



Adult CNS Tumor Pathology – WHO 2021 Brief Update and Case Examples

Grey Matters Symposium

Sahara Cathcart, MD

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I have NO financial relationships or conflicts of interests relating to the subject matter of this presentation.

ICD-O coding of central nervous system tumours

ICD-O-3.2 ICD-O label (subtypes are indicated in grey text, with the label indented);

Please note that the WHO classification of tumour types is more readily reflected in the table of contents

Gliomas, glioneuronal tumours, and neuronal tumours

Adult-type diffuse gliomas

	Astrocytoma, IDH-mutant
9400/3	Astrocytoma, IDH-mutant, grade 2
9401/3	Astrocytoma, IDH-mutant, grade 3
9445/3	Astrocytoma, IDH-mutant, grade 4
	Oligodendroglioma, IDH-mutant and 1p/19q-codeleted
9450/3	Oligodendroglioma, IDH-mutant and 1p/19q-codeleted, grade 2
9451/3	Oligodendroglioma, IDH-mutant and 1p/19q-codeleted, grade 3
9440/3	Glioblastoma, IDH-wildtype

Paediatric-type diffuse low-grade gliomas

9421/1	Diffuse astrocytoma, <i>MYB</i> - or <i>MYBL1</i> -altered [†]
9431/1	Angiocentric glioma
9413/0	Polymorphous low-grade neuroepithelial tumour of the young [†]
9421/1	Diffuse low-grade glioma, MAPK pathway-altered [†]

Paediatric-type diffuse high-grade gliomas

9385/3	Diffuse midline glioma, H3 K27-altered [†]
9385/3	Diffuse hemispheric glioma, H3 G34-mutant [†]
9385/3	Diffuse paediatric-type high-grade glioma, H3-wildtype and IDH-wildtype [†]
9385/3	Infant-type hemispheric glioma [†]

Circumscribed astrocytic gliomas

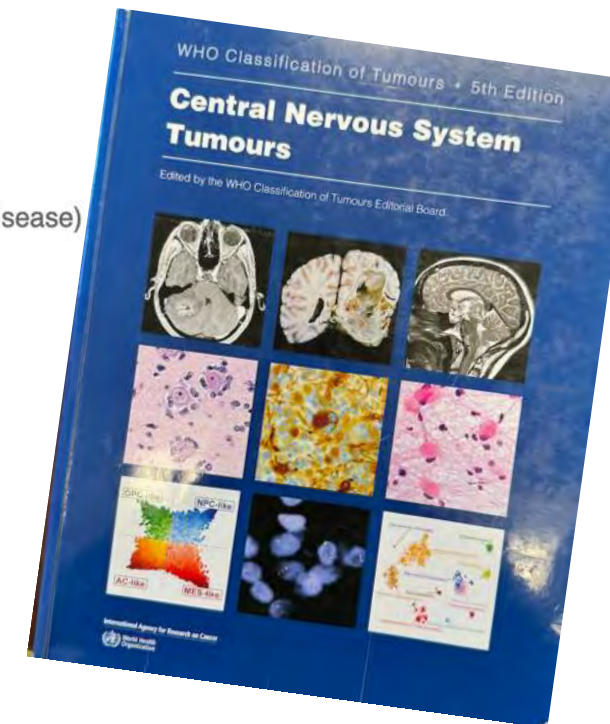
9421/1	Pilocytic astrocytoma
9421/3*	High-grade astrocytoma with piloid features
9424/3	Pleomorphic xanthoastrocytoma
9384/1	Subependymal giant cell astrocytoma
9444/1	Chordoid glioma
9430/3	Astroblastoma, <i>MN1</i> -altered [†]

