

INTEGRATING DNA MOLECULAR TESTING INTO THE ROUTINE EVALUATION OF PANCREATIC CYSTS

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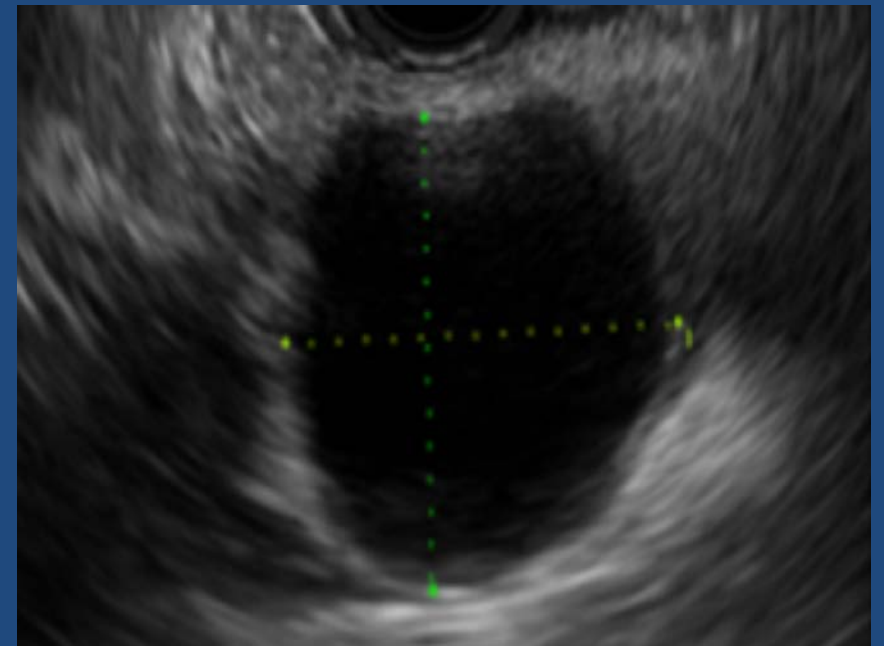
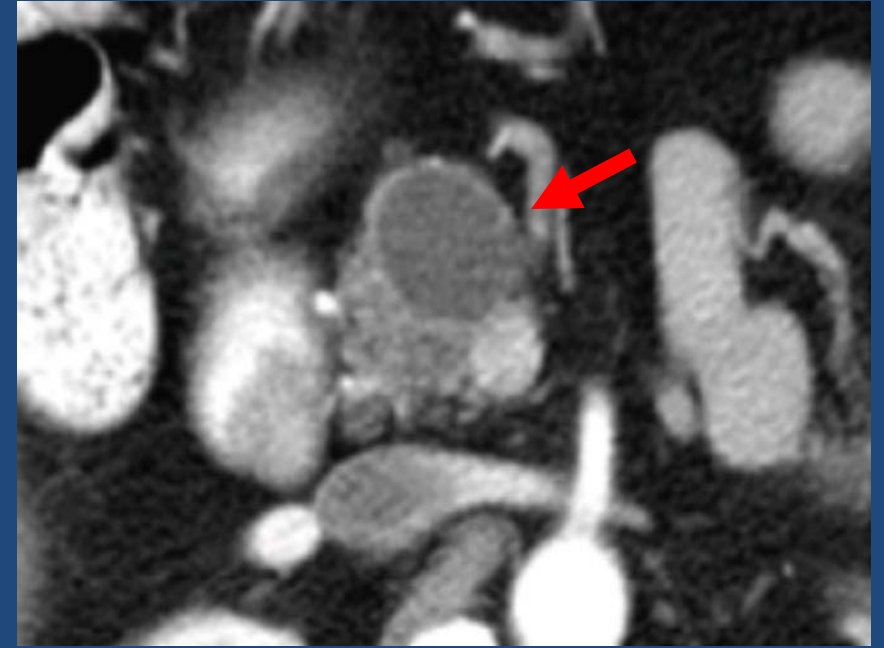
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Disclosures

- **None of the presenters have financial relationships or interests to disclose.**

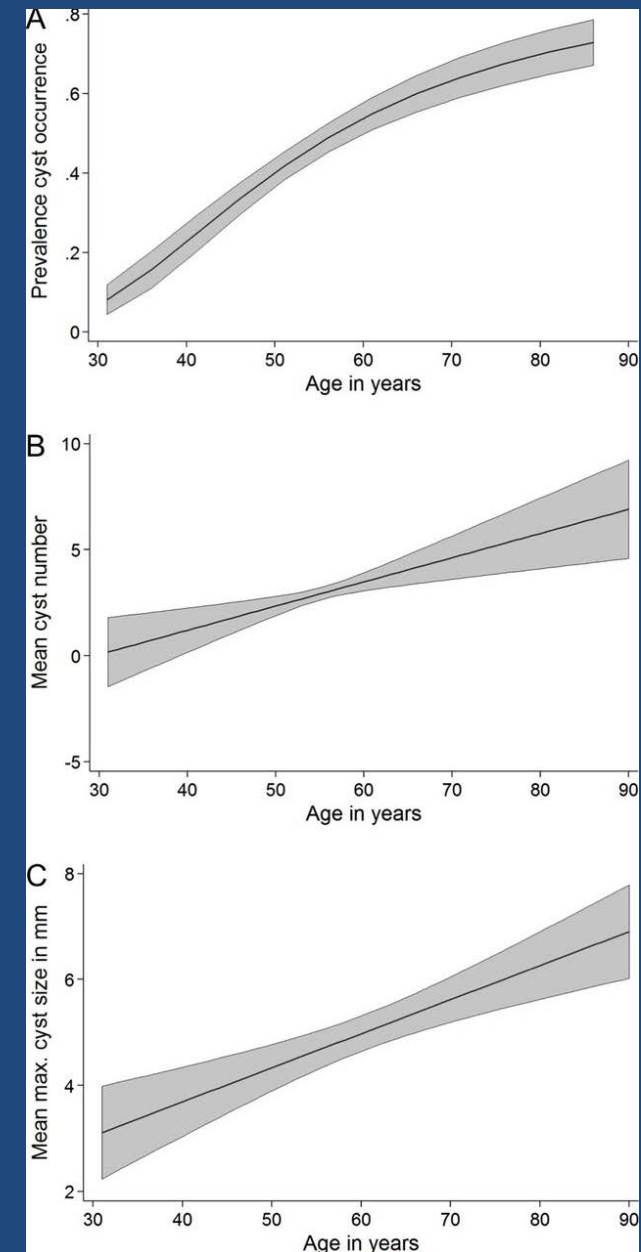
Clinical Case

- 61 yo WM with 3 cm incidental pancreatic head cyst
- No associated main duct dilation or mural nodule
- EUS-FNA cytology: atypical cells, no mucin
- CEA 157 ng/ml
- DNA analysis
 - *GNAS* (Allele Freq. 51%)
 - *PIK3CA* (Allele Freq. 50%)



Pancreatic Cyst Prevalence

- ~ 70% discovered incidentally
- MRI: 13.5%-19.6%
- Autopsy: up to 50% in elderly
- Meta-analysis: 49K pts, 17 studies
 - Pooled prevalence: 8%
- Study of Health in Pomerania
 - 1077 pts underwent MRI
 - 49.1% prevalence (cysts > 2 mm)
 - 12.9% incidence during 5 yr f/u
 - > 80 yrs: 75.7%, mean size 4.3 mm
 - 30-39 yrs: 17.1%, mean size 6.8 mm
 - 0.7% had cysts > 2 cm



Zerboni et al. Pancreatology 2019;19:2-9

Kromrey et al. Gut 2018;67:138-145

Pancreatic Cyst Classification

• Mucinous

- Mucinous cystic neoplasm (**MCN**)
- Intraductal papillary mucinous neoplasm (**IPMN**)

• Non-mucinous

- Serous cystadenoma (**SCA**)
- Cystic pancreatic neuroendocrine tumor
- Solid pseudopapillary neoplasm (**SPN**)
- Lymphoepithelial cyst
- Retention cyst
- Pseudocyst

Most Common Pancreatic Cystic Neoplasm?

- 376 pts
 - Resected between 2005 – 2011
 - **IPMN 49%**
 - MCN 16%
 - SCA 12%
 - SPN 5%
- Cystic pancreatic neuroendocrine tumor 8%

Guidelines for Evaluation / Management

- International Association of Pancreatology
 - Sendai guidelines 2006
 - Fukuoka guidelines 2012
 - Cyst fluid analysis is still investigational, but is recommended for evaluation of small BD-IPMNs without “worrisome features” in centers with expertise in EUS-FNA and cytological interpretation
 - Revised Fukuoka guidelines 2017

Tanaka et al. Pancreatology 2006;6:17-32

Tanaka et al. Pancreatology 2012;12:183-97

Tanaka et al. Pancreatology 2017;17:738-53

Revised Fukuoka Guidelines

- High risk stigmata

- Obstructive jaundice in a pt with a pancreatic head cyst
- Enhanced mural nodule ≥ 5 mm
- MPD size ≥ 10 mm

- Worrisome features

- Cyst size ≥ 3 cm
- Enhancing mural nodule < 5 mm
- MPD size 5 - 9 mm
- Abrupt change in MPD caliber with distal pancreatic atrophy
- Lymphadenopathy
- Elevated CA 19-9
- Rapid cyst growth > 5 mm / 2 yrs

Guidelines for Evaluation / Management

- AGA
- ACG
- ASGE
- UPMC

Vege et al. Gastroenterology 2015;148:819-22

Elta et al. Am J Gastroenterol 2018;113:464-79

Jacobson et al. Gastrointest Endosc 2005;61:363-70

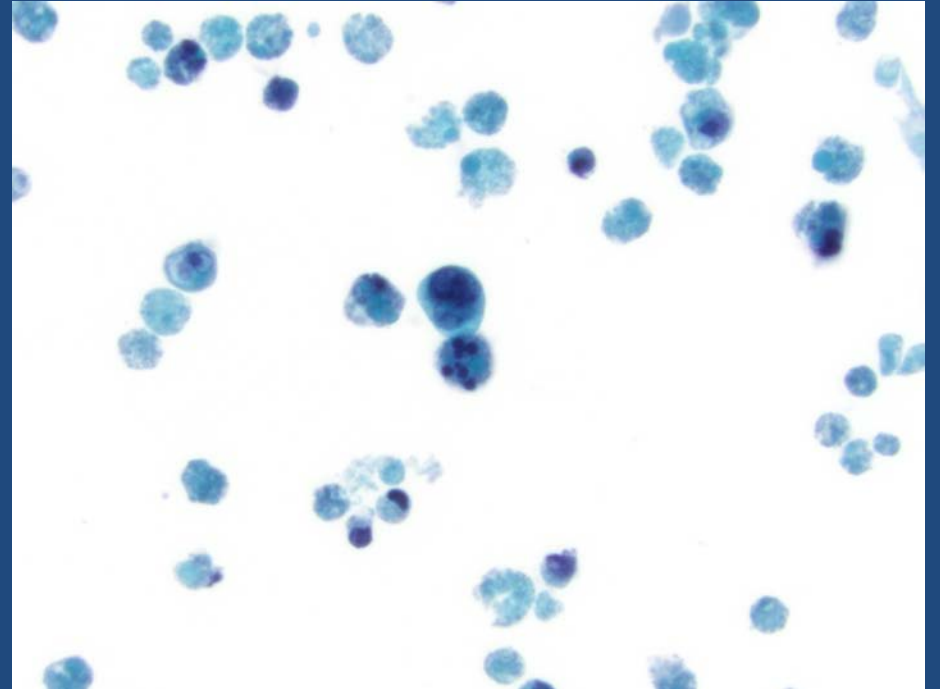
Singhi et al. Gastrointest Endosc 2016;83:1107-1117

Cytology: Mucinous vs Non-Mucinous Cysts

- Paucicellular specimen
- Viscosity dependent?

