

# **Genotyping of circulating tumor DNA in cholangiocarcinoma reveals diagnostic and prognostic information**

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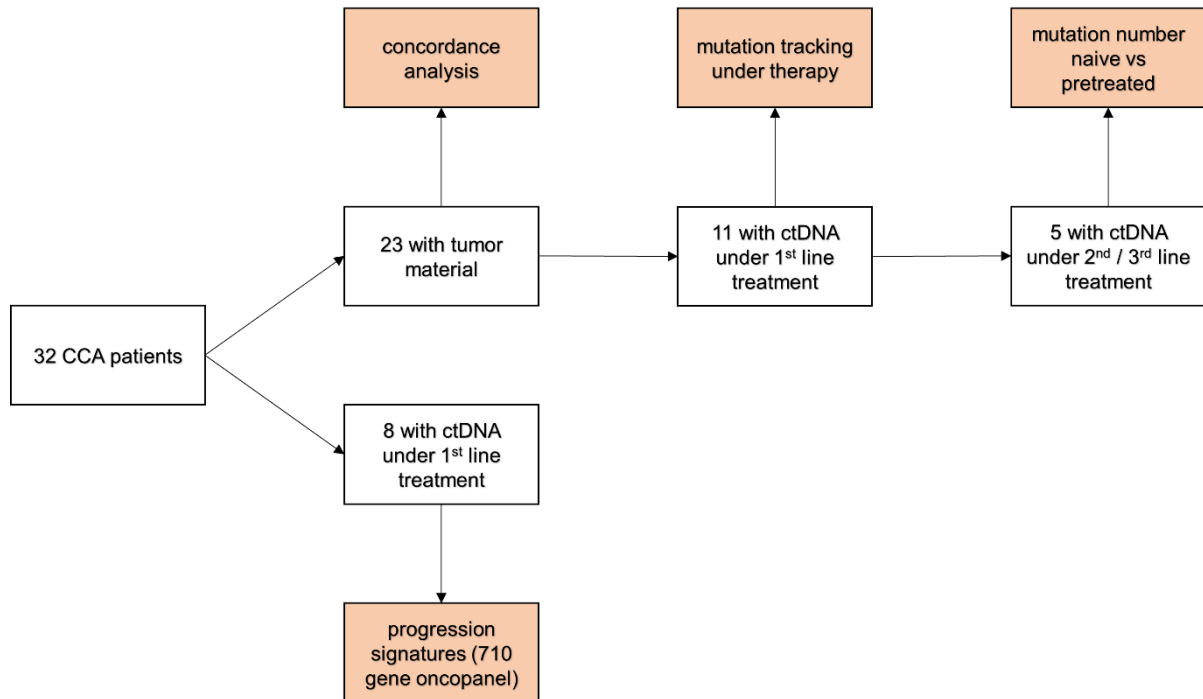
Prof. Dr. med. Thomas Seufferlein

