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Intrahepatic cholangiocarcinoma, its variants, and its precursor lesions

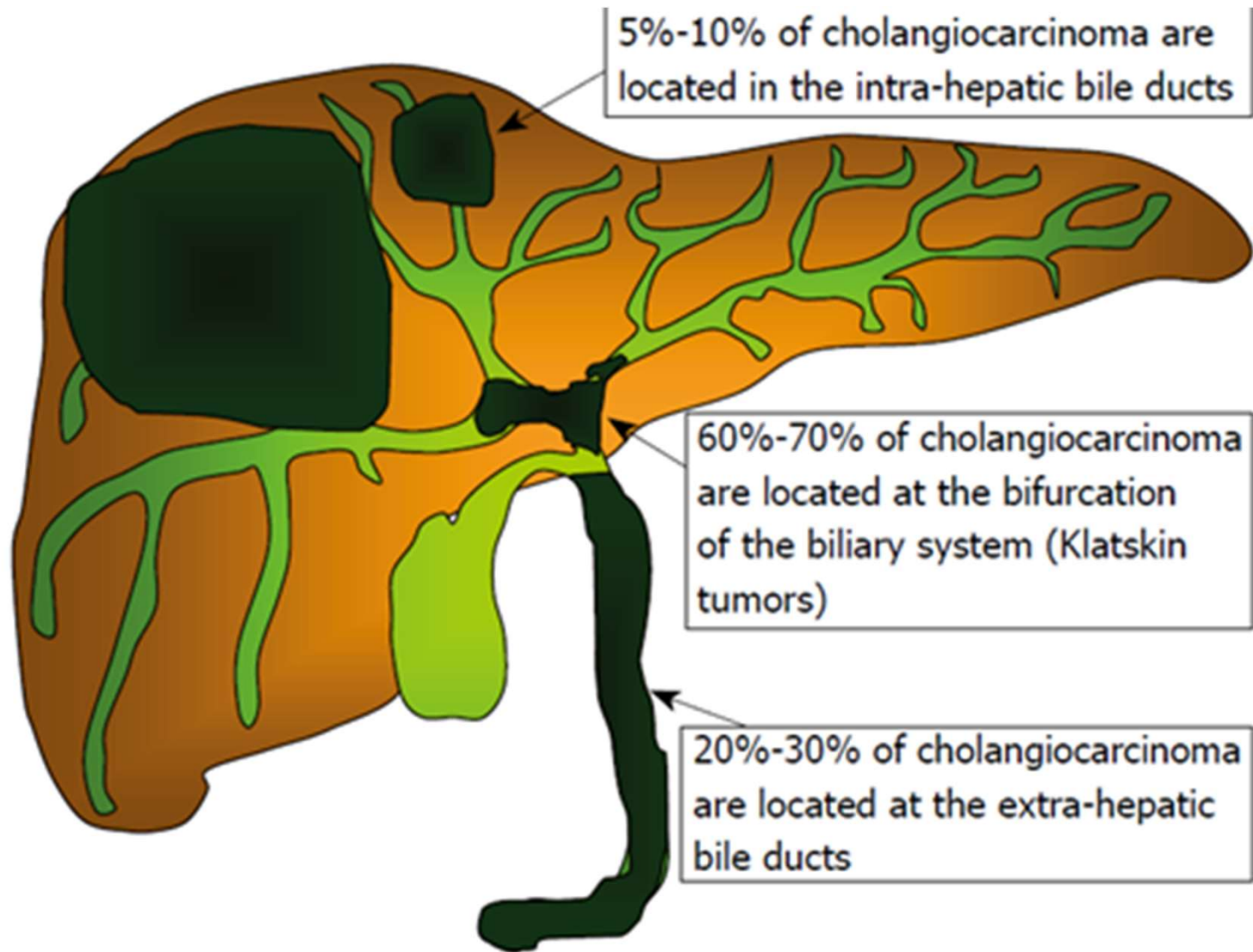
Pieter Demetter

Department of Pathology

Institut Jules Bordet

Cholangiocarcinoma

- Intrahepatic cholangiocarcinoma
- Perihilar cholangiocarcinoma (Klatskin tumour)
- Distal cholangiocarcinoma



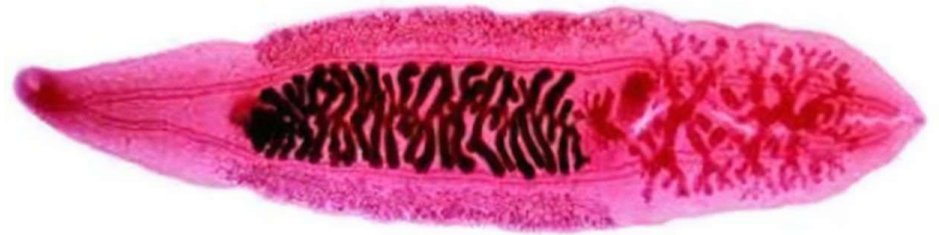
Intrahepatic cholangiocarcinoma

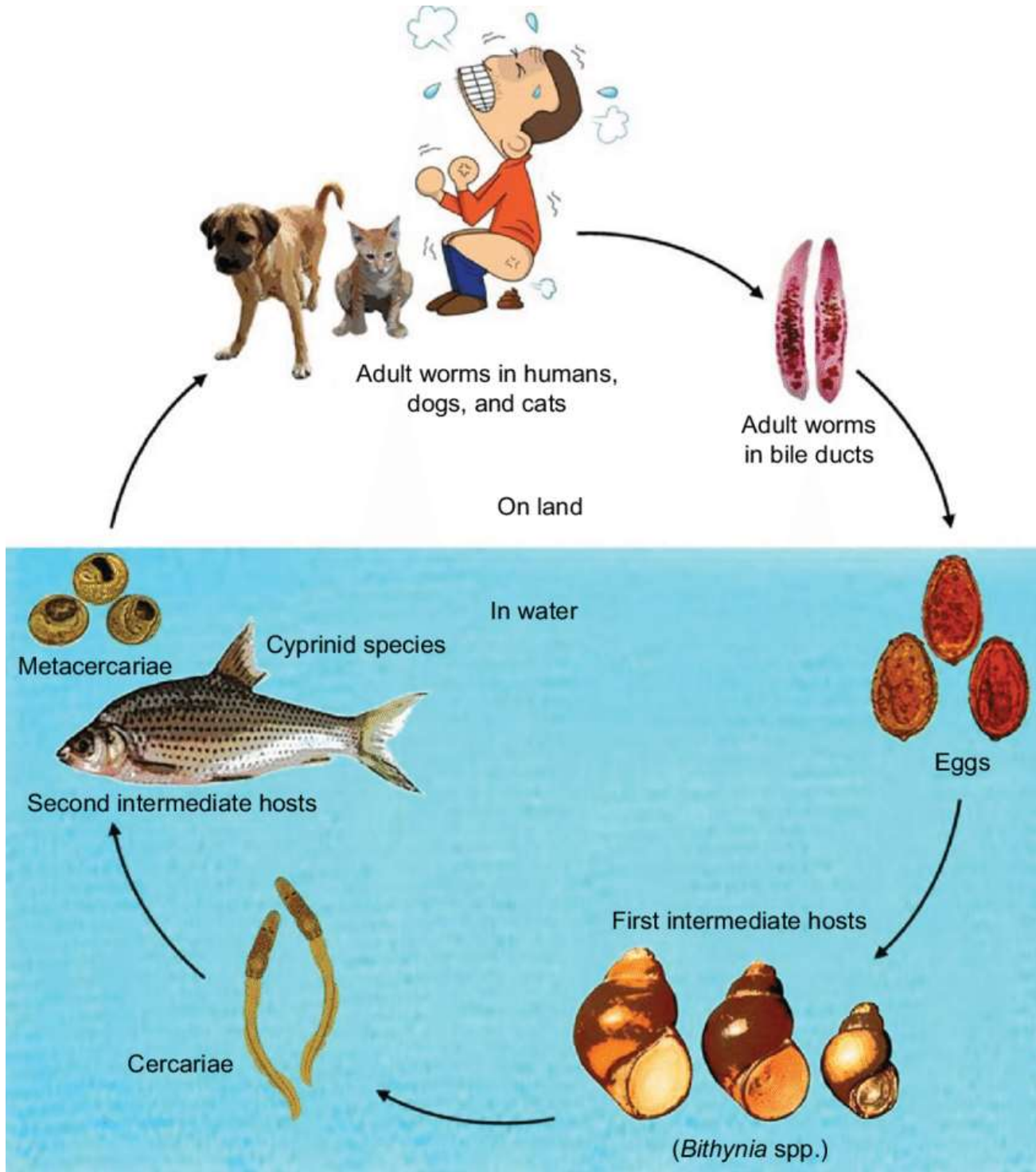
- 10-15% of primary liver malignancies
- General incidence: 1/100.000 people/year
- Proposed risk factors: choledochal cyst, anatomical anomalies, primary sclerosing cholangitis, chronic B or C viral hepatitis, non-alcoholic fatty liver disease, Lynch syndrome
- Higher incidence: South Korea (>Clonorchis sinensis) and Thailand (>Opistorchis viverrini)

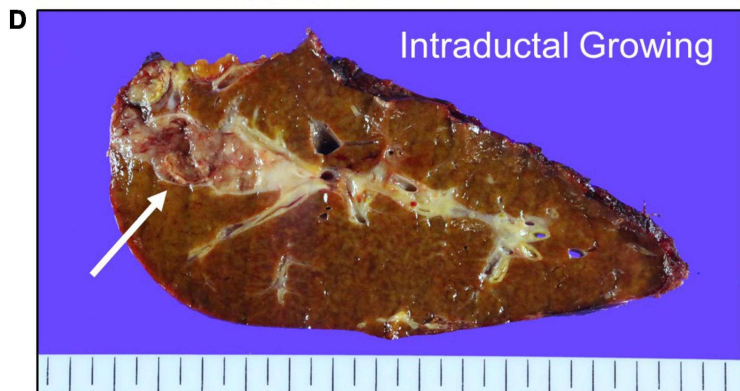
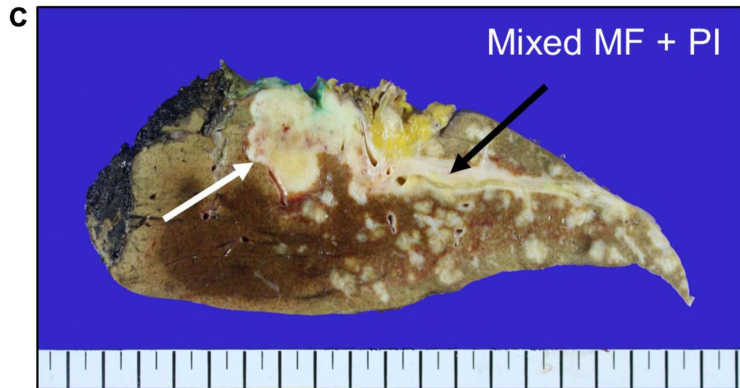
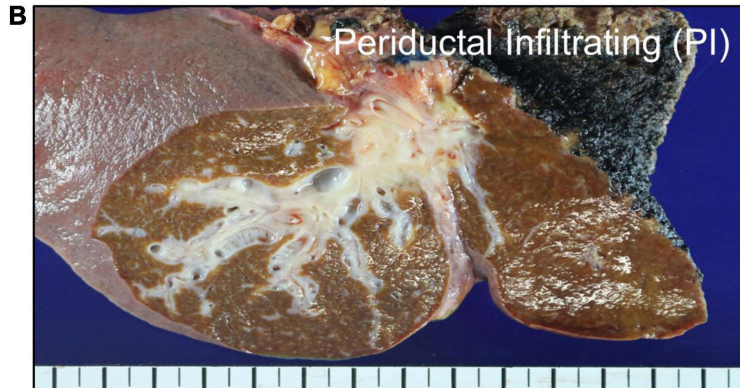
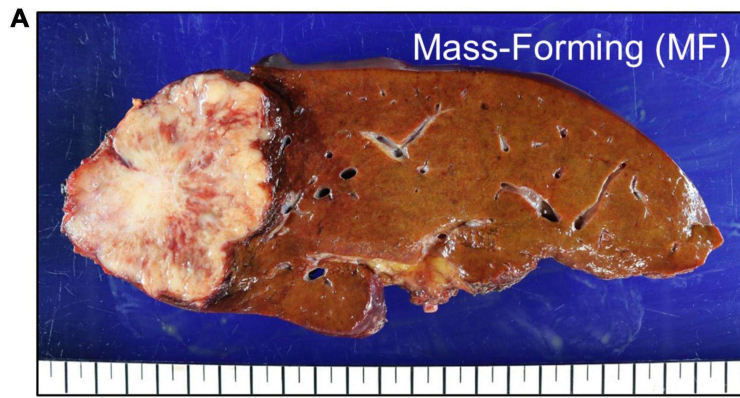
Clonorchis sinensis