Glioneuronal and neuronal tumours

9505/1	Ganglioglioma
9492/0	Gangliocytoma
9412/1	Desmoplastic infantile ganglioglioma
9412/1	Desmoplastic infantile astrocytoma
9413/0	Dysembryoplastic neuroepithelial tumour
n/a	Diffuse glioneuronal tumour with oligodendroglioma-like features and nuclear clusters (provisional entity)
9509/1	Papillary glioneuronal tumour
9509/1	Rosette-forming glioneuronal tumour
9509/1	Myxoid glioneuronal tumour [†]
9509/3*	Diffuse leptomeningeal glioneuronal tumour
9509/0*	Multinodular and vacuolating neuronal tumour
9493/0	Dysplastic cerebellar gangliocytoma (Lhermitte–Duclos disease)
9506/1	Central neurocytoma
9506/1	Extraventricular neurocytoma
9506/1	Cerebellar liponeurocytoma

Ependymal tumours

9391/3	Supratentorial ependymoma, NOS [†]
9396/3	Supratentorial ependymoma, <i>ZFTA</i> fusion-positive [†]
9396/3	Supratentorial ependymoma, <i>YAP1</i> fusion-positive [†]
9391/3	Posterior fossa ependymoma, NOS [†]
9396/3	Posterior fossa group A (PFA) ependymoma [†]
9396/3	Posterior fossa group B (PFB) ependymoma [†]
9391/3	Spinal ependymoma, NOS [†]
9396/3	Spinal ependymoma, <i>MYCN</i> -amplified [†]
9394/1	Myxopapillary ependymoma
9383/1	Subependymoma



ICD-O coding of central nervous system tumours



ICD-O-3.2 ICD-O label (subtypes are indicated in grey text, with the label indented);

Please note that the WHO classification of tumour types is more readily reflected in the table of contents

Choroid plexus tumours

9390/0	Choroid plexus papilloma
9390/1	Atypical choroid plexus papilloma
9390/3	Choroid plexus carcinoma

Embryonal tumours

Medulloblastomas, molecularly defined

9475/3	Medulloblastoma, WNT-activated
9471/3	Medulloblastoma, SHH-activated and <i>TP53</i> -wildtype
9476/3	Medulloblastoma, SHH-activated and <i>TP53</i> -mutant
9477/3	Medulloblastoma, non-WNT/non-SHH

Medulloblastomas, histologically defined

9470/3	Medulloblastoma, histologically defined
9471/3	Desmoplastic nodular medulloblastoma
9471/3	Medulloblastoma with extensive nodularity
9474/3	Large cell medulloblastoma
9474/3	Anaplastic medulloblastoma

Other CNS embryonal tumours

9508/3	Atypical teratoid/rhabdoid tumour
n/a	Cribriform neuroepithelial tumour (provisional entity)
9478/3	Embryonal tumour with multilayered rosettes
9500/3	CNS neuroblastoma, <i>FOXR2</i> -activated [†]
9500/3	CNS tumour with <i>BCOR</i> internal tandem duplication [†]
9473/3	CNS embryonal tumour, NEC/NOS

Pineal tumours

9361/1	Pineocytoma
9362/3	Pineal parenchymal tumour of intermediate differentiation
9362/3	Pineoblastoma
9395/3	Papillary tumour of the pineal region
n/a	Desmoplastic myxoid tumour of the pineal region, <i>SMARCB1</i> -mutant (provisional entity)

Cranial and paraspinal nerve tumours

9560/0	Schwannoma
9540/0	Neurofibroma
9550/0	Plexiform neurofibroma
9571/0	Perineurioma
9563/0	Hybrid nerve sheath tumour
9540/3	Malignant melanotic nerve sheath tumour
9540/3	Malignant peripheral nerve sheath tumour
8693/3	Cauda equina neuroendocrine tumour (previously paraganglioma)

Meningioma

9530/0	Meningioma
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Mesenchymal, non-meningothelial tumours involving the CNS

Fibroblastic and myofibroblastic tumours

8815/1	Solitary fibrous tumour
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Vascular tumours

