor
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C



Cell Name	Cf2
Species	<i>Homo Sapiens</i>
Sex	Male
Age	44-year-old
Diagnosis	Cholangiocarcinoma
Histological type	Papillary
Establisher	Chuaysri, C. et al. 2009
Medium	10% FBS containing DMEM
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C

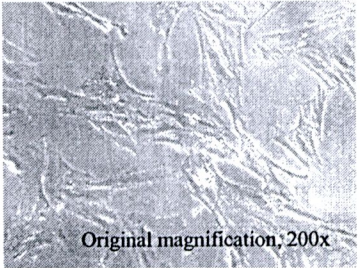


Cell Name	Lf1
Species	<i>Homo Sapiens</i>
Sex	Male
Age	56-year-old
Diagnosis	Cholangiocarcinoma
Histological type	Papillary
Establisher	Chuaysri, C. et al. 2009
Medium	10% FBS containing DMEM with 20% epidermal growth factor
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C



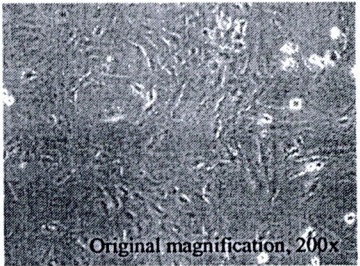
A-1 Primary culture fibroblasts (Cont.)

Cell Name	Lf2
Species	<i>Homo Sapiens</i>
Sex	Male
Age	44-year-old
Diagnosis	Cholangiocarcinoma
Histological type	Papillary
Establisher	Chuaysri, C. et al. 2009
Medium	10% FBS containing DMEM
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C



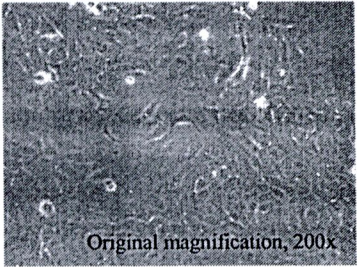
A-2 Non-tumorigenic biliary epithelial cells

Cell Name	MMNK1
Species	<i>Homo Sapiens</i>
Source	SV40T-transduced human liver OUMS-21 cells
Establisher	Maruyama et al., 2004
Medium	10% FBS containing HAM F-12
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C

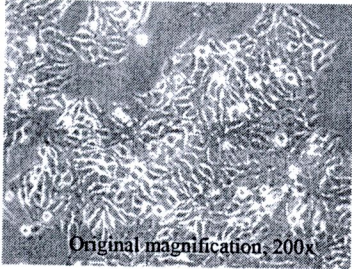
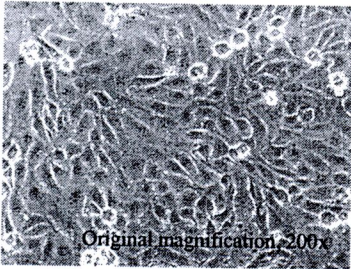
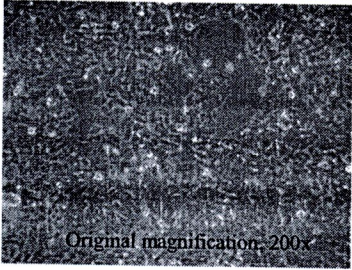


A-3 Tumorigenic biliary epithelial cells

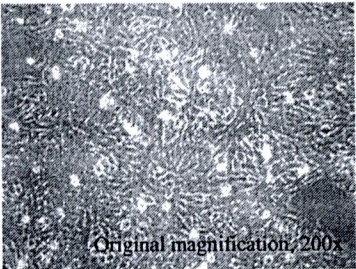
Cell Name	KKU-M213
Species	<i>Homo Sapiens</i>
Sex	Male
Age	58-year-old
Diagnosis	Cholangiocarcinoma
Histological type	Well-differentiated
Establisher	Sripa, B. et al.
Medium	10% FBS containing HAM F-12
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C

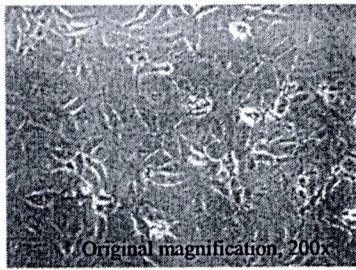


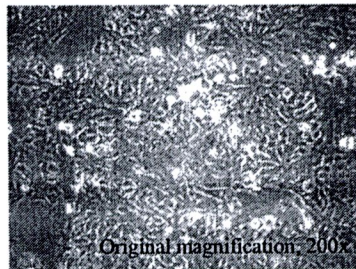
A-3 Tumorigenic biliary epithelial cells (Cont.)

Cell Name	KKU-OCA17	
Species	<i>Homo Sapiens</i>	
Sex	Male	
Age	38-year-old	
Diagnosis	Cholangiocarcinoma	
Histological type	Well-differentiated	
Establisher	Sripa, B. et al.	
Medium	10% FBS containing HAM F-12	
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C	
Cell Name	KKU-M214	
Species	<i>Homo Sapiens</i>	
Sex	Male	
Age	52-year-old	
Diagnosis	Cholangiocarcinoma	
Histological type	Well-differentiated	
Establisher	Sripa, B. et al.	
Medium	10% FBS containing HAM F-12	
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C	
Cell Name	KKU-M156	
Species	<i>Homo Sapiens</i>	
Sex	Male	
Age	68-year-old	
Diagnosis	Cholangiocarcinoma	
Histological type	Moderately-differentiated	
Establisher	Sripa, B. et al.	
Medium	10% FBS containing HAM F-12	
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C	

A-3 Tumorigenic biliary epithelial cells (Cont.)