Opistorchis viverrini







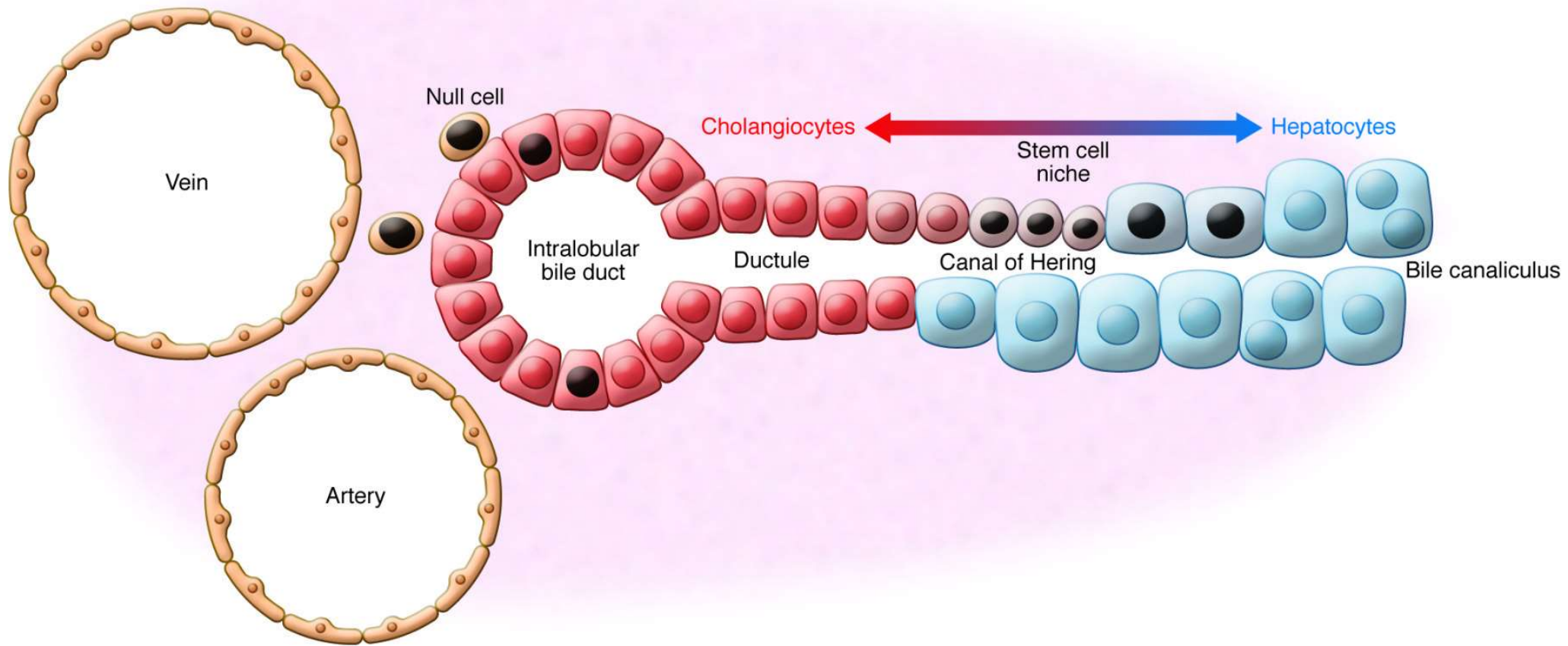
Intrahepatic cholangiocarcinoma (iCCA)

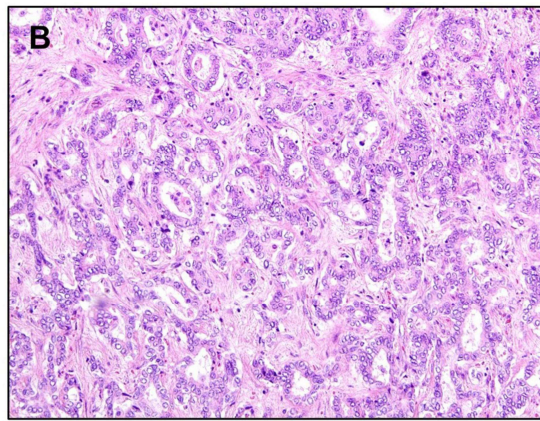
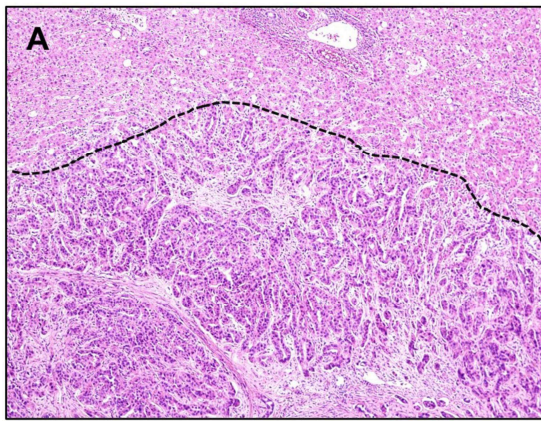
Small duct type

- Peripheral, ductular or cholangiolar type (synonyms)
- 36-84% of iCCA cases
- Background: often chronic hepatitis B or C, alcoholic hepatitis, non-alcoholic steatohepatitis
- Origin: canals of Hering, cuboidal cholangiocytes, interlobular bile ducts

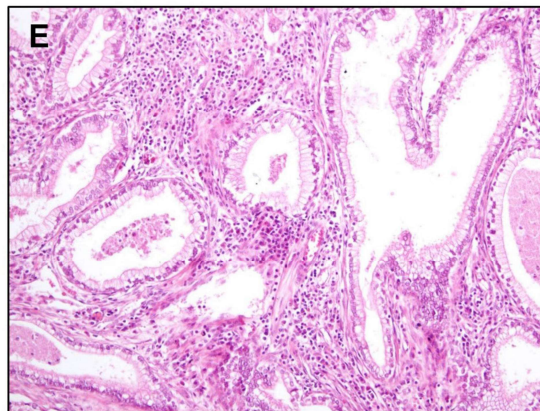
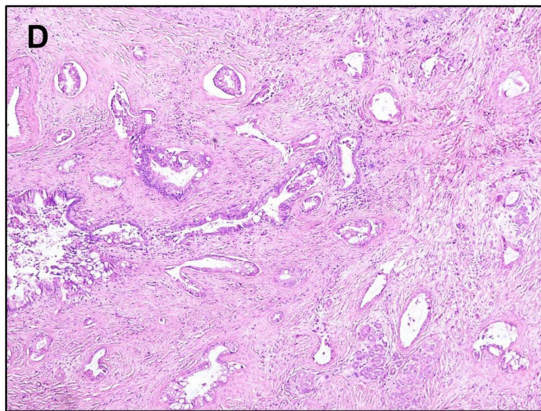
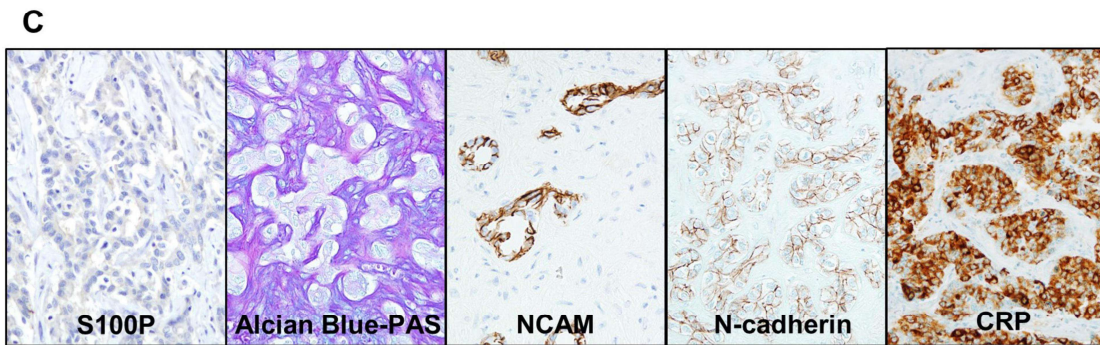
Large duct type

- Bile duct type of perihilar type (synonyms)
- 8-60% of iCCA cases
- Background: often chronic bile duct injury due to hepatolithiasis, parasites or primary sclerosing cholangitis
- Origin: columnar biliary epithelium, peribiliary glands around them

