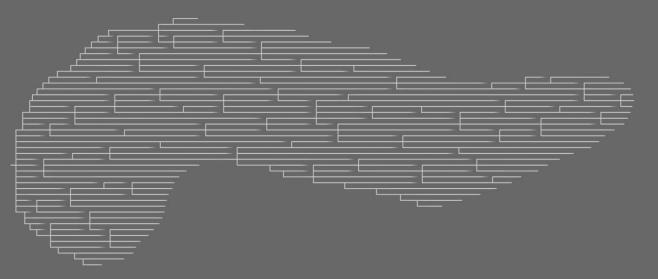
Genetics of Pancreatic Cancer



Michaël Noë

Resident, UMC Utrecht, The Netherlands

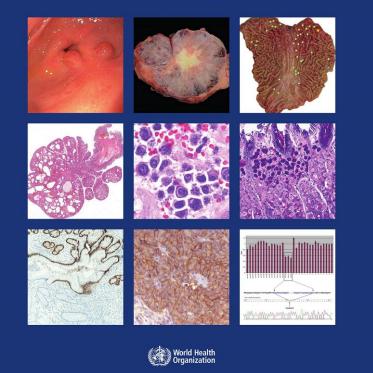
Post-doctoral Research Fellow, Johns Hopkins University, USA

WHO: 5th edition

WHO Classification of Tumours • 5th Edition

Digestive System Tumours

Edited by the WHO Classification of Tumours Editorial Board

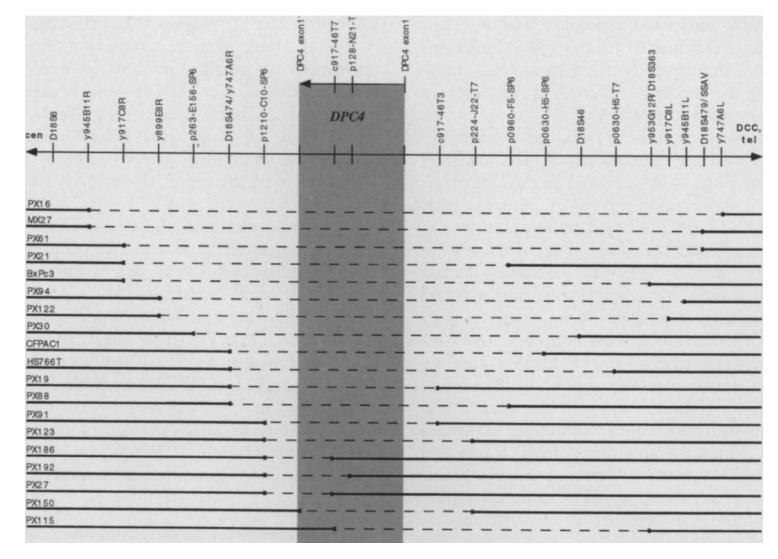


Diagnostic molecular pathology Not clinically relevant

History of the genetics of pancreatic cancer

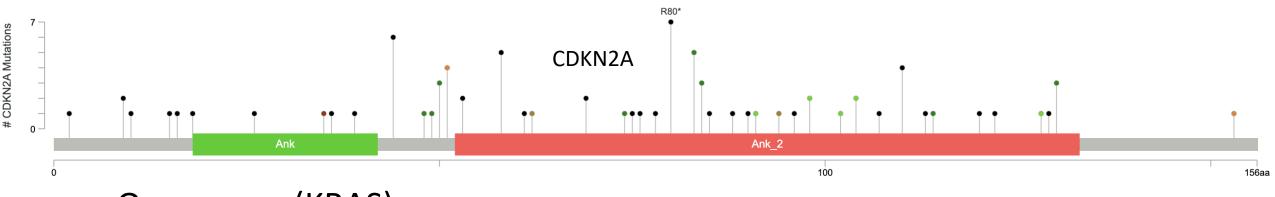
- 1982: KRAS discovered in pancreatic cancer cell lines
 - Similarities with Rous Sarcoma Virus
- 1995: DPC1 & DPC2 discovered (later: BRCA2)
 - 'Deleted in Pancreatic Cancer'
 - Finding spots of homozygous deletions in cancer cell lines (RDA)
- 1994: DPC3 discovered (later: P16, CDKN2A)
- 1996: DPC4 discovered (later: SMAD4)