Cell Name	KKU-M055	 Original magnification, 200x
Species	<i>Homo Sapiens</i>	
Sex	Male	
Age	56-year-old	
Diagnosis	Cholangiocarcinoma	
Histological type	Moderately-differentiated	
Establisher	Sripa, B. et al.	
Medium	10% FBS containing HAM F-12	
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C	

Cell Name	KKU-M139	 Original magnification, 200x
Species	<i>Homo Sapiens</i>	
Sex	Female	
Age	53-year-old	
Diagnosis	Cholangiocarcinoma	
Histological type	Moderately-differentiated	
Establisher	Sripa, B. et al.	
Medium	10% FBS containing HAM F-12	
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C	

Cell Name	KKU-100	 Original magnification, 200x
Species	<i>Homo Sapiens</i>	
Sex	Female	
Age	65-year-old	
Diagnosis	Cholangiocarcinoma	
Histological type	Poorly-differentiated	
Establisher	Sripa, B. et al.	
Medium	10% FBS containing HAM F-12	
Passage Method	Cells treated with 0.25% Trypsin-EDTA for 5 min at 37°C	

APPENDIX B
RAW DATA OF PN EXPRESSION IN CCA TISSUES

B-1 Raw data of PN expression in 52 CCA tissues

No.	Patient code	Age (yr)	Sex	Tumor size (cm)	Tumor staging	Histological type	Vascular invasion	Lymph node metastasis	Survival (d)	Status 4/10/07	Grading (IxA)	Level of PN expression*
1	M005	75	F	3	4	WD	Absence	Presence	277	Death	0	Low
2	M009	57	M	5	4B	MD	Absence	Absence	119	Death	6	High
3	M012	61	F	3	4	MD	Absence	Absence	118	Death	9	High
4	M014	70	M	2	3	WD	Absence	Absence	215	Death	4	Low
5	M030	37	F	12	3	Pap	Absence	Absence	245	Death	1	Low
6	M037	60	M	5	4B	Pap	Absence	Presence	429	Death	4	Low
7	M040	54	F	7	4	MD	Absence	Absence	425	Death	6	High
8	M043	50	M	9	4B	WD	Absence	Absence	191	Death	9	High
9	M055	56	M	7	4B	MD	Absence	Absence	34	Death	9	High
10	M057	61	M	10	4B	WD	Absence	Absence	211	Death	6	High
11	M068	54	M	5	4A	PD	Absence	Absence	395	Death	4	Low
12	M071	46	M	8	4B	WD	Absence	Presence	50	Death	6	High
13	M079	55	M	2	4A	WD	Absence	Absence	297	Death	6	High
14	M082	53	M	5	4A	Pap	Absence	Absence	637	Death	4	Low
15	M090	53	M	7	4A	Pap	Absence	Absence	294	Death	9	High
16	M097	58	M	2	2	WD	Absence	Absence	225	Death	0	Low
17	M101	64	M	9	4A	MD	Absence	Absence	51	Death	6	High
18	M102	56	M	7	4B	PD	Absence	Presence	35	Death	0	Low

B-1 Raw data of PN expression in 52 CCA tissues (Cont.)

No.	Patient code	Age (yr)	Sex	Tumor size (cm)	Tumor staging	Histological type	Vascular invasion	Lymph node metastasis	Survival (d)	Status 4/10/07	Grading (IxA)	Level of PN expression*
19	M114	61	M	3	4B	Pap	Absence	Absence	26	Death	9	High
20	M117	72	M	3.5	4B	PD	Absence	Absence	140	Death	9	High
21	M123	55	M	4	4B	WD	Absence	Absence	10	Death	6	High
22	M131	63	F	3	4B	Pap	Absence	Absence	1011	Death	0	Low
23	M136	49	F	5.5	3	WD	Absence	Absence	308	Death	9	High
24	M137	53	M	4	4	WD	Absence	Absence	3010	Death	4	Low
25	M139	53	F	5.5	4	MD	Presence	Presence	14	Death	6	High
26	M140	45	F	4.5	4B	WD	Absence	Presence	313	Death	9	High
27	M148	56	M	13	2	PD	Presence	Absence	214	Death	2	Low
28	M152	55	M	4.5	4B	MD	Presence	Absence	514	Death	0	Low
29	M155	52	M	8	4B	WD	Absence	Absence	358	Death	9	High
30	M157	72	F	8	4B	PD	Presence	Presence	139	Death	3	Low
31	M171	65	F	5	4B	WD	Absence	Presence	145	Death	6	High
32	M174	59	F	3	1	Pap	Absence	Absence	3146	Alive	0	Low
33	M189	67	M	7.5	3	Pap	Absence	Absence	2465	Death	2	Low
34	M209	61	F	5	4A	PD	Presence	Presence	157	Death	9	High
35	M213	58	M	8	4A	Pap	Presence	Presence	45	Death	9	High
36	M236	61	F	12	2	Pap	Absence	Absence	1277	Death	9	High

B-1 Raw data of PN expression in 52 CCA tissues (Cont.)

No.	Patient code	Age (yr)	Sex	Tumor size (cm)	Tumor staging	Histological type	Vascular invasion	Lymph node metastasis	Survival (d)	Status 4/10/07	Grading (IxA)	Level of PN expression*
37	M240	69	M	5	4B	WD	Absence	Presence	202	Death	1	Low
38	M250	66	M	8	4B	MD	Absence	Presence	76	Death	9	High
39	M259	63	M	4	3	WD	Absence	Absence	2505	Death	6	High
40	M266	62	M	7	4	Pap	Absence	Absence	179	Death	9	High
41	M270	55	M	5	4B	WD	Absence	Presence	149	Death	6	High
42	M273	60	F	6	4B	WD	Presence	Absence	60	Death	6	High
43	M274	47	F	5	4B	Pap	Absence	Absence	189	Death	6	High
44	M275	49	M	1.5	4B	WD	Absence	Absence	2935	Alive	6	High
45	M294	51	F	5	3	PD	Presence	Absence	2893	Alive	0	Low
46	M303	52	M	2	4B	WD	Absence	Absence	94	Death	4	Low
47	M311	41	M	3	4B	Pap	Absence	Presence	329	Death	6	High
48	M319	67	F	4.5	4B	PD	Absence	Absence	1743	Death	0	Low
49	N019	62	M	10	4B	Pap	Presence	Absence	2143	Death	9	High
50	P060	64	F	8	4B	Pap	Absence	Presence	1407	Death	1	Low
51	P078	59	F	4	4A	WD	Presence	Absence	124	Death	2	Low
52	QO66	51	F	8	4B	WD	Presence	Absence	1850	Alive	0	Low

M: Male, F: Female, WD; Well-differentiated adenocarcinoma, MD; Moderately-differentiated adenocarcinoma, PD; Poorly-differentiated adenocarcinoma and Pap; Papillary carcinoma. I: intensity, A: % positive area, * Low expression: grade 0-4; High expression: grade 5-9

