

Rotterdam, Netherlands 2024



SUMMIT

Bile extracellular vesicles hold protein biomarkers for the early diagnosis of cholangiocarcinoma in individuals with primary sclerosing cholangitis

PERUGORRIA^{1,2,11}, L. BUJANDA^{1,2,11,12}, L.AABAKKEN⁵, V.PAULSEN⁵, TH. KARLSEN^{3,4,5}, **JM. BANALES**^{1,2,6,13}, **T. FOLSERAAS**^{3,4,5}

¹Biogipuzkoa Health Research Institute, San Sebastian, Spain; ²CIBERehd, Madrid, Spain; ³Norwegian PSC Research Center, Oslo, Norway; ⁵Oslo University Hospital Rikshospitalet, Oslo, Norway; ⁴University of Oslo, Norway; ⁵Oslo University Hospital Rikshospitalet, Oslo, Norway; ⁴University of Oslo, Norway; ⁴University Norway; ⁴University of Oslo, Norway; ⁴University Norway; ⁴University of Oslo, Norway; ⁴University Norway; ⁶IKERBASQUE, Bilbao, Spain; ⁷University of Bergen, Norway; ⁸Haraldsplass Deaconess Hospital, Bergen, Norway; ¹¹University of the Basque Country, Leioa, Spain; ¹²Hospital Universitario Donostia, San Sebastian, Spain; ¹³University of Navarra, Pamplona, Spain.

		PSC	PSC to CCA	PSC-CCA			
		n = 52	n = 8	n = 14			
		71	88	86			
ampling ((years)	39.5	45.3	56.1			
es)		50	25	36			
	ERCP (%)	63	100	71			
`	PTC (%)	0	0	14			
1	TX (%)	37	0	7			
	Liver resection	0	0	7			
	(%)						
langitis	Yes (%)	2	13	0			
npling	No (%)	98	88	100			
ma	Yes (%)	19	13	14			
me or	No (%)	56	75	79			
	NA (%)	25	13	7			
	0 (%)	-	0	7			
	I (%)	-	0	7			
	II (%)	-	13	7			
	III (%)	-	25	43			
	IV (%)	-	38	14			
	NA (%)	-	25	21			
Biochemical parameters							
(% high)		71	12	21			
(% high)		62	10	15			
(% high)		81	13	27			
(% high)		33	6	19			
(% high)		0	0	4			
(% high)		2	2	4			
(% high)		4	4	17			



High-throughput proteomics of bile EVs identified 21 diagnostic biomarkers for PSC-CCA, regardless of sex, age, the presence of inflammatory bowel disease, or cirrhosis at the time of sampling. Among these, 14 biomarkers were observed to be more abundant (in red), and 7 exhibited lower levels (in blue) in patients with PSC-CCA compared to patients with isolated PSC. These 21 diagnostic biomarkers revealed a high diagnostic capacity for early CCA detection, as bile from patients with non-metastatic tumors had these biomarker levels altered.



Machine learning algorithms revealed COPA/ATP5H/VTNC/IQGA1/PRDX2 (AUC=0.996) and COPA/ATP5H/VTNC/IQGA1/CALX/PRDX2 (AUC=1.000) as highly effective in diagnosing PSC-CCA versus isolated PSC, surpassing the performance of CA19-9 alone (AUC=0.846).

A. LAPITZ^{1,2}, M. GRIMSRUD^{3,4,5}, PM. RODRIGUES^{1,2,6}, M. VESTERHUS^{3,7,8}, M. AZKARGORTA^{2,9}, K. GRZYB¹⁰, H. REIMS¹⁰, F. ELORTZA^{2,9}, L. IZQUIERDO-SANCHEZ^{1,2}, M.