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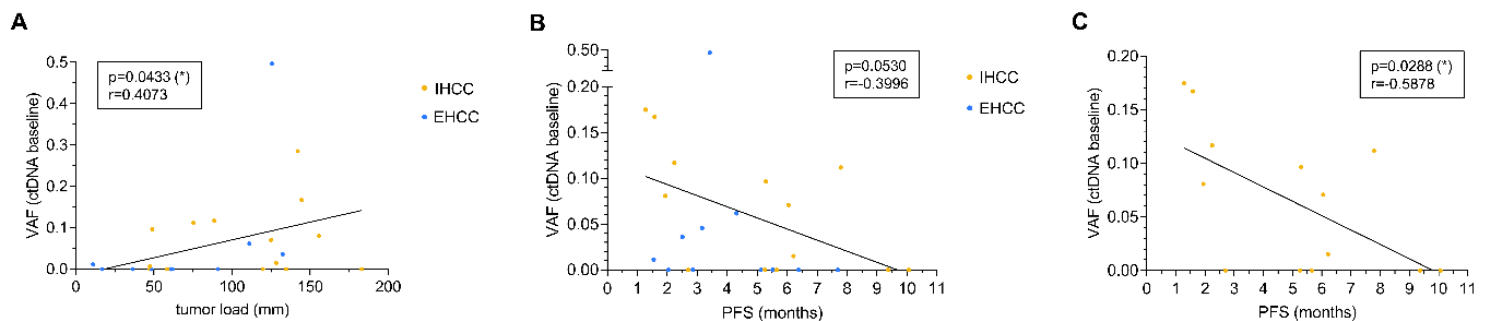
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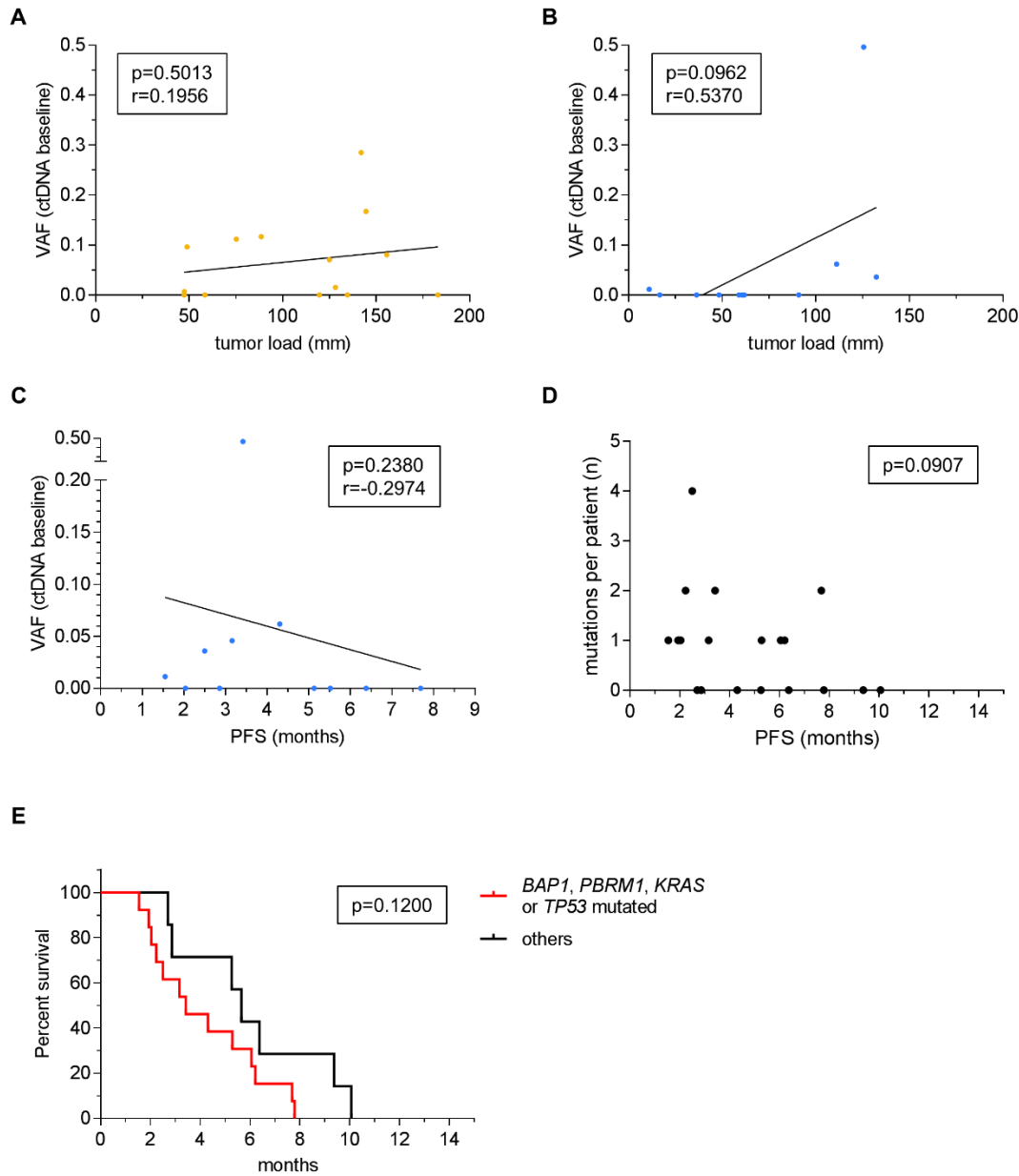
## Supplementary Materials



