

# Beyond the surface: Exploring Head and neck cancer through molecular pathology

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# CONTENTS

Clinical features  
Histopathology  
Molecular pathology

- Adenoid cystic carcinoma
- Mucoepidermoid carcinoma
- Salivary duct carcinoma
- Acinic cell carcinoma
- Secretory carcinoma
- Basal cell adenocarcinoma
- Epithelial-myoepithelial carcinoma
- Carcinoma ex pleomorphic adenoma
- Myoepithelial carcinoma
- Hyalinizing clear cell carcinoma

## 5. Salivary gland tumours

Salivary gland tumours: Introduction

*Non-neoplastic epithelial lesions*

Nodular oncocytic hyperplasia

Lymphoepithelial sialadenitis

*Benign epithelial tumours*

Pleomorphic adenoma

Basal cell adenoma

Warthin tumour

Oncocytoma

Salivary gland myoepithelioma

Canalicular adenoma

Cystadenoma of the salivary glands

Ductal papillomas

Sialadenoma papilliferum

Lymphadenoma

Sebaceous adenoma

Intercalated duct adenoma and hyperplasia

Striated duct adenoma

Sclerosing polycystic adenoma

Keratocystoma

*Malignant epithelial tumours*

Mucoepidermoid carcinoma

Adenoid cystic carcinoma

Acinic cell carcinoma

Secretory carcinoma

Microsecretory adenocarcinoma

Polymorphous adenocarcinoma

Hyalinizing clear cell carcinoma

Basal cell adenocarcinoma

Intraductal carcinoma

Salivary duct carcinoma

Myoepithelial carcinoma

Epithelial-myoepithelial carcinoma

Mucinous adenocarcinoma

Sclerosing microcystic adenocarcinoma

Carcinoma ex pleomorphic adenoma

Carcinosarcoma of the salivary glands

Sebaceous adenocarcinoma

Lymphoepithelial carcinoma

Squamous cell carcinoma

Sialoblastoma

Salivary carcinoma NOS and emerging entities

*Mesenchymal tumours specific to the salivary glands*

Sialolipoma

# Adenoid cystic carcinoma (AdCC)

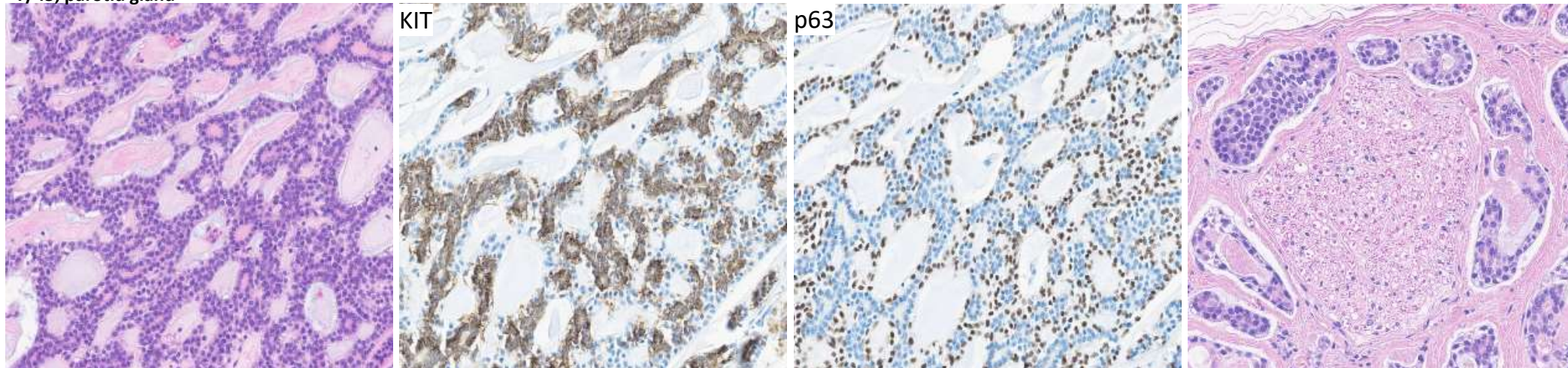
- **Clinical features**

- 25% of all primary salivary carcinomas; median age of 60 years
- 60% in major salivary gland (parotid, submandibular); 30% in minor salivary gland (palate)
- Neural symptoms are common
- Prolonged clinical course with frequent local recurrences, late onset of metastases
  - Median OS after distant metastasis : 36 months

- **Histopathology**

- Two cell component
  - Ductal cells: eosinophilic cytoplasm and uniform round nuclei (CK7+, KIT+)
  - Myoepithelial cells: clear cytoplasm and hyperchromatic angular nuclei (p63+, p40+, calponin+,  $\alpha$ -SMA+)

F/45; parotid gland

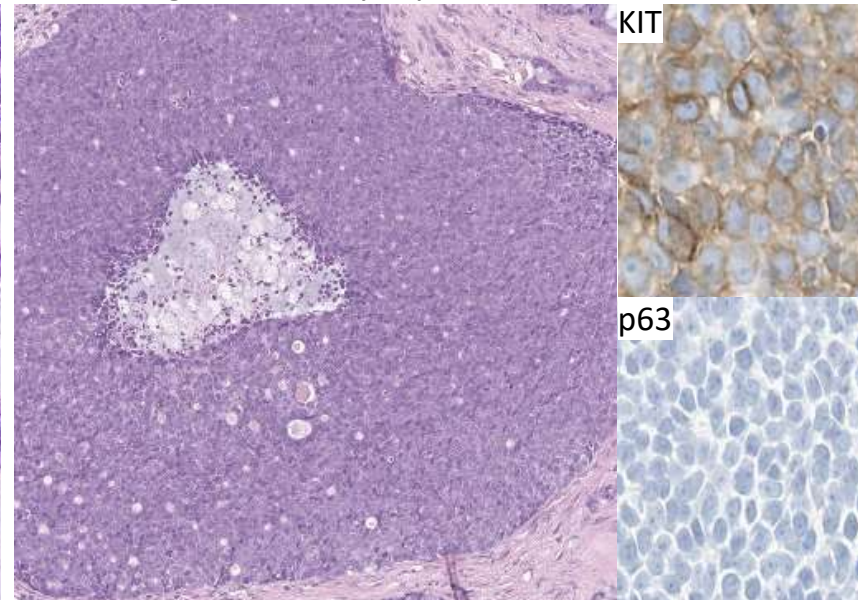
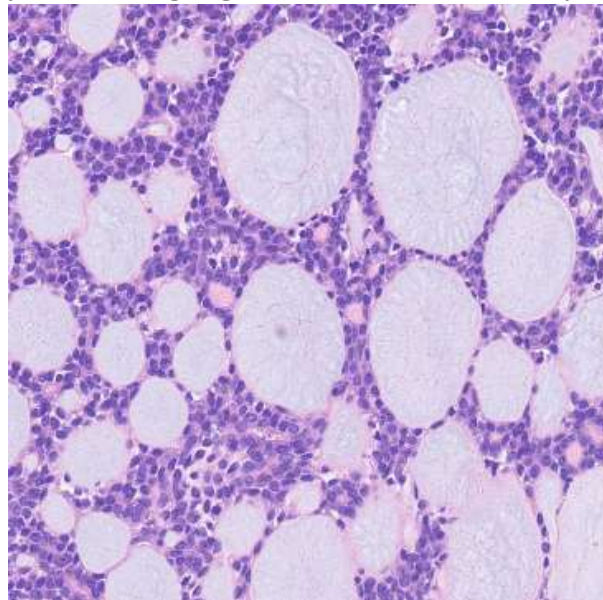
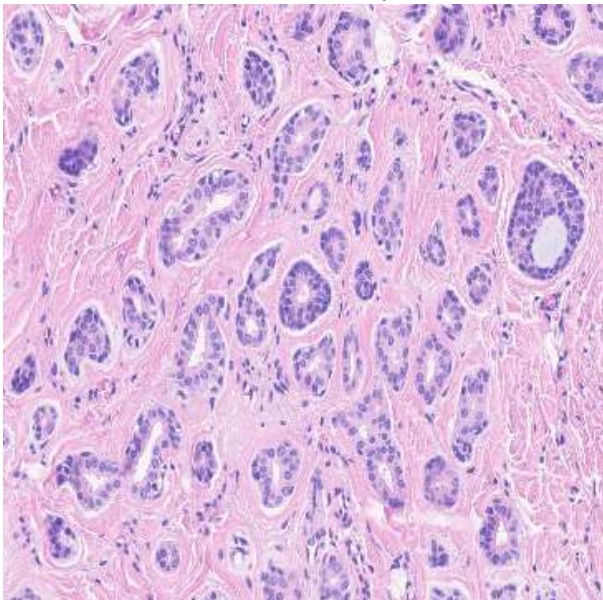




# Adenoid cystic carcinoma (AdCC)

- **Histopathology**

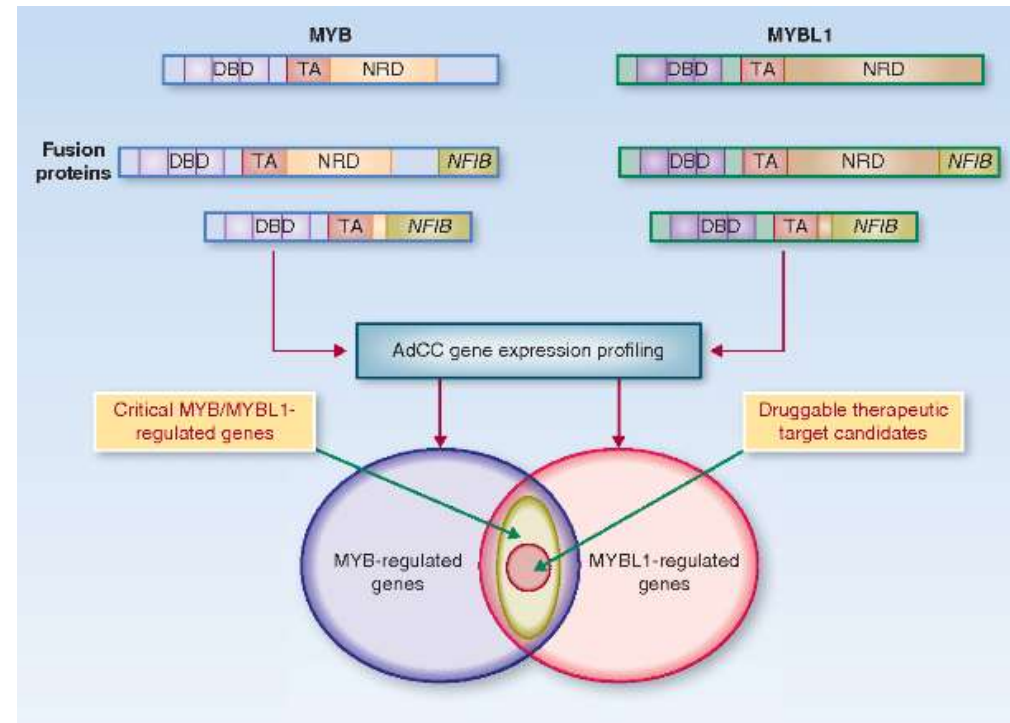
- Growth patterns
  - Tubular: well-formed ducts and tubules lined with luminal ductal and abluminal myoepithelial cells
  - Cribriform: most frequent; nests of tumor cells with microcystic spaces, filled with hyaline/basophilic material
  - Solid: tumour sheets composed of basaloid cells lacking tubular or cribriform
    - >30% solid component → more aggressive clinical course
- High-grade transformation
  - Pleomorphic, mitotically active high-grade carcinoma component → negative for myoepithelial markers



# Adenoid cystic carcinoma (AdCC)

- **Molecular pathology**

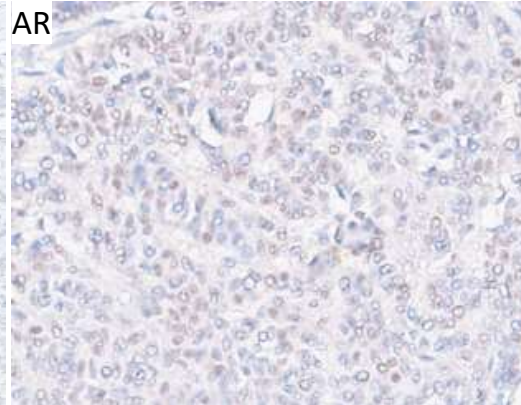
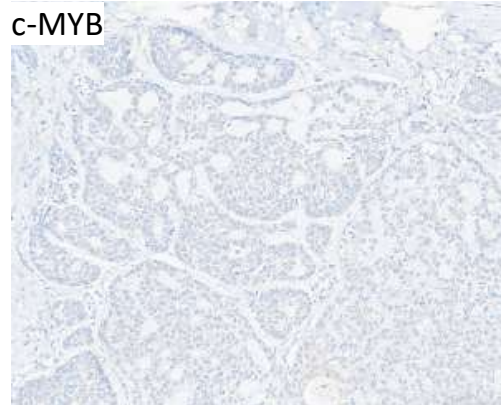
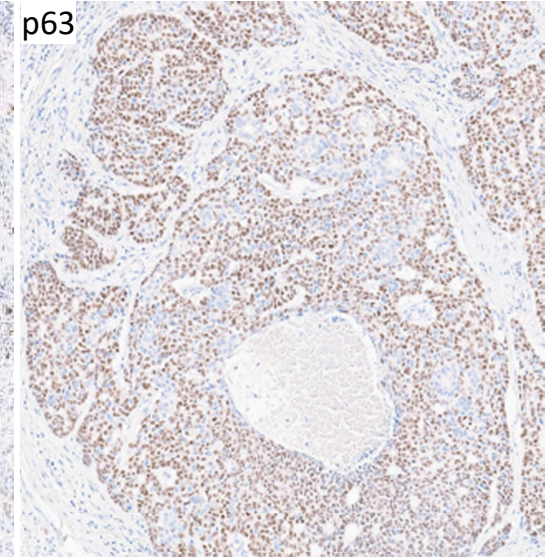
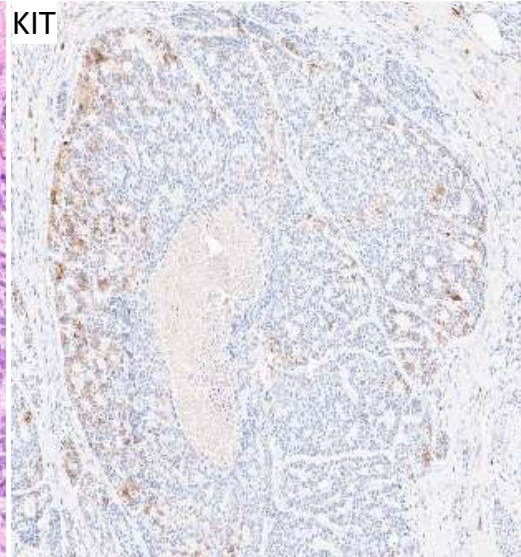
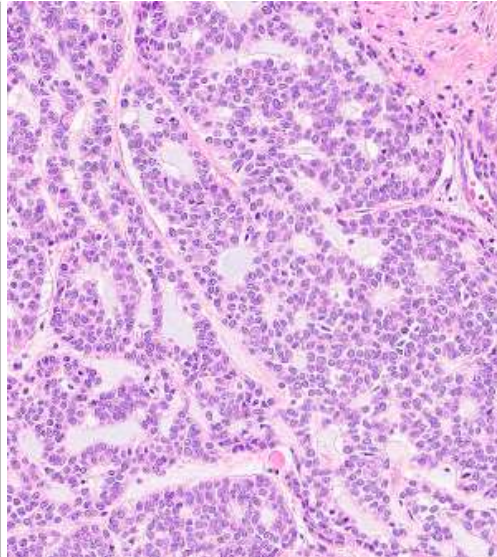
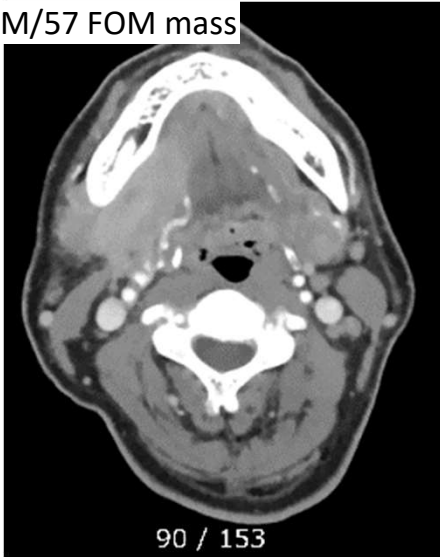
- *MYB::NFIB* fusion (29-86%); *MYBL1::NFIB* fusion (9-14%)  
→ each mutually exclusive
  - *MYB*: transcription factor regulating proliferation and differentiation of hematopoietic, colonic, and neural progenitor cells
  - *NFIB*: transcription factor nuclear factor I/B protein  
→ key regulator of hematopoietic and epithelial cells
  - *MYB::NFIB* → overexpression of truncated MYB protein d/t loss of negative regulatory elements
- Rarer fusions including *MYB::PDCD1LG2*, *MYB::EFR3A*, *MYBL1::RAD51B*, *MYBL1::YTHDF3*, *NFIB::AIG1* fusions