Figure 2. Bile EV-protein biomarkers for CCA diagnosis in patients with PSC

PSC-CCA (n=14) <i>vs</i> PSC (n=52)						PSC-CCA M0 (n=10) vs PSC (n=52)							
	ROC cu	irve		Univariable odds ratio (OR) Each protein & clinical/demographic variables [age, sex, cirrhosis, IBD]			ROC curve						
;	p-Value	SEN (%)	SPE (%)	OR (95%CI) P	-Value		OR (95%CI)	p-Value		AUC	p-Value	SEN (%)	SPE (%)
6	<0.001	92.9	61.5	20.8 (2.5 – 171.4)	<0.01		18.1 (1.3 – 249.9)	<0.05	<u>}</u> 01	0.778	<0.001	100	51.9
8	< 0.001	92.9	57.7	17.7 (2.2 – 145.8)	< 0.01		40.9 (2.2 – 756.5)	< 0.05		0.754	< 0.001	90.0	61.5
8	< 0.001	85.7	69.2	13.5 (2.7 – 67.4)	< 0.01		21.1 (1.7 – 255.7)	< 0.05		0.818	< 0.001	80.0	84.6
0	<0.01	71.4	75.0	7.5 (2.0 – 28.0)	< 0.01		8.2 (1.3 – 50.2)	<0.05		0.800	<0.001	60.0	86.5
9	< 0.001	100.0	51.9	31.3 (1.8 – 551.7)	< 0.05	⊢I	-	-		0.758	< 0.001	100.0	51.9
0	< 0.001	100.0	44.2	23.1 (1.3 – 407.8)	< 0.05		-	-		0.783	< 0.001	70.0	84.6
6	< 0.05	71.4	73.1	6.8 (1.8 – 25.2)	< 0.01		-	-		0.702	< 0.05	60.0	88.5
3	< 0.01	/1.4	/6.9	8.3 (2.2 – 31.4)	< 0.01		30.3 (2.4 – 3/4.6)	<0.01		0.712	< 0.05	70.0	/8.8
8	< 0.01	/1.4	69.2	5.6 (1.5 – 20.7)	< 0.01			-		0.771	< 0.001	70.0	80.8
7	< 0.001	85.7	53.8	7.0 (1.4 – 34.4)	< 0.05		24.2 (1.4 – 410.1)	< 0.05		0.756	< 0.001	80.0	65.4
4	< 0.01	/8.6	59.6	5.4 (1.3 – 21.8)	< 0.05		12.4 (1.4 – 108.9)	< 0.05		0.698	< 0.05	50.0	86.5
1	< 0.001	92.9	5/./	17.7 (2.2 – 145.8)	< 0.01		19.5 (1.5 -247.7)	< 0.05		0.725	< 0.001	90.0	63.5
8	< 0.01	/1.4	69.2	5.6 (1.5 – 20.7)	< 0.01		12.2 (1.5 – 98.8)	<0.05		0.701	< 0.05	70.0	69.2
2	< 0.01	64.3	76.9	6.0 (1.7 – 21.4)	< 0.01		-	-		0.729	< 0.01	90.0	53.8
7	< 0.01	5/.1	80.8	5.6 (1.6 – 19.8)	< 0.01		11.9 (1.5 – 95.3)	< 0.05		0.729	< 0.05	60.0	80.8
5	< 0.01	/1.4	65.4	4.7 (1.3 – 17.2)	< 0.05		10.4 (1.2 – 87.9)	< 0.05		0.717	< 0.05	50.0	94.2
9	< 0.01	85.7	57.7	8.2 (1.7 – 40.3)	< 0.01		1 E+4 (3.2 – 3E+7)	<0.05		0.729	<0.01	90.0	57.7
8	< 0.05	64.3	80.8	7.6 (2.1 – 27.5)	< 0.01			-		-	-	-	-
6	< 0.05	64.3	78.8	6.7 (1.9 – 24.1)	< 0.01		6.2 (1.1 – 36.5)	<0.05		-	-	-	-
5	< 0.05	78.6	/3.1	10.0 (2.4 – 41.0)	< 0.01			-		-	-	-	-
4	< 0.05	50.0	88.46	7.7 (2.0 – 29.6)	< 0.01		11.8 (1.4 – 103.1)	< 0.05		-	-	-	-
2	< 0.05	50.0	96.15	25.0 (4.3 – 145.2) <	< 0.001		45.0 (3.4 – 601.2)	<0.01		0.769	<0.05	70.0	96.2
1	< 0.05	/1.4	61.5	4.0 (1.1 – 14.5)	< 0.05		-	-		-	-	-	-
8	< 0.01	78.6	5/./	5.0 (1.2 – 20.1)	< 0.05		9.6 (1.2 - 75.2)	< 0.05		0.671	<0.05	70.0	67.3
5	< 0.05	/8.6	67.3	7.5 (1.9 – 30.7)	< 0.01		6.35 (1.1 – 36.4)	<0.05		-	-	-	-
1	< 0.05	50.0	88.46	7.7 (2.0 – 29.6)	< 0.01		-	-		-	-	-	-
0	< 0.05	/8.6	5/./	5.0 (1.2 – 20.1)	< 0.05		-	-		0.765	< 0.001	80.0	69.2
7	< 0.05	92.9	40.4	8.8 (1.1 – 72.5)	< 0.05		-	-		0.717	<0.01	80.0	55.8
5	< 0.05	92.9	40.4	8.8 (1.1 – 72.5)	< 0.05			-		-	-	-	-
4	< 0.05	50.0	88.46	7.7 (2.0 – 29.6)	< 0.01		42.0 (1.9 – 949.0)	<0.05		-	-	-	-