- CPC
 - sens 35%, spec 83%, accuracy 59%

- MGH
 - sens 43%, spec 96% , accuracy 58%



CEA: Differentiating Mucinous Cysts

- CPC
 - 112 pts
 - CEA optimal cutoff of **192** ng/mL (AUC 0.79, 73% sens, 83% spec)
- MGH
 - 198 pts
 - CEA optimal cutoff **110** ng/ml (AUC 0.93, 81% sens, 98% spec)
- Multicenter
 - 226 pts
 - CEA optimal cutoff **105** ng/ml (AUC 0.77, sens 70%, spec 63%)

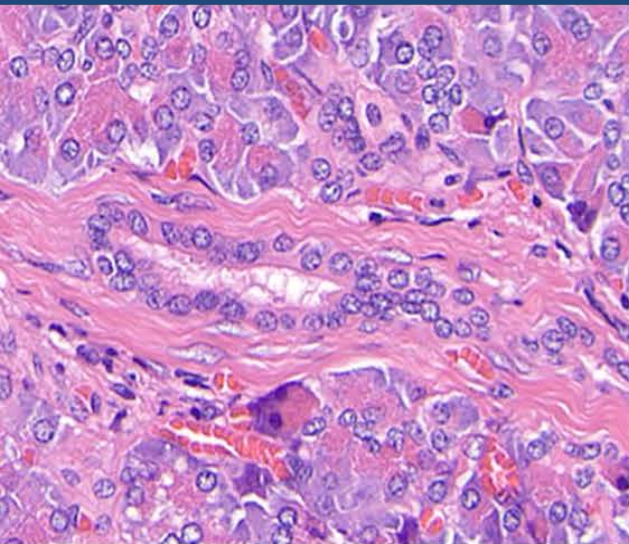
Brugge et al. Gastroenterology 2004;126:1330-6

Cizginer et al. Pancreas 2011;40:1024-8

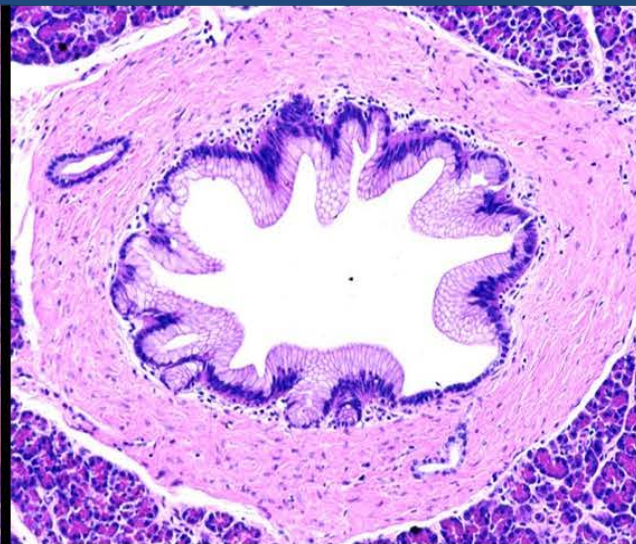
Gaddam et al. Gastrointest Endosc 2015;82:1060-9

Pancreatic Cystic Neoplasms

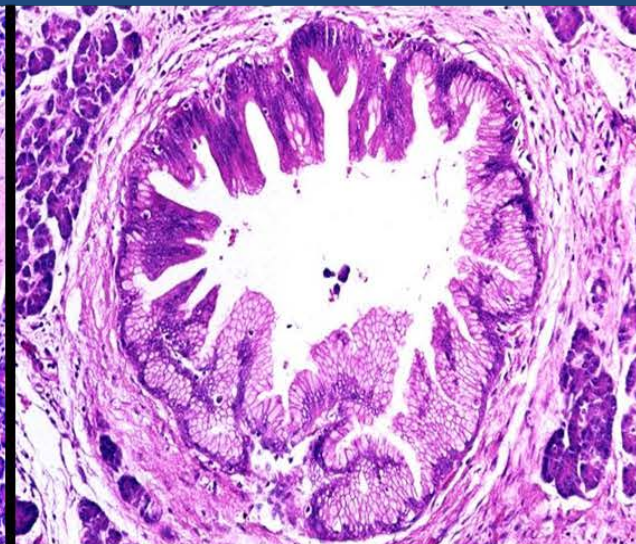
- Neoplastic transformation in cell morphology is preceded / paralleled by genetic alterations



Normal Duct



Low-Grade



High-Grade

- Hypothesis: Detection of established DNA mutations in cyst fluid may improve the yield of EUS-FNA and reflect biologic behavior

Pancreatic Cyst Fluid Molecular Testing

The role of pancreatic cyst fluid molecular analysis in predicting cyst pathology:

- 2005 pilot study
- Cyst fluid does harbor DNA for molecular analysis
- DNA quantity / quality
- *KRAS* point mutations
- Tumor suppressor gene Loss of Heterozygosity (LOH)
- Sequence – first hit *KRAS* followed by LOH predicts malignancy as does number of mutations and DNA quantity

PANDA

- Multicenter study of cyst fluid DNA analysis
 - 113 patients
 - CEA (AUC 0.74; optimal value >148 ng/mL)
 - *KRAS*: predictive of mucinous cyst
 - Predictive of malignancy:
 - Elevated DNA amount
 - High amplitude mutations
 - Sequence of mutations
- Consider DNA analysis when cytology is negative

Molecular Analysis: 10 yrs ago

- Molecular profiles may allow:
 - Differentiation of mucinous vs non-mucinous cysts
 - *KRAS*
 - Selection of high-risk lesions for surgical resection
 - DNA quantity / quality
 - LOH
- Prediction of the malignant potential of mucinous cysts

Table 1. k-ras-2 Gene and Tumor Suppressor Genes (With Associated Markers) With Chromosomal Location and Mutation Type

Proximity cancer gene ^a	Mutation type	Locus ^b	Marker 1	Marker 2
k-ras	Point mutation ^c	12p12		
CMM/RIZ	Allelic imbalance ^d	1p36–1p34	D1S407	MYCL
VHL	Allelic imbalance ^d	3p26–3p25	D3S1539	D3S2303
APC	Allelic imbalance ^d	5q23–5q23	D5S592	D5S615
P16	Allelic imbalance ^d	9p21–9p23	D9S251	D9S254
PTCH ^e	Allelic imbalance ^d	9q22	D9S252	
PTEN	Allelic imbalance ^d	10q23–10q23	D10S520	D10S1173
P53	Allelic imbalance ^d	17p13–17p13	D17S974	D17S1289

Pancreatic Cyst Molecular Testing

Whole-exome sequencing of neoplastic cysts of pancreas reveals recurrent mutations in components of ubiquitin-dependent pathways

Jian Wu^{a,1}, Yuchen Jiao^{a,1}, Marco Dal Molin^{b,1}, Anirban Maitra^b, Roeland F. de Wilde^b, Laura D. Wood^b, James R. Eshleman^b, Michael G. Goggins^{b,c}, Christopher L. Wolfgang^d, Marcia I. Canto^e, Richard D. Schulz^f, Barish H. Edil^d, Michael A. Choti^d, Volkan Adsay^e, David S. Klimstra^f, G. Johan A. Offerhaus^g, Alison P. Levy^h, Kopelovich^h, Hannah Carterⁱ, Rachel Karchinⁱ, Peter J. Allen^j, C. Max Schmidt^k, Yoshiki Naito^l, Luis Kent

A Combination of Molecular Markers and Clinical Features Improve the Classification of Pancreatic Cysts

Simeon Springer,^{1,2,*} Yuxuan Wang,^{1,2,*} Marco Dal Molin,^{2,3,*} David L. Masica,^{2,4,5,*} Yuchen Jiao,^{1,2} Isaac Kinde,^{1,2} Amanda Blackford,⁶ Siva P. Raman,⁷ Christopher L. Wolfgang,^{2,8,9} Tyler Tomita,^{4,5} Noushin Niknafs,^{4,5} Christopher Douville,^{4,5} Janine Ptak,^{1,2} Lisa Dobbyn,^{1,2} Peter J. Allen,¹⁰ David S. Klimstra,¹¹ Mark A. Schattner,¹² C. Max Schmidt,¹³ Michele Yip-Schneider,¹⁴ Oscar W. Cummings,¹⁴ Randall E. Brand,¹⁵ Herbert J. Zeh,¹⁶ Aatur D. Singhi,¹⁷ Aldo Scarpa,^{18,19} Roberto Salvia,²⁰ Giuseppe Malleo,²⁰ Giuseppe Zamboni,^{19,21} Massimo Falconi,²² Jin-Young Jang,²³ Sun-Whe Kim,²³ Wooil Kwon,²³ Seung-Mo Hong,²⁴ Ki-Byung Song,²⁵ Song Cheol Kim,²⁵ Niall Swan,²⁶ Jean Murphy,²⁶ Justin Geoghegan,²⁷ William Brugge,²⁸ Carlos Fernandez-Del Castillo,²⁹ Mari Mino-Kenudson,³⁰ Richard Schulick,³¹ Barish H. Edil,³¹ Volkan Adsay,³² Jorge Paulino,³³ Jeanin van Hooft,³⁴ Shinichi Yachida,³⁵ Satoshi Nara,³⁵ Nobuyoshi Hiraoka,³⁵ Kenji Yamao,³⁶ Susuma Hijioka,³⁶ Schalk van der Merwe,³⁷ Michael Goggins,^{2,9,38} Marcia Irene Canto,³⁸ Nita Ahuja,⁸ Kenzo Hirose,⁸ Martin Makary,⁸ Matthew J. Weiss,⁸ John Cameron,⁸ Meredith Pittman,^{2,3} James R. Eshleman,^{1,2} Luis A. Diaz Jr.,^{1,2,8} Nickolas Papadopoulos,^{1,2} Kenneth W. Kinzler,^{1,2} Rachel Karchin,^{2,4,5,9} Ralph H. Hruban,^{1,2,3,9} Bert Vogelstein,^{1,2} and Anne Marie Lennon^{2,8,38}

Targeted DNA Sequencing Reveals Patterns of Local Progression in the Pancreatic Remnant Following Resection of Intraductal Papillary Mucinous Neoplasm (IPMN) of the Pancreas

Antonio Pea, MD,*† Jun Yu, MD,‡ Neda Rezaee, MD,* Claudio Luchini, MD,‡§ Jin He, MD, PhD,* Marco Dal Molin, MD,‡ James F. Griffin, MD,* Helen Fedor,‡ Shahriar Fesharakizadeh,‡

Digital next-generation sequencing identifies



CrossMark

mutations in pancreatic juice from the duodenum of patients with intraductal papillary

pancreatic cysts. Masaya Suenaga,¹ Aaron Brant,¹ Haniel Borges,¹ Thomas Barkley,¹ Richard Hruban,^{1,2} Eun Ji Shin,³ Marcia I. Canto,^{2,3} Michael Goggins^{1,2,3}

from the prospective ZYSTEUS biomarker

Endris¹ | Matthias M. Gaida¹ | Jonas Leichsenring¹ | J. Hübner¹ | Moritz von Winterfeld¹ | Roland Penzel¹ | J. Hübner¹ | Olaf Neumann¹ | Holger Sültmann² | J. Hübner¹ | Daniel Schmitz^{3†} | Albrecht Stenzinger^{1,4†}

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Pancreatic Cyst Molecular Testing

	<i>KRAS</i>	<i>GNAS</i>	<i>VHL</i>	<i>CTNNB1</i>
• Intraductal papillary mucinous neoplasms (IPMN):				
IPMN	+	+	-	-
	<i>KRAS</i>	<i>GNAS</i>		
• Mucinous cystic neoplasms (MCN):				
MCN	+	-	-	-
• Serous cystadenomas (SCA):				
SCA	-	-	+	-
	<i>KRAS</i>			
• Solid pseudopapillary neoplasms (SPN):				
SPN	-	-	-	+
• Non-neoplastic cysts:				
Non	-	-	-	-
	<i>VHL</i>			

- Solid pseudopapillary neoplasms (SPN):

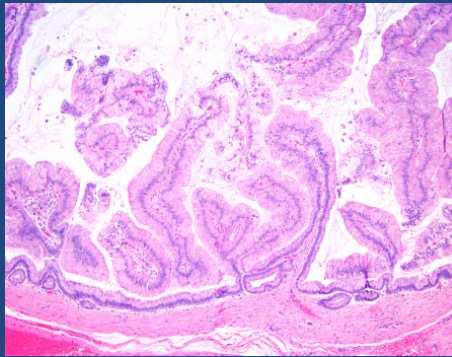
CTNNB1

- Non-neoplastic cysts:

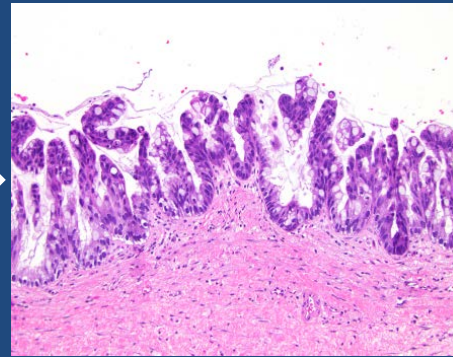
Absent

Pancreatic Cyst Molecular Testing

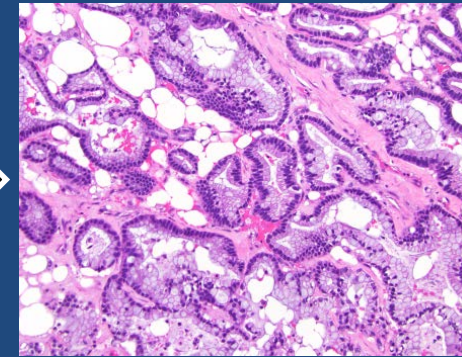
	<i>KRAS</i>	<i>GNAS</i>	<i>VHL</i>	<i>CTNNB1</i>
IPMN	+	+	-	-
MCN	+	-	-	-



Low-grade
Dysplasia



High-grade
Dysplasia



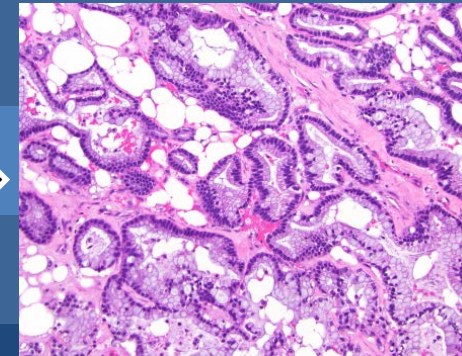
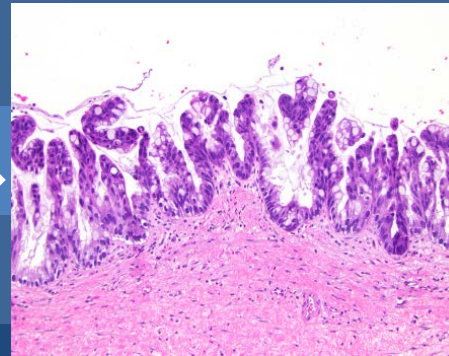
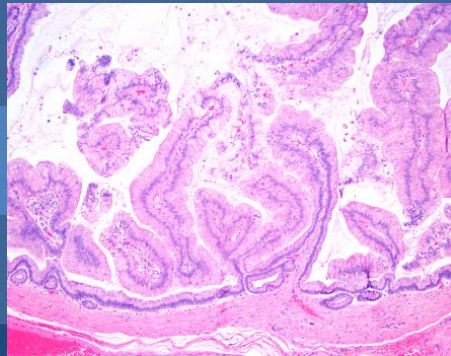
AdenoCA