B-2 Raw data of PN expression in benign and HCC tissues

No.	Patient code	Age (yr)	Sex	Diagnosis	Grading (IxA)	Level of PN expression*
1	Q067	44	F	Cavernous haemangioma	3	Low
2	R082	51	M	Chronic inflammation	2	Low
3	T075	32	F	Liver necrosis with fibrosis	3	Low
4	U115	59	M	Calcified necrotic tissue with granulomatous reaction	1	Low
5	X070	60	F	possible dead parasite Chronic cholecystitis.	0	Low
6	X086	39	F	No tumor is identified	0	Low
7	X128	66	M	Biliary cystadenoma	1	Low
8	X180	54	M	Cavernous haemangioma	4	Low
9	R140	49	M	HCC	1	Low
10	R141	33	M	HCC	2	Low
11	W001	55	M	HCC	4	Low
12	W088	47	M	HCC	0	Low

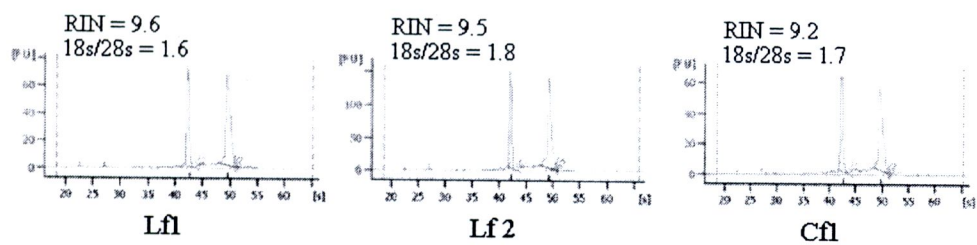
M: Male, F: Female, WD; Well-differentiated adenocarcinoma, MD; Moderately-differentiated adenocarcinoma, PD; Poorly-differentiated adenocarcinoma and Pap; Papillary carcinoma .I: intensity, A: % positive area, * Low expression: grade 0-4; High expression: grade 5-9

APPENDIX C
RNA EXTRACTION AND MELTING CURVE ANALYSIS
OF REAL TIME PCR

C-1 Quality analysis of RNA extracted from fibroblasts

Total RNA was extracted from $3\text{--}5 \times 10^5$ cells of Cf, Lf1 and Lf2, using RNeasy Micro Kit as the manufacturer’s instruction. The quality of RNA was assessed by an Agilent RNA 6000 Nano Kit. RNA integrity number (RIN) of all 3 samples showed more than 9.0 (9.6 for Lf1, 9.5 for Lf2 and 9.2 for Cf1) (Fig C-1A) and 18S/28S ratio of ribosomal RNA showed nearly to 1.8 (1.6 for Lf1, 1.8 for Lf2 and 1.7 for Cf1). Herein, these RNA samples were accepted for the good quality of RNA to perform gene expression analysis.