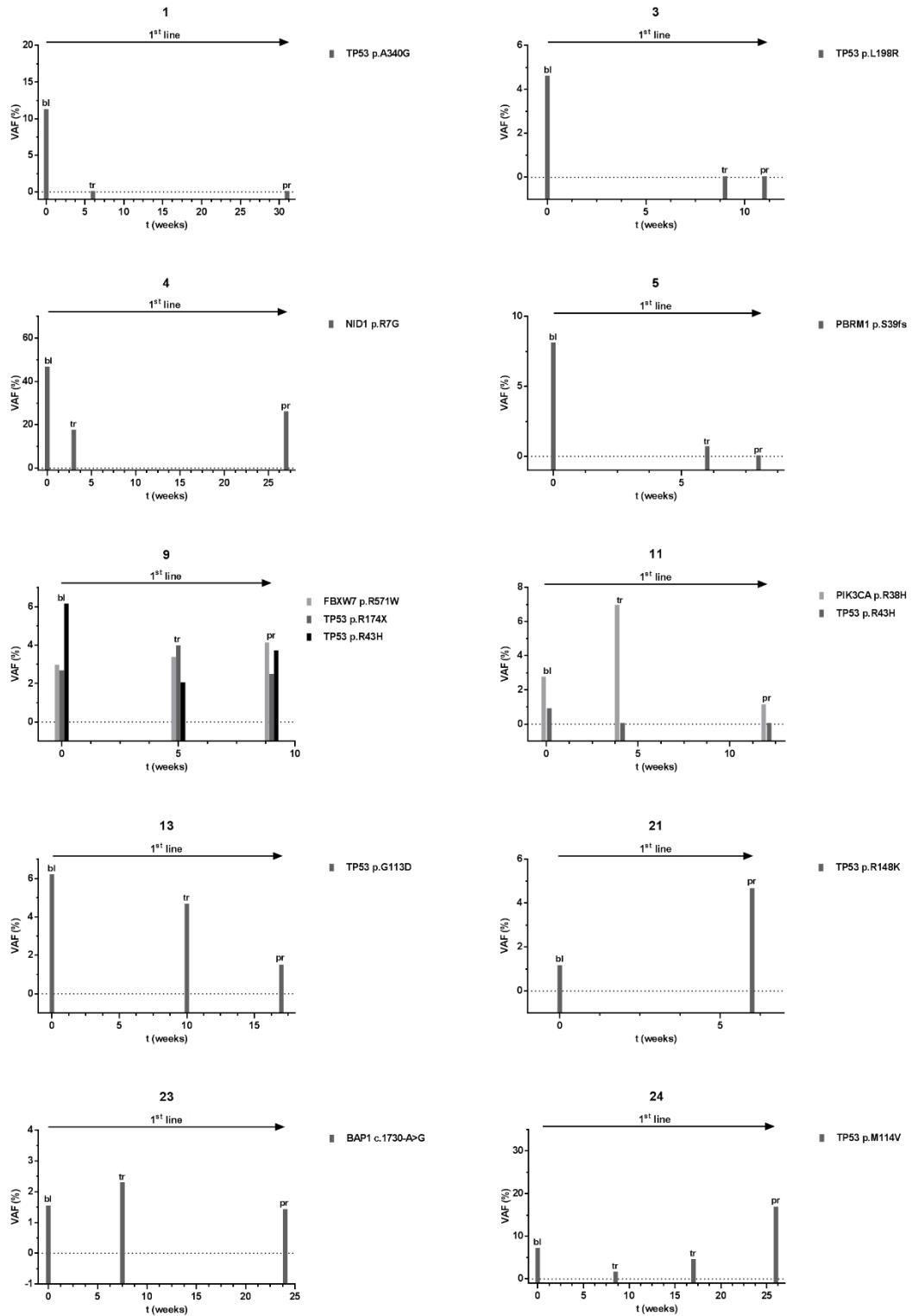
**Supplementary Figure 1.** Patient groups and applied methods.



**Supplementary Figure 2.** Correlation of ctDNA VAF baseline values and clinical parameters. **A.** VAF at baseline correlates with tumor load (RECIST 1.1) at baseline ( $P=0.0433$ ,  $r=0.4037$ , Spearman). **B.** Lower VAF at baseline shows a trend towards prolonged progression-free survival (PFS) ( $P=0.0530$ ,  $r=-0.3996$ , Spearman). **C.** VAF in the group of IHCC patients correlates with PFS ( $P=0.0288$ ,  $r=-0.5878$ , Spearman). IHCC - intrahepatic cholangiocarcinoma, EHCC - extrahepatic cholangiocarcinoma.



**Supplementary Figure 3.** Correlation of ctDNA VAF baseline values with **A.** tumor load (RECIST 1.1) in the group of IHCC patients ( $P=0.5013$ ,  $r=0.1956$ , Spearman), **B.** tumor load (RECIST 1.1) in the group of EHCC patients ( $P=0.0962$ ,  $r=0.5370$  Spearman), **C.** progression-free survival (PFS) in the group of EHCC patients ( $P=0.2380$ ,  $r=-0.2974$ , Spearman). **D.** Correlation of the number of mutations per patient with PFS ( $P=0.0907$ ,  $r=-0.3988$ , Spearman). **E.** Kaplan-Meier plot of PFS of patients with mutations in *BAP1*, *PBRM1*, *KRAS* or *TP53* vs. patients without mutations in these genes ( $P=0.1200$ , Mantel-Cox). IHCC - intrahepatic cholangiocarcinoma, EHCC - extrahepatic cholangiocarcinoma.



**Supplementary Figure 4.** Variant allele frequencies of detected mutations of all patients receiving first line chemotherapy from which cfDNA during the course of treatment was available (one patient was wild-type for the applied gene panel). bl - baseline, tr - treatment, pr - progression.