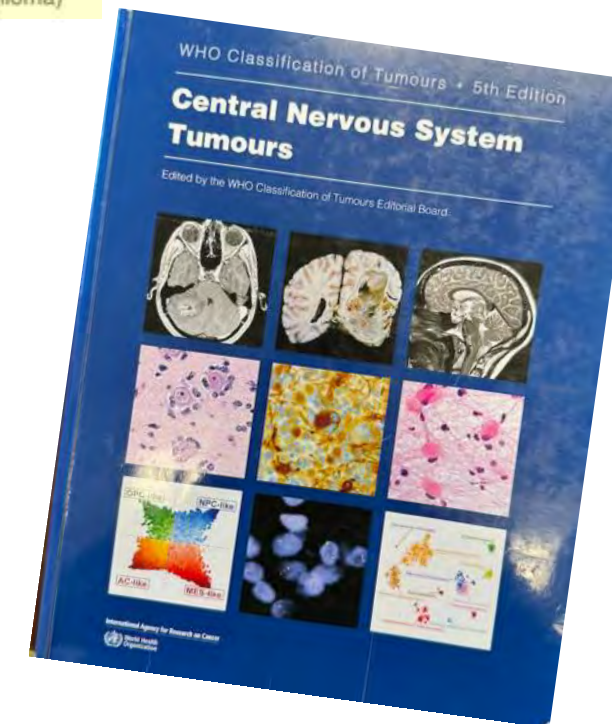
9121/0	Cavernous haemangioma
9131/0	Capillary haemangioma
9123/0	Arteriovenous malformation
9161/1	Haemangioblastoma

Skeletal muscle tumours

8910/3	Embryonal rhabdomyosarcoma
8920/3	Alveolar rhabdomyosarcoma
8901/3	Rhabdomyosarcoma, pleomorphic-type
8912/3	Spindle cell rhabdomyosarcoma