A



B

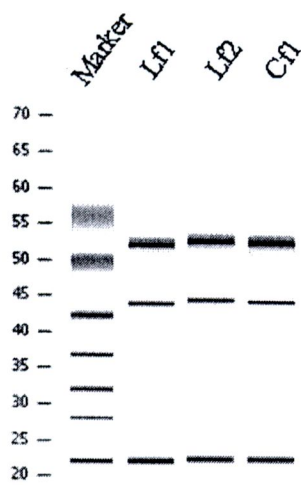
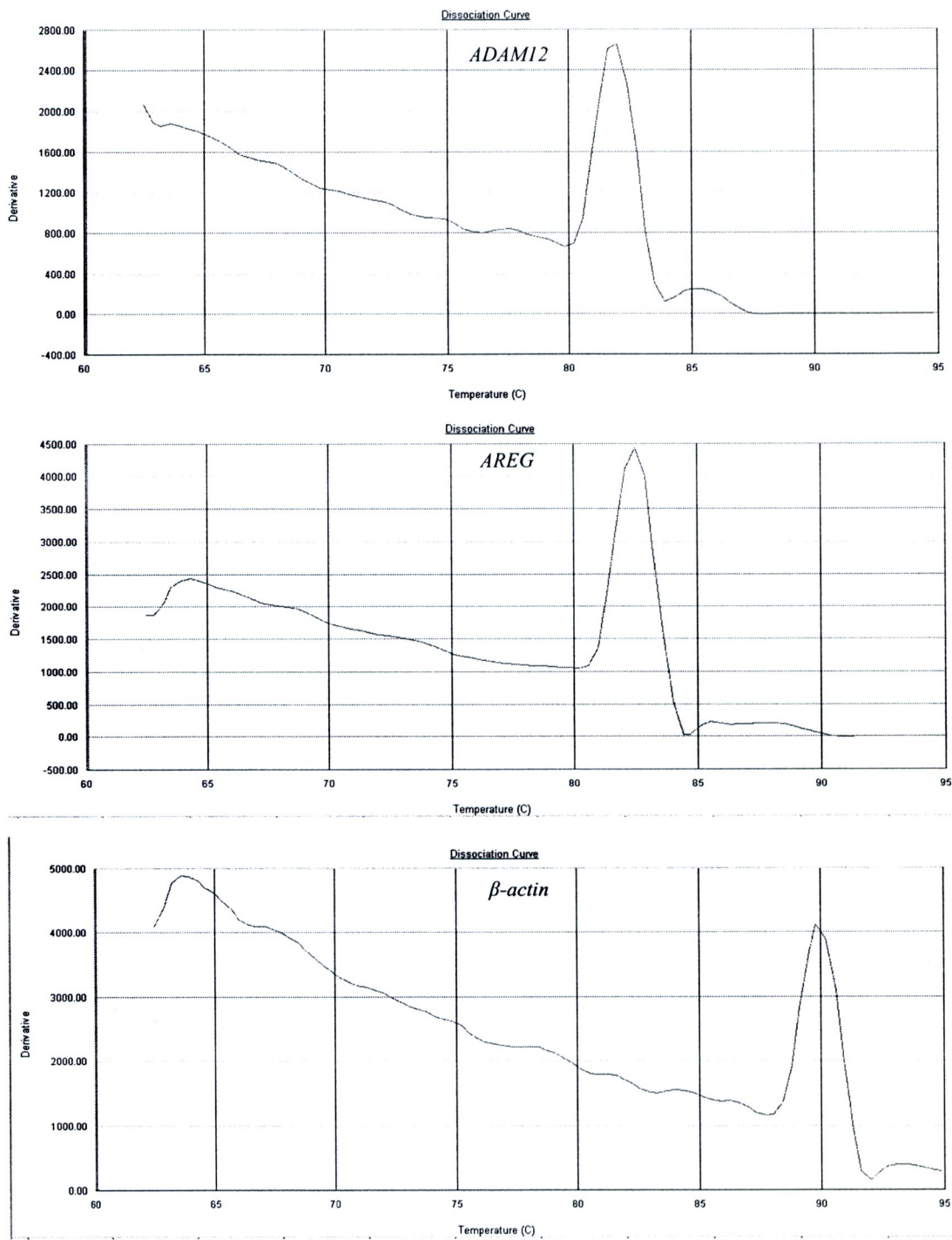
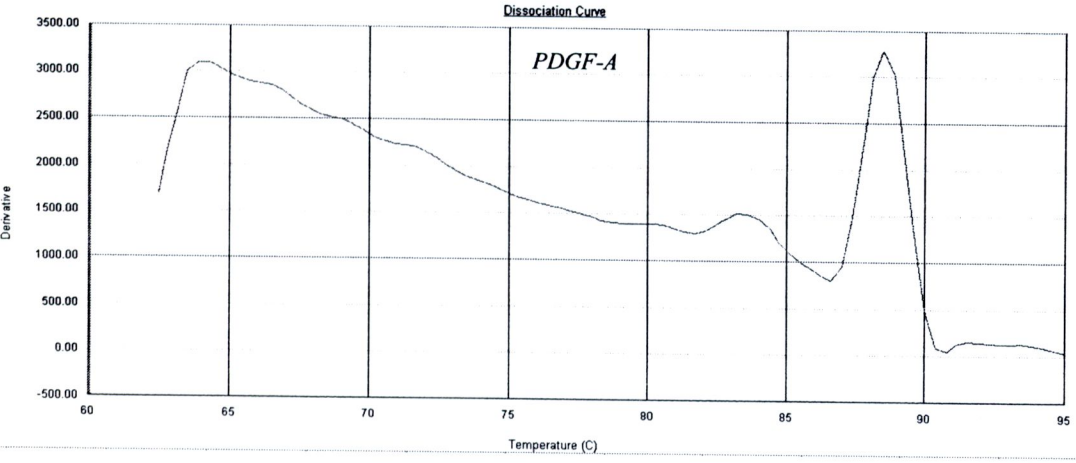
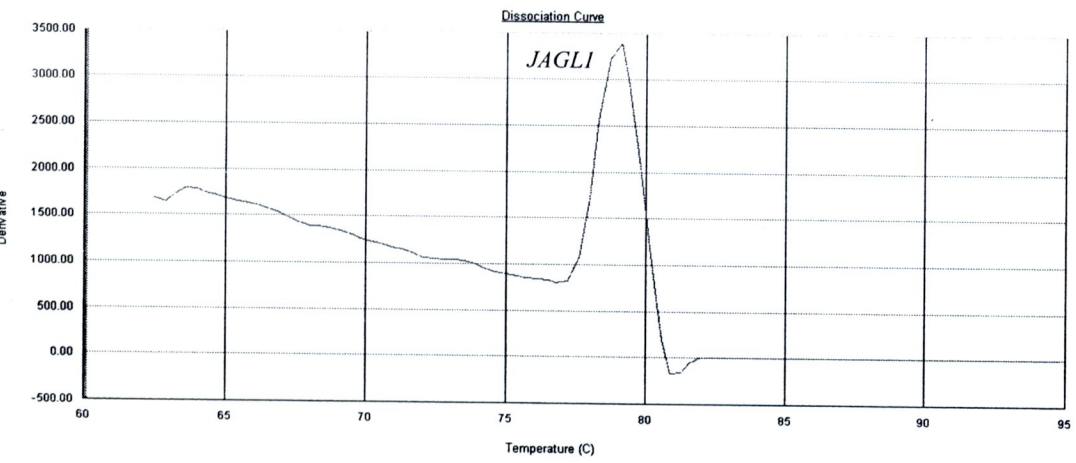
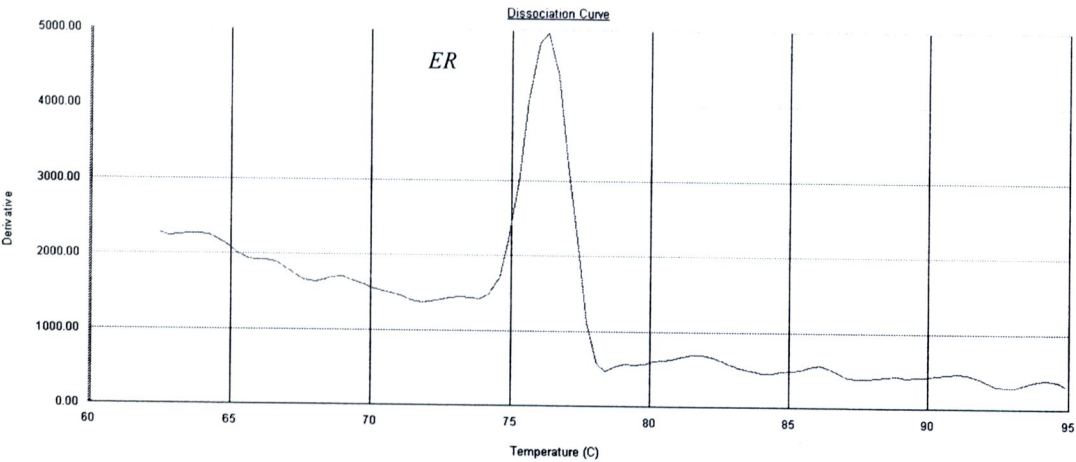


Figure C-1 Quality analysis of RNA extracted from fibroblasts. The RIN value and 18S:28S ratio of all 3 samples are shown (A). Gel electrophoresis shows bands of 18S and 28S ribosomal RNA compared to marker (B).