# Adenoid cystic carcinoma (AdCC)

M/57 FOM mass



# Adenoid cystic carcinoma (AdCC)

## Cancer panel - Report

FIRST-Cancer Panel / Molecular Genetic Testing

### Result of Actionable Gene Mutation Analysis

-SNV/INDEL

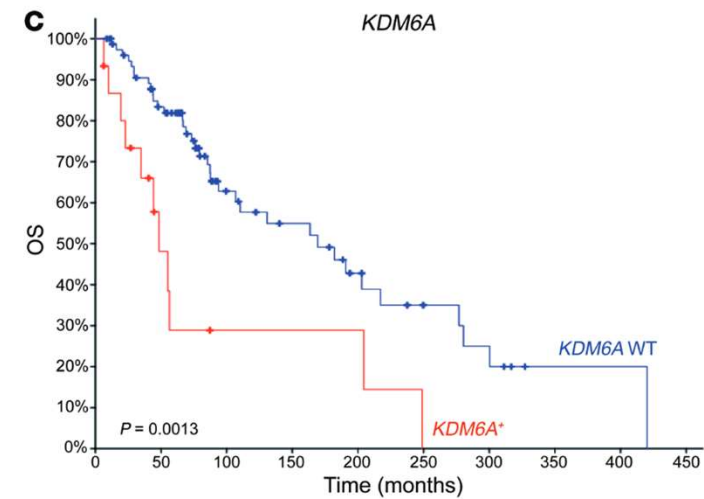
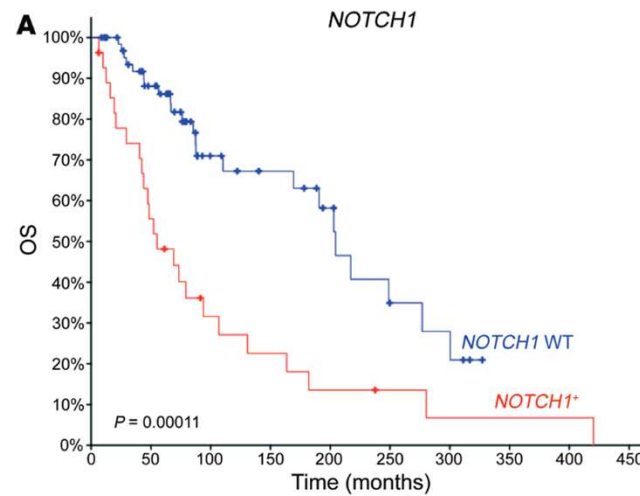
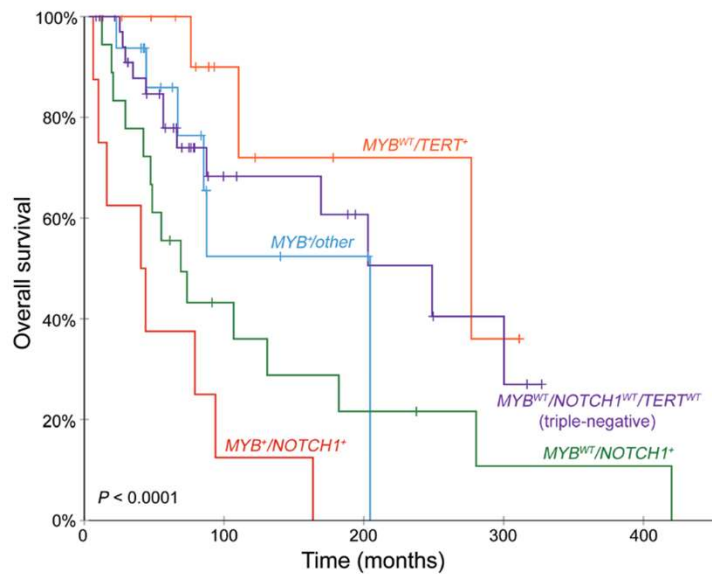
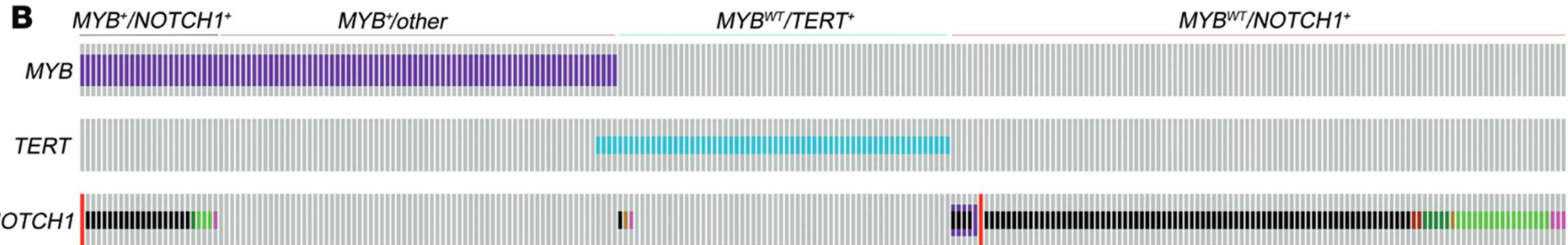
Gene	Chr	Position	Refseq	Exon	AA	CDS	% VAF	Alt	Total	Tier
Not detected										

- Fusion

Fusion Gene	Cytoband1	Cytoband2	DNA.Evidence (Span/Split)	RNA.Evidence (Span/Split)	Transcript1	Transcript2	Last Observed	First Observed	Tier
MYB->NFIB	6q23.3	9p23-p22.3	0/0	0/107	NM_005375	NM_005596	EXON_14	EXON_9	D

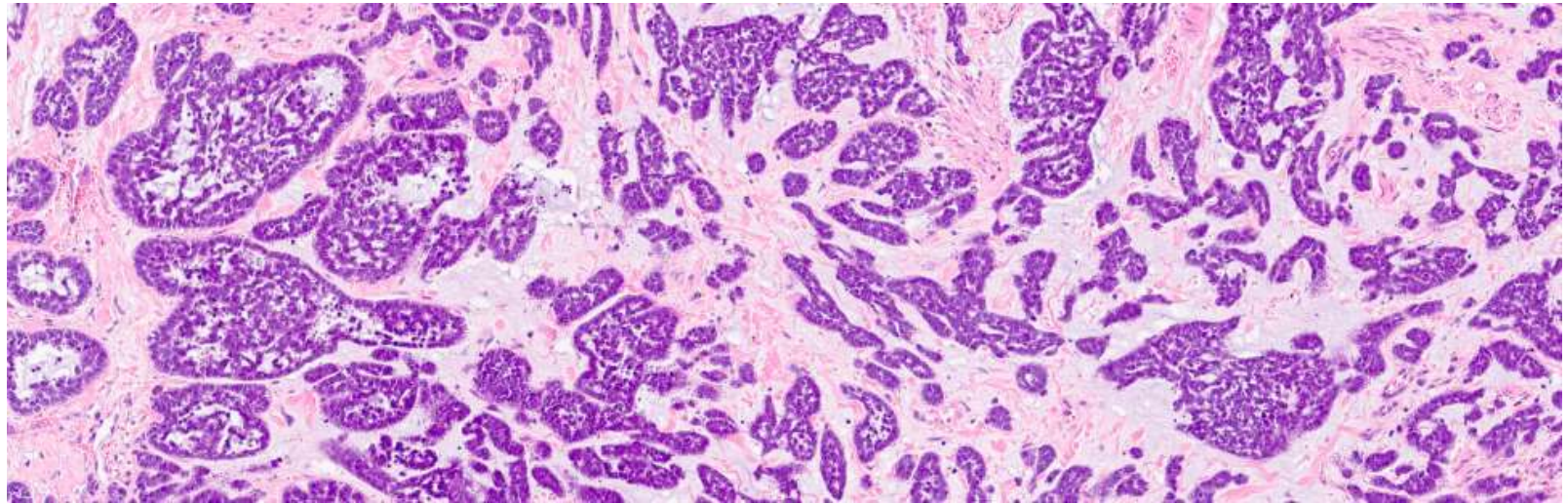


# Adenoid cystic carcinoma (AdCC)



# Adenoid cystic carcinoma (AdCC)

F/39; lacrimal gland



- SNV/INDEL

Gene	Chr	Position	Refseq	Exon	AA	CDS	% VAF	Alt	Total	Tier
NOTCH1	chr9	139390793	NM_017617.3	34	p.Ser2467fs	c.7363_7397dupACIATTCGTGC CCCAGGAGAGCCCCGCCCT GCCCAC	24.50	172	702	D
NOTCH1	chr9	139401189	NM_017617.3	23	p.Glu1294fs	c.3857_3879dupAGCGCGTCA ATGACTTCCACTGC	11.54	239	2071	D
CREBBP	chr16	3817868	NM_004380.2	16	p.Glu1035*	c.3103G>T	33.59	174	518	D

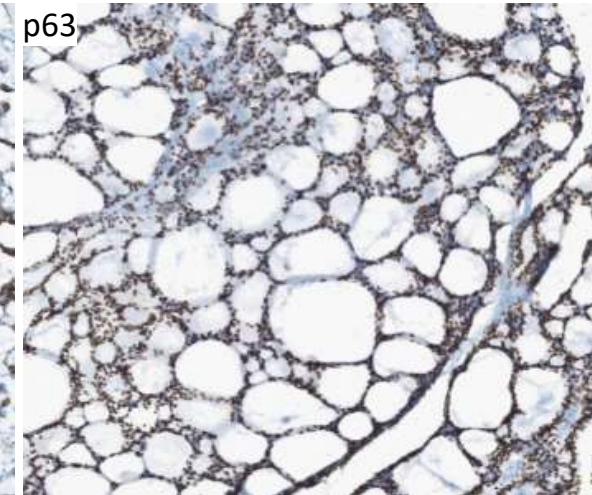
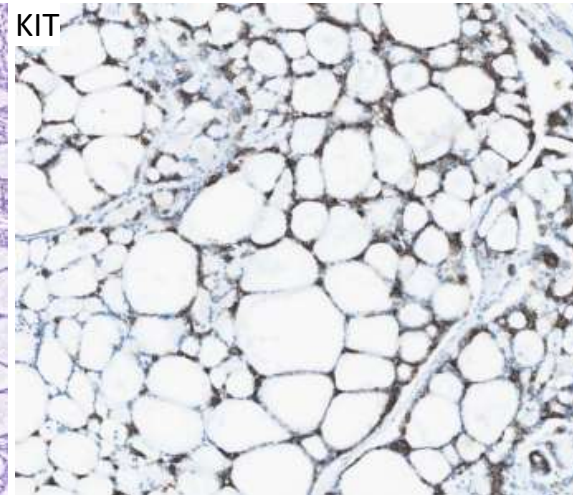
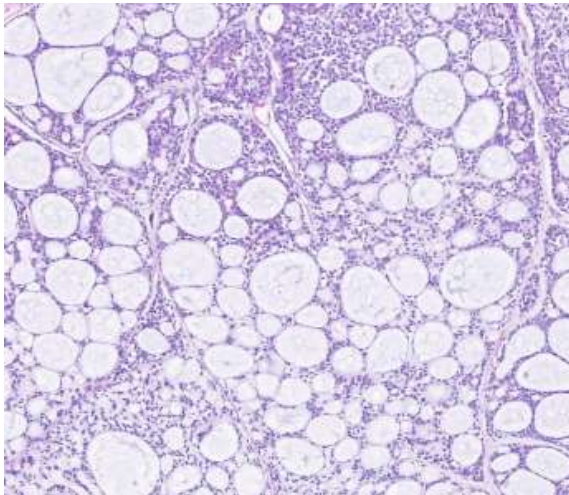
- Fusion

Fusion Gene	Cytoband1	Cytoband2	DNA.Evidence (Span/Split)	RNA.Evidence (Span/Split)	Transcript1	Transcript2	Last Observed	First Observed	Tier
MYB->NFIB	6q23.3	9p23-p22.3	0/0	9/777	NM_005375	NM_001369461	EXON 14	EXON 10	D



# Adenoid cystic carcinoma (AdCC)

M/65  
FOM mass



-SNV/INDEL

Gene	Chr	Position	Refseq	Exon	AA	CDS	%VAF	Alt	Total	Tier
Not detected										

- Fusion

Fusion Gene	Cytoband1	Cytoband2	DNA.Evidence (Span/Split)	RNA.Evidence (Span/Split)	Transcript1	Transcript2	Last Observed	First Observed	Tier
Not detected									

-CNV

Gene	Chr	Start	End	CNV	AvgL2R	DelR	NeutralR	AmpR	CN	Tier
PDGFRA	chr4	55124713	55161605	Amplification	1.957	0.0	0.0	1.0	11	D
KIT	chr4	55523974	55604878	Amplification	2.112	0.0	0.0	1.0	11	D
KDR	chr4	55945938	55991666	Amplification	2.081	0.0	0.0	1.0	10	D



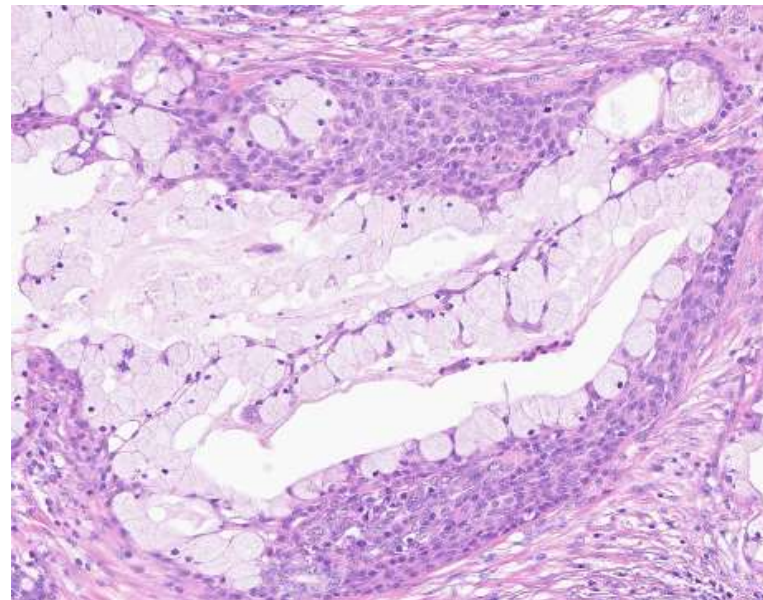
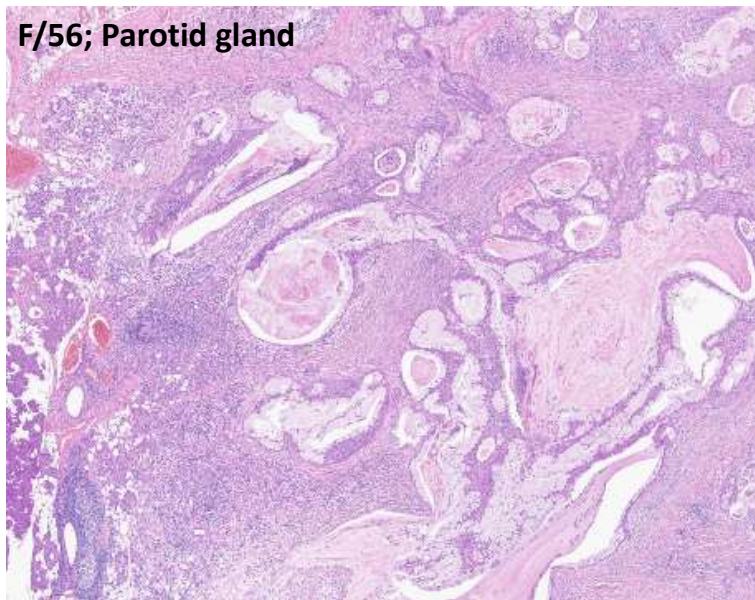
# Mucoepidermoid carcinoma (MEC)

- **Clinical features**

- Childhood to elderly (median age of 45) / M:F 1:1.1-1.5
- About 50% in major salivary gland (parotid > submandibular gland > sublingual gland)

- **Histopathology**

- Malignant salivary gland neoplasm characterized by mucous, intermediate and epidermoid (squamoid) tumor cells forming cystic and solid growth patterns
  - Mucous cells: lining cystic spaces; intracytoplasmic mucicarmine or PAS staining with diastase resistance
  - Overt keratinization is rare