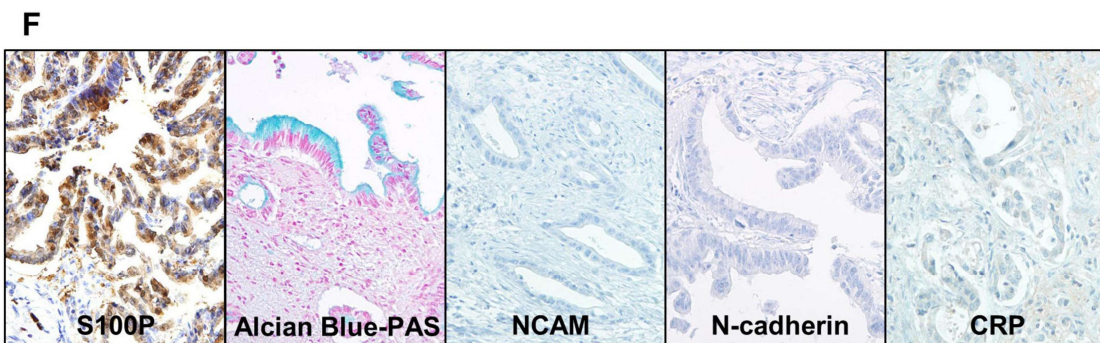


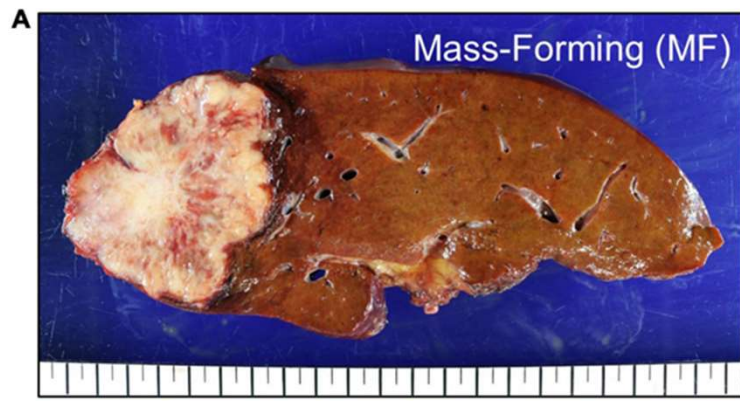


A, B and C: small duct type

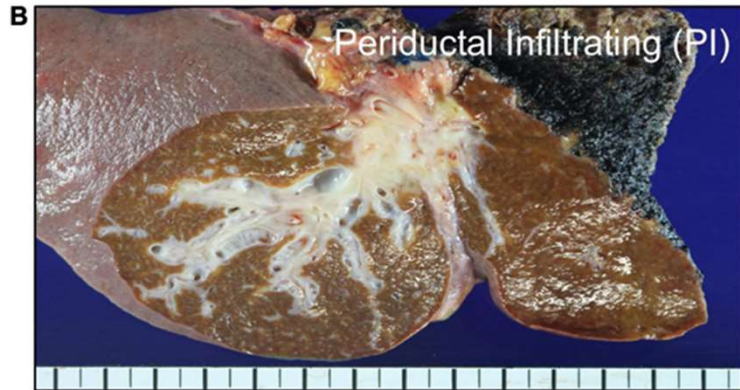


D, E and F: large duct type

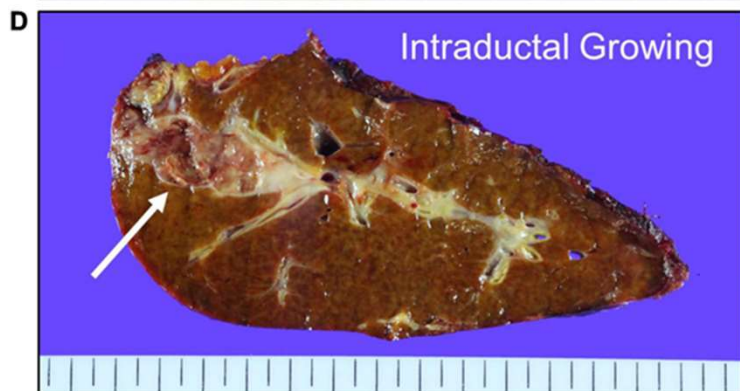
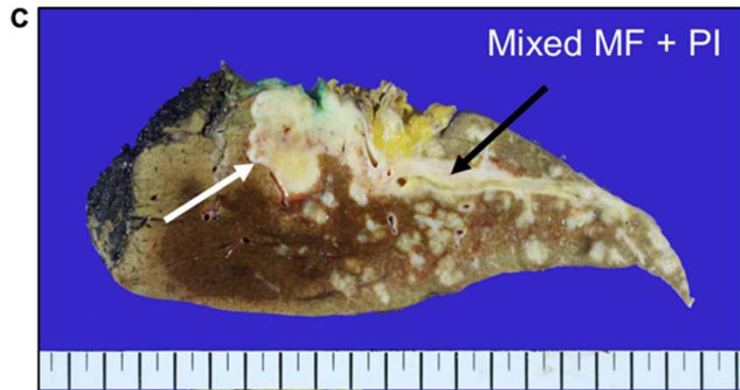




> Small duct and large duct types



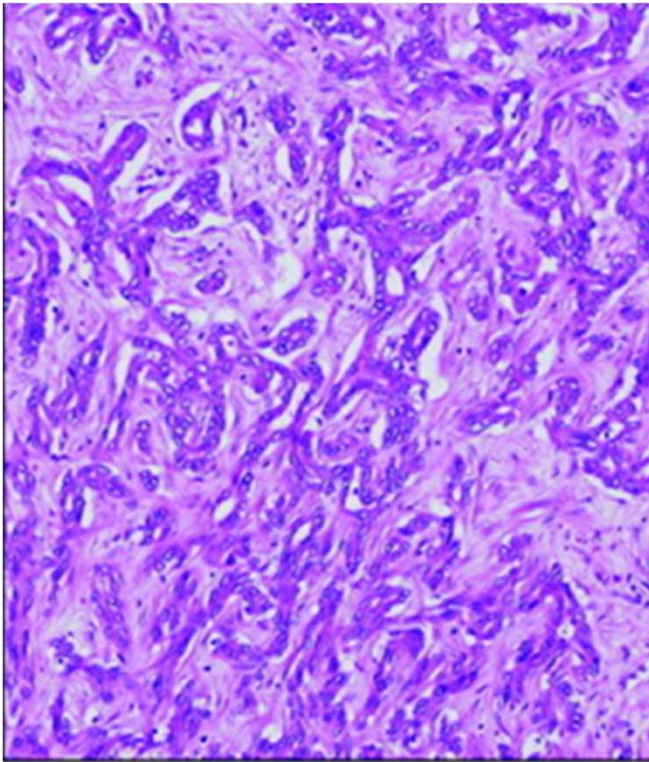
> Exclusively large duct type



Variants of intrahepatic cholangiocarcinoma

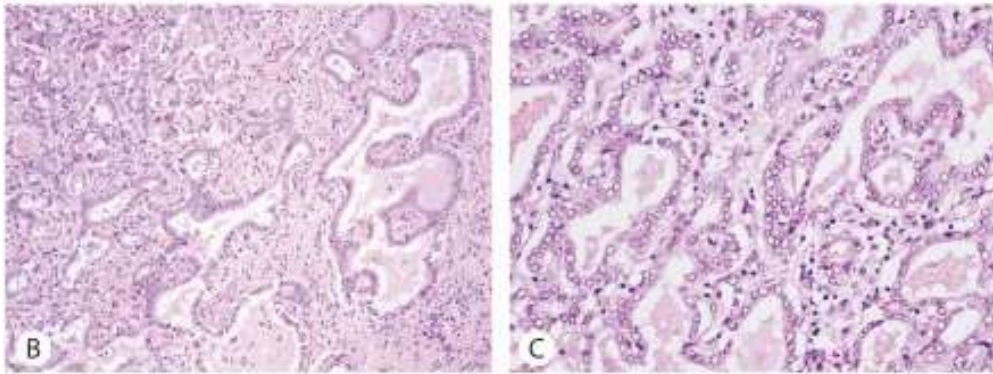
- Cholangiolocarcinoma
- Intrahepatic cholangiocarcinoma with ductal plate malformation pattern
- Adenosquamous carcinoma/squamous carcinoma
- Mucinous carcinoma/signet ring cell carcinoma
- Clear cell carcinoma
- Mucoepidermoid carcinoma
- Lymphoepithelioma-like carcinoma
- Sarcomatous intrahepatic carcinoma

Cholangiolocarcinoma



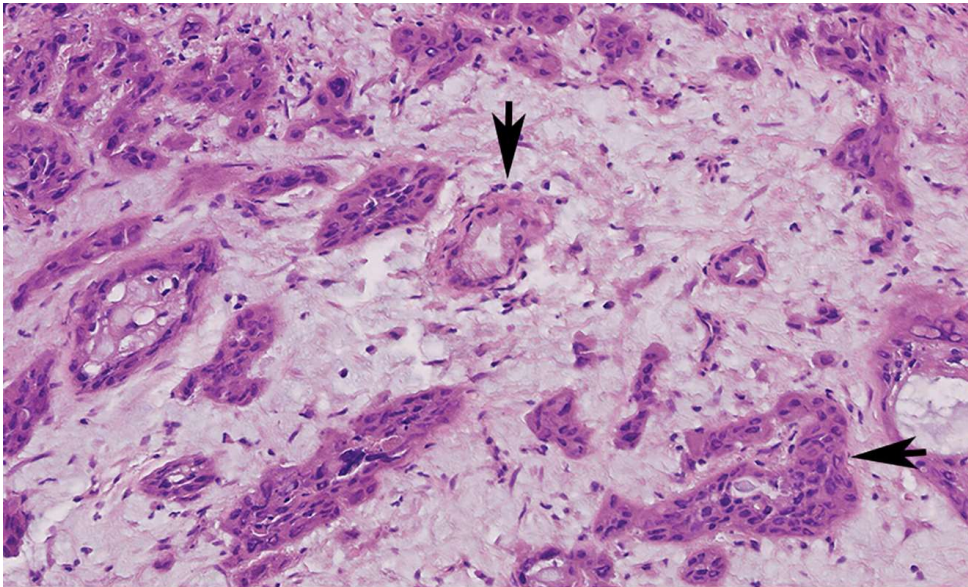
- Belongs to the small duct type
- iCCA with more than 80% of tumour area showing cholangiolocellular differentiation
- Excellent outcome

Intrahepatic cholangiocarcinoma with ductal plate malformation pattern



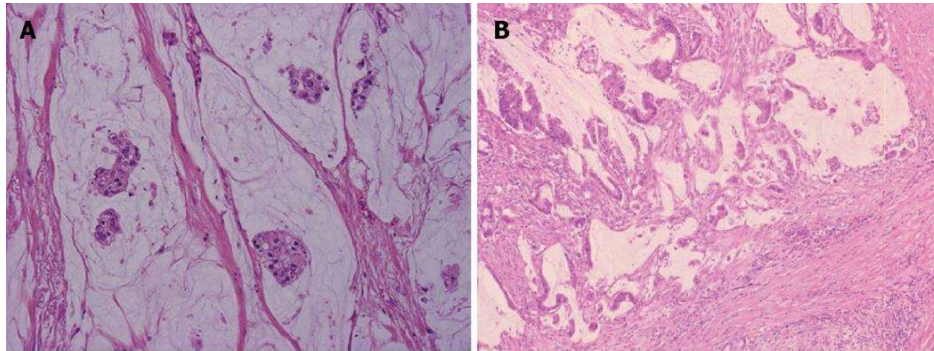
- Ductal plates: pathologically existing embryonic bile duct structures
- 3% of iCCA cases
- Survival better than that of conventional small duct ICC