KRAS & *GNAS*



Pancreatic Cyst Molecular Testing

	<i>KRAS</i>	<i>GNAS</i>	<i>VHL</i>	<i>CTNNB1</i>	<i>TP53</i>	<i>PIK3CA</i>	<i>PTEN</i>
IPMN	+	+	-	-	HR	HR	HR
MCN	+	-	-	-	HR	HR	HR



Low-grade
Dysplasia

HR = High-risk of malignancy

High-grade
Dysplasia

AdenoCA

KRAS & *GNAS*



TP53, *PIK3CA*
& *PTEN*

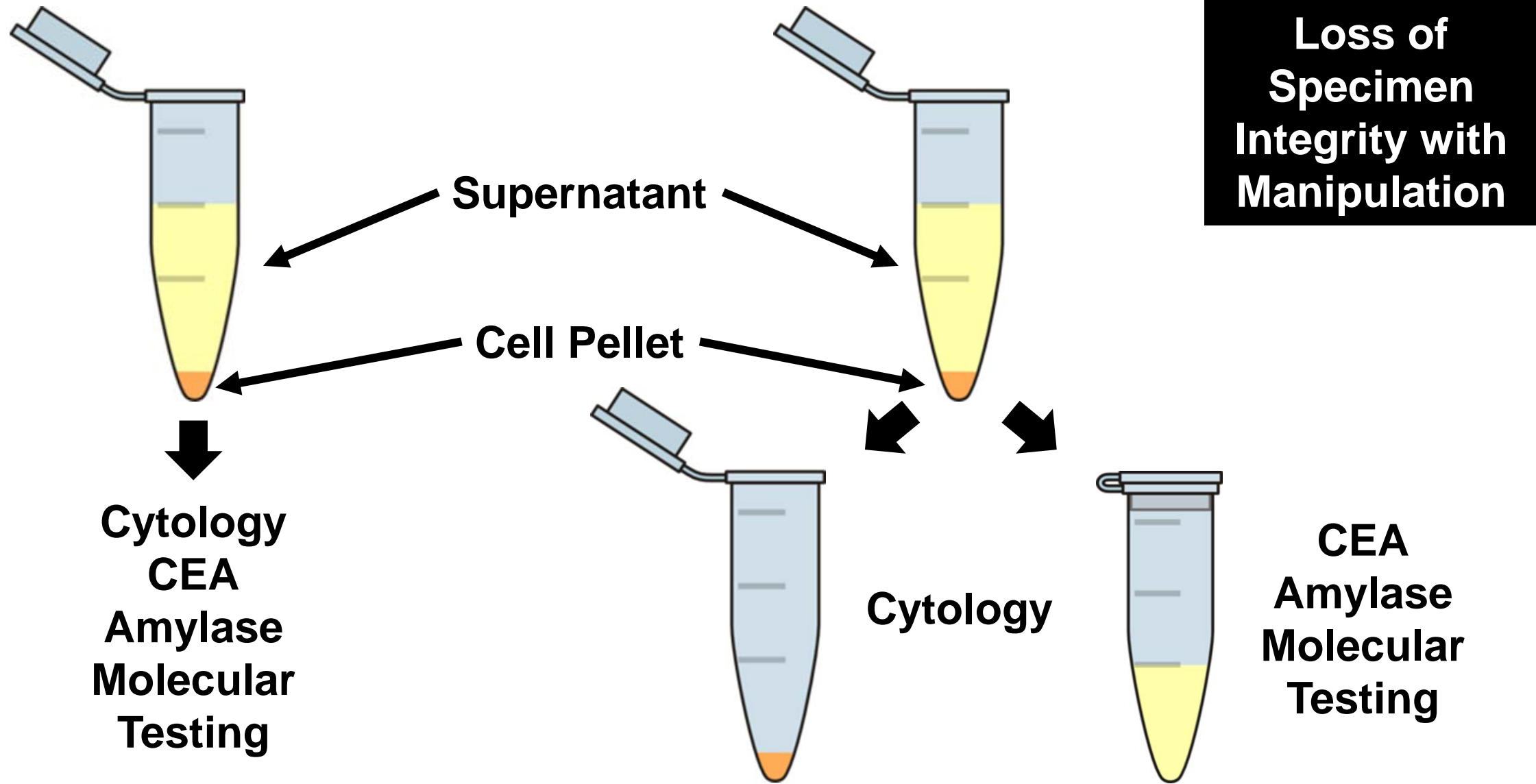


Pancreatic Cyst Molecular Testing

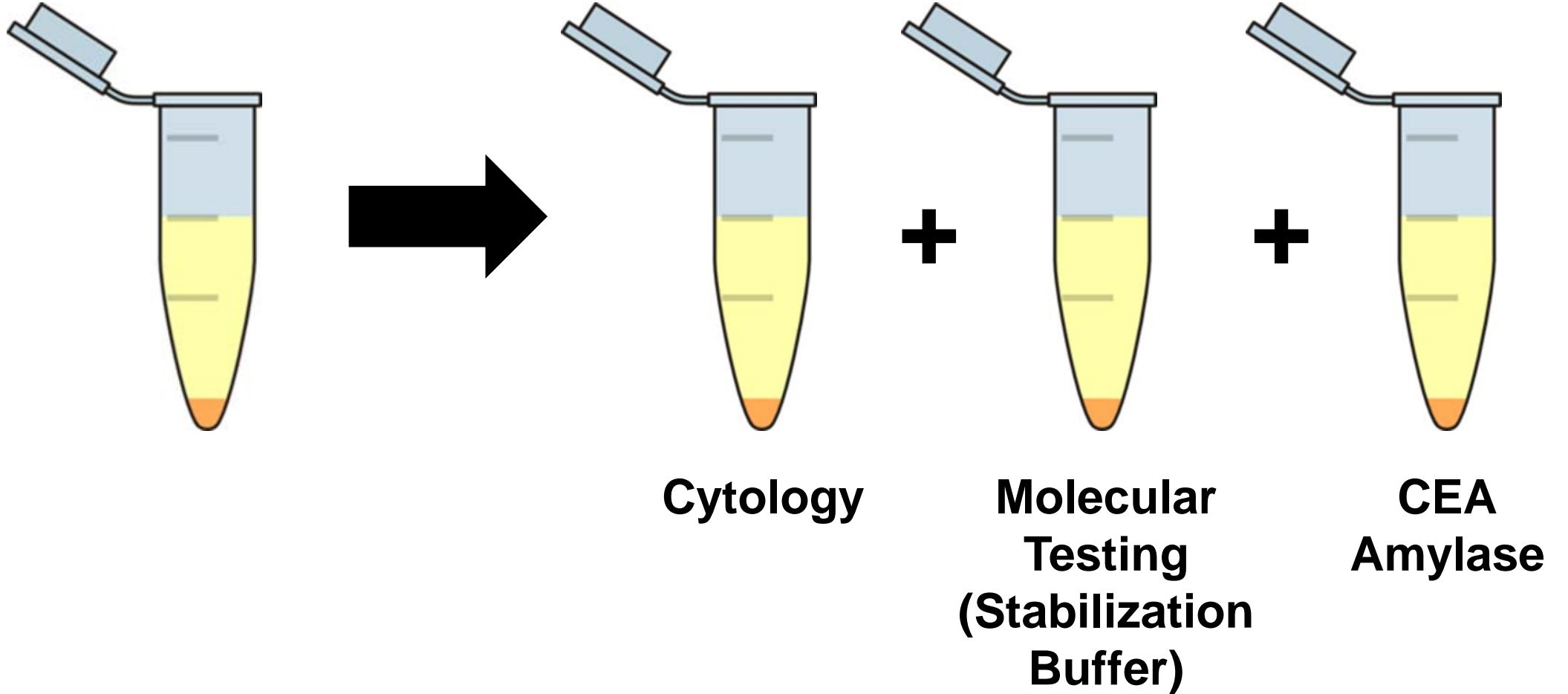
KRAS GNAS VHL CTNNB1 TP53 PIK3CA PTEN

- Next-generation sequencing panel (**PancreaSeq**) to assess preoperative EUS-FNA obtained pancreatic cyst fluid designed in **2013**.
- Exons 1 through 3 of *VHL* were assessed by Sanger sequencing, limit of detection 10-20%.
- >1,000 hot spot mutations with over 1000x to 500x depth of coverage, corresponding to a limit of detection of 3% to 5%, respectively.
- Samples below 500x were not interpreted.

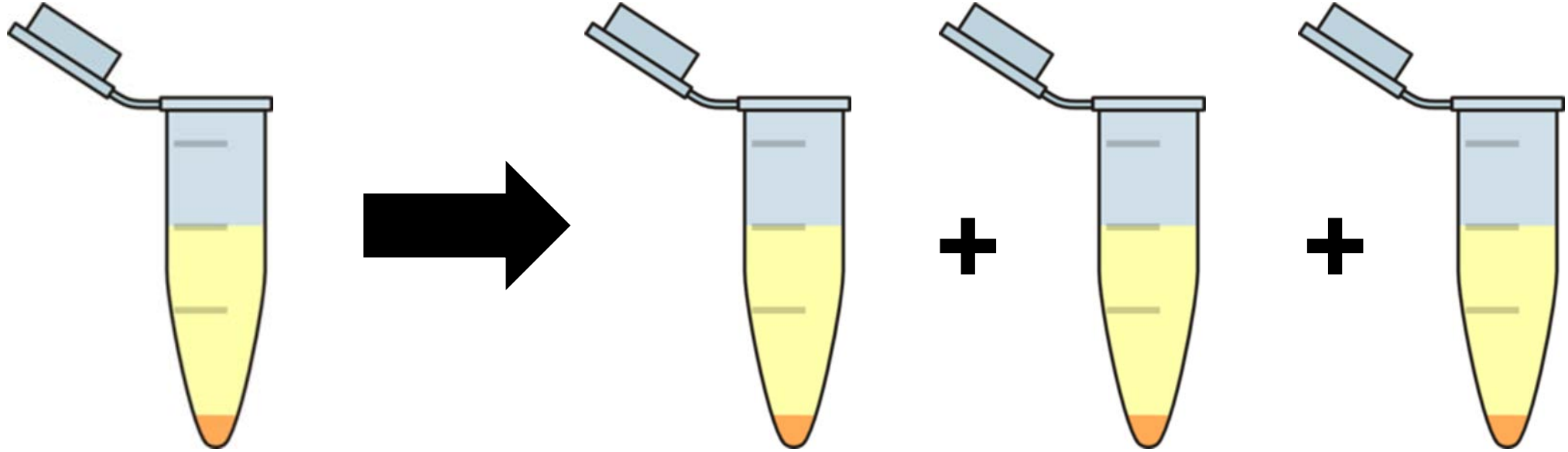
Pancreatic Cyst Fluid: Triaging



Pancreatic Cyst Fluid: Triaging



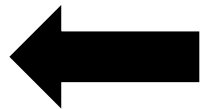
Pancreatic Cyst Fluid: Triaging



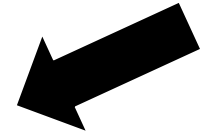
Report



10 days



Cytology



Molecular Testing (Stabilization Buffer)

CEA Amylase

PancreaSeq: Pancreatic Cyst Fluid

KRAS GNAS VHL CTNNB1 TP53 PIK3CA PTEN

- Over a 43-month period, **673 EUS-FNA** pancreatic cyst fluid specimens from 642 patients were prospectively analyzed for genetic alterations.
- Among the 673 specimens, **626 (93%)** pancreatic cysts were satisfactory for molecular analysis (**PancreaSeq**).
- In comparison, **452 (72%)** pancreatic cysts were sufficient for CEA analysis and **251 (40%)** pancreatic cysts were satisfactory for cytopathologic evaluation.
- Follow-up was available for **102 (18%)** patients.