**Supplementary Table 1. Clinical patient characteristics.**

ID	sex	age	tumor localization	stage	metastases	chemotherapeutic regimen	
						therapy line	chemotherapy
1*	F	59	Intrahepatic	UICC IV	ADR, HEP, LYM, PER, OTH	1st, 2nd	Gem/Cis, FOLFIRI
2	F	78	Extrahepatic	UICC III	-	-	-
3	M	70	Extrahepatic	UICC IV	ADR, PLE	1st	Gem/Cis
4*	M	74	Extrahepatic	UICC IV	HEP, LYM, PUL	1st, 2nd, 3rd	Gem/Cis, FOLFIRI, Doce
5	M	70	Intrahepatic	UICC III	-	1st	Gem
6	F	55	Intrahepatic	UICC IV	ADR, LYM	1st	Gem/Cis
7	M	75	Intrahepatic	UICC III	-	1st	Gem/Cis
8	F	80	Intrahepatic	UICC IV	HEP, OSS, PUL	1st	Gem
9	M	84	Intrahepatic	UICC IV	LYM	1st	Gem
10	F	70	Extrahepatic	UICC IV	PER	1st	Gem/Ox
11	M	62	Extrahepatic	UICC IV	LYM	1st	Gem/Cis
12	M	75	Extrahepatic	UICC IV	PER	1st	Gem/Carbo
13	M	62	Extrahepatic	UICC IV	LYM	1st	Gem/Ox
14	M	74	Extrahepatic	UICC IV	PER	1st	Gem
15	F	54	Intrahepatic	UICC IV	LYM	1st	Gem/Cis
16	F	42	Intrahepatic	UICC IV	HEP, LYM	1st	Gem/Cis
17	M	39	Extrahepatic	UICC IV	LYM, OSS	1st	Gem/Cis
18*	F	59	Intrahepatic	UICC IV	HEP	2nd	FOLFIRI
19	F	49	Intrahepatic	UICC IV	LYM	1st	Gem/Ox
20	M	86	Extrahepatic	UICC IV	LYM, PUL	1st	Gem
21	M	73	Extrahepatic	UICC IV	PER, PUL, HEP, LYM	1st	Gem/Cis
22	M	72	Intrahepatic	UICC IV	LYM	1st	Gem/Ox
23	M	65	Intrahepatic	UICC IV	HEP, OSS, PUL	1st	Gem/Cis
24*	M	59	Intrahepatic	UICC IV	PUL	1st, 2nd	Gem/Cis, FOLFIRI

\* = analyzed prior advanced therapy line(s); F = female; M = male; UICC = Union Internationale Contre le Cancer; ADR = adrenal; HEP = hepatic; LYM = lymphoid; OSS = osseous; PER = peritoneal; PUL = pulmonary; OTH = other; Gem = Gemcitabine; Cis = Cisplatin; Doce = Docetaxel; Ox = Oxaliplatin; Carbo = Carboplatin; FOLFIRI = folinic acid/5-fluorouracil/irinotecan



**Supplementary Table 3.** Baseline laboratory findings of patients in the study

<b>Baseline laboratory findings</b>	<b>Median Value EHCC (min-max)</b>	<b>Median Value IHCC (min-max)</b>	<b>Unit</b>
Hemoglobin	12.8 (11.6-15.8)	12.2 (9.7-16.1)	g/dl
Platelets	282 (161-511)	240 (173-335)	Giga/l
Leucocytes	7 (6-9.7)	6.7 (4-21.6)	Giga/l
Bilirubin	17 (2-54)	7 (5-171)	μmol/l
Aspartate Aminotransferase (AST)	53 (19-266)	45 (14-329)	U/l
Alanine Aminotransferase (ALT)	56 (12-378)	29 (17-292)	U/l
Gamma-Glutamyl Transferase (GGT)	371 (18-1525)	122 (22-1103)	U/l
Alkaline Phosphatase (AP)	226 (72-682)	133 (39-268)	U/l
Lactate Dehydrogenase (LDH)	238 (170-501)	213 (149-1114)	U/l
CA19-9	114 (5.4-1415)	103 (1.2-10000)	IU/ml
Carcinoembryonic antigen (CEA)	4 (1.2-451)	2.7 (1.3-31)	μg/l
Alpha-fetoprotein (AFP)	6.5 (1-105)	5 (2-41)	μg/l

**Supplementary Table 4.** Patient risk factors for cholangiocarcinoma in the study cohort

<b>Characteristics</b>	<b>N (%)</b>
<b>Positive risk factors</b>	<b>16/24 (66.7)</b>
Choledocholithiasis / Cholecystolithiasis	3 (12.5)
Cholecystectomy	2 (8.3)
Hepatitis C	1 (4.2)
Smoker	5 (20.8)
Chronic alcohol abuse	1 (4.2)
Type 2 diabetes	4 (16.7)
<b>Negative risk factors</b>	<b>8/24 (33.3)</b>