4	< 0.05	100.0	42.3	21.4 (1.2 – 3/7.8)	< 0.05		-	-		0.665	<0.05	100.0	42.3
3	<0.05	18.0	01.3	7.5 (1.9 – 30.7)	<0.01		11.0 (1.3 – 91.0)	<0.05		- 0.700	-	-	- 74.0
5	< 0.001	80.7 71.4	72 4	14.0 (3.0 - 74.2)	<0.01		10.1(1.9 - 134.9)	<0.05		0.790	< 0.001	90.0	71.Z
5	<0.001	71.4	76.0	0.0(1.0 - 20.2) 9.2(2.2 - 21.4)	<0.01		30.0 (4.2 - 2230.2)	<0.01		0.707	<0.001	00.0	61.5
5 7	< 0.01	05.7	10.9 EE 0	7.6 (2.2 - 31.4) 7.6 (1.5 27.2)	<0.01		33.0(2.0-404.0)	<0.01		0.733	< 0.01	90.0	01.0 EE 0
י 2	<0.01	79.6	67.2	7.0(1.3 - 37.2) 7.5(1.0 - 30.7)	<0.05		10.2(1.3 - 133.2)	<0.05		0.744	<0.01	900	67.2
<u>د</u>	<0.01	50.0	01.3	163 (2,4 79.2)	<0.01		86 8 (4 5 1690 0)	<0.03		0.721	S0.01	00.0	07.5
2 2	<0.05	02.0	55 Q	16.3 (3.4 - 70.3)	<0.001		66.6 (4.5 - 1060.5)	NU.U1		0 734	<0.05	90.0	- 55 8
5	<0.01	92.9 85.7	51.0	65(13-310)	<0.01		360 (26 - 500 8)	<0.01		0.734	<0.05	100.0	34.6
ă	<0.01	71.4	67.3	51 (1.4 – 18.8)	<0.05		90(12-658)	<0.01		0.000	-0.00	-	
6	< 0.03	71.4	71.2	6 2 (1 7 – 22 8)	<0.00		-	-0.00		0 692	<0.05	80.0	712
6	<0.01	100.0	46.2	24.9 (1.4 – 439.9)	< 0.05		-	-		0.665	<0.05	100.0	46.2
6	< 0.05	50.0	84.6	5.5 (1.5 – 20.0)	< 0.01		9.3 (1.0 – 84 7)	< 0.05		-	-	-	-
4	< 0.05	92.9	48.1	12.1 (1.5 – 98.8)	< 0.05		47.9 (1.8 – 1308.8)	< 0.05		0.708	< 0.05	90.0	50.0
0	< 0.05	42.9	88.5	5.8 (1.5 – 22.4)	< 0.05		-	-		-	-	-	-
7	< 0.05	64.3	76.9	3.8 (1.2 – 11.8)	< 0.05		-	-		0.698	< 0.05	70.0	76.9
4	< 0.05	92.9	40.4	8.8 (1.1 – 72.5)	<0.05		-	-		0.716	< 0.05	60.0	84.6
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Figure 3. Logistic models (LM) combining bile EV

n=14) <i>vs</i> PSC (n=52)						
	p-Value	SEN (%)	SPE (%)			
	<0.0001	100.0	100.0			
	<0.0001	100.0	98.1			
	<0.001	75.0	94.3			

Figure 4. Logistic models combining bile EV proteins for the prediction of CCA development in patients with PSC



Machine learning algorithm TM9S4/RS18/LPPRC/NHRF1 demonstrated predictive capacity for CCA development in PSC before any clinical evidence of malignancy with 100% of sensitivity and specificity (AUC=1.000), whereas serum CA19-9 exhibited no significant predictive capacity for CCA development (AUC=0.596).

Conclusion

Bile EVs harbor valuable protein biomarkers for predicting the development of CCA and enabling early diagnosis in individuals with PSC.

Given the ease of bile collection during stenting for dominant strictures in individuals with PSC, this innovative liquid biopsy may be of significant value for monitoring disease progression and aiding access of potentially curative treatment options.

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Contact Information

ainhoa.lapitz@biodonostia.org jesus.banales@biodonostia.org trine.folseraas@medisin.uio.no bio construction to the second BIOGIPUZKOA NOPSC Norwegian PSC Research Center