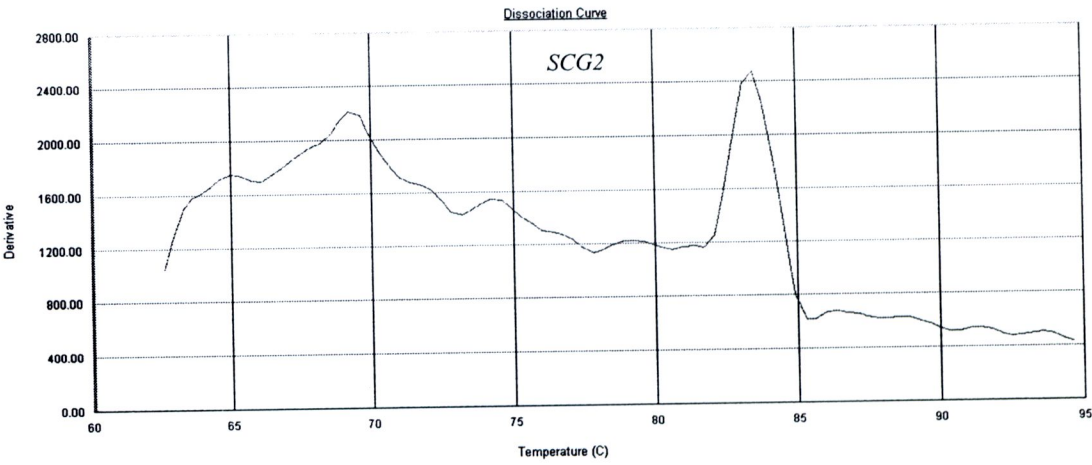
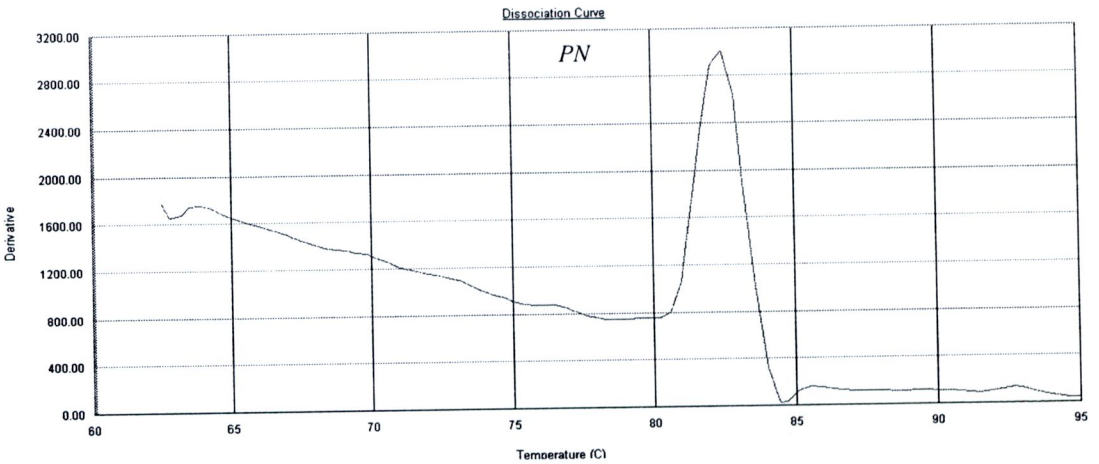
C-2 Melting curve analysis by ABI 7500 machine



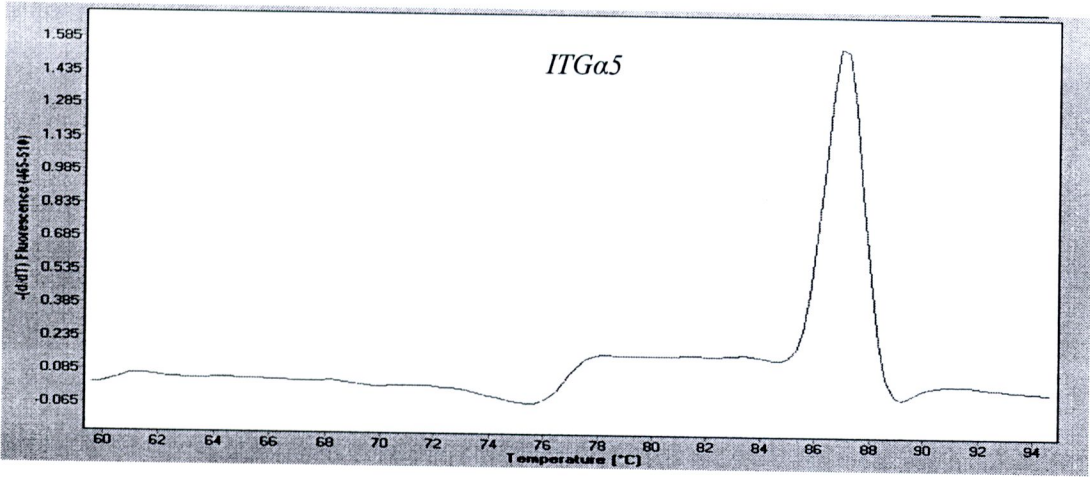
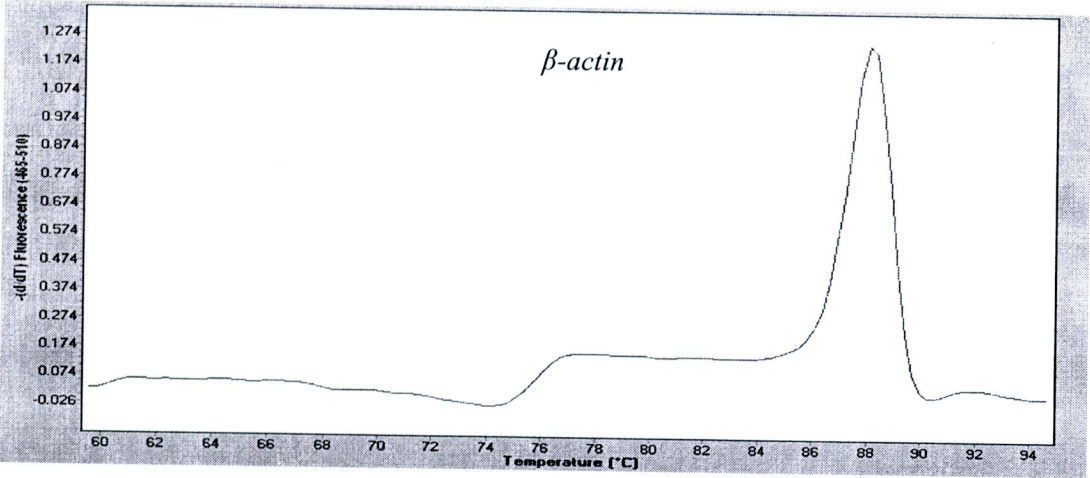
C-2 Melting curve analysis by ABI 7500 machine (Cont.)