Adenosquamous carcinoma/ squamous carcinoma



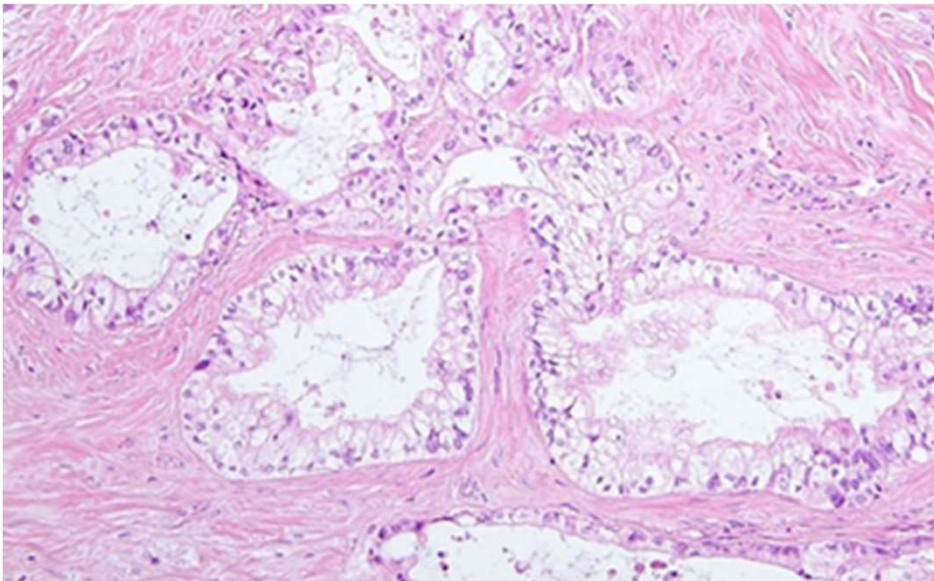
- Correlated with chronic cholangitis caused by liver flukes or hepatolithiasis
- Poor prognosis, median survival 6 months

Mucinous carcinoma/ signet ring cell carcinoma



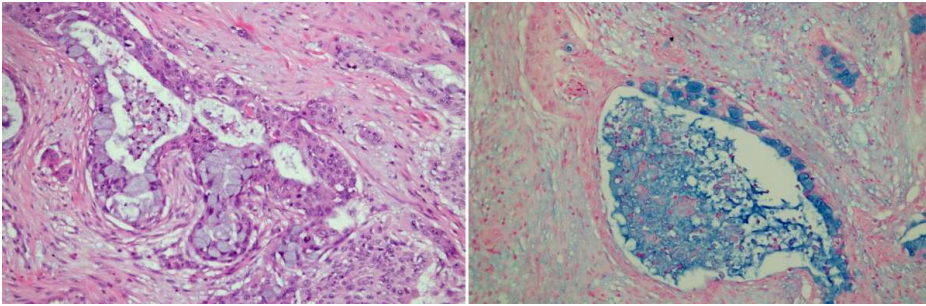
- Belongs to large duct type iCCA
- By convention at least 50% of tumour volume
- Usually due to malignant transformation of intraductal papillary neoplasm of the bile duct (IPNB)

Clear cell carcinoma



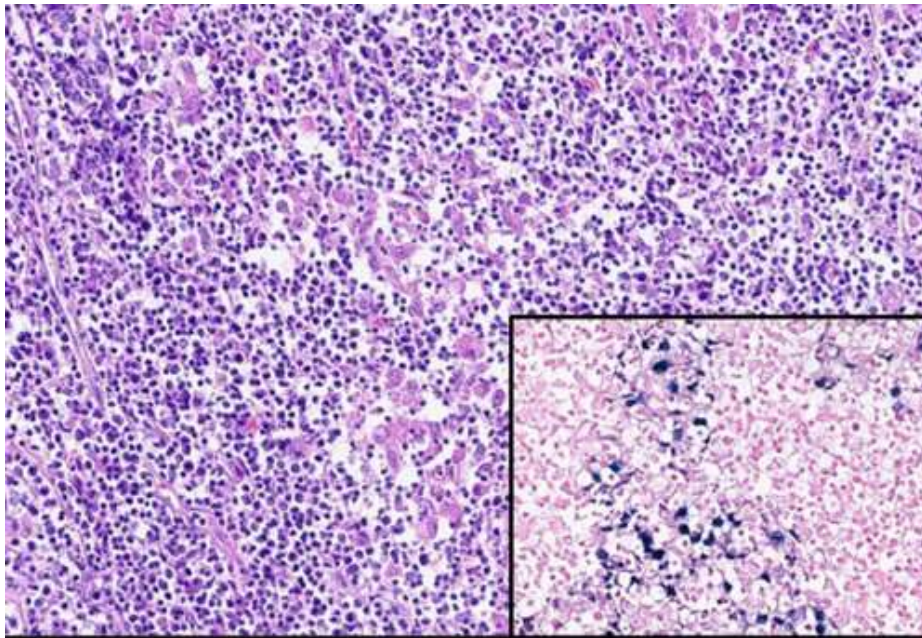
- Bulky cytoplasm clearing
- Eccentrically located nuclei
- Glandular and/or trabecular growth patterns
- DD: HCC with clear cell change, metastatic clear cell carcinoma of the kidney, metastasis of other GI tract tumours

Mucoepidermoid carcinoma



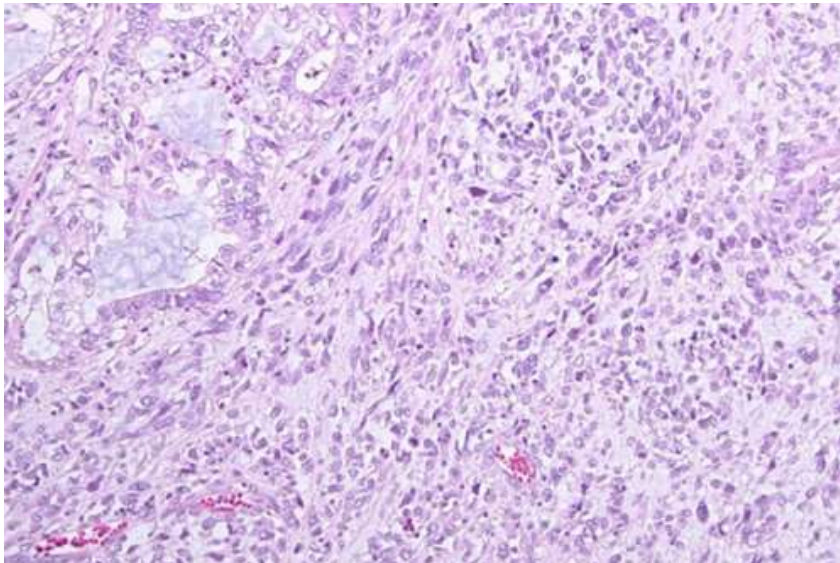
- Mixture of epidermoid/squamous and mucin-secreting elements
- Only a few case reports
- Poor prognosis

Lymphoepithelioma-like carcinoma



- Dense lymphoid stroma
- Undifferentiated or gland-forming tumour cells
- Almost all cases are Epstein-Barr encoded small RNA (EBER) positive
- Usually favourable outcome

Sarcomatous iCCA



- Mixed features of conventional iCCA and undifferentiated components with spindle cell features
- Sarcomatoid component often negative for epithelial markers
- Worse prognosis than conventional iCCA

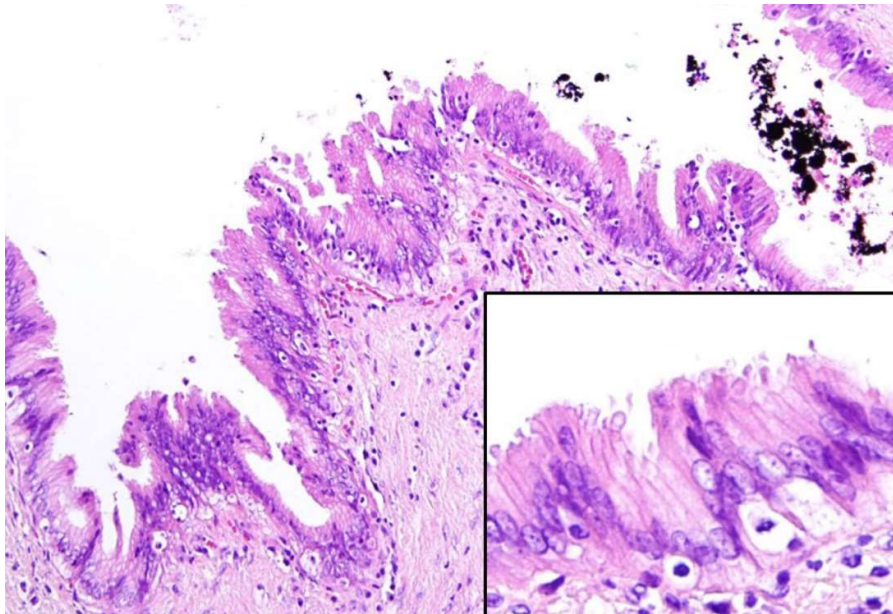
Precursor lesions of iCCA

- Biliary intraepithelial neoplasia (BillIN)
 - Low-grade BillIN
 - High-grade BillIN
- Intraductal papillary neoplasm of the bile duct (IPNB)
 - Low-grade IPNB
 - High-grade IPNB

Biliary intraepithelial neoplasia (BiIN)

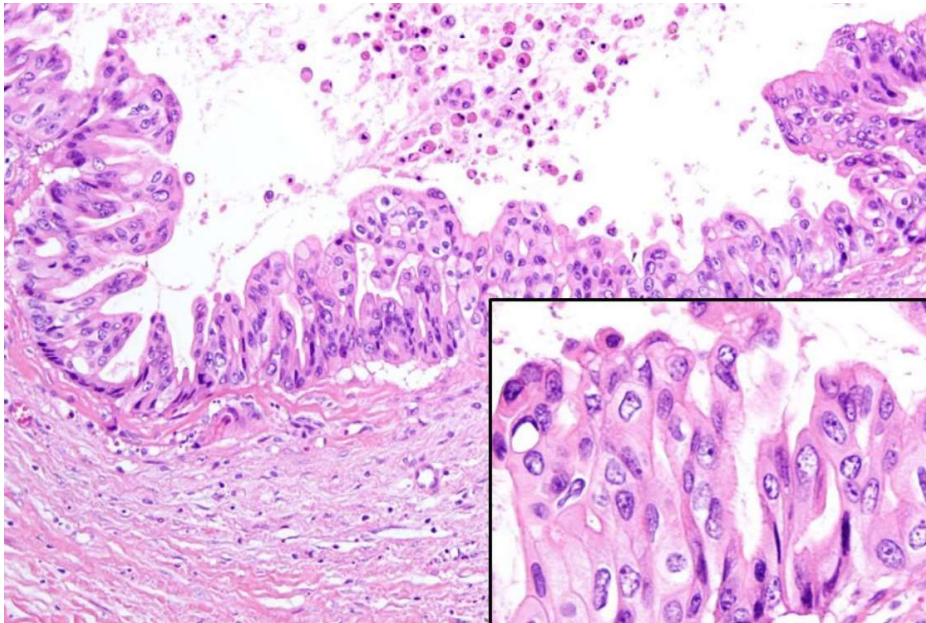
- Often accompanying large duct type iCCA, but not small duct type
- Invisible upon gross examination (or associated with subtle mucosal thickening)

Low-grade BiIN



- Mild cytoarchitectural atypia (flat pseudopapillary and/or micropapillary growth)
- Hyperchromatic nuclei
- Increase nuclear-cytoplasmic ratio
- Nuclear polarity is preserved
- p53 usually negative
- p16 relatively preserved