DPC: Deleted in Pancreatic Cancer

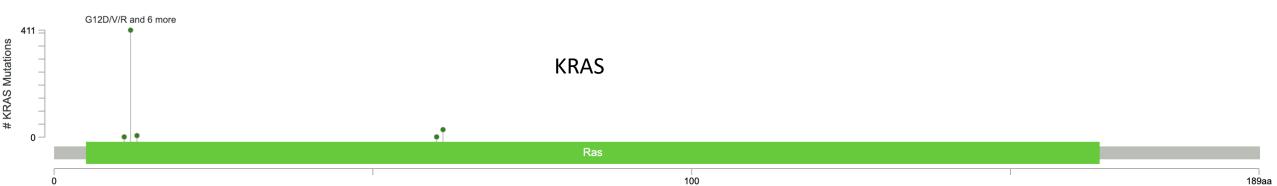


Tumor Suppressors vs. Oncogenes

- Tumor suppressors (CDKN2A, TP53, SMAD4)
 - many ways to break them (chaos = more entropy)



- Oncogenes (KRAS)
 - Only a couple of ways to make it work better / more (structure = less entropy)

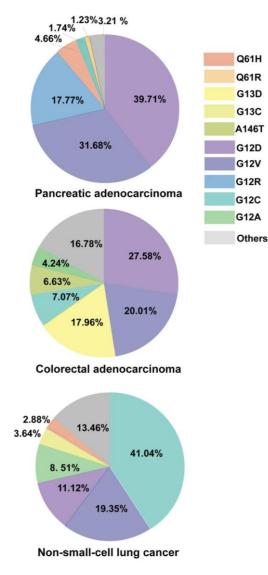


Pancreatic cancer: driver genes

- KRAS
- CDKN2A (P16)
- TP53
- SMAD4 (immunohistochemistry if it works)

Pancreatic cancer: driver genes

- KRAS
- CDKN2A (P16)
- TP53
- SMAD4

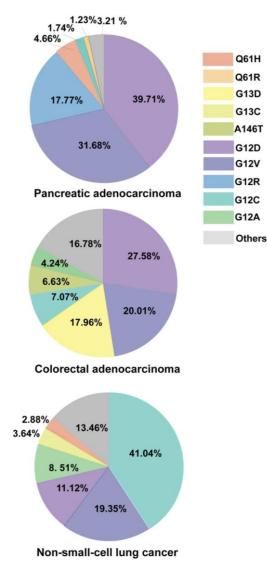


Different KRAS-mutations in different cancers: reflection of different impact of carcinogens

- Smoking (C>A)
- Bacteria
- Aging
- APOBEC
- etc...