PancreaSeq: Pancreatic Cyst Fluid

Surgical Resection Dx	Total, n = 102 (18%)
AdenoCA arising in an IPMN	13
IPMN with HGD	4
MCN with HGD	2
IPMN with LGD	39
MCN with LGD	8
Serous cystadenoma	3
Cystic PanNET	9
Acinar cell cystadenoma	1
Pseudocyst	17
Retention cyst	2
Lymphoepithelial cyst	2
Epidermoid cyst	1
Squamoid cyst	1

PancreaSeq: Pancreatic Cyst Fluid

Surgical Resection Dx	Total, n = 102 (18%)
AdenoCA arising in an IPMN	13
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Pseudocyst	17
Retention cyst	2
Lymphoepithelial cyst	2
Epidermoid cyst	1
Squamoid cyst	1

66 Mucinous Cysts:
56 IPMNs
10 MCNs

PancreaSeq: Pancreatic Cyst Fluid

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Pseudocyst	17
Retention cyst	2
Lymphoepithelial cyst	2
Epidermoid cyst	1
Squamoid cyst	1

66 Mucinous Cysts:
56 IPMNs
10 MCNs

**36 Non-Mucinous
Cysts**

PancreaSeq: Pancreatic Cyst Fluid

Surgical Resection Dx	Total, n = 102 (18%)	<i>KRAS</i> / <i>GNAS</i> wildtype	<i>KRAS</i> / <i>GNAS</i> mutant
AdenoCA arising in an IPMN	13	0 (0%)	13 (100%)
IPMN with HGD	4	0 (0%)	4 (100%)
MCN with HGD	2	0 (0%)	2 (100%)
IPMN with LGD	39	0 (0%)	39 (100%)
MCN with LGD	8	7 (87%)	1 (13%)
Serous cystadenoma	3	3 (100%)	0 (0%)
Cystic PanNET	9	9 (100%)	0 (0%)
Acinar cell cystadenoma	1	1 (100%)	0 (0%)
Pseudocyst	17	17 (100%)	0 (0%)
Retention cyst	2	2 (100%)	0 (0%)
Lymphoepithelial cyst	2	2 (100%)	0 (0%)
Epidermoid cyst	1	1 (100%)	0 (0%)
Squamoid cyst	1	1 (100%)	0 (0%)

PancreaSeq: Pancreatic Cyst Fluid

Surgical Resection Dx	Total, n = 102 (18%)	<i>KRAS</i> / <i>GNAS</i> wildtype	<i>KRAS</i> / <i>GNAS</i> mutant
AdenoCA arising in an IPMN	13	0 (0%)	13 (100%)
IPMN with HGD	4	0 (0%)	4 (100%)
MCN with HGD	2	0 (0%)	2 (100%)
IPMN with LGD	39	0 (0%)	39 (100%)
MCN with LGD	8	7 (87%)	1 (13%)
Se		3 (100%)	0 (0%)
C		9 (100%)	0 (0%)
A		1 (100%)	0 (0%)
P		17 (100%)	0 (0%)
R		2 (100%)	0 (0%)
L		2 (100%)	0 (0%)
E		1 (100%)	0 (0%)
Sq		1 (100%)	0 (0%)

IPMNs & MCNs

***KRAS* &/or *GNAS* mutations**

- Sensitivity: 89%
- Specificity: 100%

Elevated **CEA***

- Sensitivity: 57%
- Specificity: 80%

PancreaSeq: Pancreatic Cyst Fluid

Surgical Resection Dx	Total, n = 102 (18%)	<i>KRAS</i> / <i>GNAS</i> wildtype	<i>KRAS</i> / <i>GNAS</i> mutant
AdenoCA arising in an IPMN	13	0 (0%)	13 (100%)
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MCN with HGD	2	0 (0%)	2 (100%)
IPMN with LGD	39	0 (0%)	39 (100%)
MCN with LGD	8	7 (87%)	1 (13%)

IPMNs & MCNs

***KRAS* &/or *GNAS* mutations**

- Sensitivity: 89%
- Specificity: 100%

Elevated CEA

- Sensitivity: 57%
- Specificity: 80%

IPMNs

***KRAS* &/or *GNAS* mutations**

Sensitivity: 100%

MCNs

***KRAS* mutations**

Sensitivity: 30%

PancreaSeq: Pancreatic Cyst Fluid

Surgical Resection Dx	Total, n = 102 (18%)	<i>TP53/PIK3CA/PTEN</i> wildtype	<i>TP53/PIK3CA/PTEN</i> mutant
AdenoCA arising in an IPMN	13	0 (0%)	13 (100%)
IPMN with HGD	4	2 (50%)	2 (50%)
MCN with HGD	2	2 (100%)	0 (0%)
IPMN with LGD	39	36 (92%)	3 (8%)
MCN with LGD	8	8 (100%)	0 (0%)

Alterations in ***TP53/PIK3CA/PTEN*** were preoperatively detected in all 13 (100%) adenocarcinomas.

Epidermoid cyst	1	1 (100%)	0 (0%)
Squamoid cyst	1	1 (100%)	0 (0%)

PancreaSeq: Pancreatic Cyst Fluid

Surgical Resection Dx	Total, n = 102 (18%)	<i>TP53/PIK3CA/PTEN</i> wildtype	<i>TP53/PIK3CA/PTEN</i> mutant
AdenoCA arising in an IPMN	13	0 (0%)	13 (100%)
IPMN with HGD	4	2 (50%)	2 (50%)
MCN with HGD	2	2 (100%)	0 (0%)
IPMN with LGD	39	36 (92%)	3 (8%)
MCN with LGD	8	8 (100%)	0 (0%)

Alterations in ***TP53/PIK3CA/PTEN*** were detected in 2 (50%) IPMNs with HGD

Epidermoid cyst	1	1 (100%)	0 (0%)
Squamoid cyst	1	1 (100%)	0 (0%)

PancreaSeq: Pancreatic Cyst Fluid

Surgical Resection Dx	Total, n = 102 (18%)	<i>TP53/PIK3CA/PTEN</i> wildtype	<i>TP53/PIK3CA/PTEN</i> mutant
AdenoCA arising in an IPMN	13	0 (0%)	13 (100%)
IPMN with HGD	4	2 (50%)	2 (50%)
MCN with HGD	2	2 (100%)	0 (0%)
IPMN with LGD	39	36 (92%)	3 (8%)
MCN with LGD	8	8 (100%)	0 (0%)

Alterations in ***TP53/PIK3CA/PTEN*** were detected in 2 (50%) IPMNs with HGD and 3 (8%) IPMNs with LGD (*PIK3CA* mutations).

Epidermoid cyst	1	1 (100%)	0 (0%)
Squamoid cyst	1	1 (100%)	0 (0%)

PancreaSeq: Pancreatic Cyst Fluid

Surgical Resection Dx	Total, n = 102 (18%)	TP53/PIK3CA/PTEN wildtype	TP53/PIK3CA/PTEN mutant
AdenoCA arising in an IPMN	13	0 (0%)	13 (100%)
IPMN with HGD	4	2 (50%)	2 (50%)
MCN with HGD	2	2 (100%)	0 (0%)
IPMN with LGD	39	36 (92%)	3 (8%)
MCN with LGD		8 (100%)	0 (0%)
IPMN with F1		3 (100%)	0 (0%)
IPMN with F2		9 (100%)	0 (0%)
IPMN with F3		1 (100%)	0 (0%)
IPMN with F4		17 (100%)	0 (0%)
IPMN with F5		2 (100%)	0 (0%)
IPMN with F6		2 (100%)	0 (0%)
IPMN with F7		1 (100%)	0 (0%)
Sq		1 (100%)	0 (0%)

Advanced Neoplasia

KRAS and/or **GNAS** and **TP53**, **PIK3CA**, and/or **PTEN**

- Sensitivity: 79%
- Specificity: 96%

Cytology

- Sensitivity: 32%
- Specificity: 98%

PancreaSeq: Pan

Surgical Resection Dx	Total, n = 102 (18%)
AdenoCA arising in an IPMN	13
IPMN with HGD	4
MCN with HGD	2
IPMN with LGD	39

Advanced Neoplasia

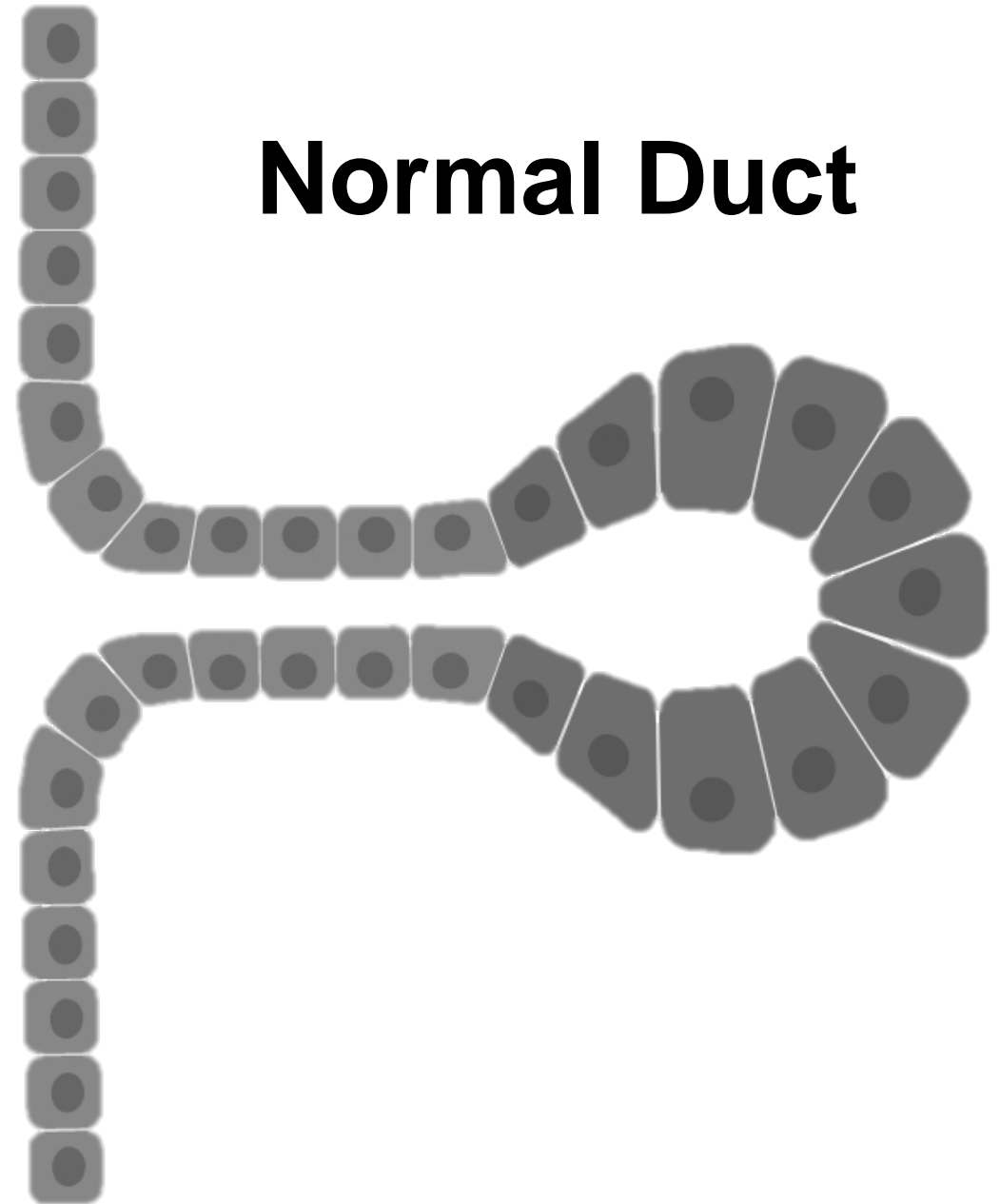
KRAS and/or **GNAS** and
TP53, **PIK3CA**, and/or **PTEN**

- Sensitivity: 79%
- Specificity: 96%

Cytology

- Sensitivity: 32%
- Specificity: 98%

Normal Duct



PancreaSeq: Pan

Surgical Resection Dx	Total, n = 102 (18%)
AdenoCA arising in an IPMN	13
IPMN with HGD	4
MCN with HGD	2
IPMN with LGD	39

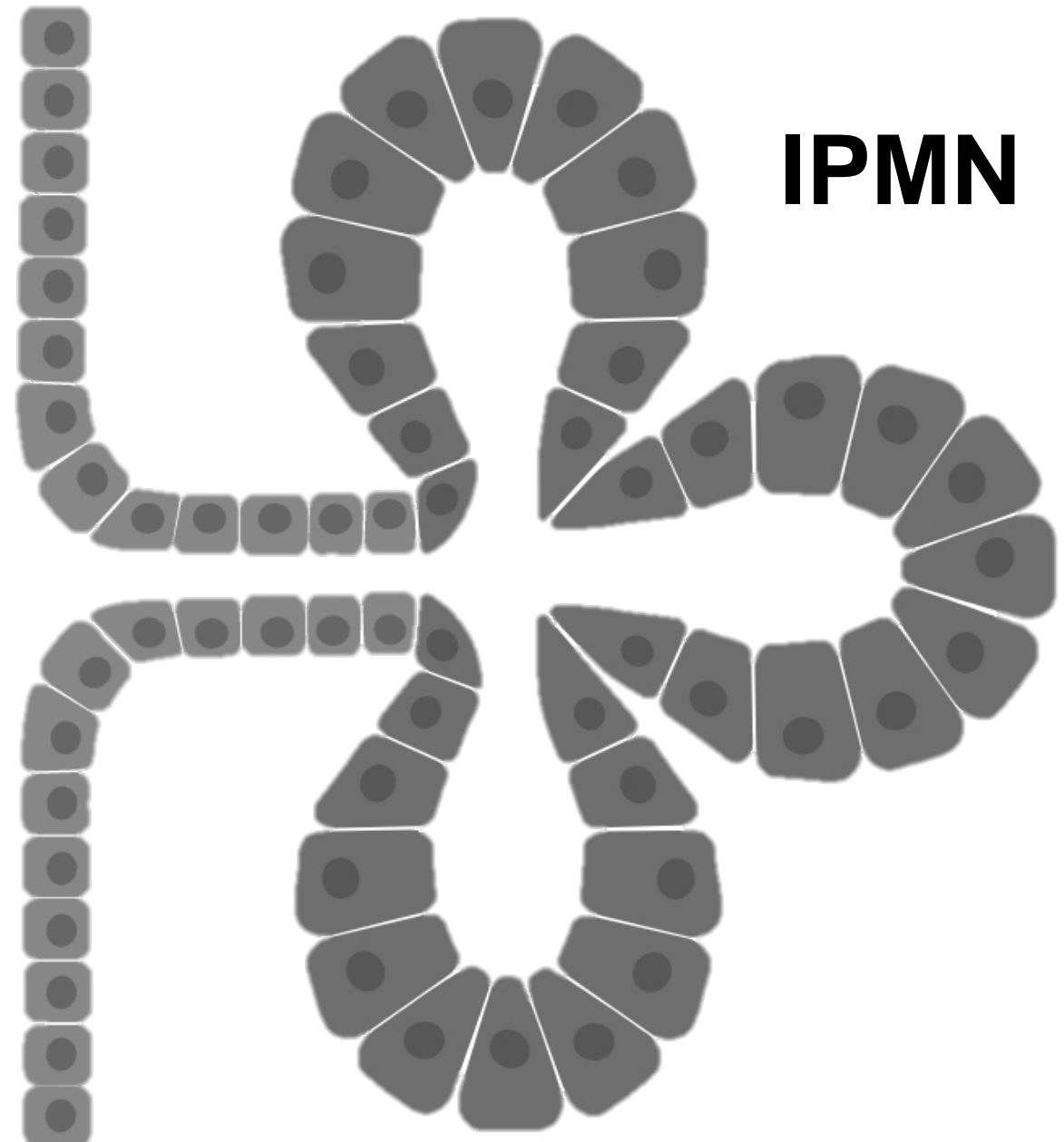
Advanced Neoplasia

KRAS and/or **GNAS** and
TP53, **PIK3CA**, and/or **PTEN**

- Sensitivity: 79%
- Specificity: 96%

Cytology

- Sensitivity: 32%
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PancreaSeq: Pan