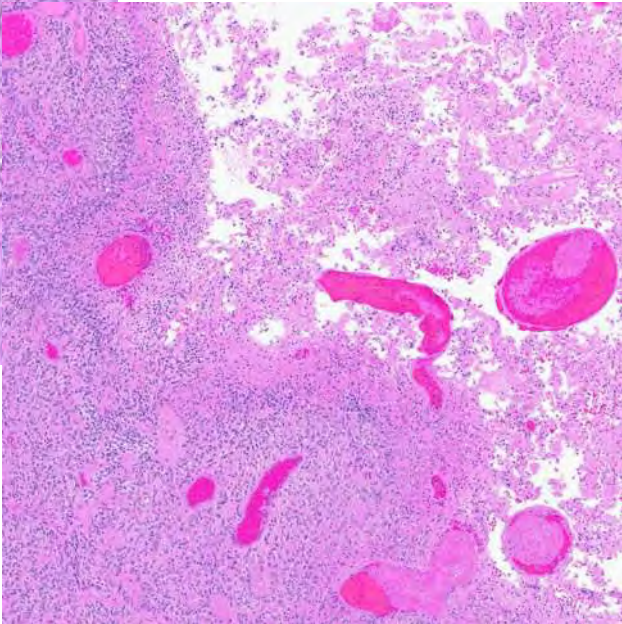
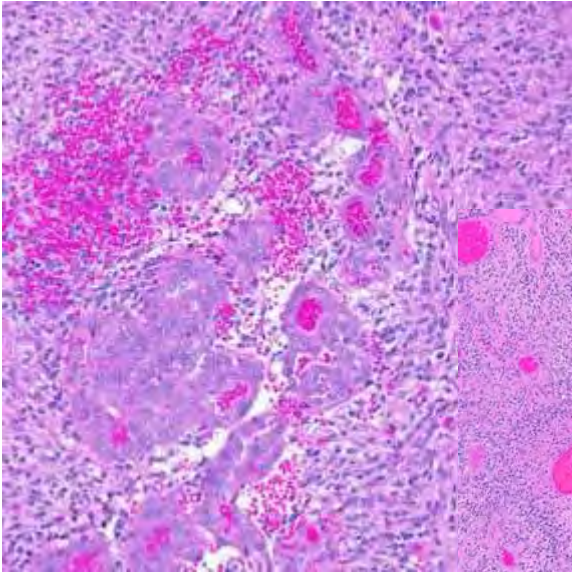
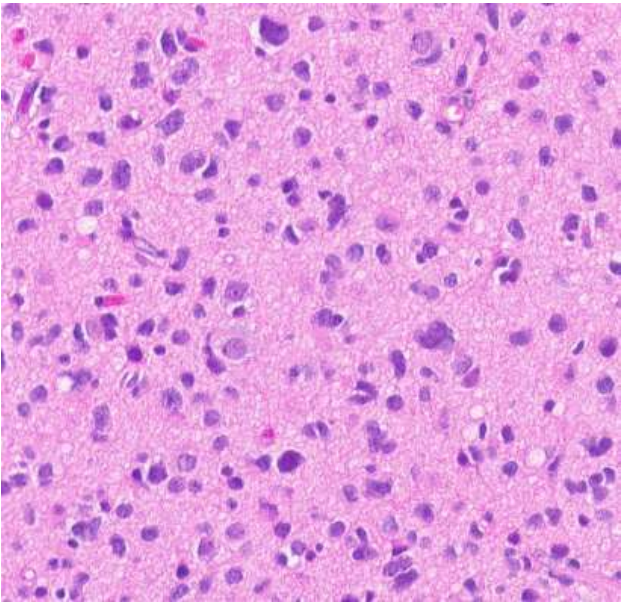
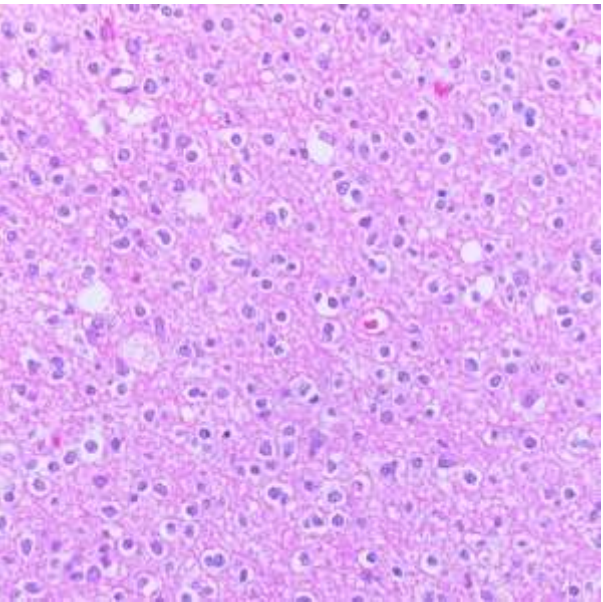
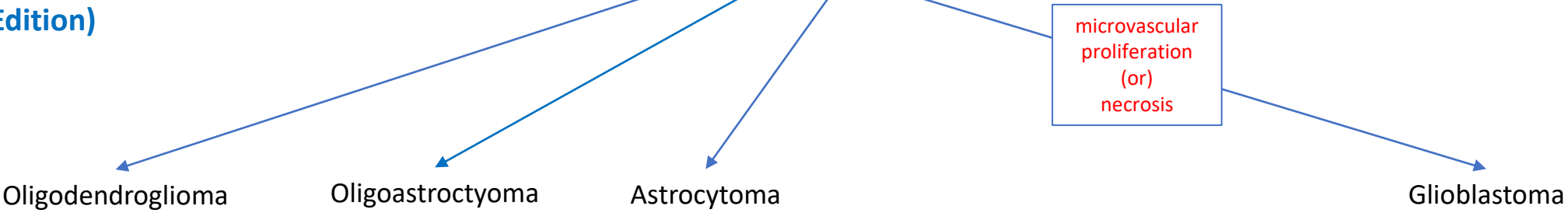
Tumours of uncertain differentiation

n/a	Intracranial mesenchymal tumour, FET:: <i>CREB</i> fusion-positive (provisional entity)
9367/3	<i>CIC</i> -rearranged sarcoma
9480/3	Primary intracranial sarcoma, <i>DICER1</i> -mutant [†]
9364/3	Ewing sarcoma