Pancreatic cancer: driver genes

- KRAS
- CDKN2A (P16)
- TP53
- SMAD4

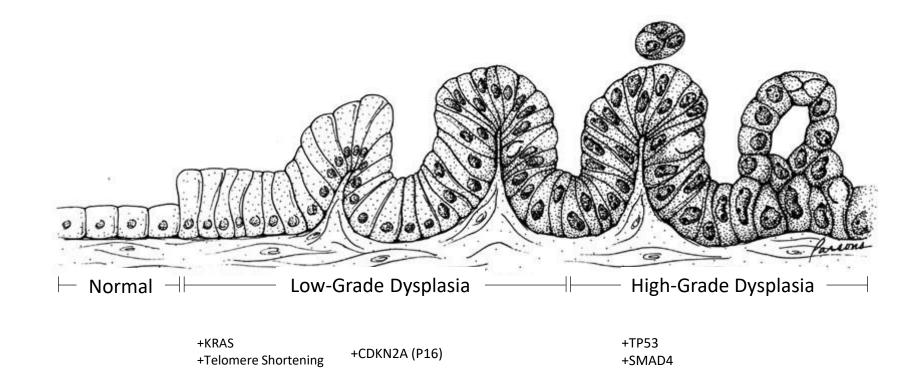


→ Sotorasib: irreversible inhibitor of KRAS G12C (1-2%)

- ASCO 2022: CodeBreaK100 study showed effect of the therapy in pancreatic cancer patients
 - Partial response: 21%
 - Disease control: 84%

From precursor to carcinoma (PDAC)

• Deciphering the pathogenesis:



Precursors to pancreatic ductal adenocarcinoma (PDAC)

Precursor	Malignant	Mutations
PanIN	PDAC	KRAS, CDKN2A, TP53, SMAD4
IPMN	PDAC-light	KRAS, <u>GNAS</u> , CDKN2A, <u>RNF43</u> , TP53, SMAD4
IOPN	PDAC-light	Fusions between ATP1B1 / DNAJB1 and PRKACA / PRKACB
MCN	PDAC-light	KRAS, CDKN2A, <u>RNF43</u> , TP53, SMAD4
ITPN	PDAC	PIK3CA, PTEN, KMT2A, KMT2B, KMT2C, FGFR2 fusions
SCAdenoma	SCAdenocarcinoma?	VHL
	SPN	CTNNB1
	Acinar Cell Carcinoma	BRCA2, CHEK2, FAT

PanIN: Pancreatic Intraepithelial Neoplasia PDAC: Pancreatic Ductal Adenocarcinoma IPMN: Intraductal Papillary Mucinous Neoplasm IOPN: Intraductal Oncocytic Papillary Neoplasm MCN: Mucinous Cystic Neoplasm ITPN: Intraductal Tubulopapillary Neoplasm SCAdenoma: Serous Cystadenoma SPN: Solid Pseudopapillary Neoplasm

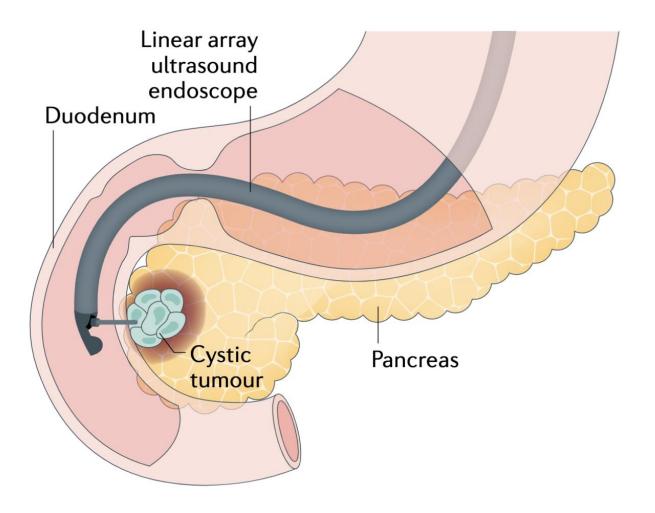
DNAJB1 - PRKACA fusions: also seen in fibrolamellar hepatocellular carcinoma

PDAC is molecular heterogenic

Small amounts of targetable driver-mutations

- Microsatellite instability (immunotherapy)
- ERBB2 amplifications (trastuzumab)
- Targets for Kinase Inhibitors (BRAF, FGFR2, FGFR3,...)
- Homologous Repair Deficiency (BRCA2, ATM,...)