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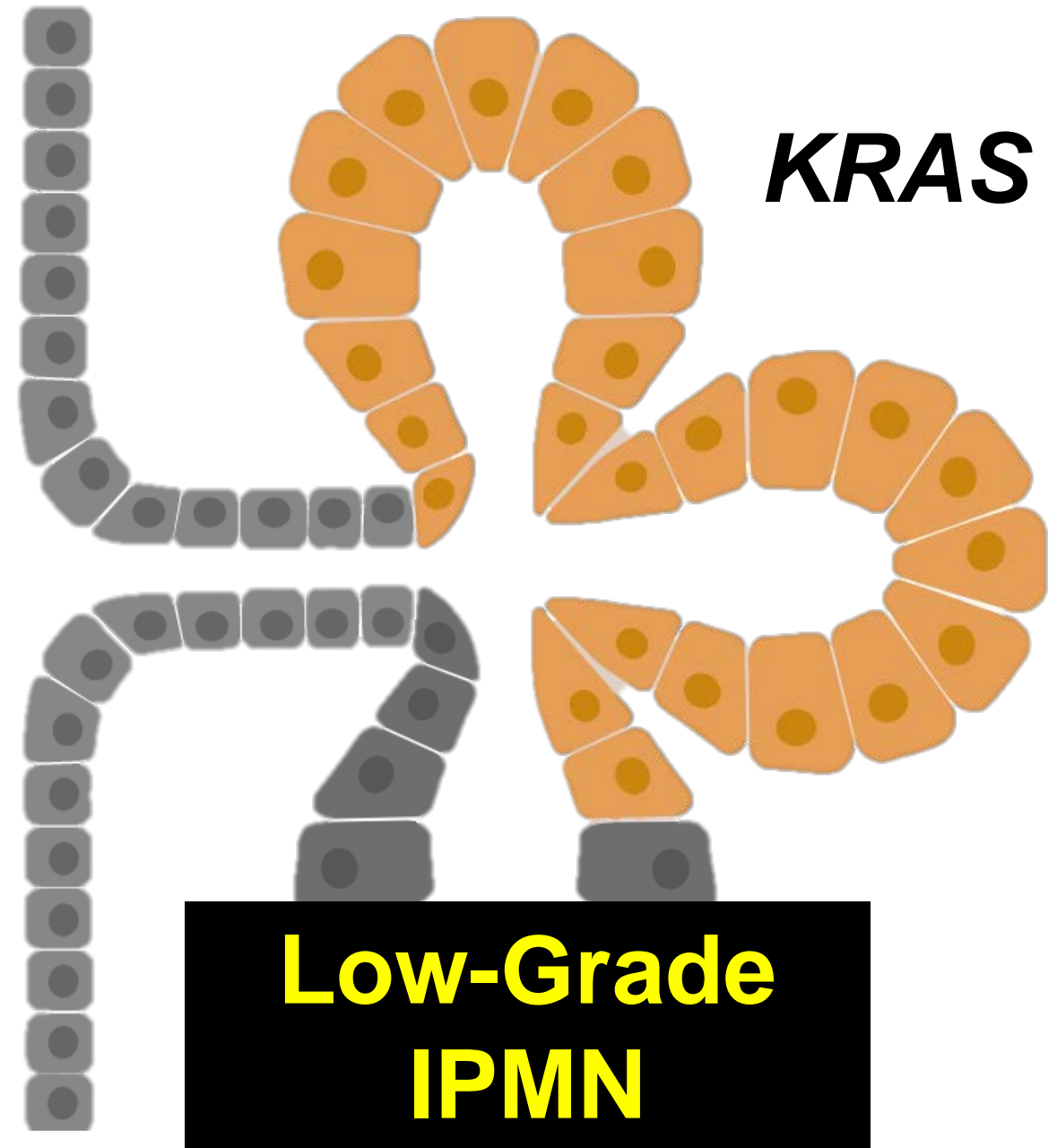
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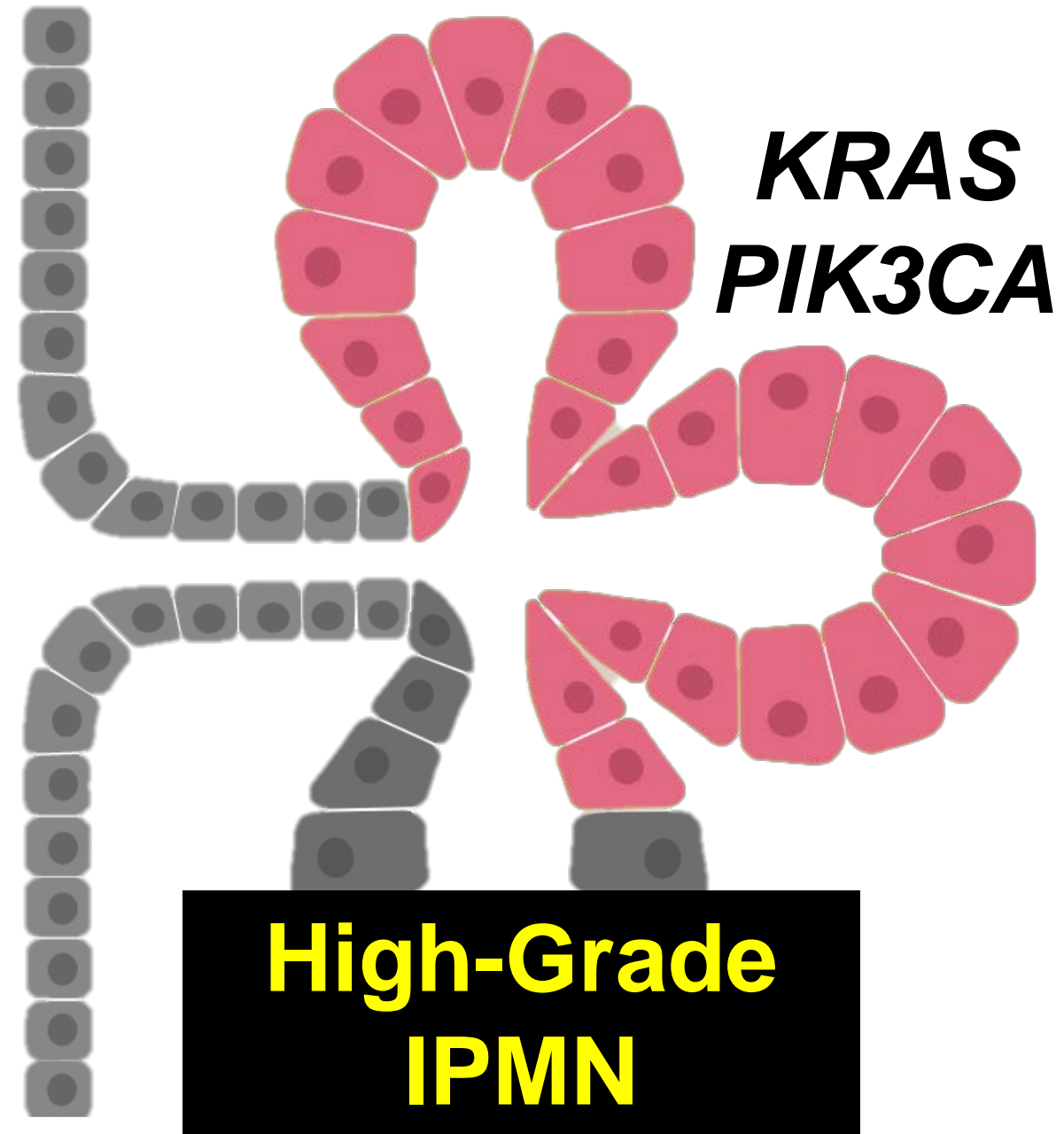
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- Sensitivity: 79%
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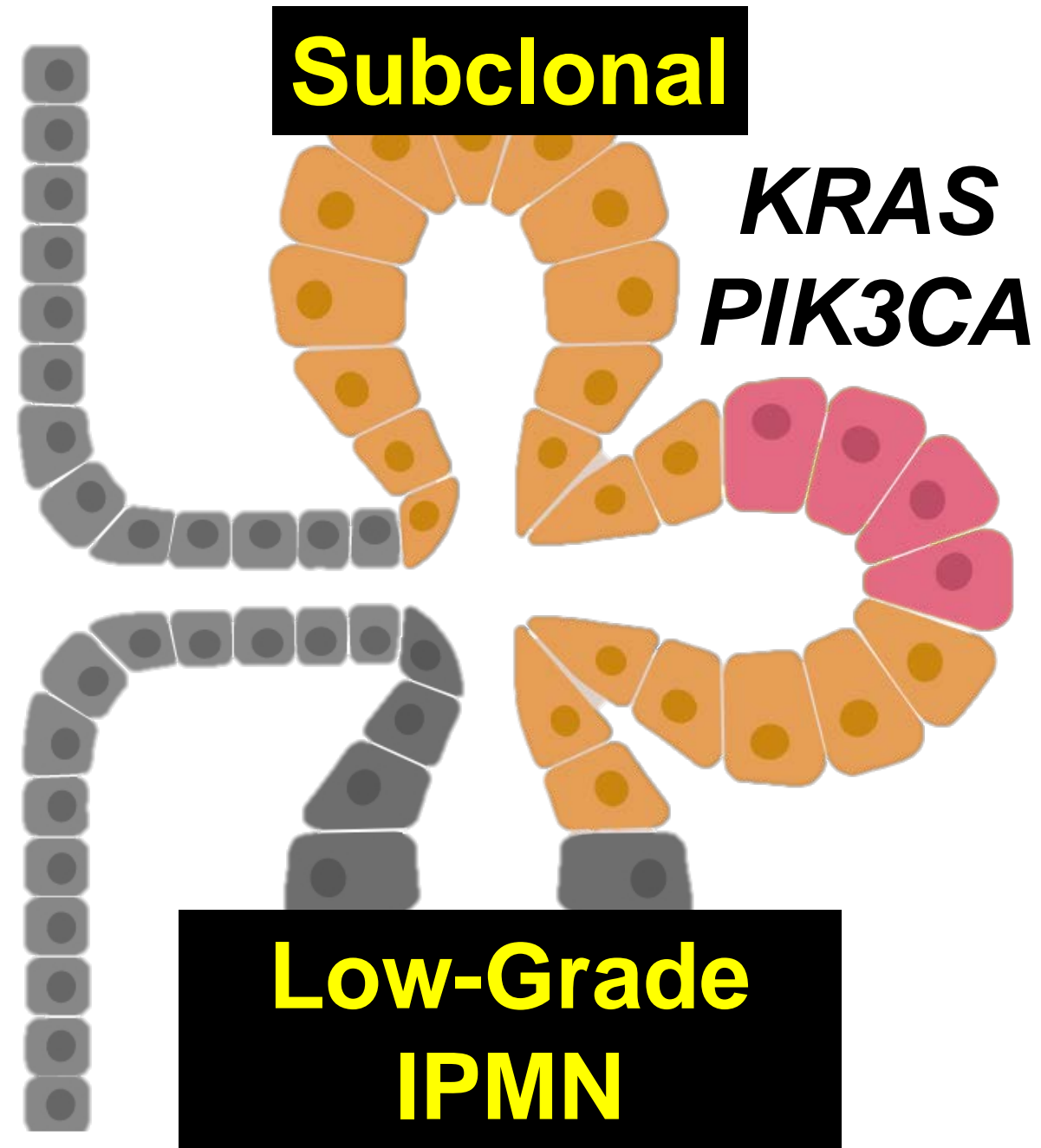
Advanced Neoplasia

KRAS and/or **GNAS** and
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- Sensitivity: 79%
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PancreaSeq: Pan