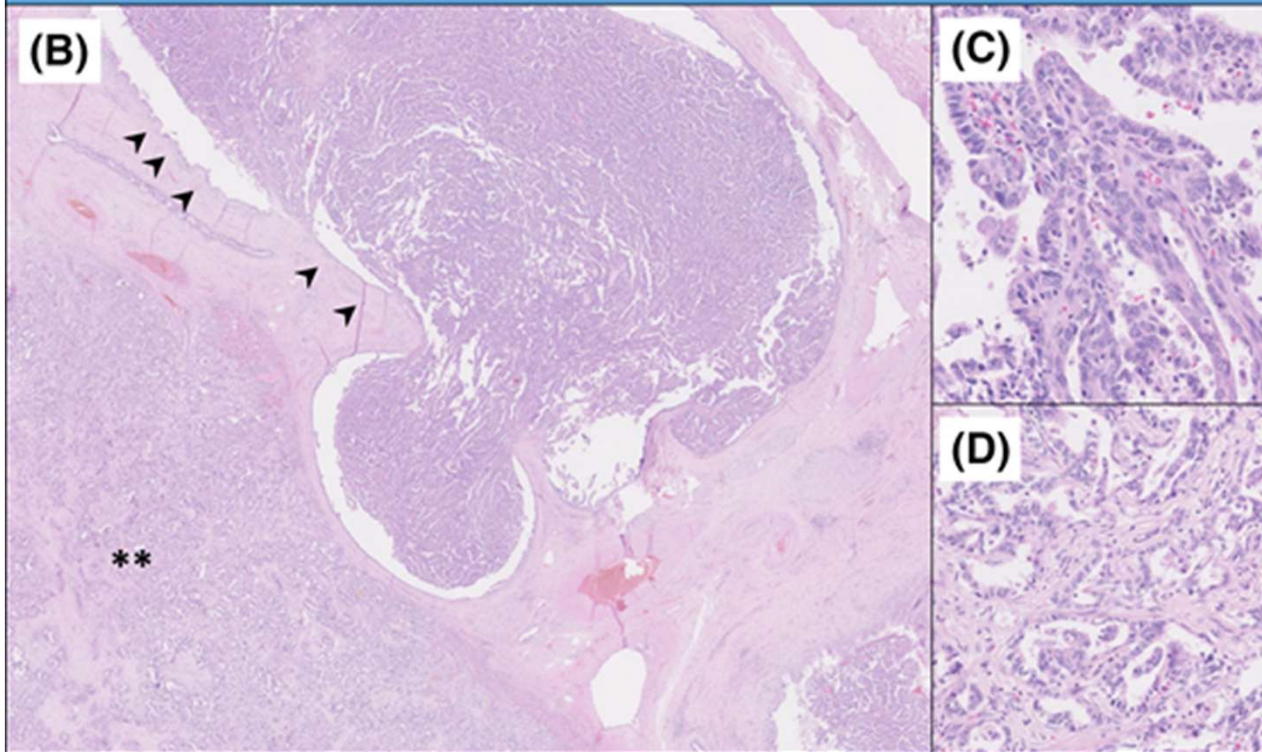
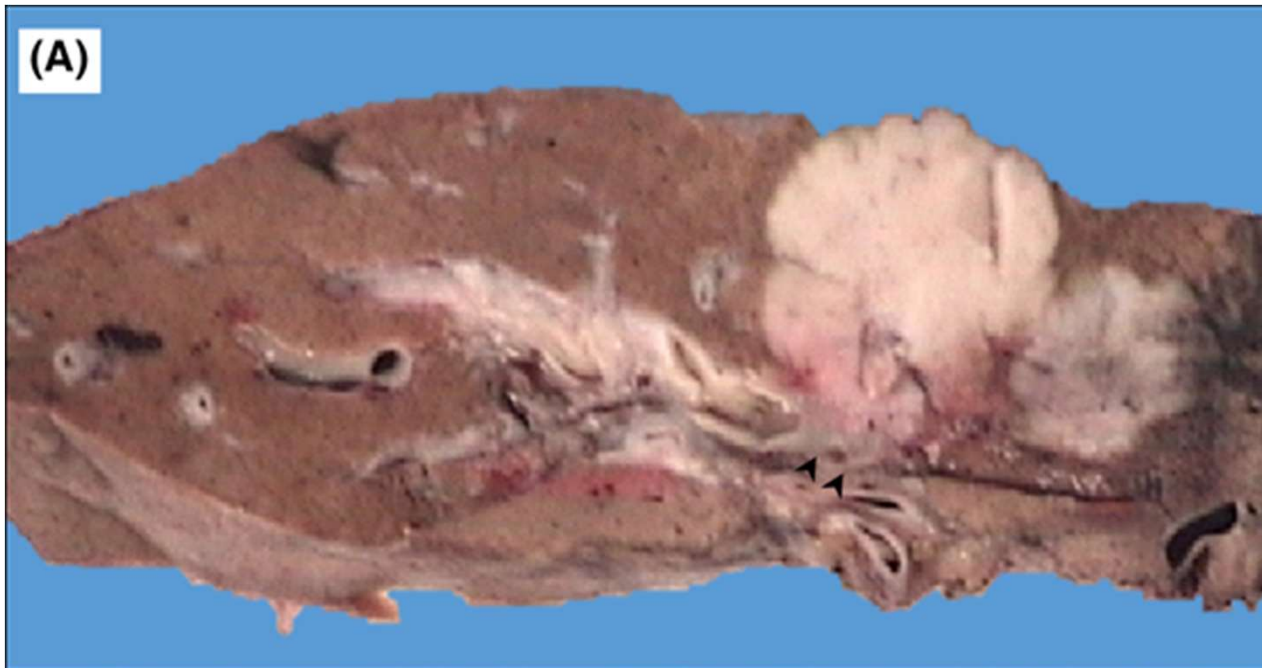
High-grade BiIN



- Moderate to severe cytoarchitectural atypia (more complex patterns, complete loss of polarity, marked nuclear atypia)
- Frequent mitosis
- p53 often overexpressed
- p16 decreased

Intraductal papillary neoplasm of the bile duct (IPNB)

- Grossly visible premalignant neoplasm
- Intraductal papillary or villous growth of biliary-type epithelium
- Four subtypes: pancreatobiliary, intestinal, gastric and oncocytic
- When invasive carcinoma develops in this lesion, it should be diagnosed as IPNB with associated invasive carcinoma



Intraductal papillary neoplasm of the bile duct (IPNB)

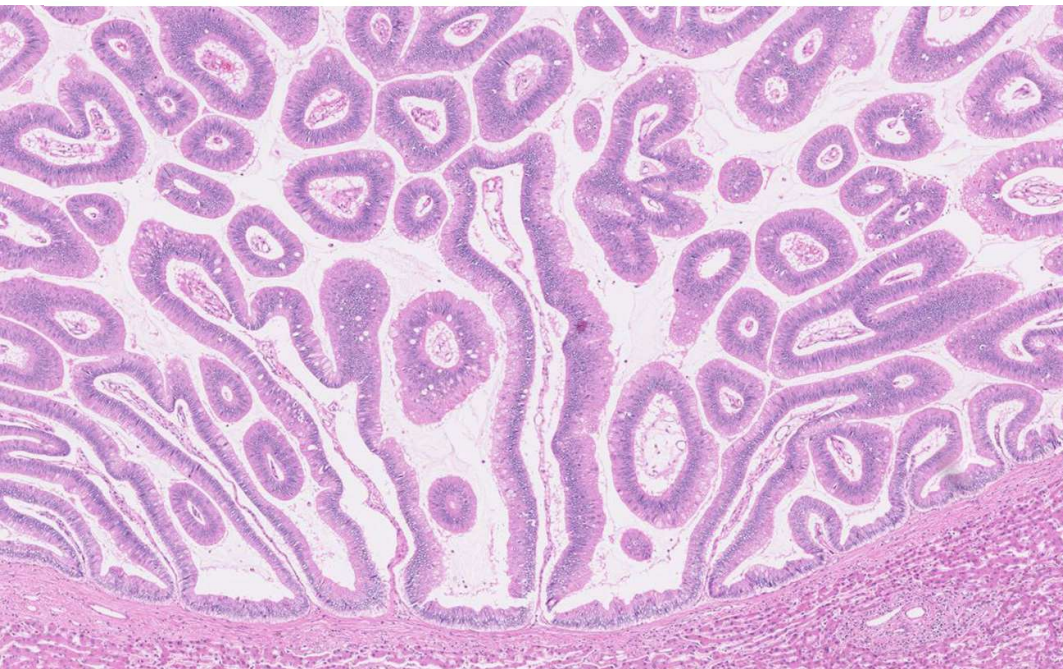
Type 1

- More homogeneous appearance
- Regular villous, papillary or tubular structures
- Usually low-grade dysplasia
- Frequently mucin overproduction
- Stromal invasion uncommon
- Most common in intrahepatic bile ducts
- Higher mutation rates of *KRAS*, *GNAS* and *RNF43*

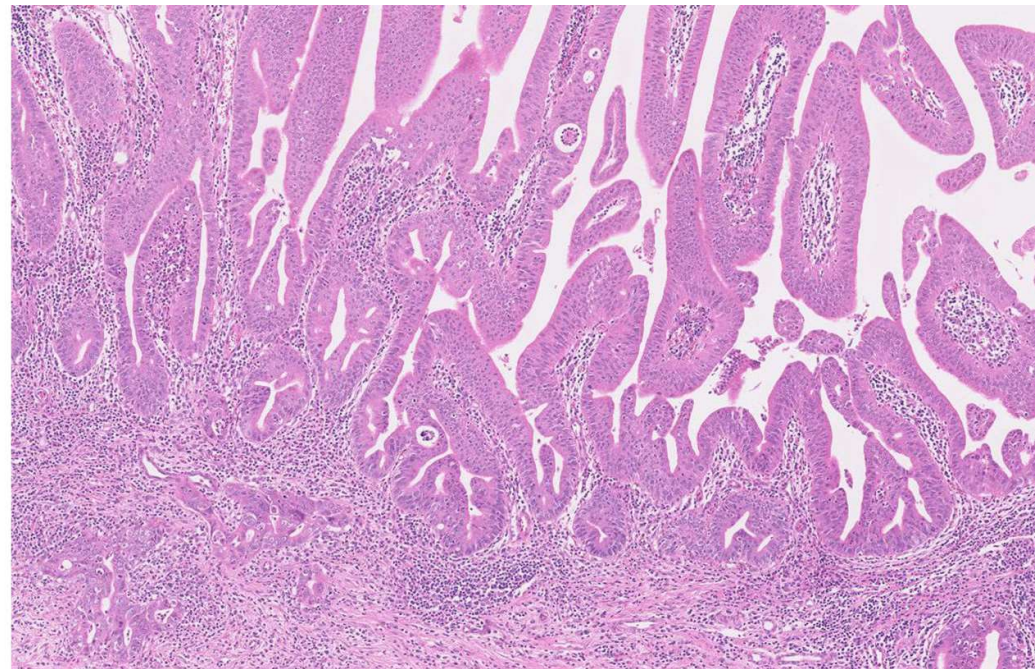
Type 2

- Heterogeneous appearance
- Irregular villous, papillary or tubular structures
- Usually high-grade dysplasia
- Rarely mucin overproduction
- Stromal invasion more common
- Arises throughout the biliary tree
- Higher mutation rates of *TP53* and *SMAD4*