# Mucoepidermoid carcinoma (MEC)

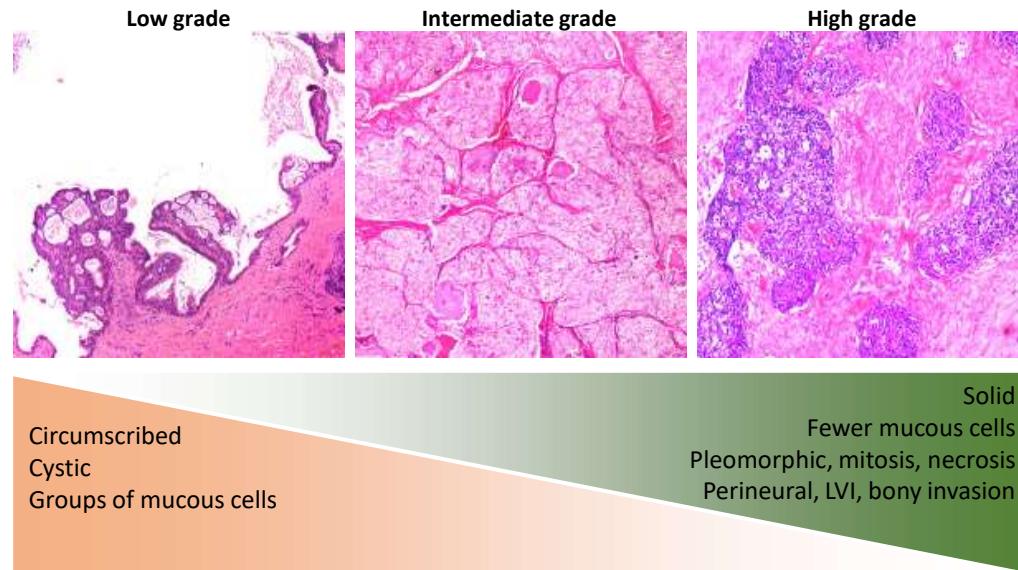
## Histopathology – grading

**Table 1.** AFIP, Brandwein, and modified Healy grading systems (modified from Seethala)<sup>5</sup>

AFIP		Brandwein		Modified Healy
Intracystic component <20%	+2	Intracystic component <25%	+2	Low grade: • Macrocysts and microcysts • Differentiated mucin with 1/1 mucin/epidermoid with minimal to moderate intermediate cells • Absent to minimal pleomorphism and rare mitoses • Broad front or circumscribed borders • Pools of extravasated mucin
Neural invasion	+2	Neural invasion	+2	
		Tumour invades in small nests	+2	
Mitosis (≥4/10HPFs)	+3	Mitosis (≥4/10HPFs)	+3	
Necrosis	+3	Necrosis	+3	Intermediate grade: • Solid nests, few microcysts, no macrocysts • Mild to moderate pleomorphism, few mitoses, prominent nuclei and nucleoli • Invasive, well defined and uncircumscribed with fibrosis separating tumour nests • Peripheral chronic inflammation
		Vascular invasion	+3	
Anaplasia	+4	Nuclear atypia	+3	
		Bone invasion	+3	
Grade	Score	Grade	Score	High grade: • Predominately solid with no macrocysts • Tumour cells range from poorly differentiated to epidermoid and intermediate cells to ductal cells adenocarcinoma • Marked pleomorphism and easily found mitoses • Perineural and vascular invasions, soft tissue invasion, and desmoplasia • Less common peripheral chronic inflammation
Low grade	0–4	Low grade	0	
Intermediate grade	5–6	Intermediate grade	2–3	
High grade	>7	High grade	≥4	

**Table 2.** Our own (Memorial Sloan-Kettering Cancer Center) grading system

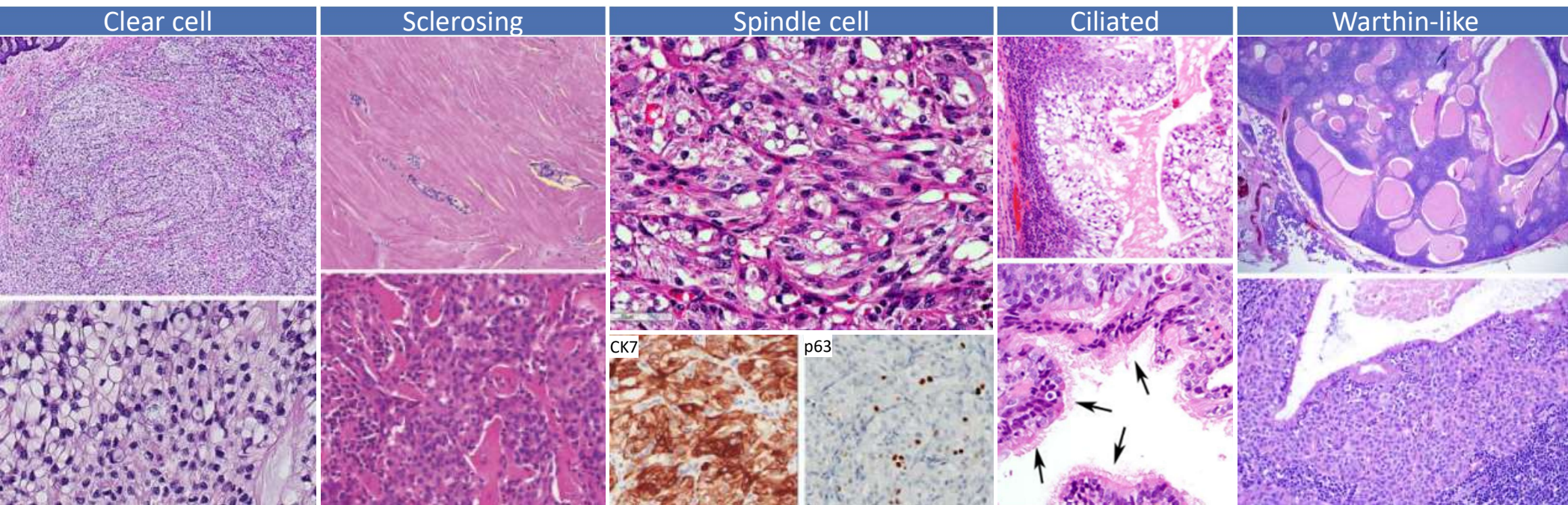
	Low grade	Intermediate grade	High grade
Predominant growth pattern	Cystic	Predominantly solid	Any (usually solid)
Infiltration	Well circumscribed borders	Well circumscribed or infiltrative borders	Any (usually infiltrative borders)
Mitosis	0–1/10 HPF	<4/10 HPFs	≥4/10 HPFs
Tumour necrosis	Absent	Absent	Present





# Mucoepidermoid carcinoma (MEC)

- Histopathology – variants



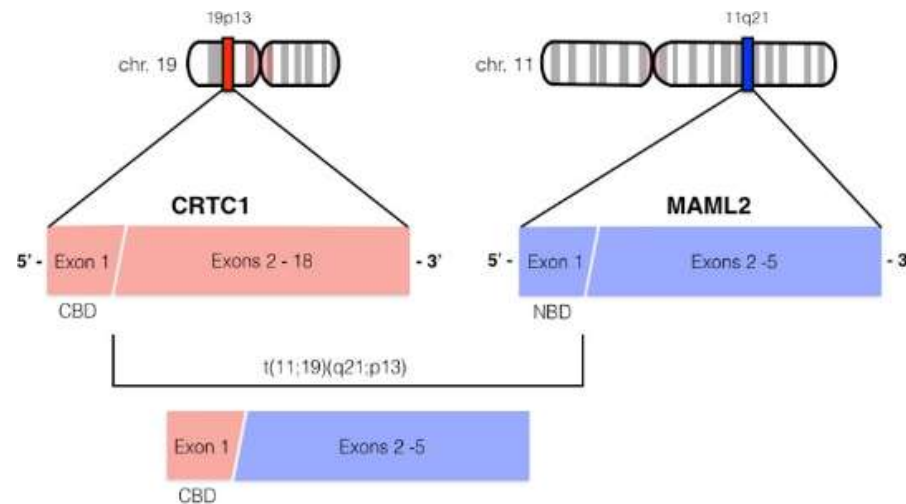
Tajima, Med Mol Morphol. 2015  
Goh, Int J Surg Pathol. 2017  
Yabuki, Int J Surg Pathol. 2017  
Bishop, Am J Surg Pathol JSP, 2019  
Skalova, Am J Surg Pathol. 2020



# Mucoepidermoid carcinoma (MEC)

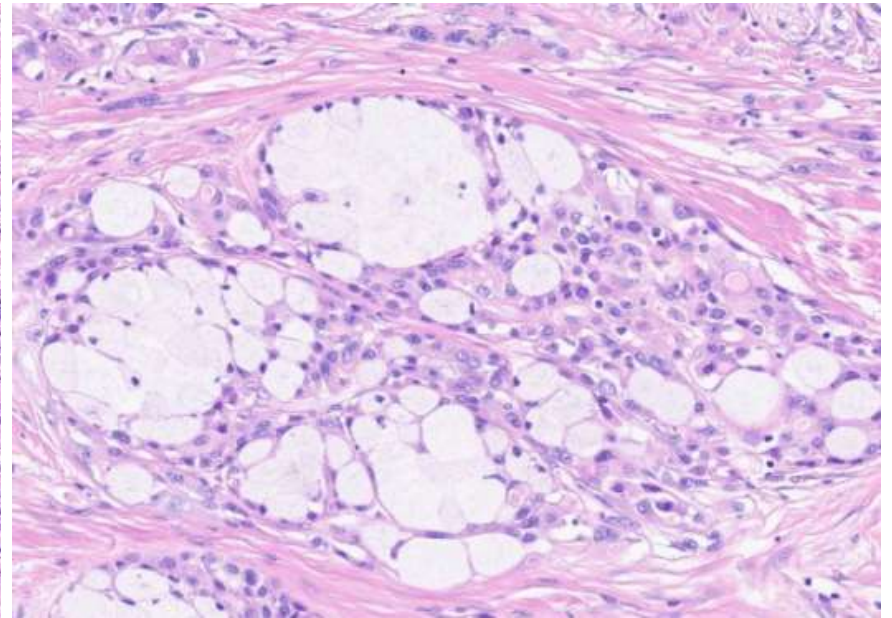
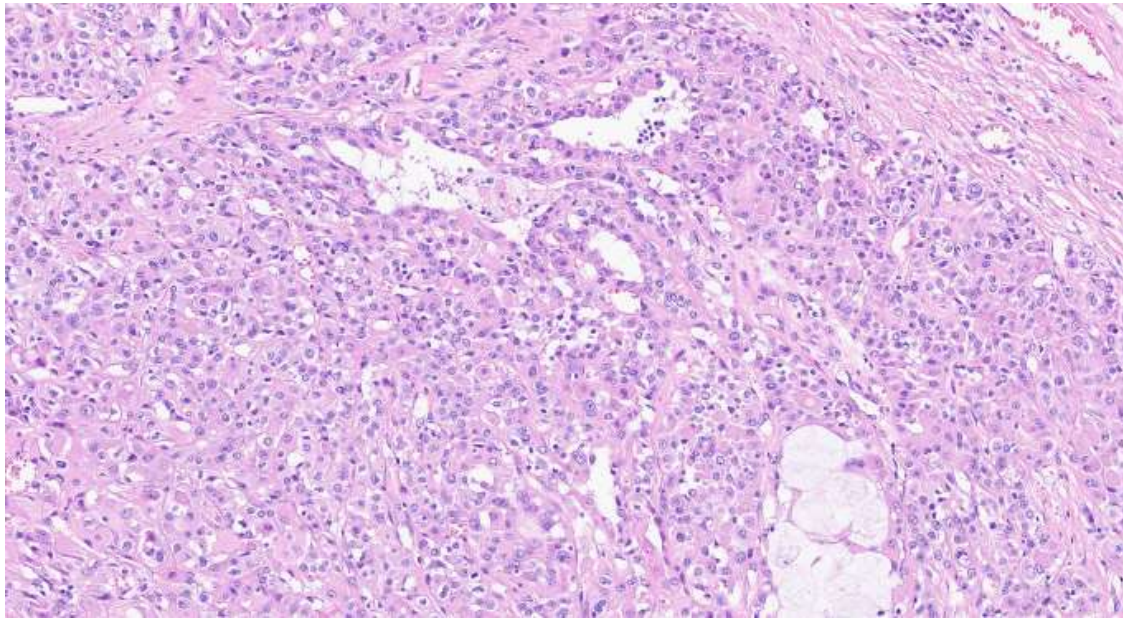
- **Molecular pathology**

- t(11;19)(q21;p13) *CRTC1::MAML2* fusion
  - Present in 55-88% of MECs
  - *CRTC1* exon 1 ~ *MAML2* exon 2 to 5
  - CREB-binding domain of *CRTC1* replace Notch-binding domain of *MAML2*
- Minor fusions including *CRTC1::MAML2*; *EWSR1::POU5F1*
- ***CRTC1/3-MAML2* as prognostic biomarker**
  - Favorable prognosis
  - Mostly seen in younger patients with low-intermediate grade MEC



# Mucoepidermoid carcinoma (MEC)

F/72; Tongue



-SNV/INDEL

Gene	Chr	Position	Refseq	Exon	AA	CDS	%VAF	Alt	Total	Tier
BAP1	chr3	52437795	NM_004656.3	13	p.Gln456*	c.1366C>T	5.11	97	1900	D

- Fusion

Fusion Gene	Cytoband1	Cytoband2	DNA.Evidence (Span/Split)	RNA.Evidence (Span/Split)	Transcript1	Transcript2	Last Observed	First Observed	Tier
CRTC1->MAML2	19p13.11	11q21	0/0	1/14	NM_001098482	NM_032427	EXON_1	EXON_2	D

# Salivary duct carcinoma (SDC)

- **Clinical features**

- Rapidly growing **high-grade** tumor, mostly involving major salivary gland
- Distinct **male predilection** with peak incidence in the fifth to seventh decades of life

- **Histopathology**

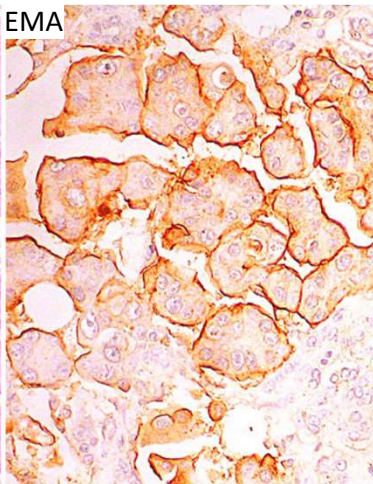
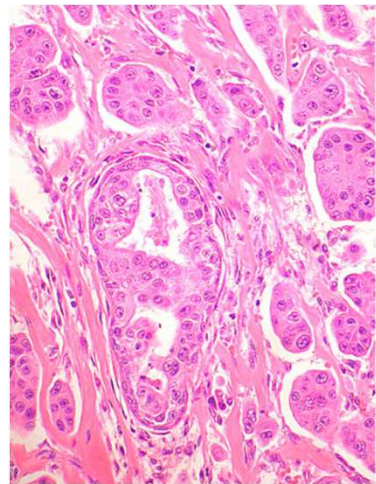
- Complex solid, cribriform, and papillary-cystic architecture with frequent comedo-necrosis
- Large pleomorphic nuclei with coarse chromatin and prominent nucleoli
- Abundant eosinophilic, typically apocrine cytoplasm
- Lymphovascular and perineural invasion are common
- IHC
  - **AR expression in 90%**
  - Diffuse and strong immunoreactivity for **ERBB2/HER2 is identified in about 30%**
  - CK7+, S100-, SOX10-



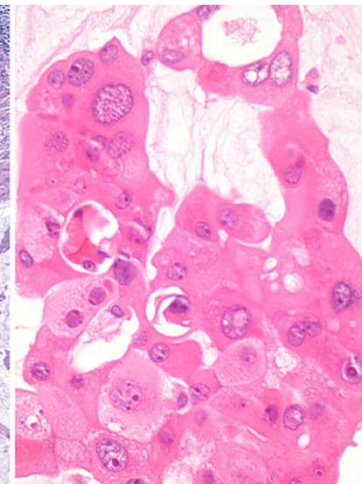
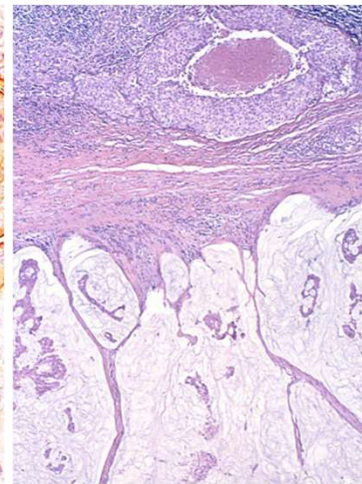
# Salivary duct carcinoma (SDC)