C-2 Melting curve analysis by ABI 7500 machine (Cont.)

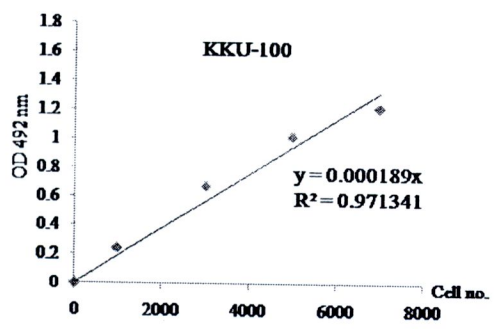
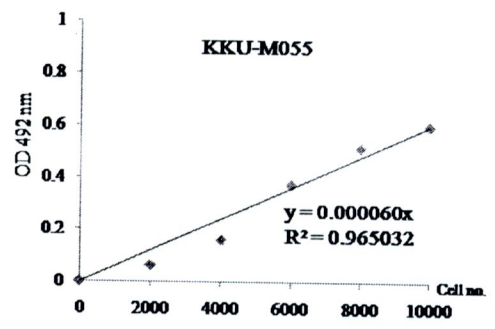
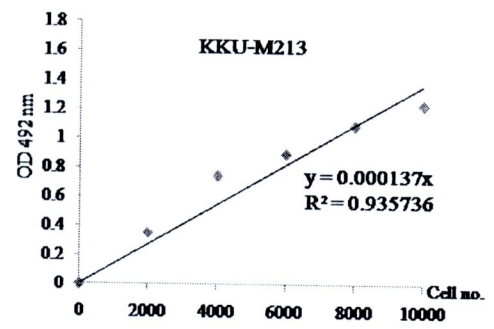
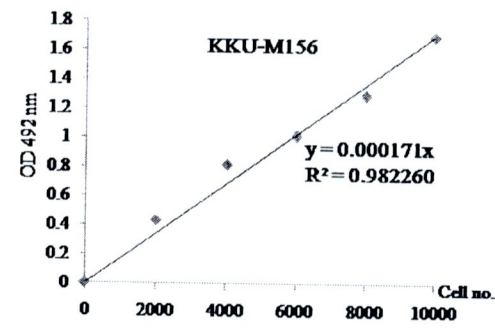


C-3 Melting curve analysis by LightCycler® 480 system



APPENDIX D
STANDARD CURVE OF CELL PROLIFERATION ASSAY

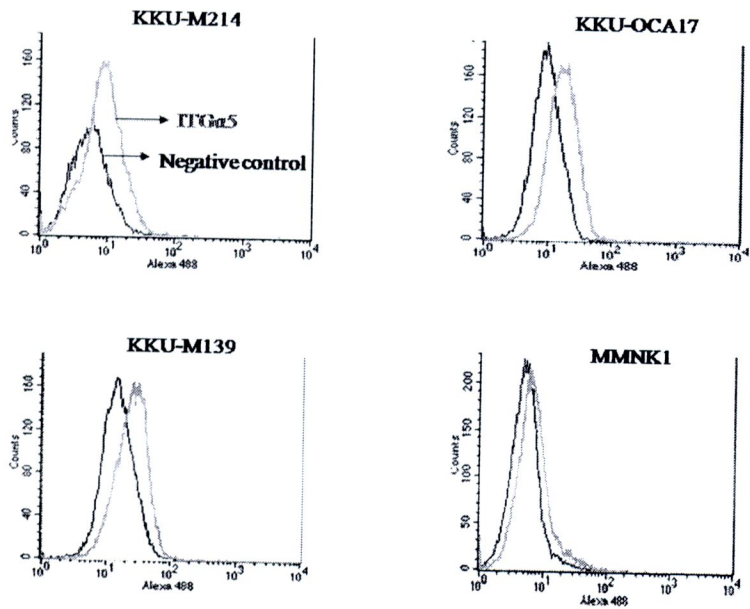
D-1 Standard curve of CCA cell lines measured by MTS assay



APPENDIX E

**RAW DATA OF ITG α 5 DETECTION ON BILIARY EPITHELIAL
CELLS USING FLOW CYTOMETRY**

E-1 Histogram plots of cell numbers in channel FL-1 (Alexa 488)



E-2 Raw data of MFI measured for ITGa5 signal

Biliary epithelial cell lines	Relative MFI (Normalized by negative control)
KKU-M213	2.0 ± 0.57
KKU-M156	1.4 ± 0.09
KKU-M055	1.6 ± 0.26
KKU-100	1.5 ± 0.21
KKU-M214	1.7 ± 0.22
KKU-OCA17	1.5 ± 0.24
KKU-M139	1.3 ± 0.42
MMNK1	1.5 ± 0.23

APPENDIX F
SECRETOME ANALYSIS OF CCA-ASSOCIATED
FIBROBLASTS

Introduction

In addition to whole gene expression analysis of CCA-associated fibroblasts, the proteomic analysis of the substances released from these fibroblasts were performed in order to understand the possible effect that fibroblasts act paracrinely on the neighboring cancer cells or even autocrinely on themselves.

Material and methods

Cf2 and Lf2 were cultured in DMEM supplemented with 10% FBS until reach 80-90% confluence and then switched to 1% FBS containing DMEM for 48 h. The 48-h conditioned media (CM) was collected and centrifuged to get rid of cell debris.