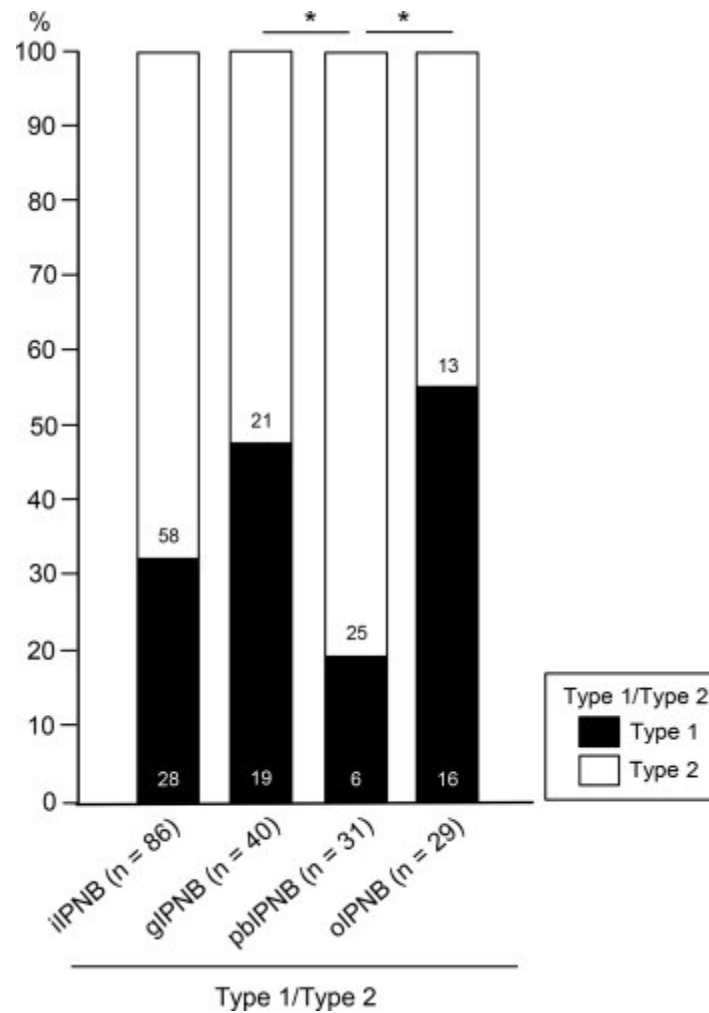
Type 1



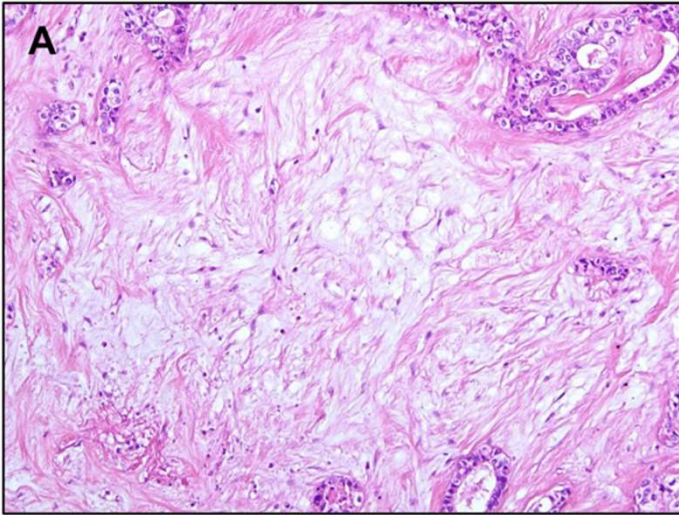
Type 2



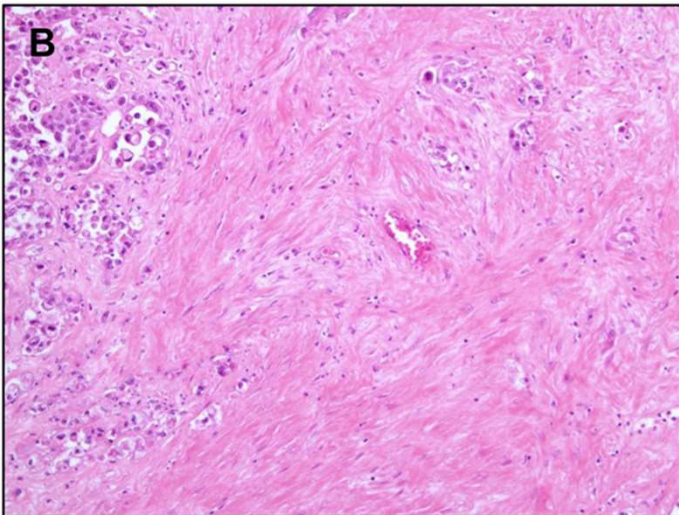
Proportion of types 1 and 2 in the four subtypes