Surgical Resection Dx	Total, n = 102 (18%)
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IPMN with HGD	4
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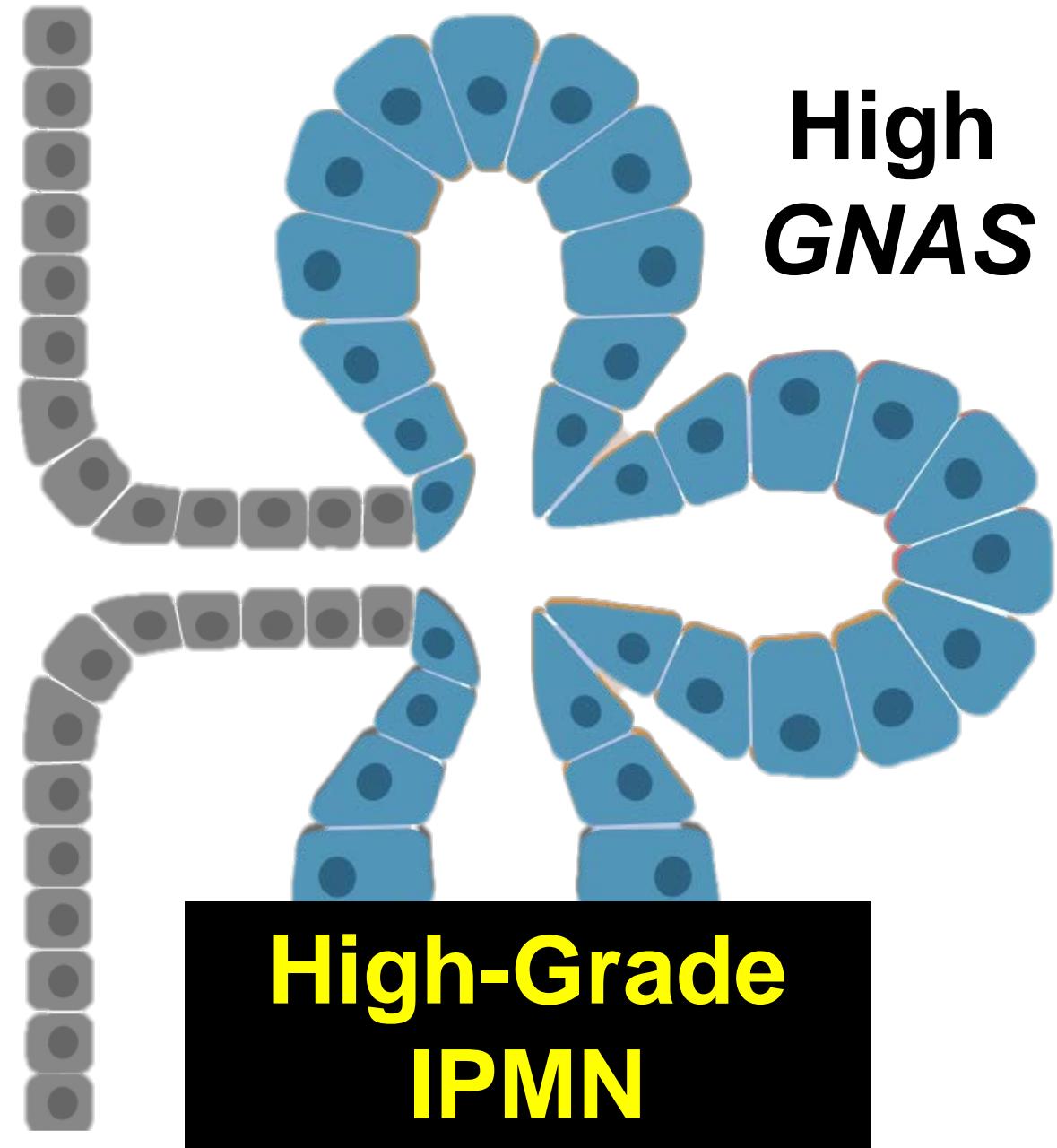
Advanced Neoplasia

KRAS and/or **GNAS** and
TP53, **PIK3CA**, and/or **PTEN**

- Sensitivity: 79%
- Specificity: 96%

Cytology

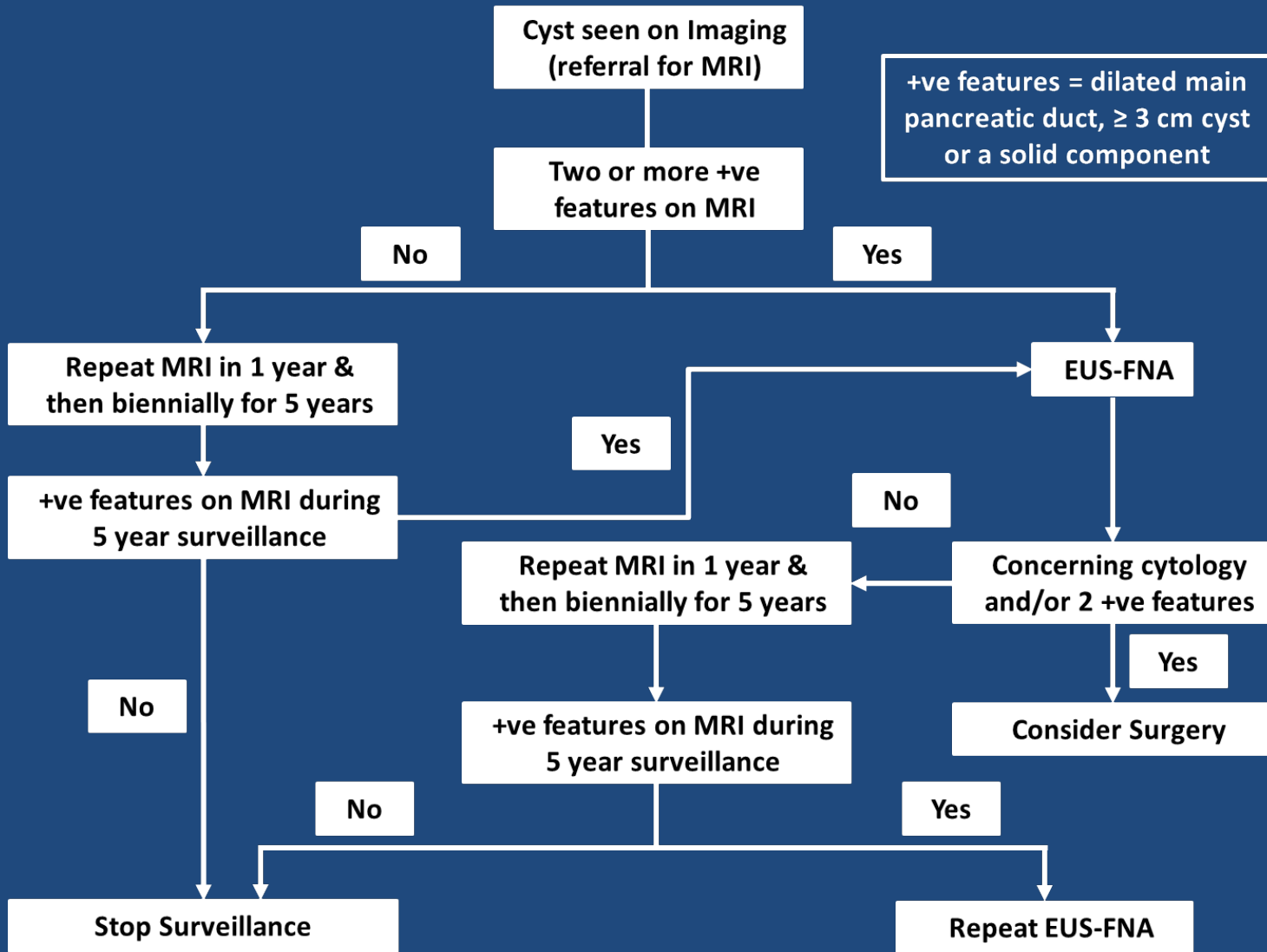
- Sensitivity: 32%
- Specificity: 98%



PancreaSeq: Pancreatic Cyst Fluid

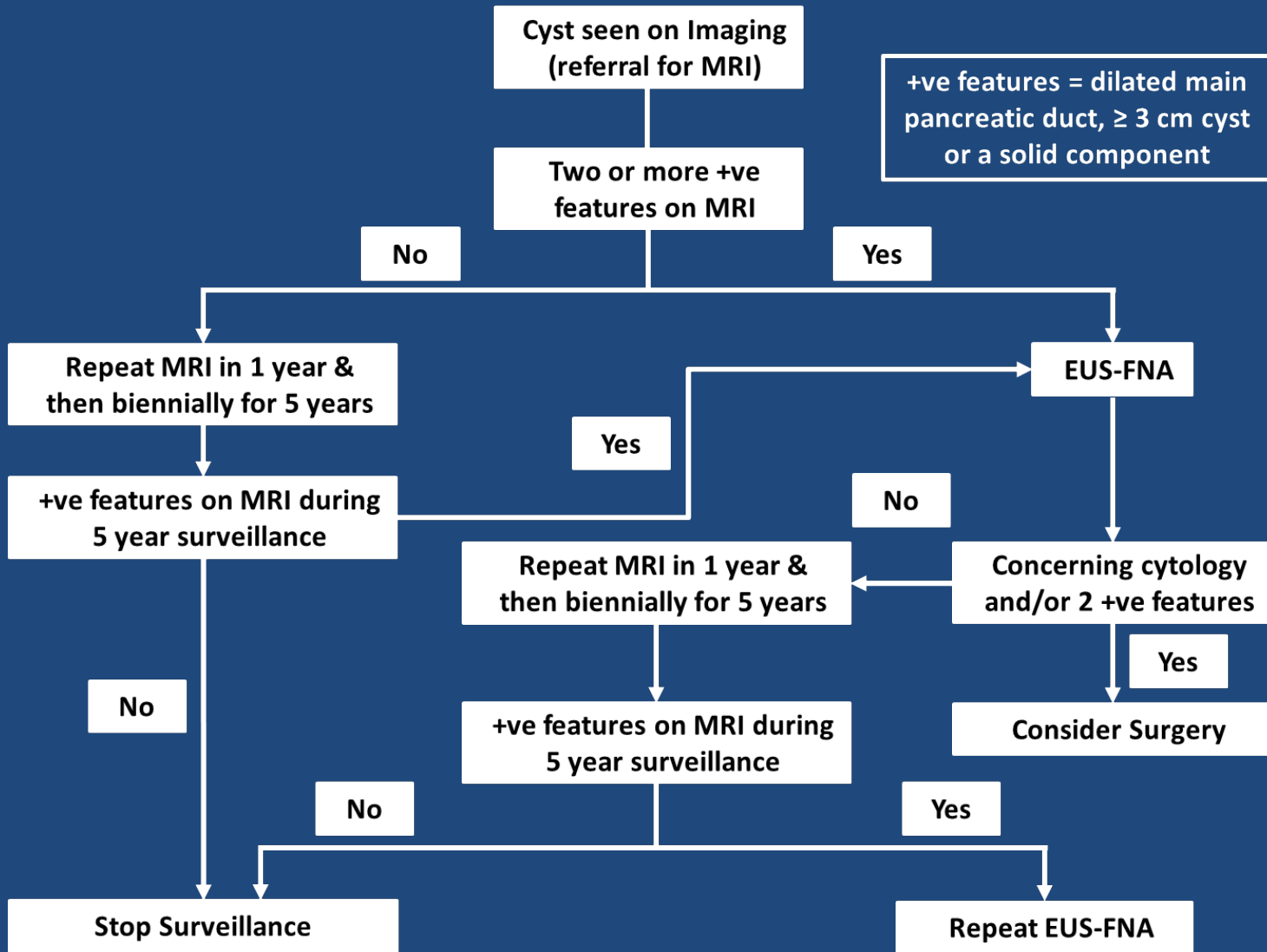
Surgical Resection Dx	Total, n =	TP53/PIK3CA/PTEN	TP53/PIK3CA/PTEN
<u>Advanced Neoplasia</u> Main duct dilation <ul style="list-style-type: none">• Sensitivity: 42%• Specificity: 74%		<u>Advanced Neoplasia</u> Mural nodule <ul style="list-style-type: none">• Sensitivity: 32%• Specificity: 94%	
<u>Advanced Neoplasia</u> KRAS and/or GNAS and TP53, PIK3CA, and/or PTEN <ul style="list-style-type: none">• Sensitivity: 79%• Specificity: 96% Cytology <ul style="list-style-type: none">• Sensitivity: 32%• Specificity: 98%		<u>Advanced Neoplasia</u> GNAS AF >55% or KRAS and/or GNAS AF = TP53, PIK3CA, and/or PTEN AF <ul style="list-style-type: none">• Sensitivity: 89%• Specificity: 100%	