Application: cyst fluid analysis



Diagnosis of cysts:

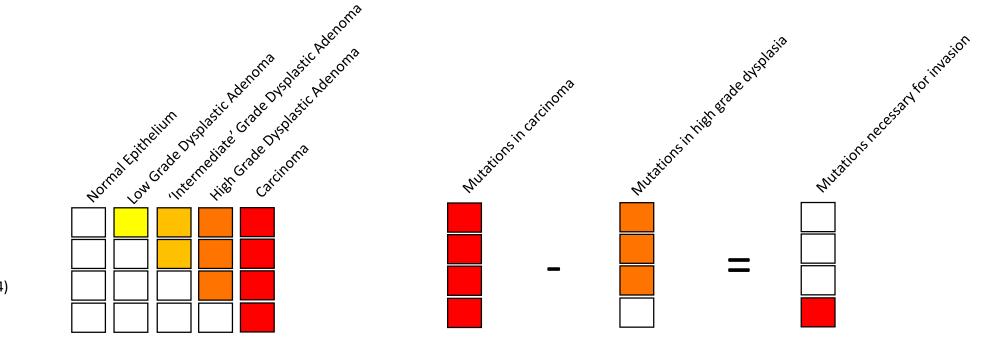
- Enzymes (CEA and amylase)
- Driver mutation analysis

Risk stratification of IPMNs / MCNs:

• TP53, SMAD4: invasion

Discovering the driver mutations for invasion

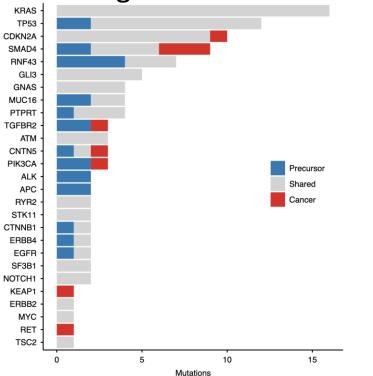
- Vogelstein et al., New England Journal of Medicine, 1988:
 - FAP patients
 - Try to detect the previously discovered molecular aberrations of colorectal cancers in adenomatous polyps



Loss of APC KRAS mutations Loss of 18q (SMAD4) Loss of 17p (TP53)

IPMN to PDAC: driving invasion

- Deciphering the pathogenesis (whole-exome sequencing):
 - 4/18 cases had cancer specific SMAD4/TGFBR2 mutations
 - Other cancer-specific driver-genes were CDKN2A, CNTN5, PIK3CA, KEAP1 and RET

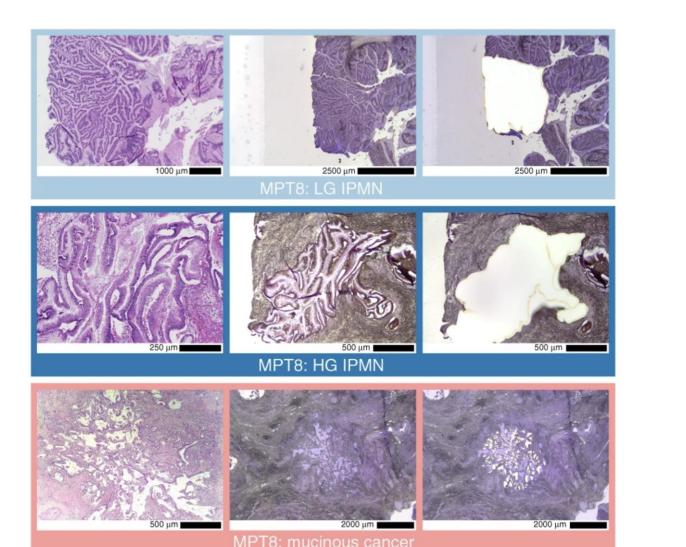


Noë M. et al. Genomic characterization of malignant progression in neoplastic pancreatic cysts. Nat Commun. 2020