Immature versus mature stroma in intrahepatic cholangiocarcinoma



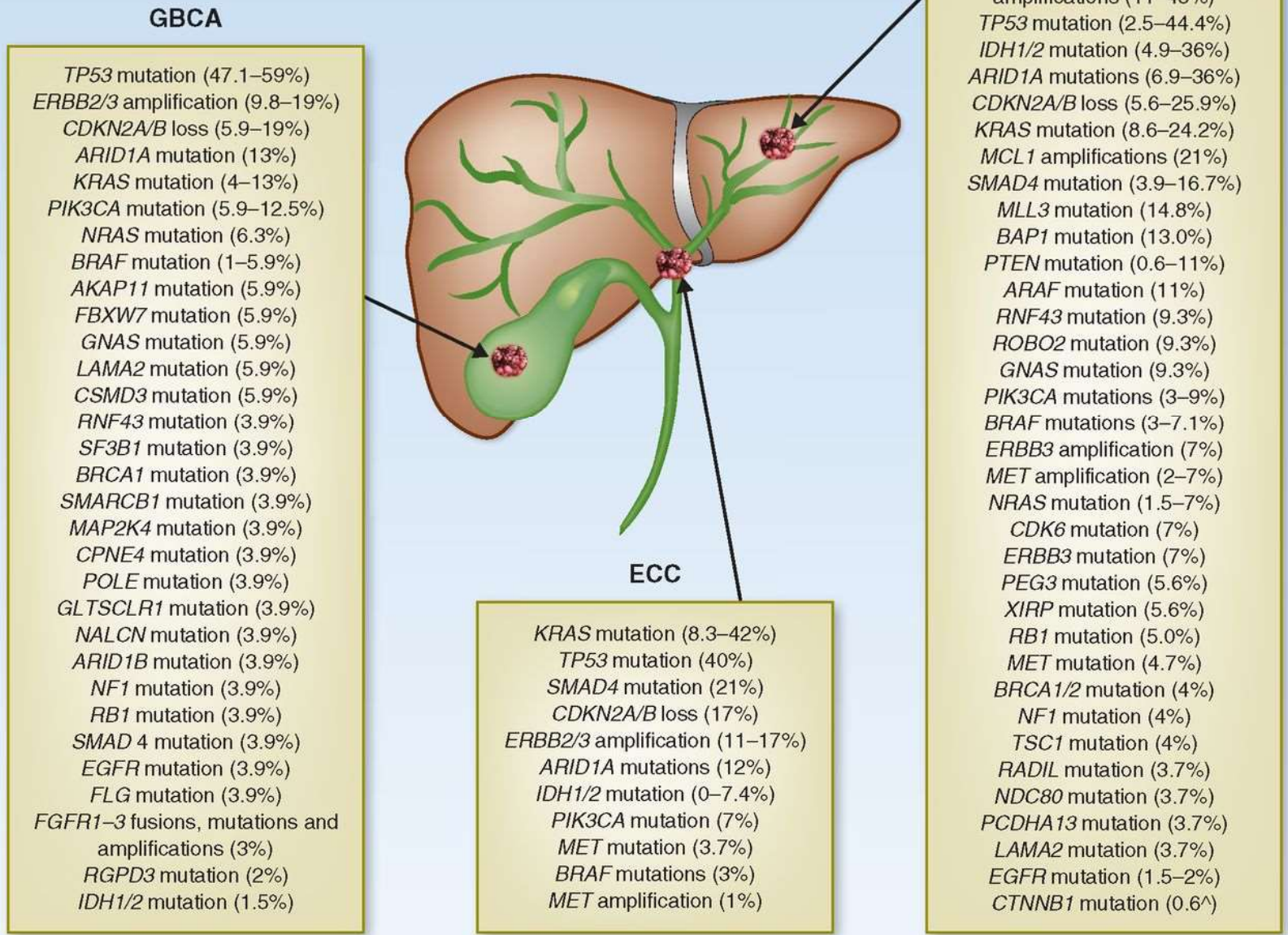
> Immature stroma; poorer prognosis



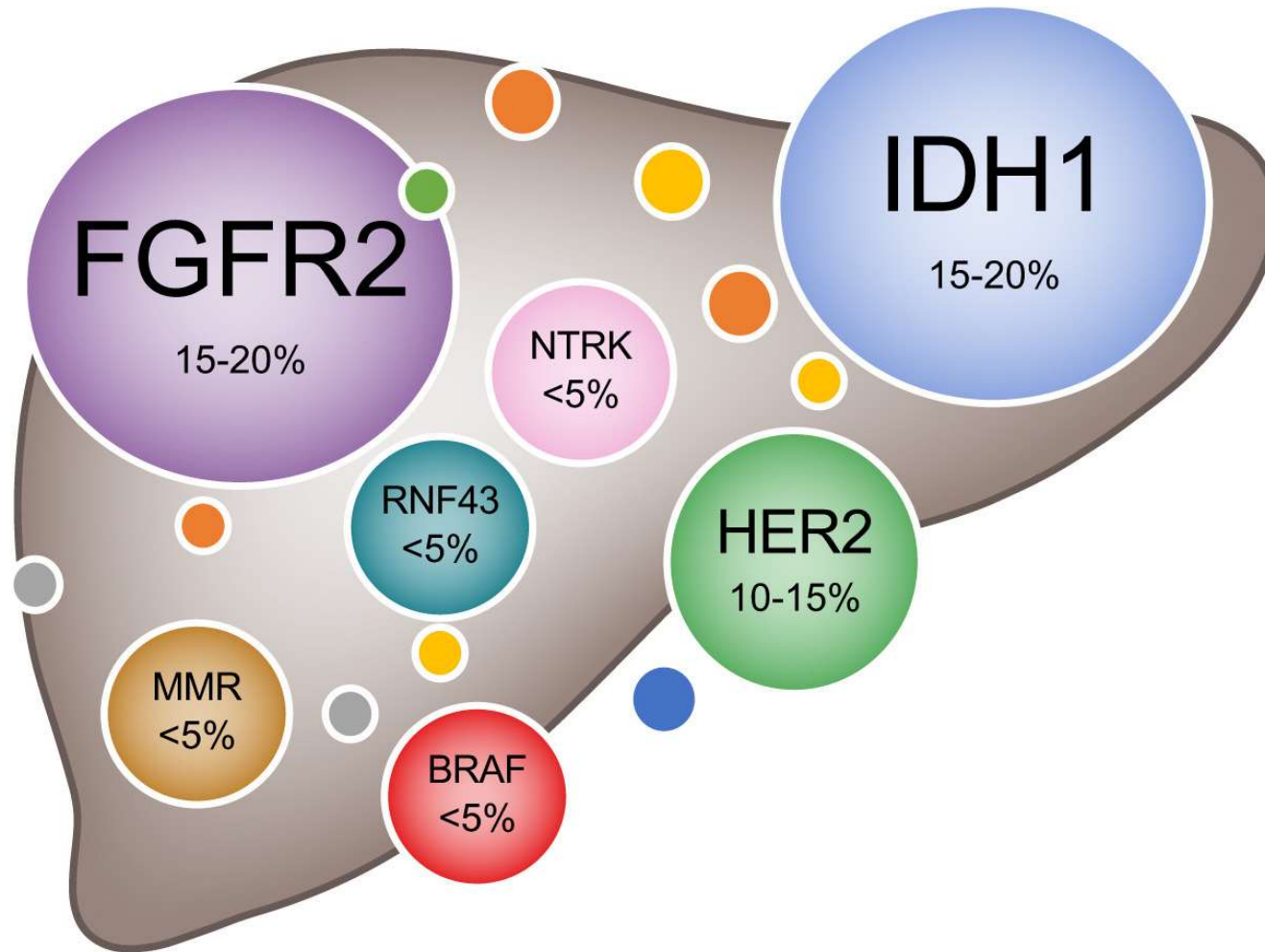
> Mature stroma; better prognosis

Zhang XF, *Hum Pathol* 2017
Kojima S, *Anticancer Res* 2020
Chung T, *Front Oncol* 2022

Molecular genetics of BTC



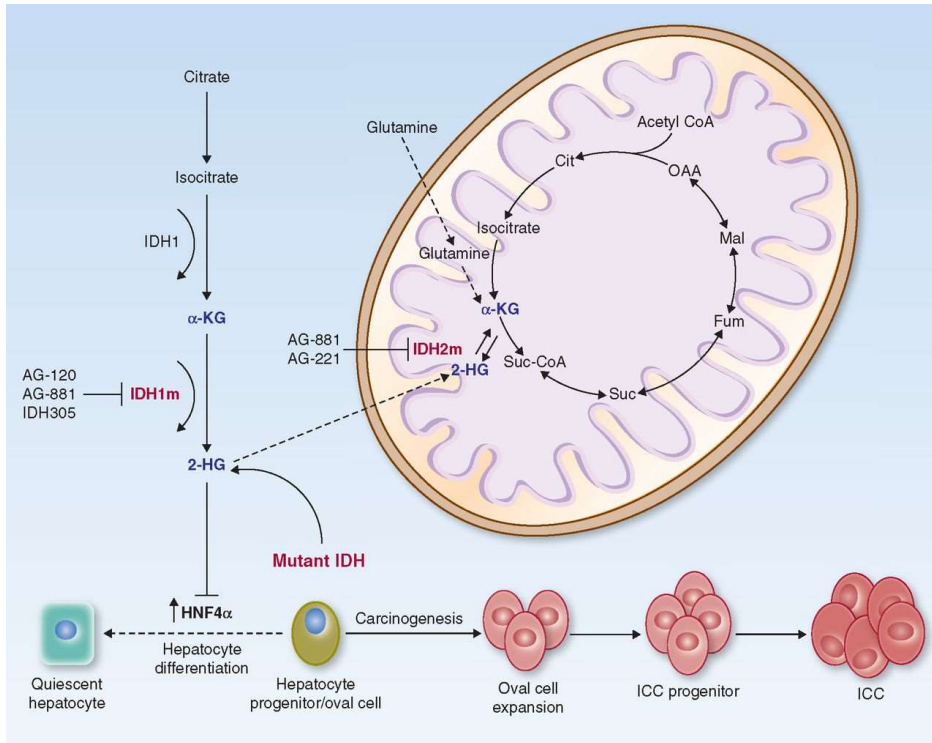
Molecular subgroups of intrahepatic cholangiocarcinoma



FGFR2 fusions

- *FGFR2* fusions/rearrangements in 15-20% of intrahepatic cholangiocarcinoma
- Associated with better prognosis
- Tyrosine kinase inhibitors
 - Pemigatinib: oral FGFR 1-3 TK inhibitor
 - Derazantinib: oral pan-TK inhibitor
 - Infigratinib: oral FGFR 1-3 TK inhibitor

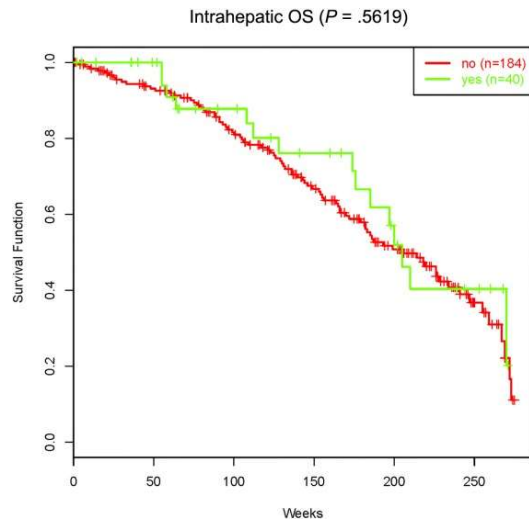
IDH 1 / 2 mutations



IDH : isocitrate dehydrogenase

IDH 1 or 2 mutations in 20% of intrahepatic cholangiocarcinoma

2-hydroglutarate: oncometabolite

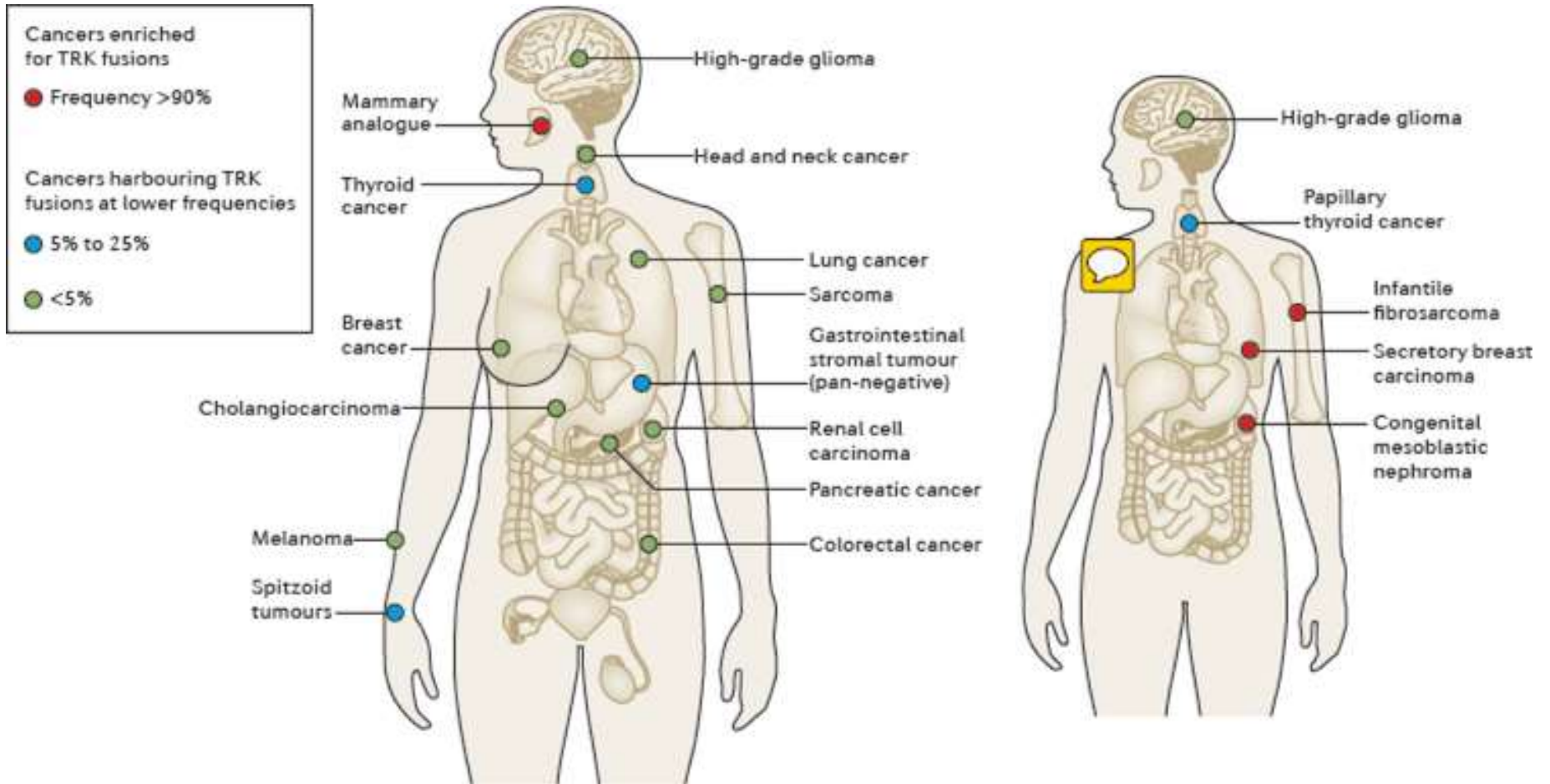


IDH 1 mutations have no prognostic value

Ivosidenib: IDH1 inhibitor

Javle M, *Cancer* 2016

Valle JW, *Cancer Discov* 2017



ESMO recommendations on the standard methods to detect NTRK fusions in daily practice and clinical research.

Marchiò C^{1,2}, Scaltriti M^{3,4}, Ladanyi M⁵, Iafrate AJ^{5,6}, Bibeau F⁷, Diotel M⁸, Hechtman JF³, Troiani T⁹, López-Ríos F¹⁰, Douillard JY¹¹, André F¹², Reis-Filho JS³.

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- 8 Institute of Pathology, Charité, University Medicine Berlin, Berlin, Germany.
- 9 Medical Oncology, Department of Precision Medicine, University of Campania "Luigi Vanvitelli", Naples, Italy.
- 10 Pathology & Targeted Therapies Laboratory, HM Sanchinarro University Hospital, Madrid, Spain.
- 11 European Society for Medical Oncology, Lugano, Switzerland.
- 12 Department of Medical Oncology, INSERM Unit 981, Institut Gustave Roussy, Villejuif, France.

Abstract

BACKGROUND: NTRK1, NTRK2 and NTRK3 fusions are present in a plethora of malignancies across different histologies. These fusions represent the most frequent mechanism of oncogenic activation of these receptor tyrosine kinases, and biomarkers for the use of TRK small molecule inhibitors. Given the varying frequency of NTRK1/2/3 fusions, crucial to the administration of NTRK inhibitors is the development of optimal approaches for the detection of human cancers harbouring activating NTRK1/2/3 fusion genes.