PancreaSeq versus AGA Guidelines



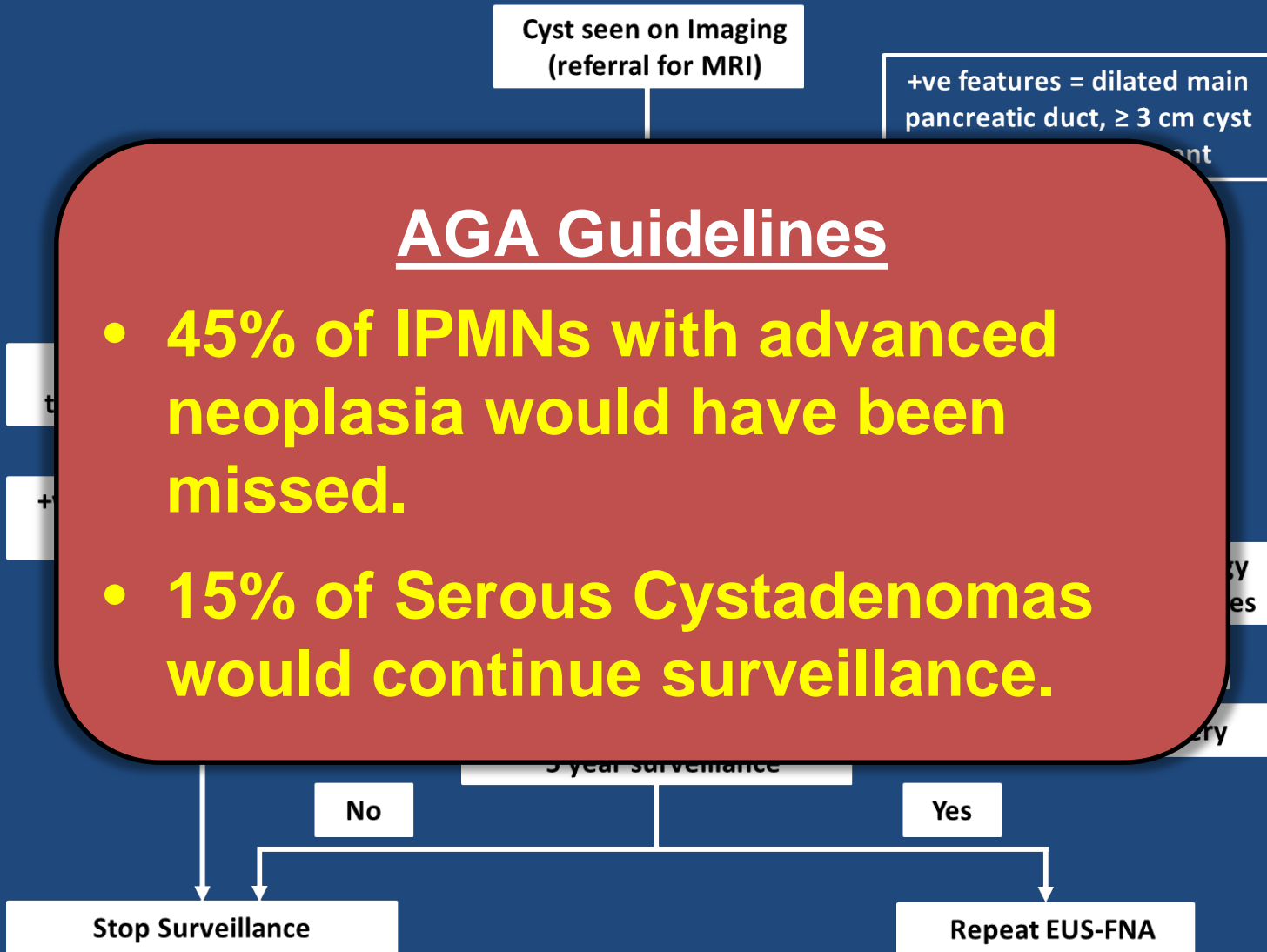
- **Study cohort:**
 - **225 patients with corresponding diagnostic pathology for 41 patients.**
 - **EUS-FNA with PancreaSeq testing.**

PancreaSeq versus AGA Guidelines



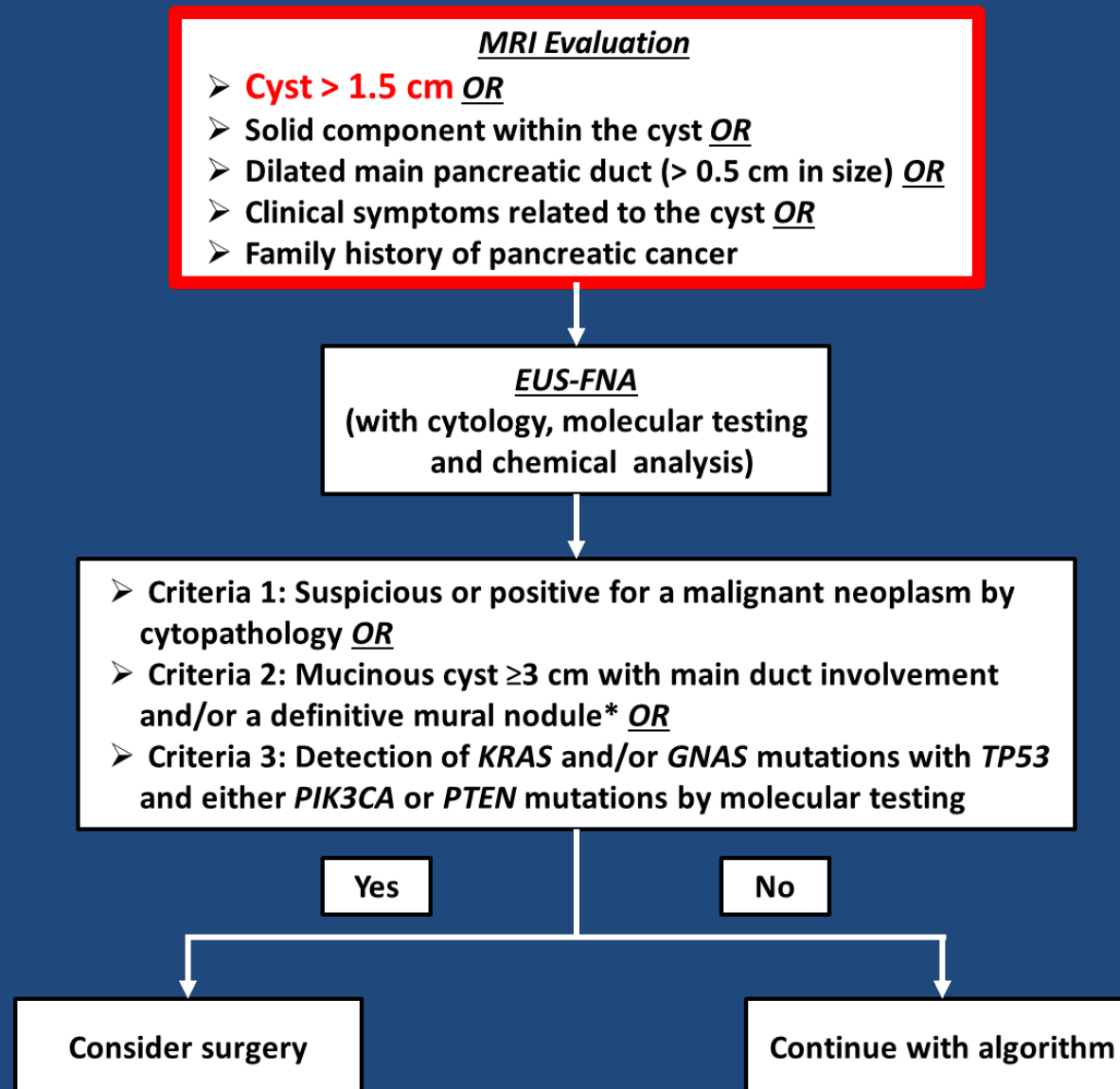
- Study cohort:
 - 225 patients with corresponding diagnostic pathology for 41 patients.
 - EUS-FNA with PancreaSeq testing.
- **AGA Guidelines:**
 - **Sensitivity: 62%**
 - **Specificity: 79%**
- Modified Fukuoka with PancreaSeq (UPMC):
 - Sensitivity: 100%
 - Specificity: 90%