The 1.4 mg Cf-CM and Lf-CM were subjected to RayBio® Biotin Label-based Human Antibody Array I (RayBiotech, Inc, Norcross, GA, USA) containing 507 human proteins followed the company's instruction. The signal was scanned using Axon GenePix scanner (RayBiotech, Inc, Norcross, GA, USA) and the data was analyzed by GeneSpring G.X. 7.3 (Agilent Technologies, Waldbronn, Germany).

Results

Among 507 proteins, 123 proteins were up-regulated for 2-fold or more in the Cf-CM compared to those of Lfs. These proteins were categorized into several biological functions including cytokine (63.4%), receptor (17.1%), growth factor (16.3%), matrix metalloproteinases (1.6%) and phosphorylated protein (1.6%) (Table F-1). Whereas 70 secreted proteins were down-regulated with less than 1-fold in Cf-CM compared to that of Lf-CM (Table E-2).

Table F-1 List of up-regulated proteins in Cf-CM

No	Biological function	No. of protein (%)	List of protein	Fold change (Cf/Lf)
1	Cytokine	63.4	- Endoglin / CD105	8.4
			- Lymphotactin/ XCL1	6.2
			- CXCL14 / BRAK	3.8
			- VEGF	3.7
			- Cripto-1	3.6
			- Pentraxin3 / TSG-14	3.3
			- WIF-1	3.1
			- Orexin A	3.0
			- TRAIL R4 / TNFRSF10D	3.0
			- GREMLIN	2.9
			- DcR3 / TNFRSF6B	2.9
			- MIF	2.9
			- Tarc	2.8
			- NeuroD1	2.8
			- Siglec-5 (CD170)	2.8
			- ENA-78	2.8
			- TRAIL/ TNFSF10	2.7
			- Activin A	2.7
			- ICAM-1	2.6
			- 6Ckine	2.6
			- uPAR	2.6
			- P-selectin	2.6
			- SMDF / NRG1Isoform	2.6
			- Activin RIB / ALK-4	2.5
			- Osteoactivin / GPNMB	2.5
			- FAM3B	2.4
			- RANK / TNFRSF11A	2.4
			- ALCAM	2.4
			- Thymopoietin	2.4
			- Thrombospondin-4	2.4
			- IL-31	2.4
			- MCP-2	2.4
			- MIP-3 beta	2.3
			- IL-16	2.3

Table F-1 List of up-regulated proteins in Cf-CM (Cont.)

No	Biological function	No. of protein (%)	List of protein	Fold change (Cf/Lf)
1	Cytokine (Cont.)	63.4	- Dkk-4	2.3
			- GRO-a	2.3
			- Galectin-3	2.3
			- VCAM-1 (CD106)	2.3
			- WISP-1 / CCN4	2.3
			- LECT2	2.3
			- Prolactin	2.2
			- IL-1 ra	2.2
			- Inhibin A	2.2
			- Sonic Hedgehog (Shh N- terminal)	2.2
			- PECAM-1 (CD31)	2.2
			- MDC	2.2
			- MIG	2.2
			- TACI / TNFRSF13B	2.2
			- IL-1 sRI	2.1
			- VEGF-B	2.1
			- Insulin	2.1
			- SDF-1 / CXCL12	2.1
			- IL-31 RA	2.1
			- Endostatin	2.1
			- IL-22 BP	2.1
			- L-Selectin (CD62L)	2.1
			- PDGF-C	2.1
			- FGF R5	2.1
			- MPIF-1 / CCL23	2.1
			- TSLP	2.1
			- Cerberus 1	2.1
			- IGFBP-1	2.1
			- BCMA / TNFRSF17	2.1
			- sgp130	2.1
			- Dkk-3	2.1
			- Kremen-1	2.1
			- ICAM-2	2.1

Table F-1 List of up-regulated proteins in Cf-CM (Cont.)

No	Biological function	No. of protein (%)	List of protein	Fold change (Cf/Lf)
1	Cytokine (Cont.)	63.4	- IL-6	2.0
			- CTLA-4 (CD152)	2.0
			- Latent TGF-beta bp1	2.0
			- sFRP-3	2.0
			- Soggy-1	2.0
			- TRAIL R3 / TNFRSF10C	2.0
			- M-CSF	2.0
			- CRIM 1	2.0
			- TRAIL R1 / DR4 /	2.0
			- TNFRSF10A	2.0
			- TMEFF1 / Tomoregulin-1	2.0
			- NT-3	2.0
			-	
2	Receptor	17.1	- VEGF R2(KDR)	3.1
			- SIGIRR	2.9
			- IL-13 R alpha 2	2.9
			- CCR6	2.8
			- IL-22 R	2.6
			- FGF R3	2.5
			- TMEFF2	2.4
			- TLR3	2.4
			- IL-1 R6 / IL-1 Rrp2	2.3
			- IL-2 R beta(CD122)	2.3
			- Frizzled-3	2.3
			- FGF R4	2.2
			- IL-1 R8	2.2
			- IL-17R	2.2
			- EGF R / ErbB1	2.2
			- Frizzled-7	2.2
			- 17 IFN-alpha / beta R2	2.1
			- Spinesin	2.1
			- TCCR / WSX-1	2.1
			- LFA-1 alpha	2.0
			- IL-1 R9	2.0

Table F-1 List of up-regulated proteins in Cf-CM (Cont.)

No	Biological function	No. of protein (%)	List of protein	Fold change (Cf/Lf)
3	Growth factor	16.3	- NRG1 Isoform GGF2	4.1
			- NRG3	3.3
			- GDF5	3.2
			- HRG-alpha	3.2
			- GDF9	3.2
			- GDF3	2.9
			- GDF8	2.7
			- GDF1	2.6
			- FGF-2 (b FGF)	2.6
			- HRG-beta1	2.6
			- TGF-alpha	2.6
			- NOV / CCN3	2.5
			- GDF11	2.4
			- EDG-1	2.4
			- NRG1-alpha / HRG1-alpha	2.3
			- GFR alpha-4	2.2
			- TGF-beta 1	2.1
			- FGF-21	2.0
			- TGF-beta 5	2.0
			- FGF-21	2.0
4	Matrix metallo-proteinase	1.6	- MMP-1	5.2
			- MMP-12	2.1
5	Phosphorylated protein	1.6	- TGF-beta RI / ALK-5	2.5
			- Smad 8	2.4