MATERIALS AND METHODS: Experts from several Institutions were recruited by the European Society for Medical Oncology (ESMO) Translational Research and Precision Medicine Working Group (TR and PM WG) to review the available methods for the detection of NTRK gene fusions, their potential applications, and strategies for the implementation of a rational approach for the detection of NTRK1/2/3 fusion genes in human malignancies. A consensus on the most reasonable strategy to adopt when screening for NTRK fusions in oncologic patients was sought, and further reviewed and approved by the ESMO TR and PM WG and the ESMO leadership.

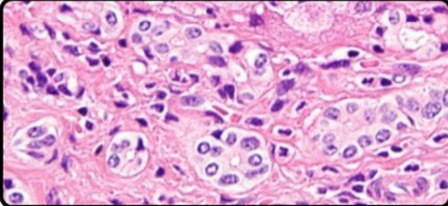
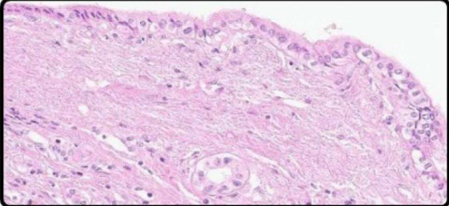
RESULTS: The main techniques employed for NTRK fusion gene detection include immunohistochemistry, fluorescence in situ hybridization (FISH), RT-PCR, and both RNA-based and DNA-based next generation sequencing (NGS). Each technique has advantages and limitations, and the choice of assays for screening and final diagnosis should also take into account the resources and clinical context.

CONCLUSION: In tumours where NTRK fusions are highly recurrent, FISH, RT-PCR or RNA-based sequencing panels can be used as confirmatory techniques, whereas in the scenario of testing an unselected population where NTRK1/2/3 fusions are uncommon, either front-line sequencing (preferentially RNA-sequencing) or screening by immunohistochemistry followed by sequencing of positive cases should be pursued.

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Molecular subclassification of intrahepatic cholangiocarcinoma

Inflammation subclass (38%)	Proliferation subclass (62%)
Activation of inflammatory pathways	Activation of oncogenic signalling pathways
Overexpression of cytokines (IL-10/IL-6)	RAS, MAPK
STAT3 activation	C-MET, BRAF, mutations in <i>KRAS</i>
Cholangiolar differentiation	Genomic resemblance to poor-prognosis hepatocellular carcinoma
	Aggressive clinical behaviours and poor prognosis
	Hepatic stem cell-like features
	Chromosomal instability
	IDH mutations
	Moderate/poor differentiation and intraneural invasion

Intrahepatic Cholangiocarcinoma		
Classification	Small Duct Type	Large Duct Type
Gross Type	Mass-forming	Mixed Periductal Infiltrating
Cell of Origin		
	Canal of Hering Bile ductule	Columnar cholangiocytes Peribiliary glands
Main Etiology	Chronic hepatitis HBV / HCV Alcoholic / Metabolic	Hepatolithiasis Liver fluke PSC
Immuno- histochemistry & Mucin stain	NCAM N-cadherin CRP	S100P Mucin
Frequent Mutations	<i>BAP1</i> <i>IDH1/2</i> <i>FGFR2 fusion</i>	<i>KRAS</i> <i>TP53</i> <i>SMAD4</i>
Suggested Molecular Classification*	Inflammation Class	Proliferation Class
Patient Outcome	Favorable	Poor

Combined hepatocellular- cholangiocarcinoma

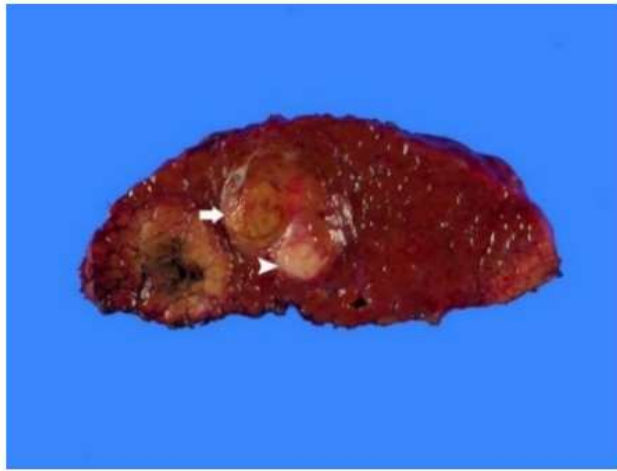
Table 1. Evolution of the WHO classification of cHCC-CCA.

	2000 WHO Classification (3rd Edition)	2010 WHO Classification (4th Edition)	2019 WHO Classification (5th Edition)
Tumor category	Malignant epithelial tumors	Malignancies of mixed or uncertain origin	Malignant biliary tumors
Tumor entities or subtypes	cHCC-CCA	cHCC-CCA, classical type cHCC-CCA with stem cell features ^(a) , typical subtype cHCC-CCA with stem cell features, intermediate-cell type cHCC-CCA with stem cell features, cholangiolocellular type	cHCC-CCA ^(b) Intermediate cell carcinoma ^(c) Cholangiolocarcinoma ^(d)

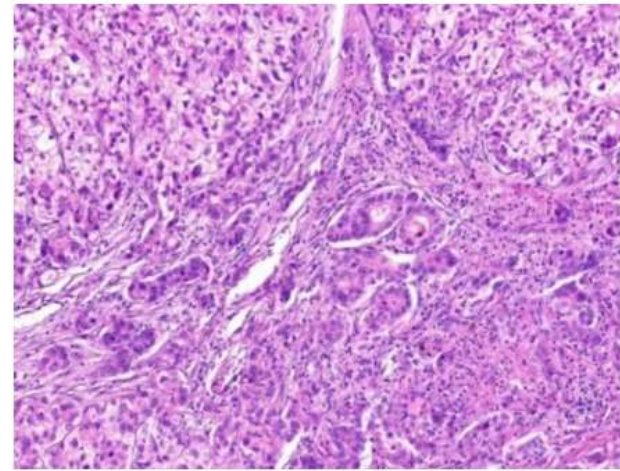
Combined hepatocellular-cholangiocarcinoma

- Unequivocal presence of both hepatocytic and cholangiocytic differentiation
- 2-5% of primary liver carcinomas
- Synonyms: mixed HCC-CCA, mixed hepatobiliary carcinoma, hepatocholangiocarcinoma, biphenotypic primary liver cancer

Combined hepatocellular- cholangiocarcinoma

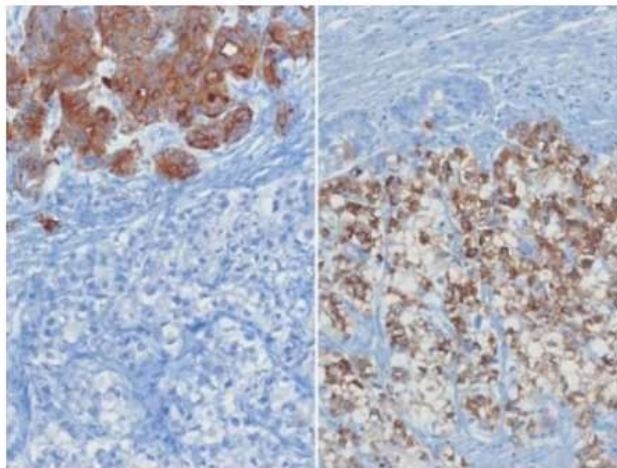


(A)



(B)

CK19



(C)

HepPar-1

Combined hepatocellular-cholangiocarcinoma: pathogenesis

- Three possible pathogenetic processes:
 - HCC and CCA may arise independently and separately
 - cHHC-CCA may originate from stem/progenitor cells that differentiate into both hepatocytic and cholangiocytic lines
 - HCC may arise first and transform into CCA at varying degrees

Theise ND, *Histopathology* 2003

Li L, *Am J Pathol* 2018

Choi JH, *Biomedicines* 2022

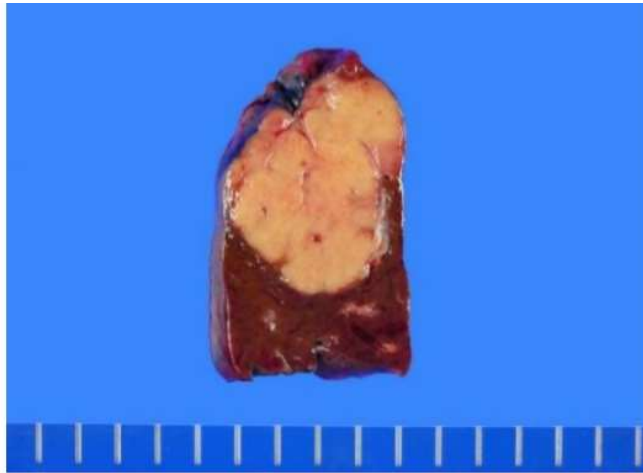
Combined hepatocellular-cholangiocarcinoma: histopathology

- Two components are either close to each other or extensively intermingled
- No consensus regarding a cutoff for each component
- Transitional are between the two components often exhibits mixed features with intermediate morphology
- Distant metastasis can show both components, or one single component

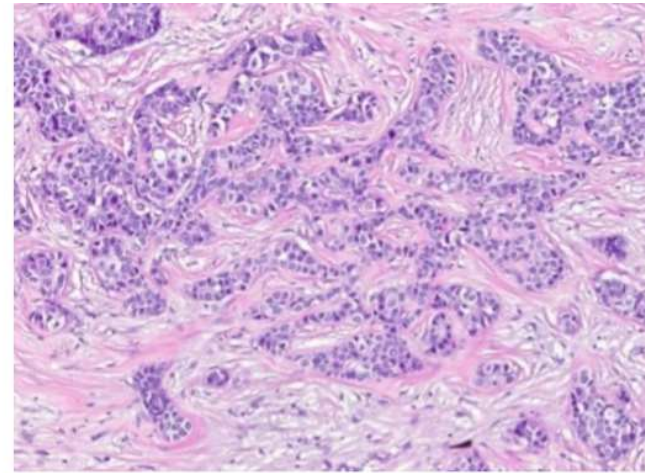
Intermediate cell carcinoma

- 2010, WHO classification: 'cHCC-CCA with stem cell features, intermediate-cell subtype'
- 2018, Brunt *et al.* : 'intermediate cell carcinoma'
- 2019, WHO classification: 'intermediate cell carcinoma'

Intermediate cell carcinoma

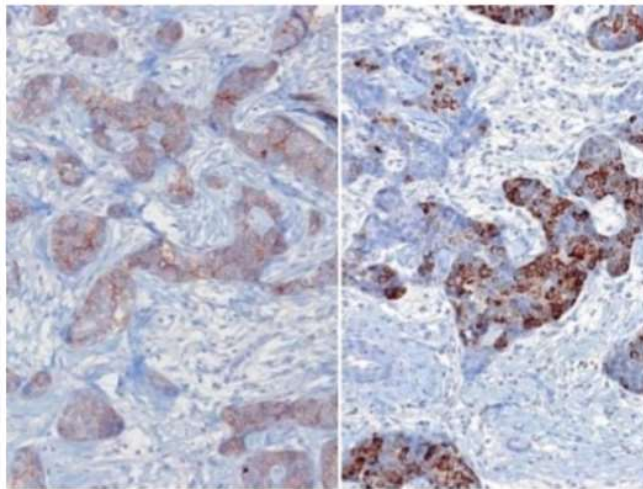


(A)



(B)

CK19



(C)

HepPar-1