- Histopathology – variants

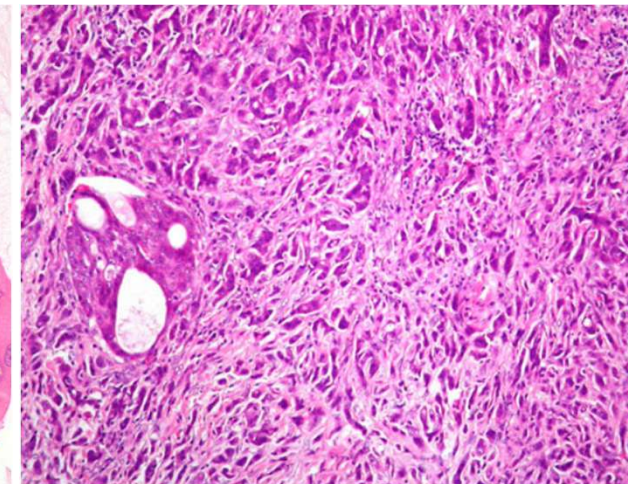
Micropapillary



Mucin-rich



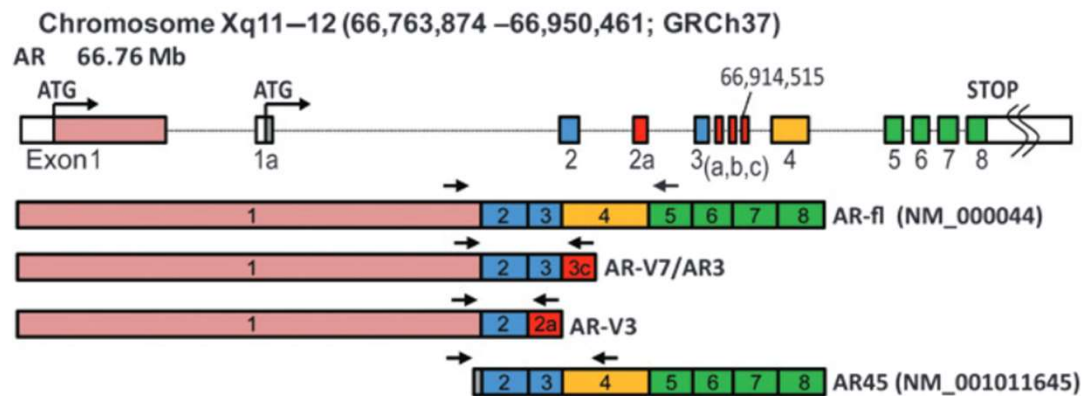
Mucin-rich



# Salivary duct carcinoma (SDC)

- **Molecular pathology**

- AR
  - chromosome Xq11-12
  - Copy number gain, mutation, alternative splicing



- Others

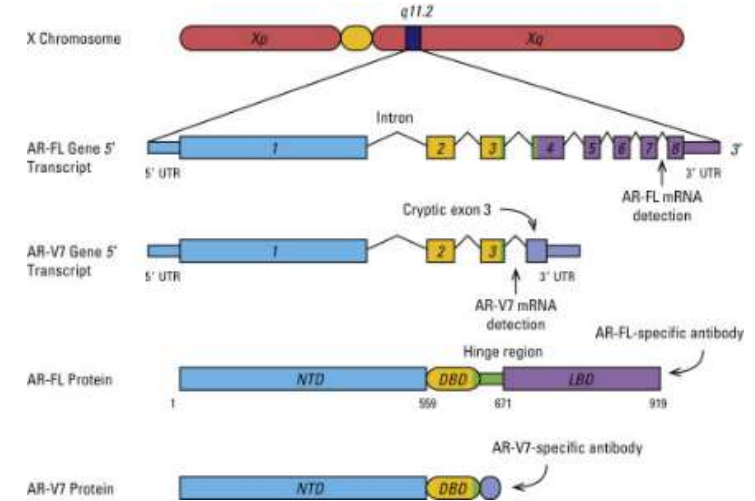
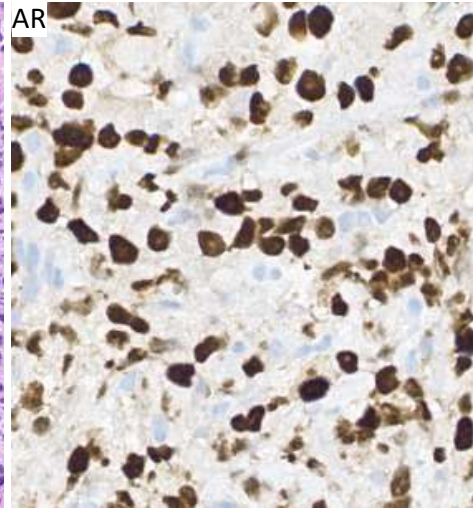
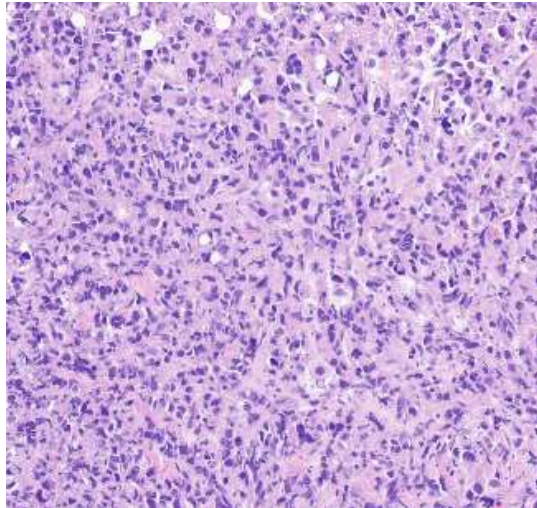
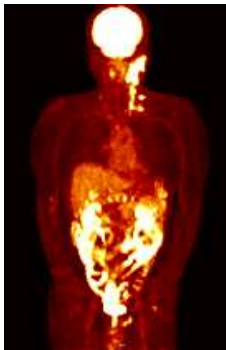
**TABLE 4.** Key Gene Mutations in Salivary Duct Carcinoma

Genetic alterations	Frequency (%)
<i>AR</i> alteration	40-70
<i>TP53</i> mutation	55-68
<i>ERBB2</i> amplifications	29-35
<i>PIK3CA</i> mutation	18-37
<i>PTEN</i> mutation	10-50
<i>NF1</i> mutation	16-21
<i>HRAS</i> mutation	4-27
<i>KMT2C</i> mutation	11
<i>EGFR</i> mutation	9
<i>ALK</i> mutation	7
<i>CDKN2A</i> mutation	7
<i>NOTCH1</i> mutation	7
<i>KDM5C</i> mutation	7
<i>NRAS</i> mutation	4
<i>BRAF</i> mutation	3-7
<i>AKT</i> mutation	2



# Salivary duct carcinoma (SDC)

M/67  
Parotid mass  
M/LN, brain



## Result of Actionable Gene Mutation Analysis

-SNV/INDEL

Gene	Chr	Position	Refseq	Exon	AA	CDS	%VAF	Alt	Total	Tier
ERBB2	chr17	37881000	NM_004448.3	20	p.Val777Leu	c.2329G>T	53.94	1492	2766	C

- Fusion

Fusion Gene	Cytoband1	Cytoband2	DNA.Evidence (Span/Split)	RNA.Evidence (Span/Split)	Transcript1	Transcript2	Last Observed	First Observed	Tier
AR-V7	Xq12	Xq12	0/0	0/50	NM_000044	NM_000044	EXON_3	EXON_CE3	D

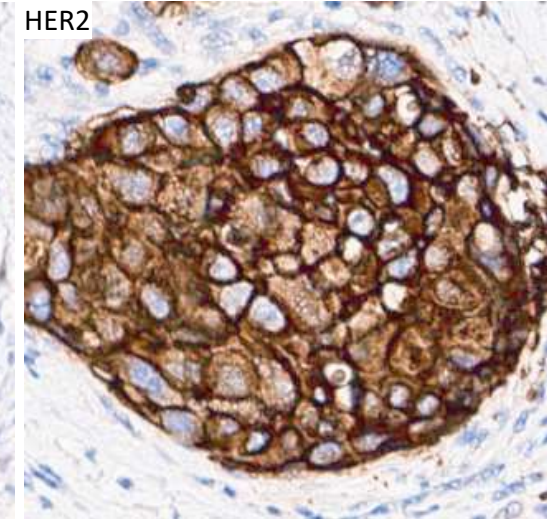
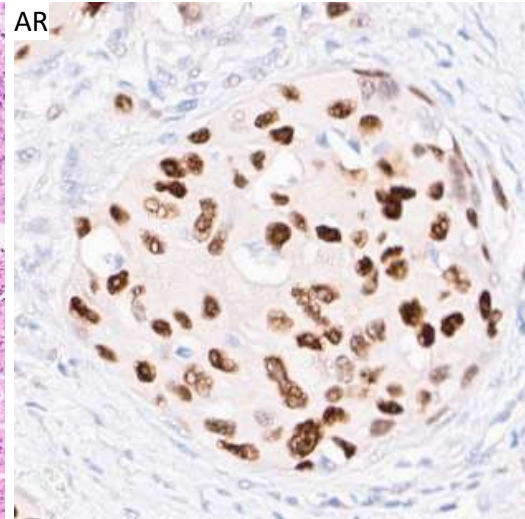
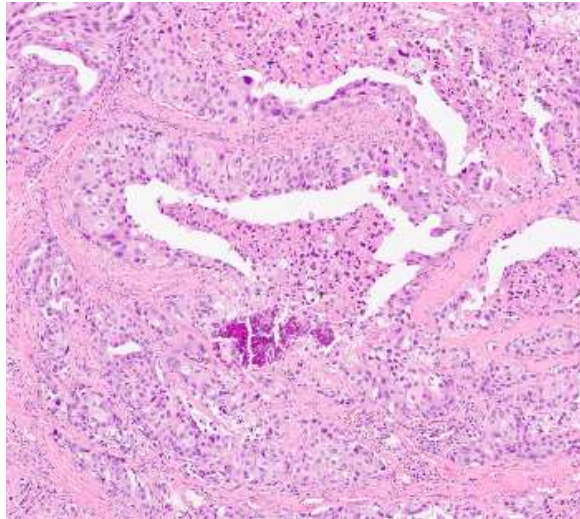
## AR-V7

Splice variant of the androgen receptor mRNA resulting in the truncation of the ligand-binding domain



# Salivary duct carcinoma (SDC)

M/51  
SMG mass



-SNV/INDEL

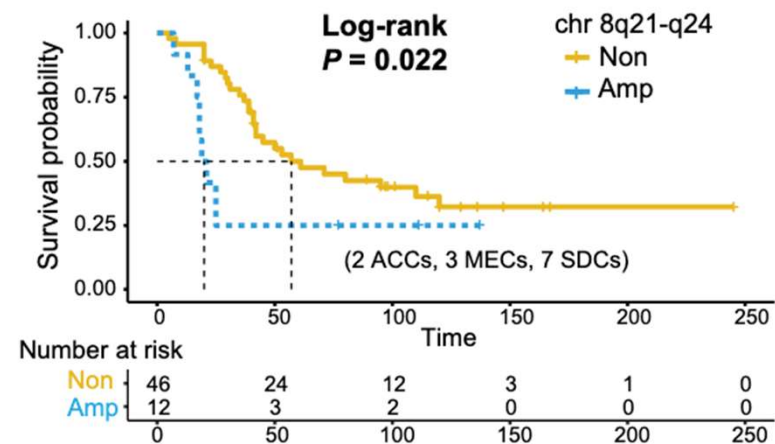
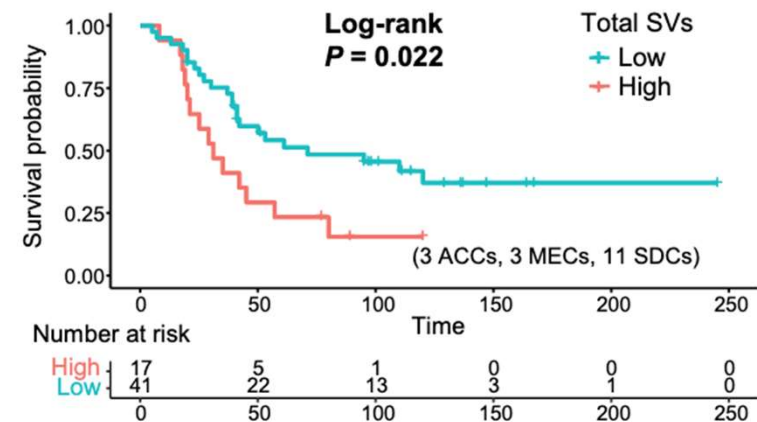
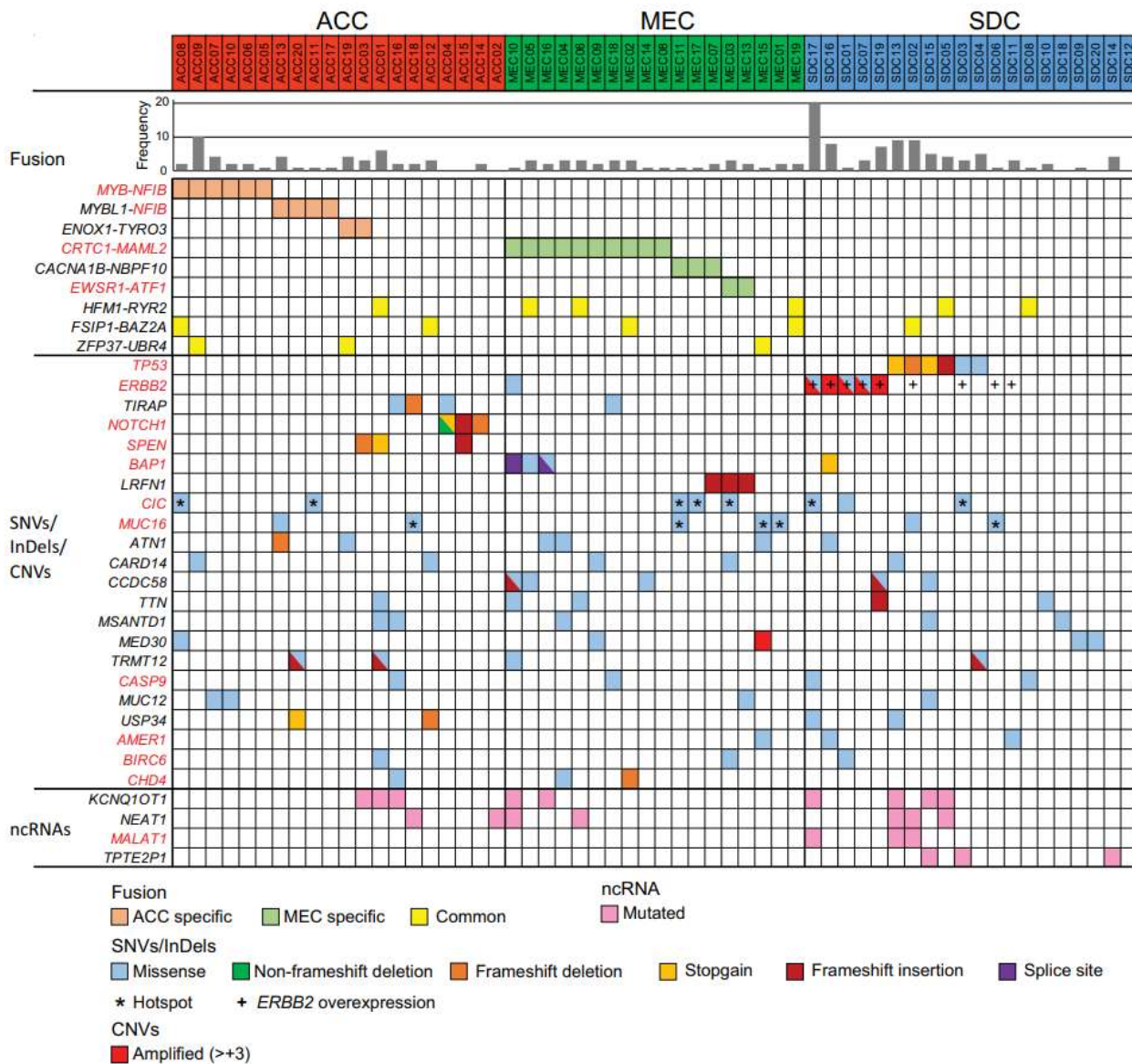
Gene	Chr	Position	Refseq	Exon	AA	CDS	%VAF	Alt	Total	Tier
NRAS	chr1	115258744	NM_002524.4	2	p.Gly13Asp	c.38G>A	1.27	26	2047	C
PBRM1	chr3	52668619	NM_018313.4	12	p.Arg434*	c.1300C>T	18.98	26	137	D
TP53	chr17	7578553	NM_000546.5	5	p.Tyr126Cys	c.377A>G	28.56	347	1215	D