Table F-2 List of 70 down-regulated proteins in Cf-CM

Rank	Protein	Fold change (Cf/Lf)
1	Smad 4	0.9
2	Frizzled-6	0.9
3	FGF-16	0.9
4	FGF-10 / KGF-2	0.9
5	Activin RIA / ALK-2	0.9
6	CNTF R alpha	0.9
7	Axl	0.9
8	OSM	0.9
9	Hepassocin	0.9
10	Thrombospondin-1	0.9
11	Dkk-1	0.9
12	MMP-24 / MT5-MMP	0.9
13	IL-24	0.8
14	Glypican 3	0.8
15	Angiopoietin-like Factor	0.8
16	CCL14 / HCC-1 / HCC-3	0.8
17	IGFBP-rp1 / IGFBP-7	0.8
18	Pref-1	0.8
19	CXCR6	0.8
20	Smad 7	0.8
21	NCAM-1 / CD56	0.8
22	CCR2	0.8
23	CD40 / TNFRSF5	0.8
24	NRG1-beta1 / HRG1-beta1	0.8
25	IL-1 F6 / FIL1 epsilon	0.8
26	RELT / TNFRSF19L	0.8
27	CLC	0.8
28	BMP-7	0.8
29	LIF R alpha	0.8

Table F-2 List of 70 down-regulated proteins in Cf-CM (Cont.)

Rank	Protein	Fold change (Cf/Lf)
30	B7-1 (CD80)	0.8
31	TRANCE	0.8
32	Heregulin	0.7
33	MSPbeta-chain	0.7
34	BMP-5	0.7
35	CCR3	0.7
36	IL-3 R alpha	0.7
37	IL-4 R	0.7
38	MMP-8	0.7
39	MMP-10	0.7
40	Thrombospondin (TSP)	0.7
41	IL-19	0.7
42	TIMP-2	0.7
43	Progranulin	0.7
44	Dtk	0.7
45	Luciferase	0.7
46	Artemin	0.6
47	Neurturin	0.6
48	sFRP-4	0.6
49	BTC	0.6
50	Angiopoietin-1	0.6
51	IGFBP-6	0.6
52	E-Selectin	0.6
53	CV-2 / Crossveinless-2	0.6
54	BMP-3b / GDF-10	0.6
55	FGF-18	0.6
56	BD-1	0.6
57	CCR9	0.6
58	Osteoprotegerin / TNFRSF11B	0.5

Table F-2 List of 70 down-regulated proteins in Cf-CM (Cont.)

Rank	Protein	Fold change (Cf/Lf)
59	Siglec-9	0.5
60	Angiostatin	0.5
61	MMP-7	0.4
62	SCF	0.4
63	APRIL	0.4
64	BMP-15	0.4
65	Angiopoietin-like 1	0.4
66	Angiopoietin-like 2	0.3
67	GDF-15	0.3
68	ErbB3	0.3
69	HGF	0.1
70	uPA	0.1

APPENDIX G

***O. VIVERRINI* EXCRETORY/SECRETORY PRODUCT
INDUCED *PN* EXPRESSION IN NON-TUMORIGENIC
LIVER FIBROBLAST**

Introduction

PN has been proposed as a TGF- β inducible gene. Recently, our group has reported the induction effect of *O. viverrini* excretory/secretory product (ESP) on cell proliferation of fibroblasts possibly through TGF- β signaling pathway (Thuwajit et al., 2006). It is of great interest to explore whether *O. viverrini* ESP when reach fibroblasts underlying the bile duct epithelium can activate PN expression.

Material and method

To investigate the capability of *O. viverrini* ESP to induce PN expression in non-tumorigenic liver fibroblast, Lf was subjected to treat with the parasitic product containing serum-free medium for 48 h and PN expression was measured using real time PCR. Lf cultured in serum-free medium was used as the negative control whereas Cf was served as positive control.

Results

The result showed that PN expression was markedly increased with statistical significance in Lf exposed to *O. viverrini* ESP compared the control cells without *O. viverrini* ESP treatment (Fig G-1). Though the level of PN expression in parasitic product-induced Lf was not as high as that observed in Cf, the result suggested that *O. viverrini* ESP could activate Lf to produce PN. With our study confirmed that CCA-associated fibroblasts produced PN, it may suggest *O. viverrini* infection induced CCA may, in part, activate fibroblasts to produced tumorigenic substance(s). This observation, though needs further experiment before conclusion, it supports the roles of *O. viverrini* in induction CCA through the parasitic substances.

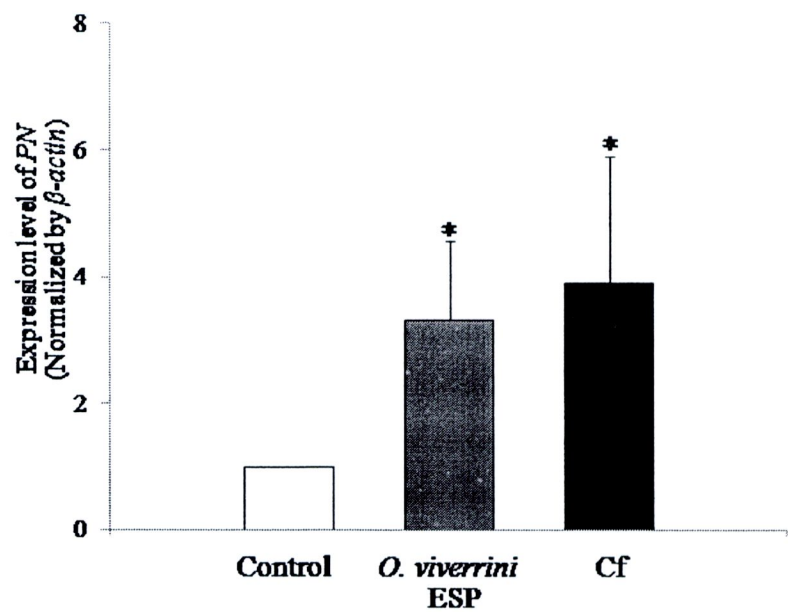


Figure G-1 *O. viverrini* ESP-induced *PN* expression in Lf. Expression level of *PN* in each condition was shown after normalized with β -actin level and assumed the level in the control condition to be 1. Bar represents mean $\pm$ SD of duplicate independent experiment. The * represents $P < 0.05$ compared with negative control.

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