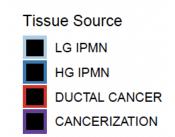
Association, but not always causation

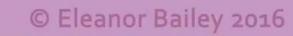
- Clonal relationship between concurrent IPMN and PDAC:
 - 1/3 of IPMNs in patients with a PDAC is not clonally related!
 - Different driver mutations
 - Probably a smaller PanIN lesion resulted in the PDAC
 - Reporting:
 - Just describe what you see (IPMN, PanIN and PDAC)
 - Use of 'Malignant IPMN' is discouraged

Cancerization: molecular evidence

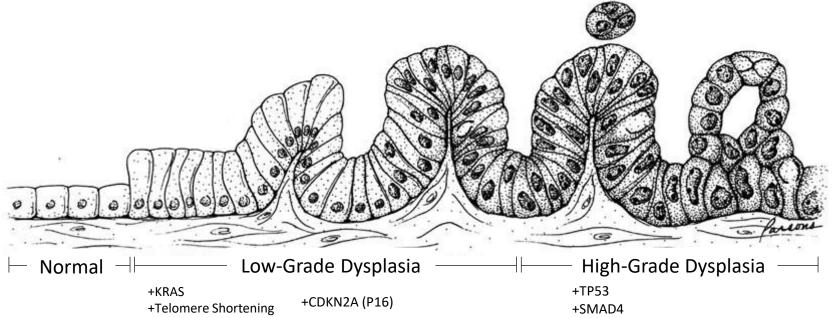


KRAS:G12R KRAS:Q61H-12:25380275 T/G KRAS:Q61H-12:25380275 T/A TMEM56-RWDD3:c.566+2T>C [SP] KCNA5:A451A C17orf98:T142M DUSP27:T652T ADAMTS12:P429L COL12A1:R933H SCAF8:G961V TMEM246:A372A MCM10:E613K PNLIPRP1:T465T C13orf35:L33L SMAD6:L179L GRIN2A:S397S AP1G1:T798T DDDDDDD TP53:E171* FCH01:c.2247+1G>C [SP] LRFN1:S632S THE PERSON NUMBER OF STREET LOH:chr17:0.01-18.26 ABR:F645S ALPK1:R873R LOH:chrX:2.75-48.42 DEL:chr9:21.8-23.77(CDKN2A) BSN:D861N SUPT6H:1104T PLEKHF1:D258D LOH:chr22:17.16-51.12 LTBR:R425G ACOT11:R306L OR6N1:Y120Y ENPP5:Y341Y ITGB3:C562* BTK:P116T SLC1A7:G465D FGF3:R104Q ZBED4:P1100L SIM1:R192C PHC1:S627C MS4A1:T41T LOH:chr7:127.25-158.72 SH2D5:R199W RNPEPL1:G413V GAL3ST2:P85P JADE2:N352S HHLA1:A97V TMEM141:Q61Q BPTF:R296H KLHL22:R603R GALE:Q261L ANKS4B:Q368R KRI1:S326S OR812:C240C LRTM2:G319V LOH:chr9:71.23-139.96 LAMB3:R887L SPTAN1:I1484F TP53:R175H





Implications of the Driver Mutations Sequence



- The sequence is important to evaluate risk from germline mutations
 - CDKN2A (P16): early driver mutation in PDAC
 - Germline mutations: Familial Atypical Multiple Mole Melanoma syndrome (FAMMM) have high risk for PDAC
 - SMAD4: late driver mutation in PDAC
 - Germline mutations: Juvenile Polyposis syndrome have normal / low risk for PDAC

Conclusions

- KRAS is targetable
- Small amount of PDACs harbour targetable driver mutations
- PDAC is heterogeneous and subgroups have different prognosis
 - Precursors have different mutations, which can be used for diagnosis on cyst fluid
 - Small amount of PDACs have targetable driver mutations
- Co-occurring IPMNs and PDACs are not always related
- PDACs can grow back in ducts and present as 'precursor' lesion

Thank you

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