-Fusion

Fusion Gene	Cytoband1	Cytoband2	DNA.Evidence (Span/Split)	RNA.Evidence (Span/Split)	Transcript1	Transcript2	Last Observed	First Observed	Tier
Not detected									

-CNV

Gene	Chr	Start	End	CNV	AvgL2R	DelR	NeutralR	AmpR	CN	Tier
ERBB2	chr17	37855608	37884459	Amplification	2.974	0.0	0.0	1.0	14	C



# Acinic cell carcinoma (AciCC)

- **Clinical features**

- 10% of all salivary gland malignancies; 90-95% of cases arise in the parotid gland; 18.7% of parotid carcinoma
- 2<sup>nd</sup> most common salivary gland malignancy in children; wide age range (average 47.7 – 52)

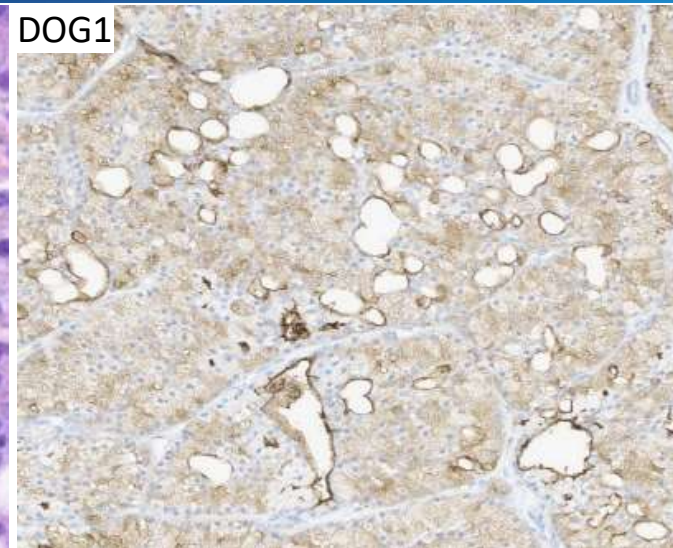
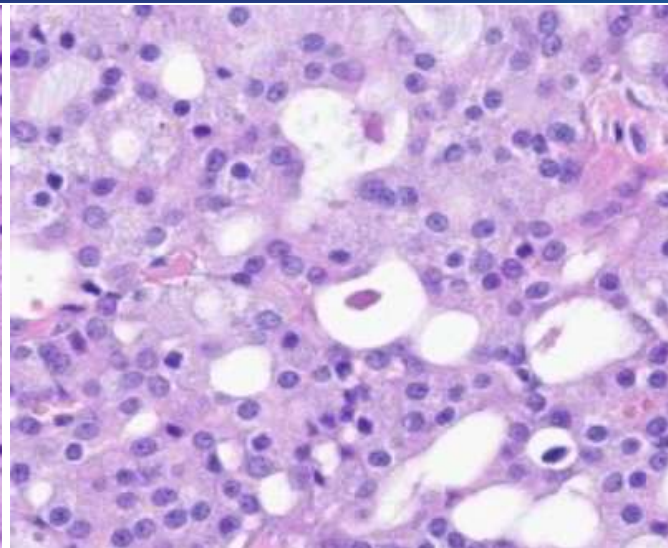
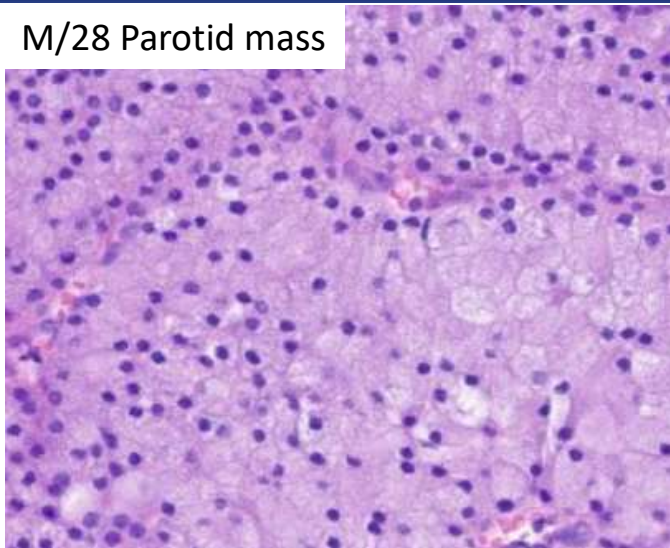
- **Histopathology**

- Solid, microcystic, follicular, papillary-cystic architectures
- Most commonly serous acinar cell proliferation
  - PAS-positive, diastase-resistant basophilic cytoplasmic zymogen granules
- Variable intercalated duct-type, nonspecific glandular, vacuolated, oncocytic, and rarely clear cells
- IHC
  - SOX10+, DOG1+ / p40-, p63-, mammaglobin-, S100-
- High grade transformation
  - nuclear pleomorphism, coarse chromatin, necrosis, ↑mitotic activity/Ki-67 index, ↑ frequent perineural / LVI

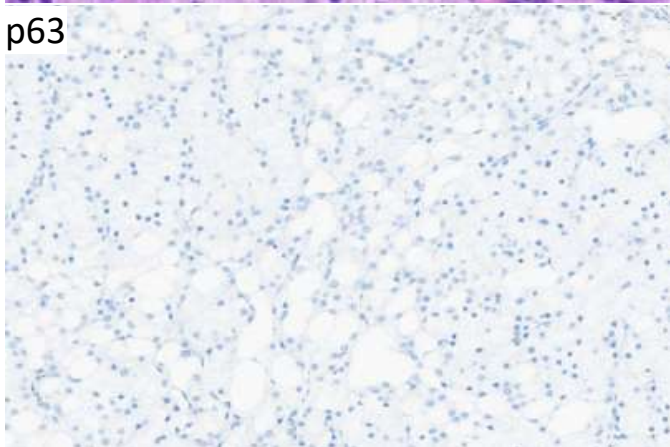


# Acinic cell carcinoma (AciCC)

M/28 Parotid mass



p63



S100



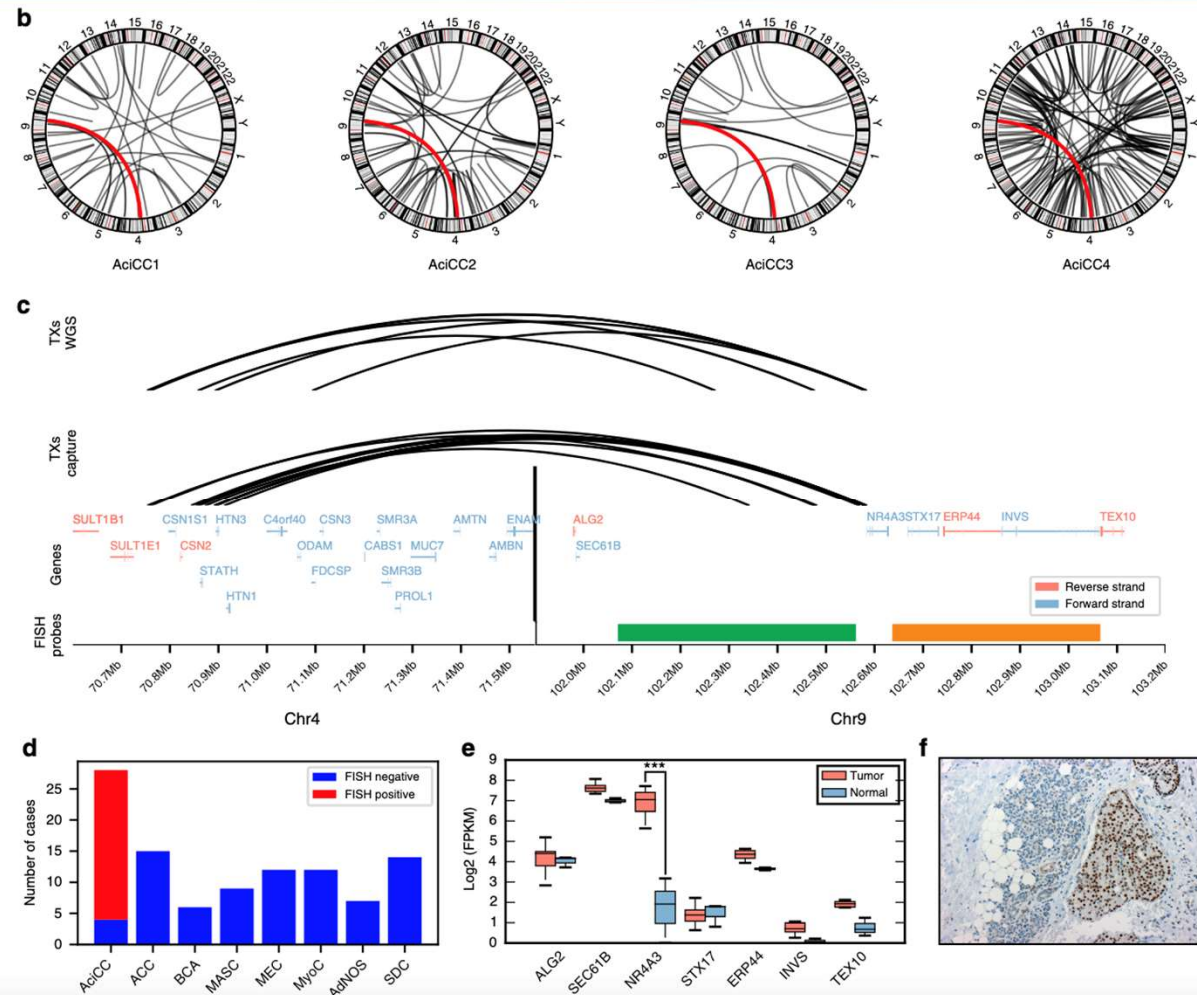
TRK



# Acinic cell carcinoma (AciCC)

## Molecular pathology

- $t(4;9)(q13;q31)$  rearrangement
  - Active enhancer of secretory Ca-binding phosphoprotein (SCPP) gene cluster at upstream of *NR4A3* gene
  - Upregulation of *NR4A3* via enhancer hijacking
- NR4A3 IHC
  - Sensitivity 94.4%; specificity 99%



Haller, Nat Commun. 2019  
Haller, Am J Surg Pathol. 2019

# Secretory carcinoma (SC)

- **Clinical features**

- Painless slowly growing mass; parotid gland (m/c) > oral cavity > submandibular gland
- Mean patient age of 46.5 (range 10-86 years)

- **Histopathology**

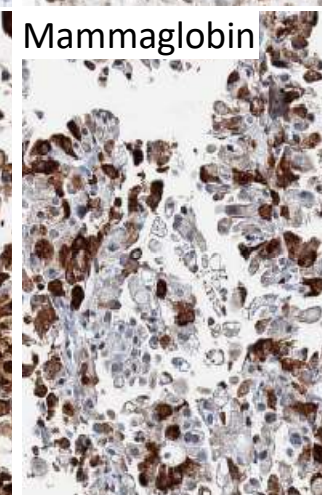
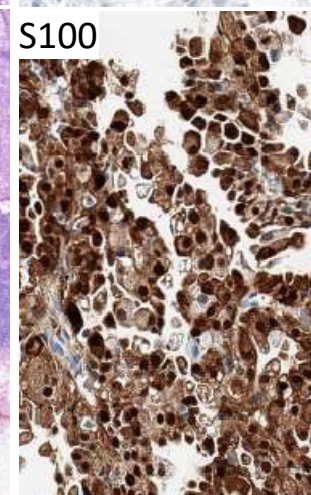
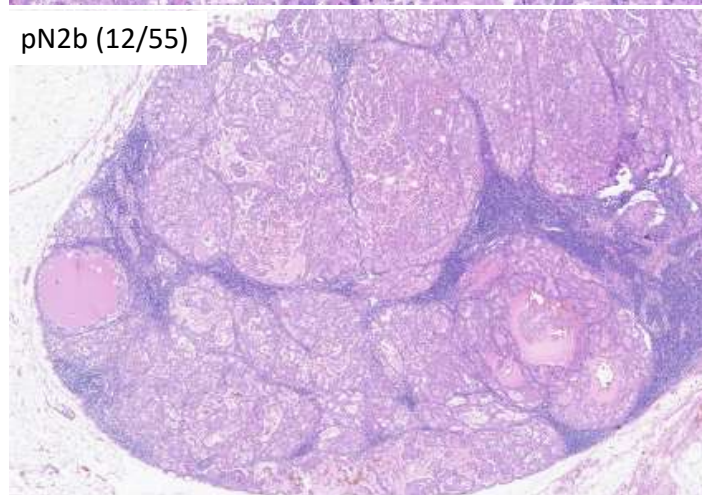
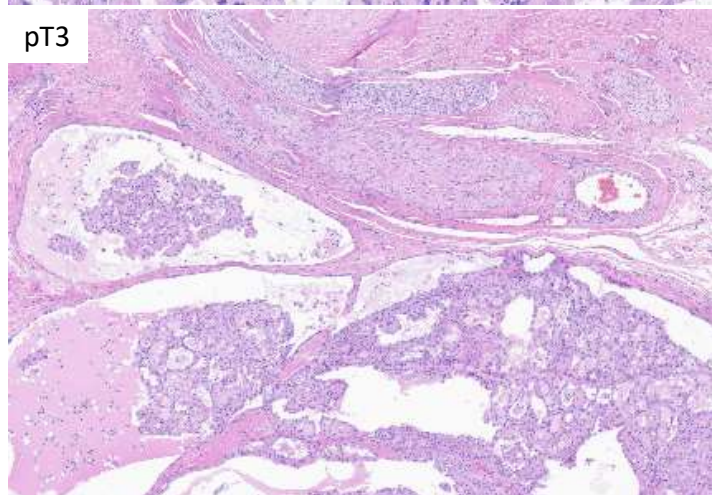
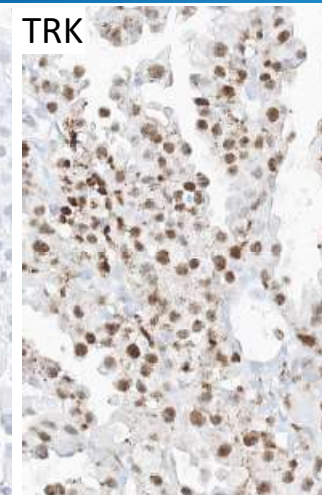
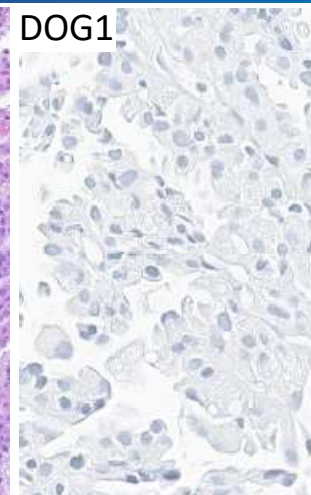
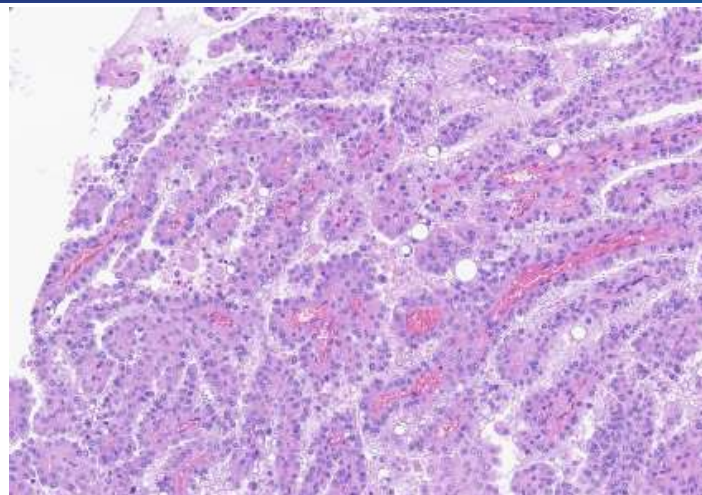
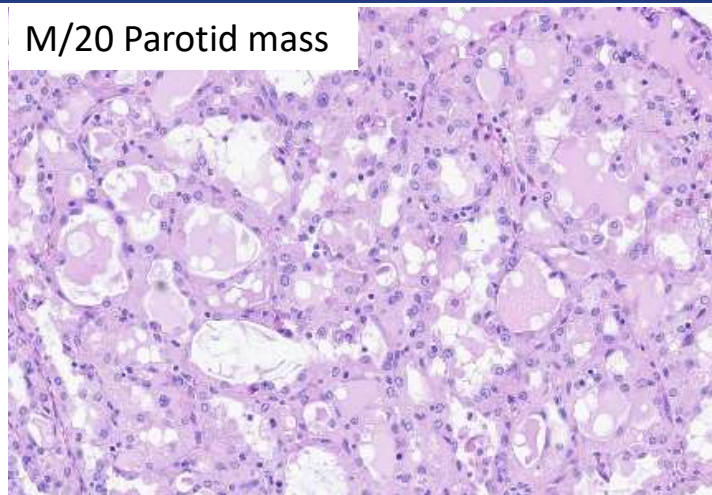
- Lobulated growth pattern separated by fibrous septa
- Microcystic/solid, tubular, follicular, and papillary-cystic structures with distinctive luminal secretions
- Low grade vesicular round-to-oval nuclei with finely granular chromatin and distinctive centrally located nucleoli
- Granular to vacuolated pale, pink cytoplasm
- Special stain/IHC
  - PAS, D-PAS, alcian-blue+ luminal secretions
  - CK7+ S100+, SOX10+, vimentin+, mammaglobin+
  - p63-, p40- DOG1-, NR4A3-