PancreaSeq versus AGA Guidelines

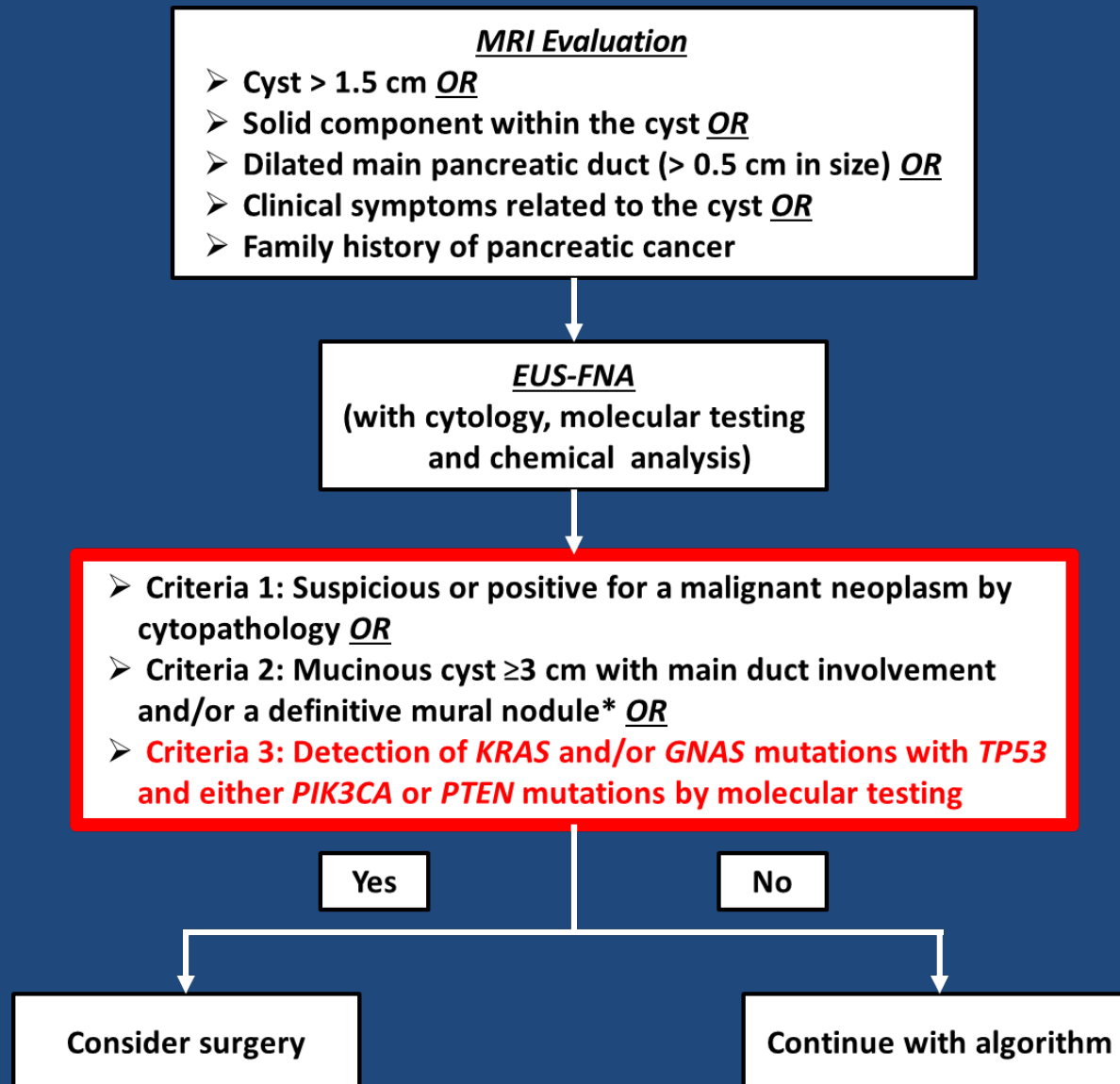


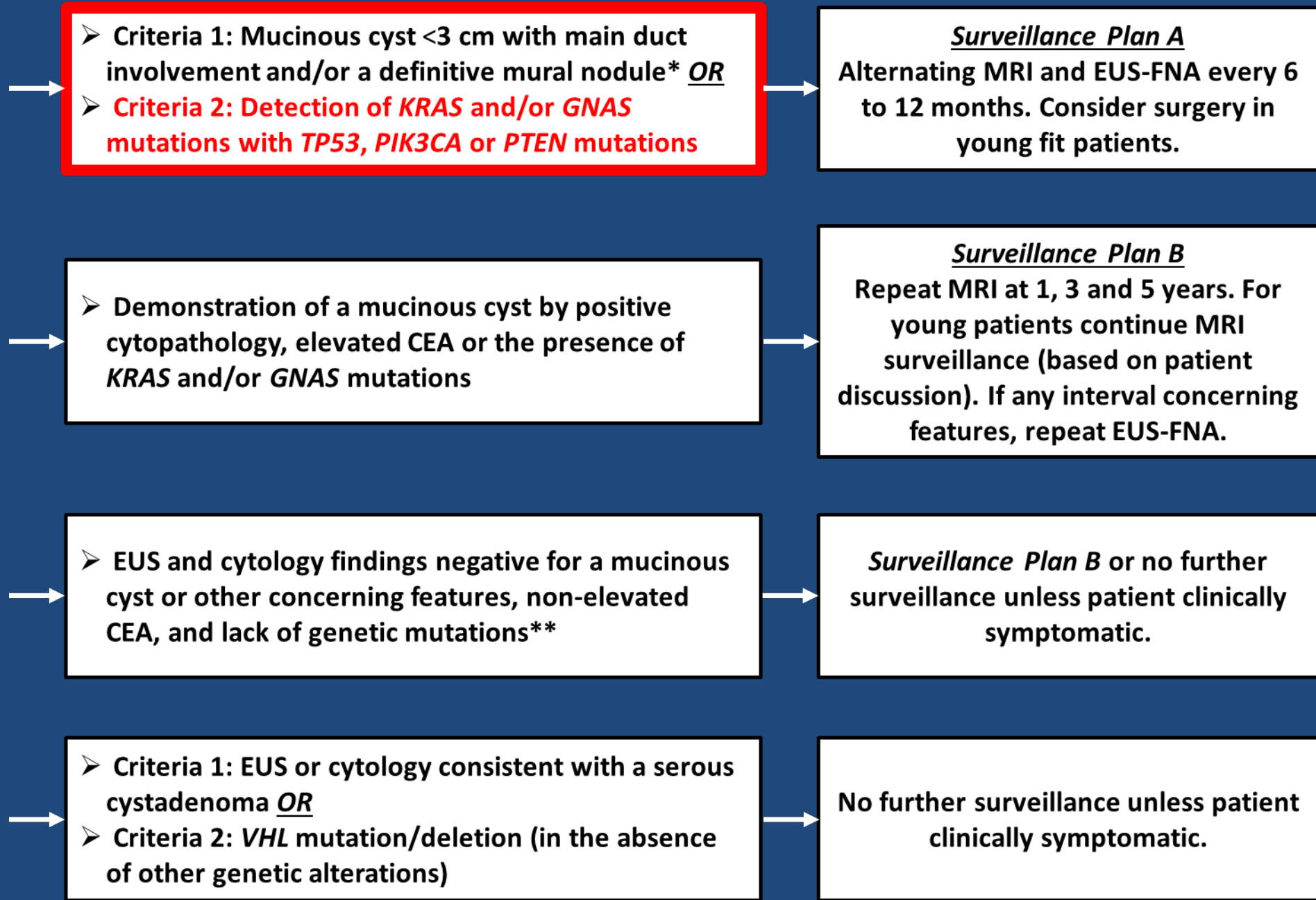
- Study cohort:
 - 225 patients with corresponding diagnostic pathology for 41 patients.
 - EUS-FNA with PancreaSeq testing.
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 - Sensitivity: 62%
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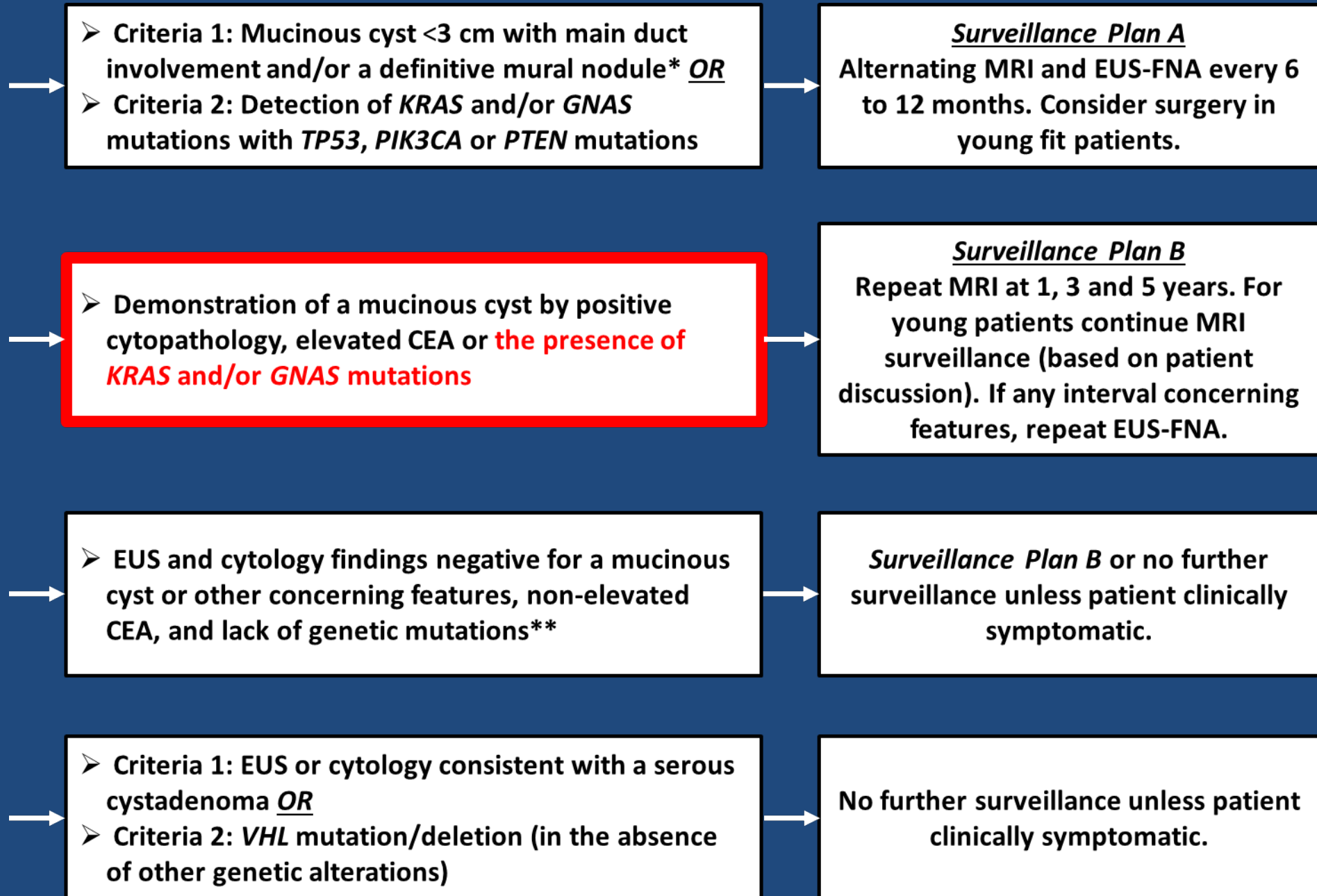
UPMC Pancreatic Cyst Workflow

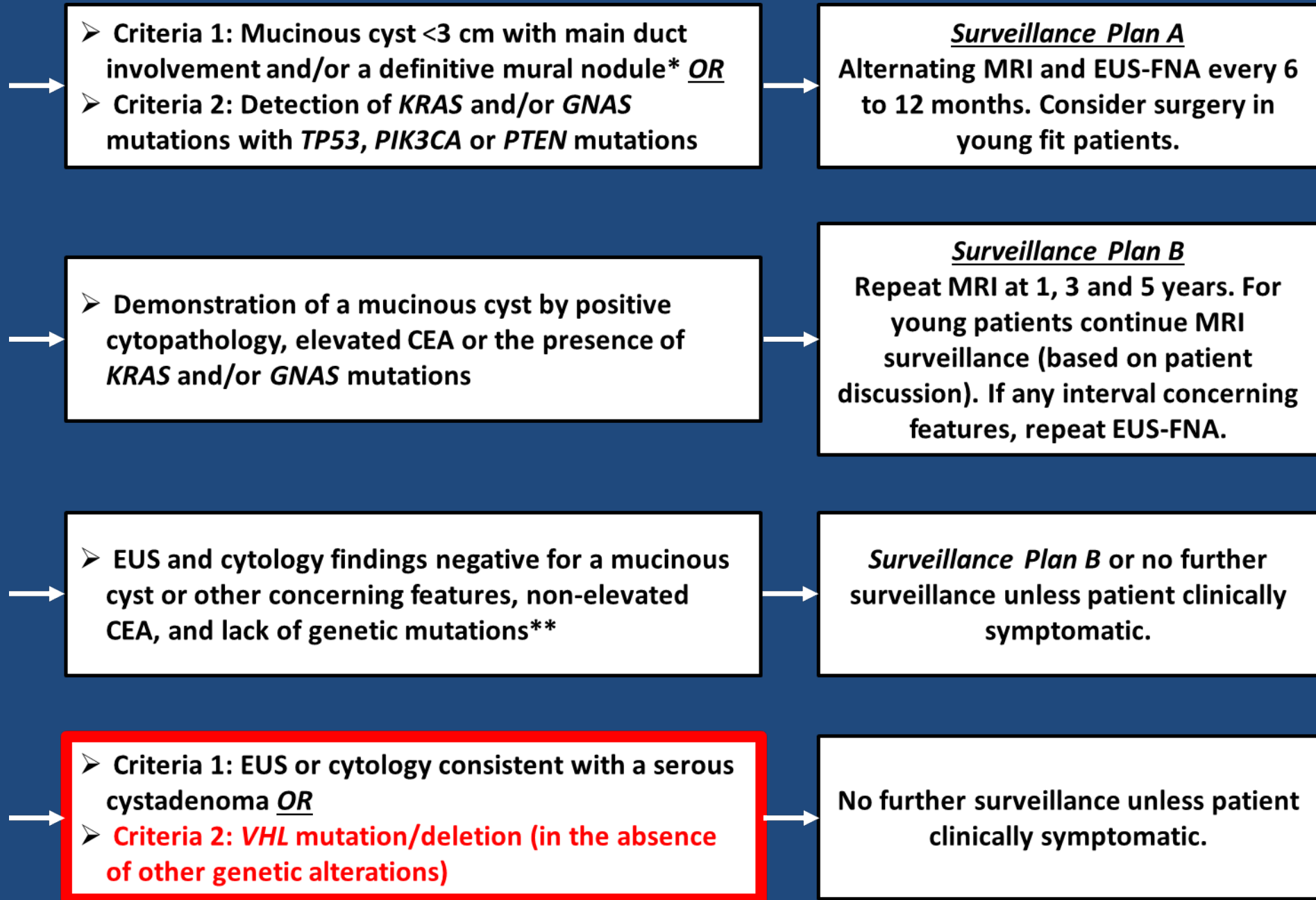


UPMC Pancreatic Cyst Workflow





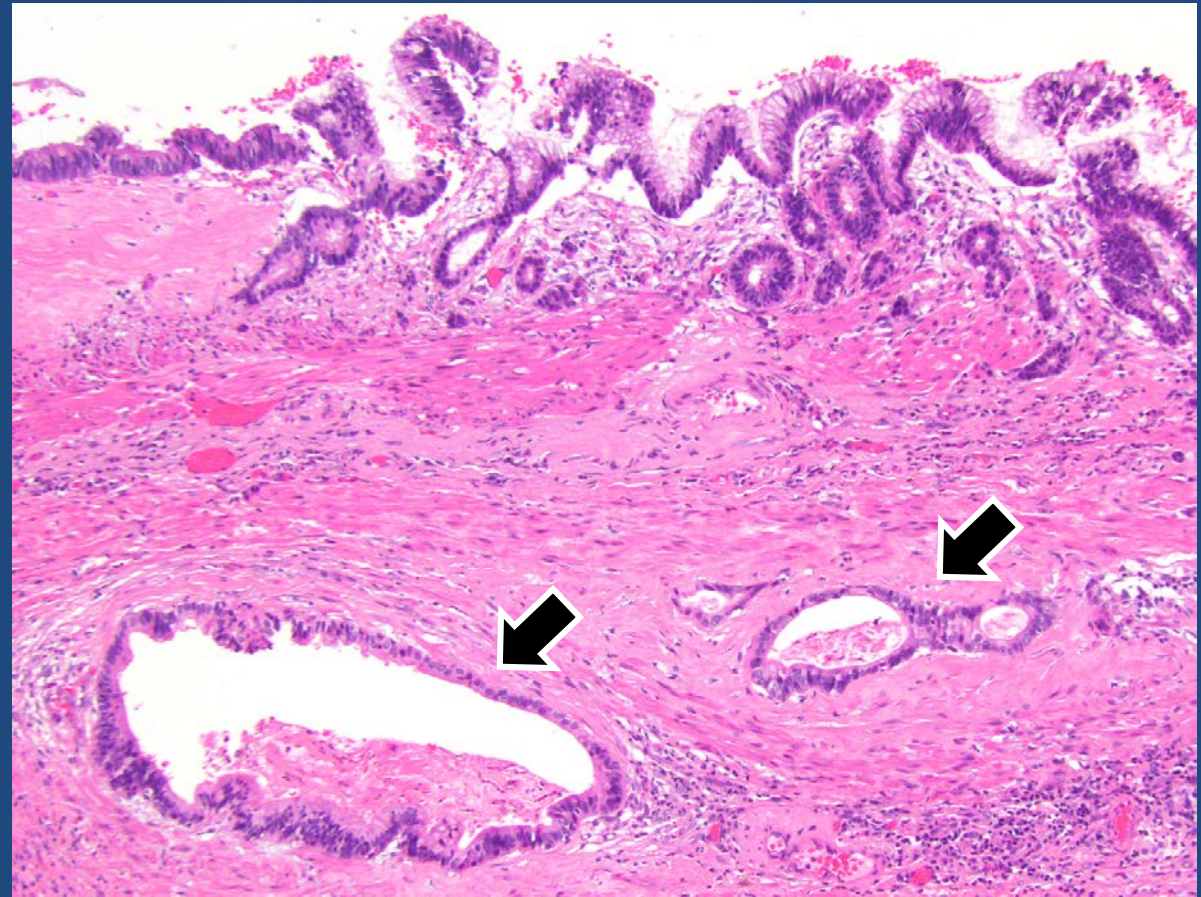




Clinical Case

Surgical Resection:

- Invasive moderately-differentiated adenocarcinoma, 0.9 cm (black arrow), arising in association with an IPMN.
- Pathologic stage (AJCC 8th edition): pT1a N0.



Summary

- **The application of molecular testing to pancreatic cyst fluid analysis can be a useful adjunct in the evaluation of pancreatic cysts.**
- **Cellular content and fluid volume of aspirated cysts are commonly suboptimal for routine ancillary studies (e.g. CEA and cytopathology).**
- **DNA from lysed or exfoliated pancreatic cyst epithelial lining shed into the cyst can be analyzed for genetic abnormalities.**

Summary

- Mutations in ***KRAS* & *GNAS*** are highly specific for branch duct IPMNs, but not MCNs.
- The presence of **high-risk alterations** (e.g. *TP53*, *PIK3CA*, *PTEN*, high *GNAS*, etc.) can predict advanced neoplasia.
- Alterations in ***VHL*** can aid in classification of **serous cystadenomas** and decrease the number of patients undergoing surveillance.

Take Away Message

	IPMN	MCN	SCA	SPN	Pseudocyst
Gender	M>F	F>>M	F>M	F>>M	M>F
Location	Head>Tail	Tail>>Head	Head>Tail	Tail>Head	Head=Tail
Viscosity	Increased	Increased	Low	Low	Low
CEA	>192 ng/mL	>192 ng/mL	<0.5 ng/mL	<192 ng/mL	<192 ng/mL
Amylase	High	Low	Low	Low	High
Cytology	Mucinous	Mucinous	Scant, Bland PAS+	Papillary & Vascular	Pigmented Histiocytes
Genetics	<i>KRAS,GNAS</i>	<i>KRAS</i>	<i>VHL</i>	<i>CTNNB1</i>	Absent

Challenges Ahead



- Continuing evolution of pancreatic cyst molecular profiling.
- **Additional biomarkers:** DNA, RNA, protein, carbohydrates and others.
- Refinements and optimization with other preoperative clinical, imaging and pathology metadata.
- Benefits versus costs: **insurance reimbursement.**
- **Multiple assays available:** academic versus industry.