- **Molecular pathology**

- 90% harbor t(12;15)(p13;q25) chromosomal rearrangement → *ETV6::NTRK3* fusion
- Rarer rearrangements include *ETV6::RET*, *ETV6::MET*, *ETV6::MAML3*, *VIM::RET*



# Secretory carcinoma (SC)



# Basal cell adenocarcinoma (BCAC)

- **Clinical features**

- Slow growing nodule in parotid gland (>90%) in 6<sup>th</sup>–7<sup>th</sup> decades
- About 15% occur in the setting of familial/multiple cylindromatosis syndromes
- Complete surgical excision is curative in >90%
- Regional lymph node and distant metastasis and disease related death are rare (<10%)

- **Histopathology**

- Tubulo-trabecular, membranous, solid growth patterns
- Tumor nests show peripheral palisading of dark cells with paler cells and centrally located ducts
- DDx BCAC from basal cell adenoma (BCA)
  - Infiltration; necrosis; mitosis (>4 per 2mm<sup>2</sup>)
- IHC
  - Epithelial and myoepithelial markers highlights the dual cell components
  - Nuclear  $\beta$ -catenin immunoexpression: BCAC (66.7%-100%); BCA (70%-100%)

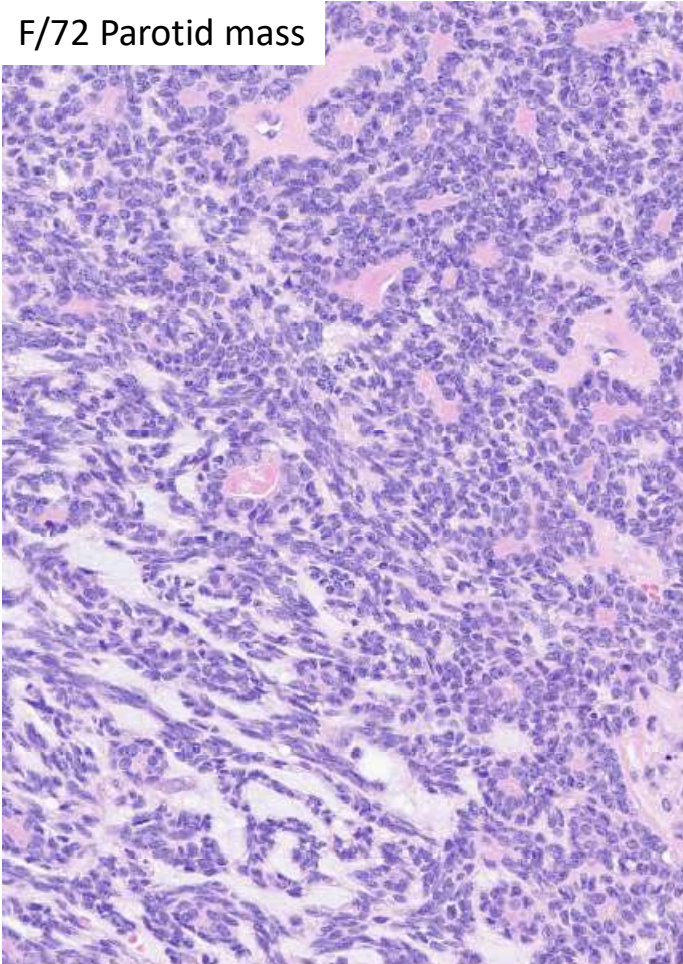
- **Molecular pathology**

- Diagnostic feature: none
- *CTNNB1* mutations: more common in BCA than BCAC
- Other genetic profiles of BCAC
  - Activating mutations in *PIK3CA*, biallelic inactivation of *NFKB1A*, focal *CYLD* deletion

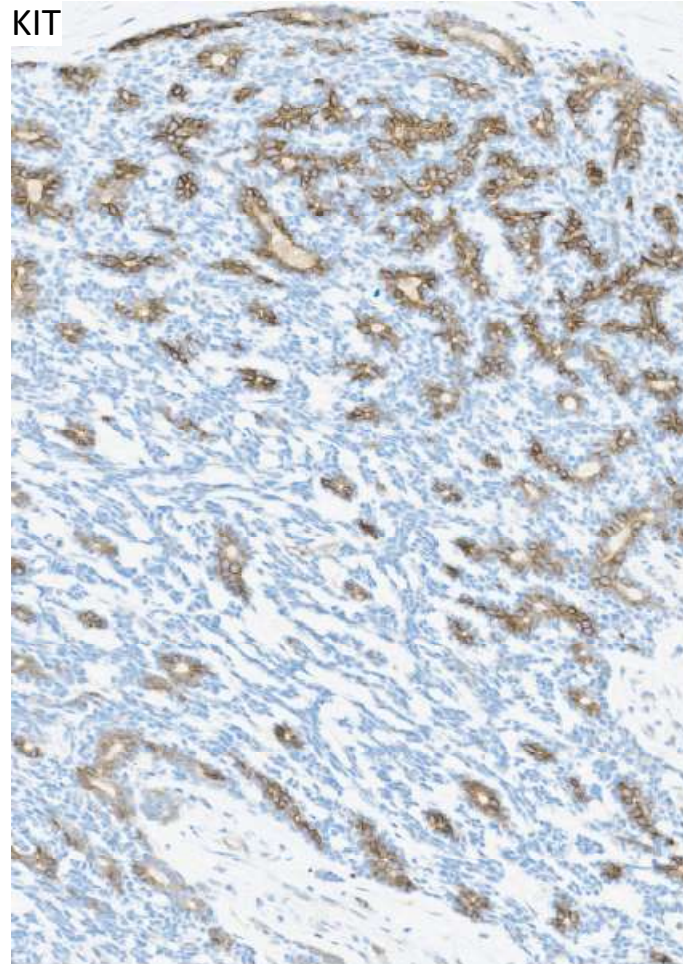


# Basal cell adenocarcinoma (BCAC)

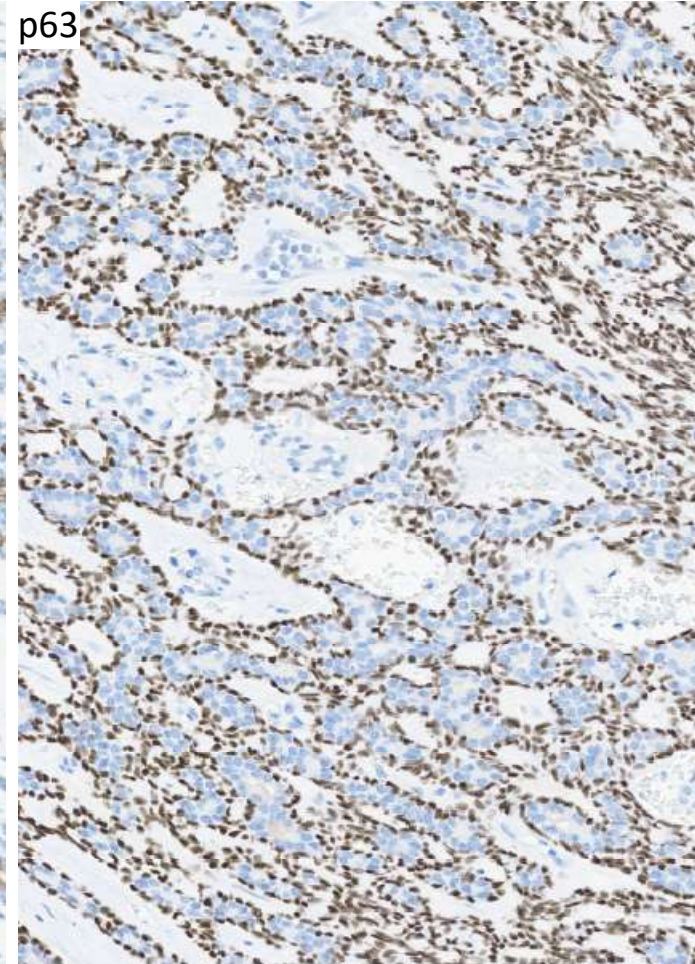
F/72 Parotid mass



KIT

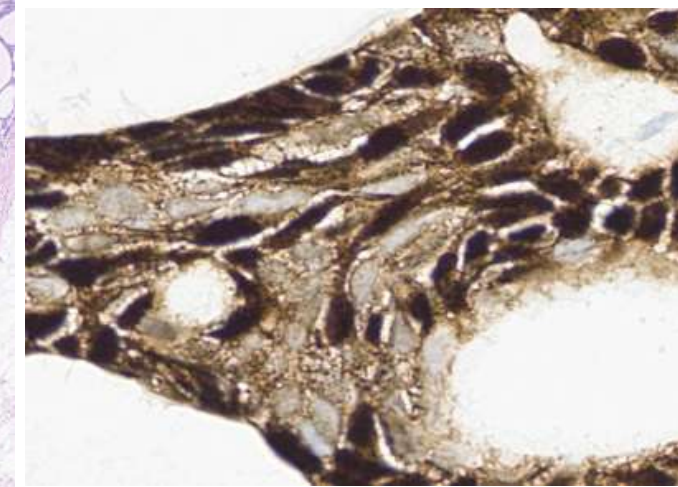
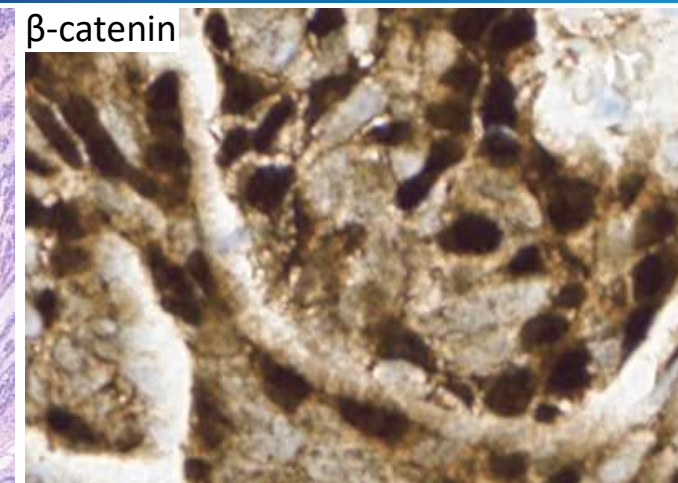
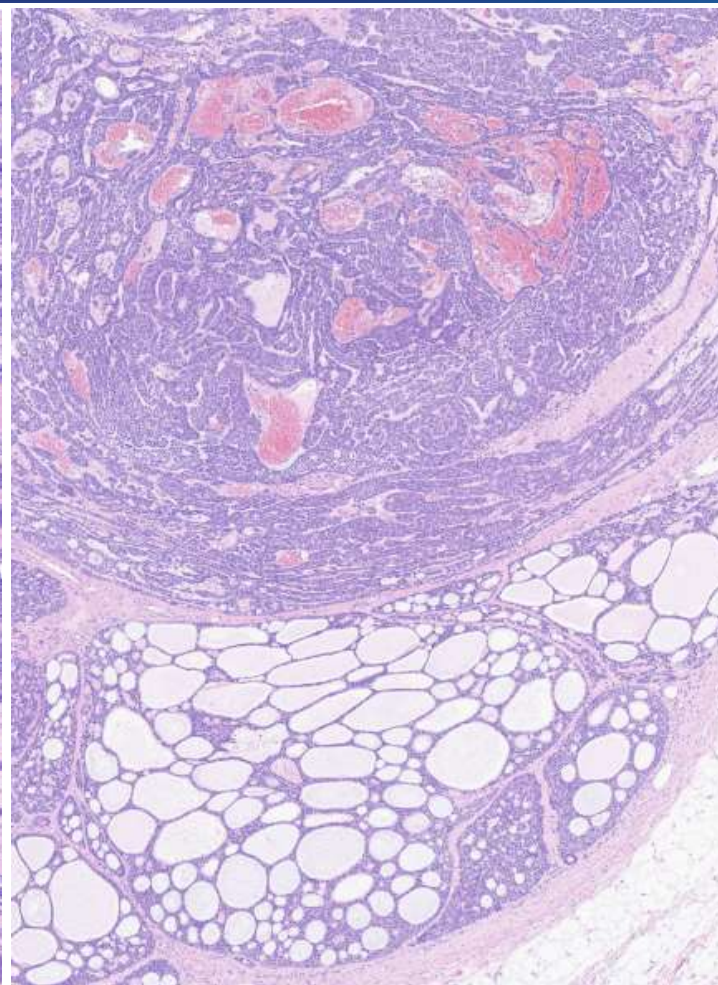
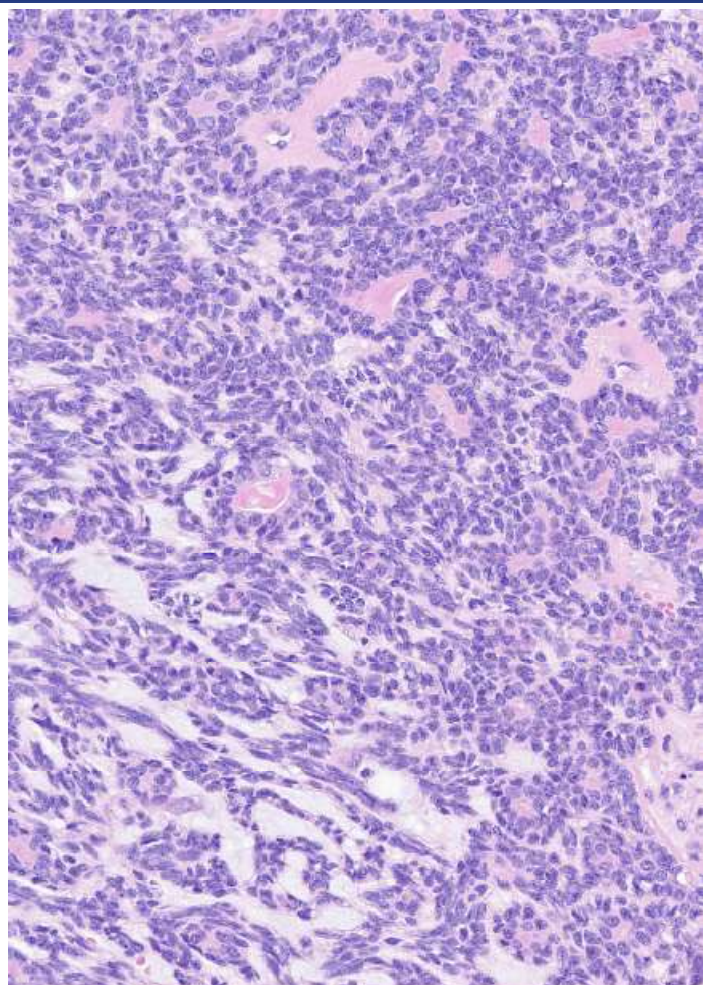


p63





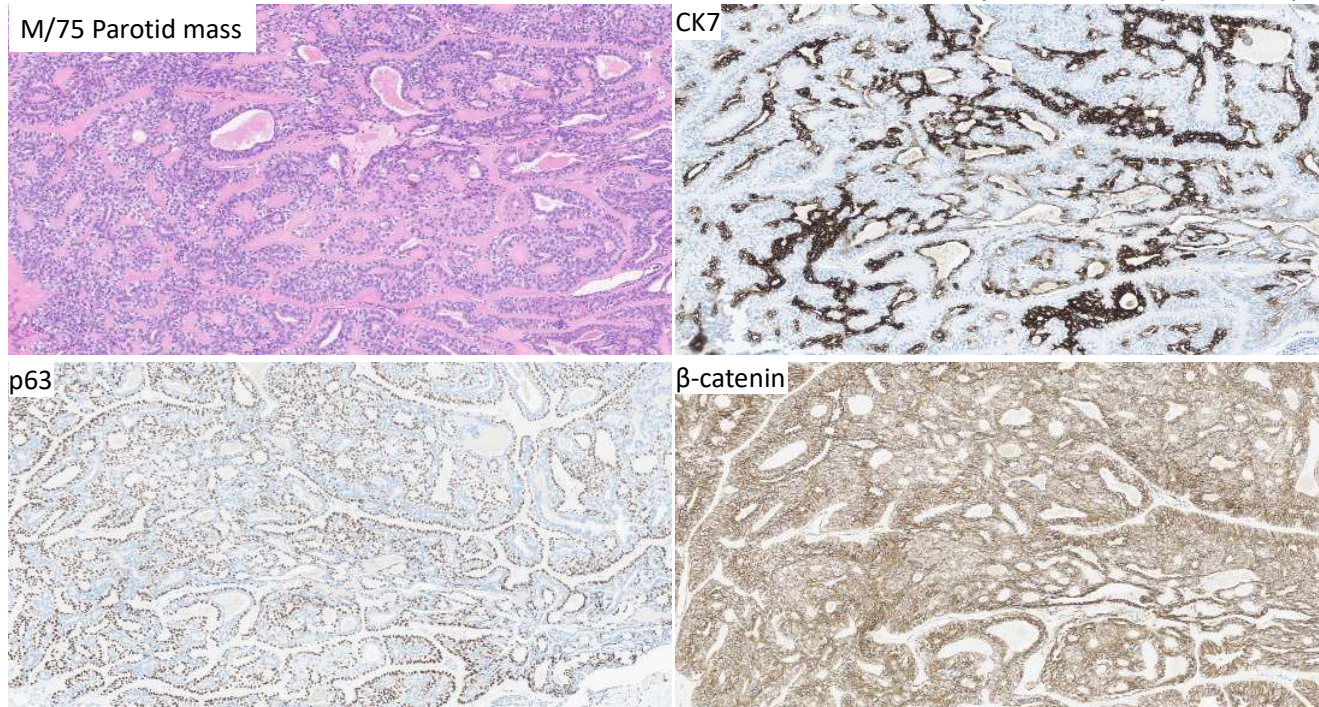
# Basal cell adenocarcinoma (BCAC)





# Epithelial-myoepithelial carcinoma (EMC)

- **Clinical features**
  - 1% of all salivary gland tumors mostly affecting parotid gland in 6<sup>th</sup>-7<sup>th</sup> decades
- **Histopathology**
  - Biphasic tubules → tightly coupled inner eosinophilic ductal and prominent outer (often clear) myoepithelial cells
- **Molecular pathology**
  - **HRAS codon 61 mutation** in 82.7%; HRAS Q61R mutation-specific IHC positivity in 65%



# Carcinoma ex pleomorphic adenoma (CXPA)

- **Clinical features**

- Epithelial and/or myoepithelial malignancy in association with a primary or recurrent pleomorphic adenoma (PA), presenting at 6<sup>th</sup>-7<sup>th</sup> decades
- Long-standing painless mass with recent rapid progression or of previous diagnosis of PA
- Most commonly at parotid gland
- Aggressive tumors with local and distant recurrence rates of 70%; 5-year survival from 25% to 75%
  - Poor prognostic factors: invasive CXPA, large size > 4 cm, multiple LN+, distant metastasis

- **Histopathology**

- PA and carcinoma components can be intermixed or appear as discrete nodules
- Histologic types of malignant components
  - Most commonly SDC > myoepithelial carcinoma > adenocarcinoma, NOS
  - Rarely carcinosarcoma : malignant epithelial and sarcomatous components arising in association with a PA

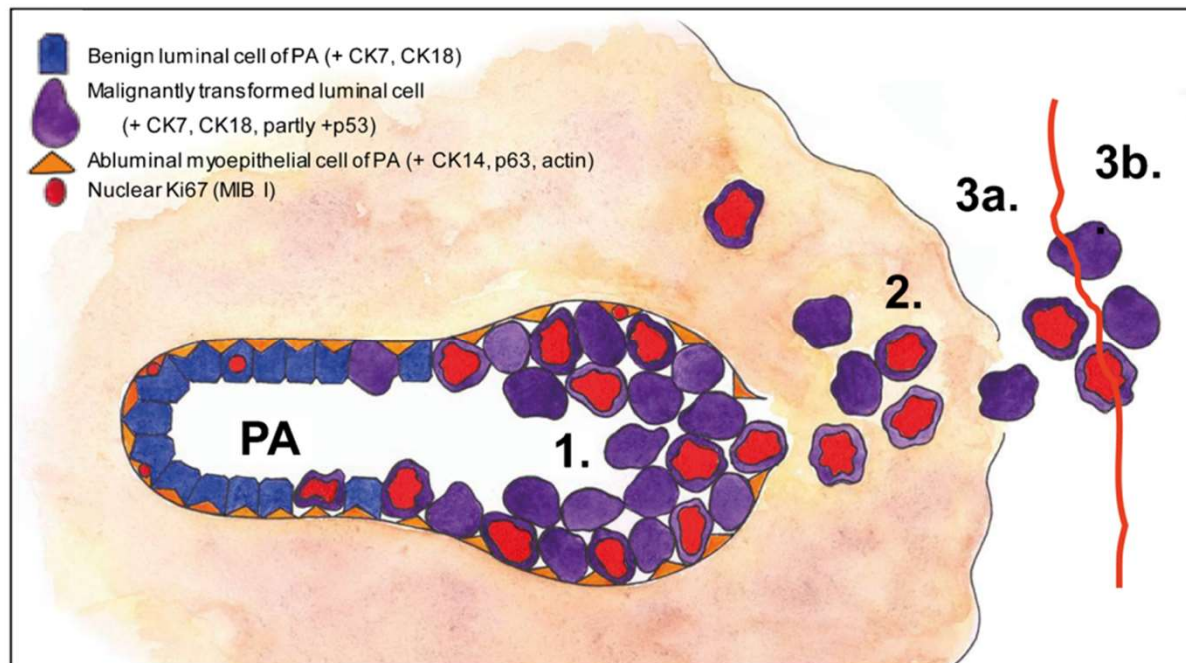


# Carcinoma ex pleomorphic adenoma (CXPA)

- **Histopathology – Subclassification**

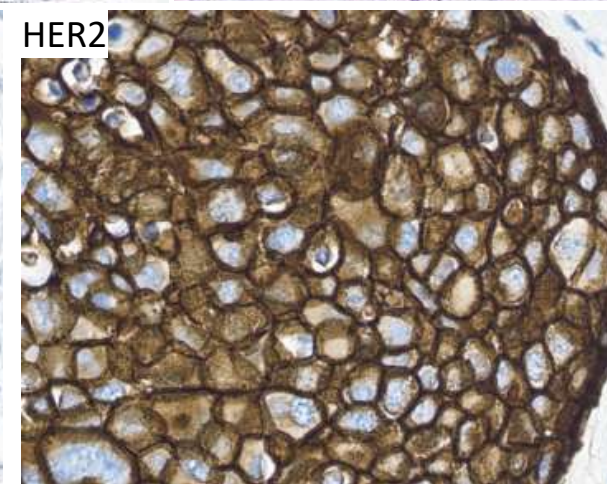
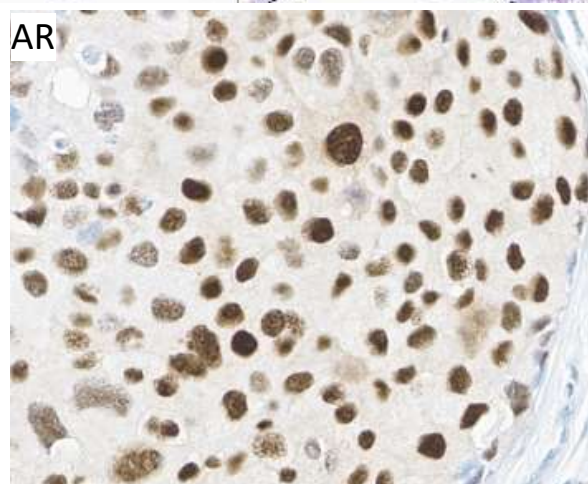
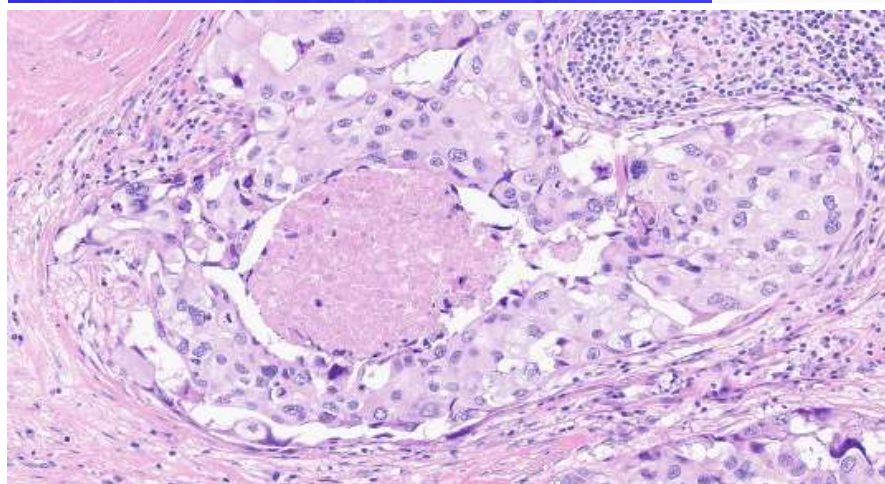
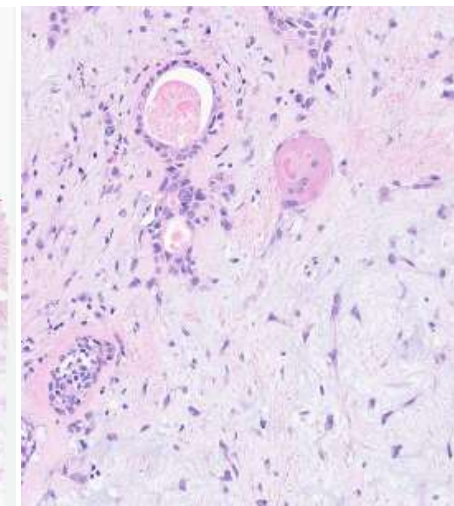
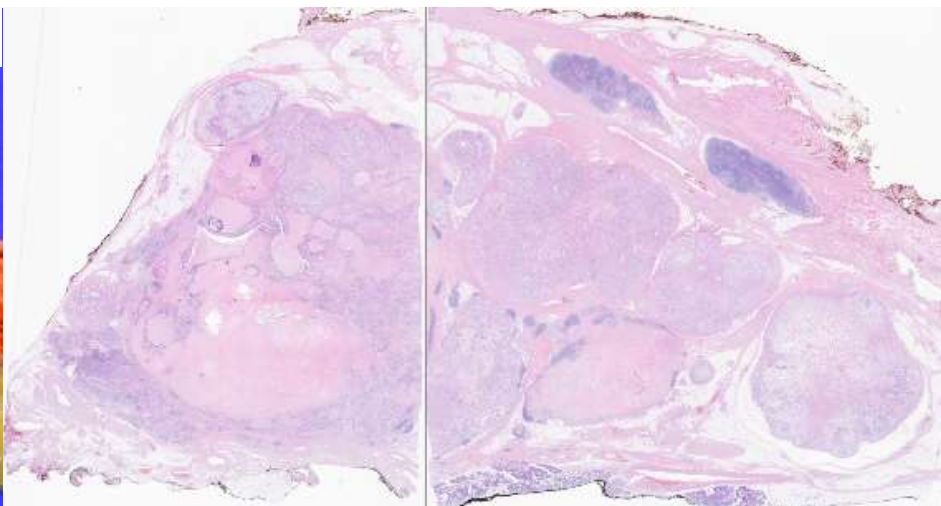
- Based on the **extent of invasion beyond PA borders**

- 1) Intracapsular: carcinoma is confined within the PA capsule
  - In situ intracapsular when the malignant tumor cells replace ductal cells with an intact myoepithelial layer
- 2) Minimally invasive: carcinoma invades < 4 - 6 mm beyond the PA borders
- 3) Invasive: invasion beyond the PA capsule measures  $\geq 6$ mm



# Carcinoma ex pleomorphic adenoma (CXPA)

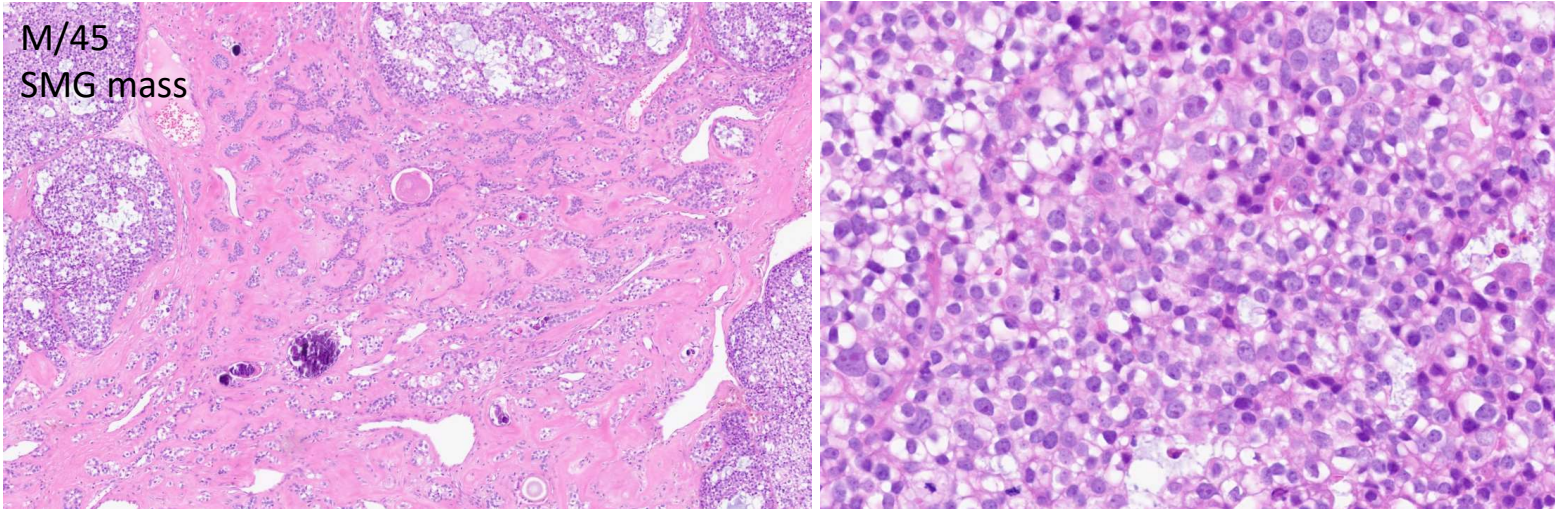
M/66 Parotid mass



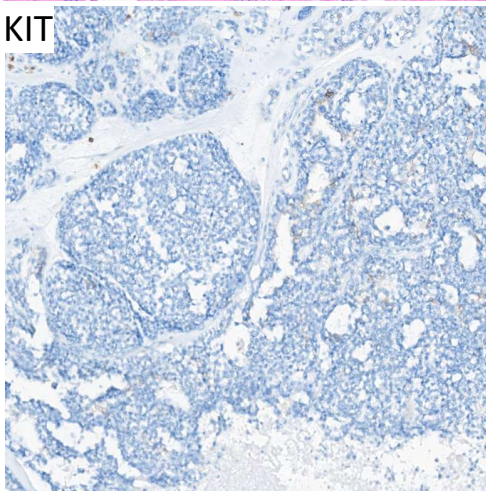


# Carcinoma ex pleomorphic adenoma (CXPA)

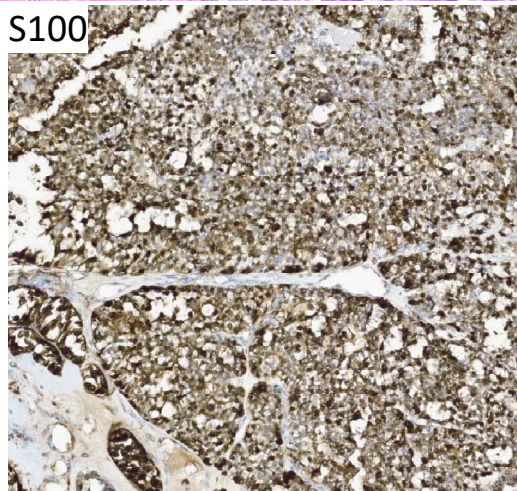
M/45  
SMG mass



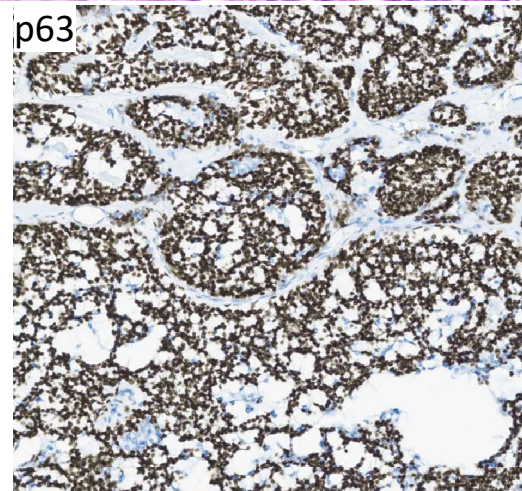
KIT



S100



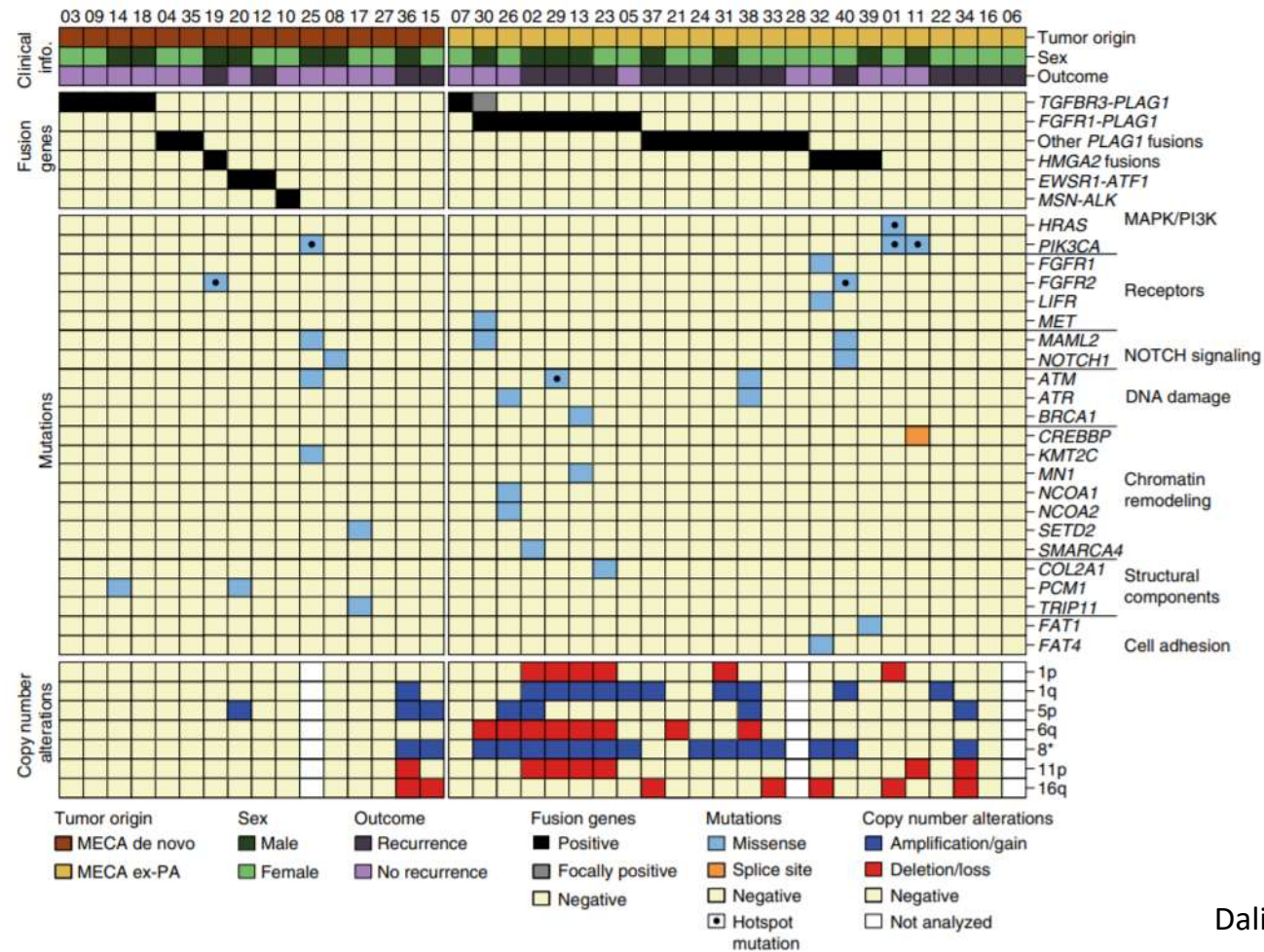
p63





# Carcinoma ex pleomorphic adenoma (CXPA)

- Molecular pathology



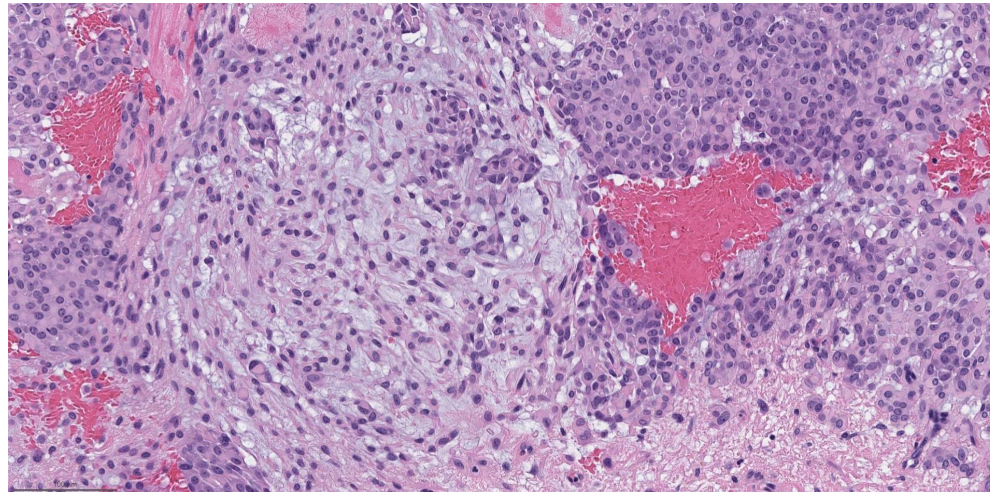
# Myoepithelial carcinoma (MC)

- **Clinical features**

- Parotid gland followed by the palate, and the submandibular gland

- **Histopathology**

- De novo or CXPA (2<sup>nd</sup> most common histologic subtype of CXPA)
- Diverse cellular morphology including spindle, epithelioid, plasmacytoid, vacuolated, and clear cells
- Solid, trabecular, and reticular growth patterns
- Myxoid, myxochondroid or hyalinized stroma
- IHC
  - SOX10+, S100+, CK+, myoepithelial markers+ (SMA, calponin, p63, p40)



# Hyalinizing clear cell carcinoma (HCCC)

- **Clinical features**

- Mostly at **oral minor salivary glands** > oropharynx, nasopharynx, parotid, sinonasal tract, larynx, and hypopharynx
- Median age at diagnosis of 56 years (range 23-87 years)

- **Histopathology**

- Nests, cords and trabeculae composed of clear or eosinophilic cells
  - Some cases completely lack clear cells
- Unencapsulated with cells infiltrating the surrounding tissue
- Densely hyalinized basement membrane-like to desmoplastic or fibrocellular stroma
- IHC
  - CK7+, CK19+, CK14+, EMA+
  - P63+, p40+, CK5/6+ → supporting squamous differentiation
  - Negative for myoepithelial markers (S100, SMA, calponin)

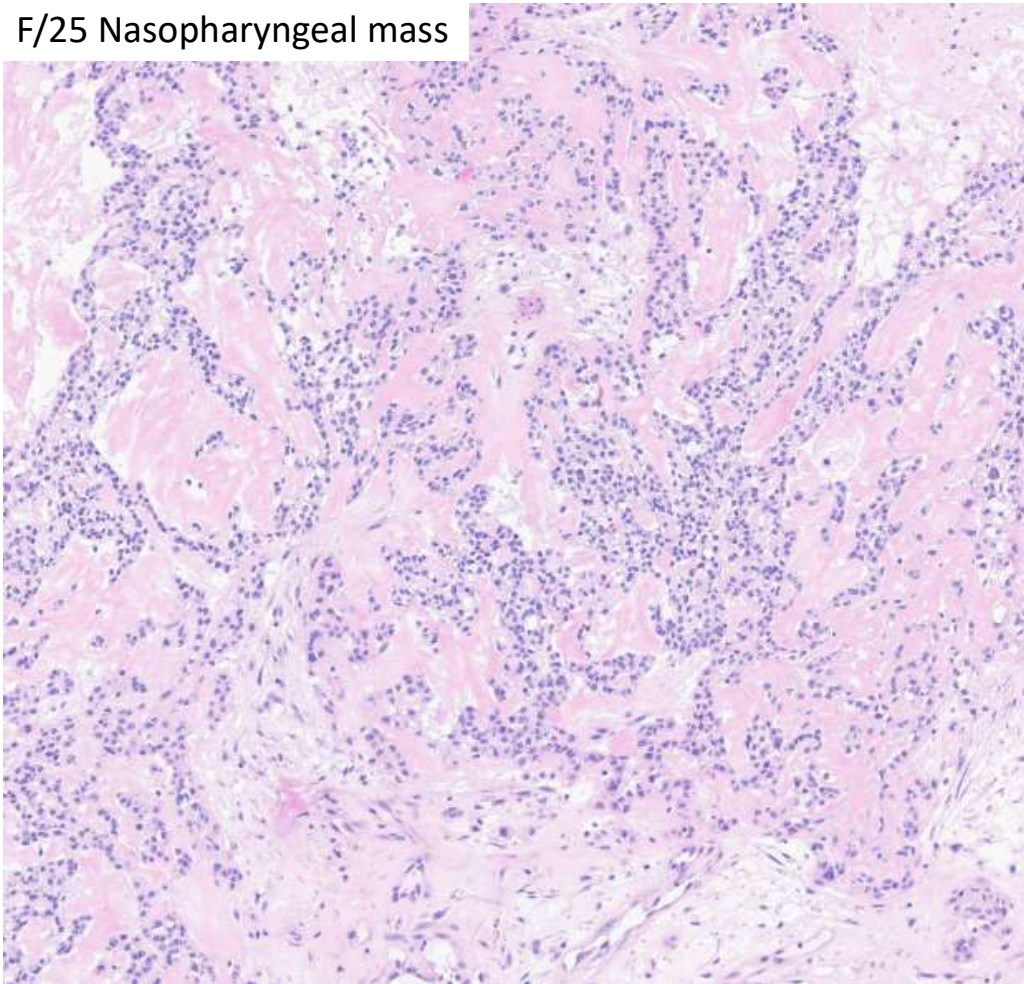
- **Molecular pathology**

- ***EWSR1::ATF1***; rarer fusion including *EWSR1::CREM*

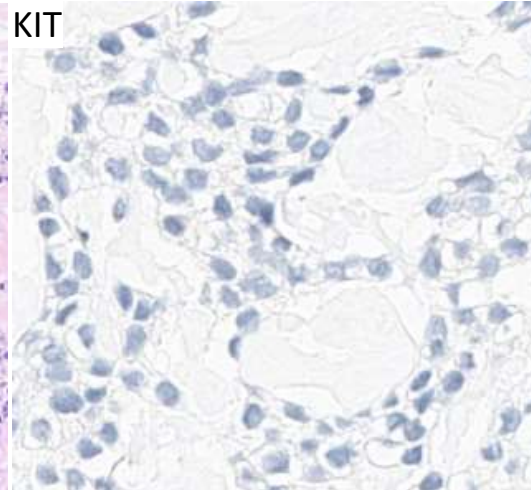


# Hyalinizing clear cell carcinoma (HCCC)

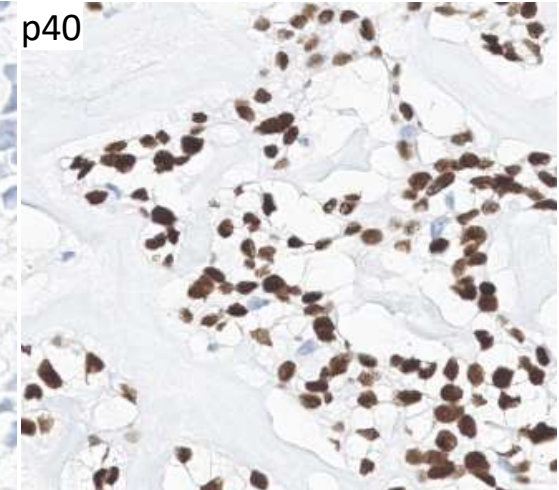
F/25 Nasopharyngeal mass



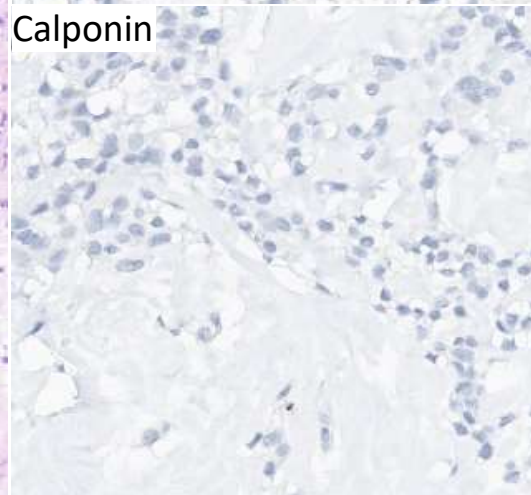
KIT



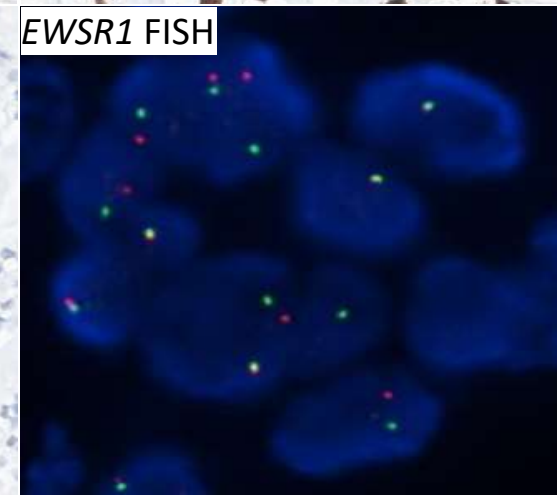
p40



Calponin



*EWSR1* FISH



# Summary

	Gene and mechanism	Prevalence
AdCC	<i>MYB</i> fusion/activation/amplification	~80%
	<i>MYBL1</i> fusion/activation/amplification	~10%
	<i>NOTCH</i> mutations	14%
MEC	<i>CRTC1::MAML2</i>	40-90%
	<i>CRTC3::MAML2</i>	6%
	<i>CDKN2A</i> deletion	25%
SDC	<i>ERBB2</i> amplification	31%
	<i>FGFR1</i> amplification	10%
	<i>TP53</i> mutation	56%
	<i>PIK3CA</i> mutation	33%
	<i>HRAS</i> mutation	33%
	<i>AR</i> copy gain	35%
	<i>PTEN</i> loss	38%
	<i>CDKN2A</i> loss	10%

	Gene and mechanism	Prevalence
AciCC	<i>NR4A3</i> fusion/activation	86%
	<i>MSANTD3</i> fusion/amplification	4%
SC	<i>ETV6::NTRK3</i> fusion	> 90%
	<i>ETV6::RET</i> fusion	2–5%
	<i>ETV6::MET</i> fusion	<1%
	<i>ETV6::MAML3</i> fusion	<1%
	<i>VIM::RET</i> fusion	<1%
BCAC	<i>CYLD</i> mutations	29%
CXPB	<i>PLAG1</i> fusions/amplification	73%
	<i>HMGA2</i> fusions/amplification	14%
	<i>TP53</i> mutations	60%
EMC	<i>HRAS</i> mutations	78%
MC	<i>PLAG1</i> fusions	38%
	<i>EWSR1::ATF1</i> fusions	13%
HCCC	<i>EWSR1::ATF1</i> fusions	93%
	<i>EWSR1::CREM</i> fusions	<5%



**Thank you for your attention**