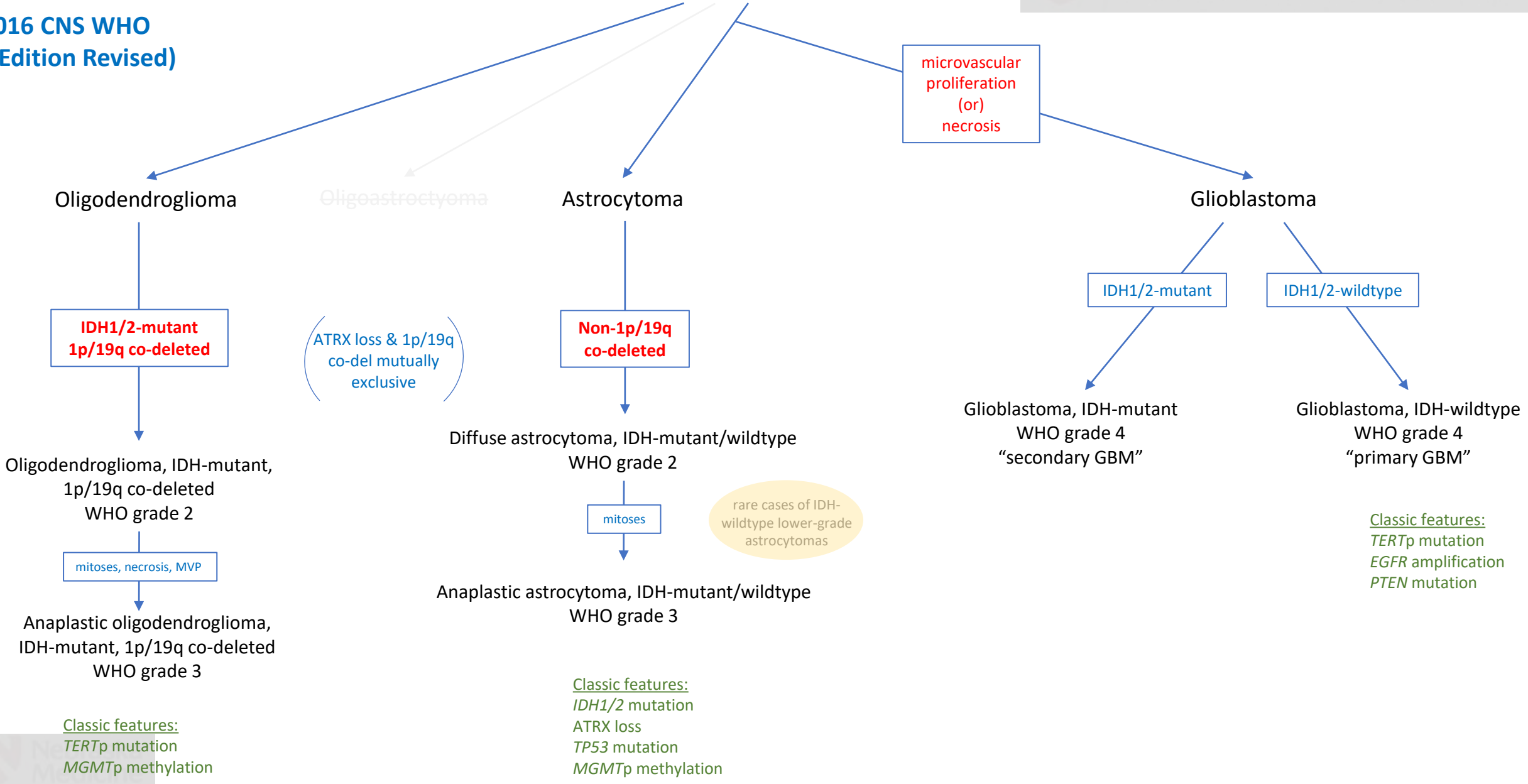
Adult Infiltrating Gliomas

2007 CNS WHO
(4th Edition)



Adult Infiltrating Gliomas

2016 CNS WHO
(4th Edition Revised)



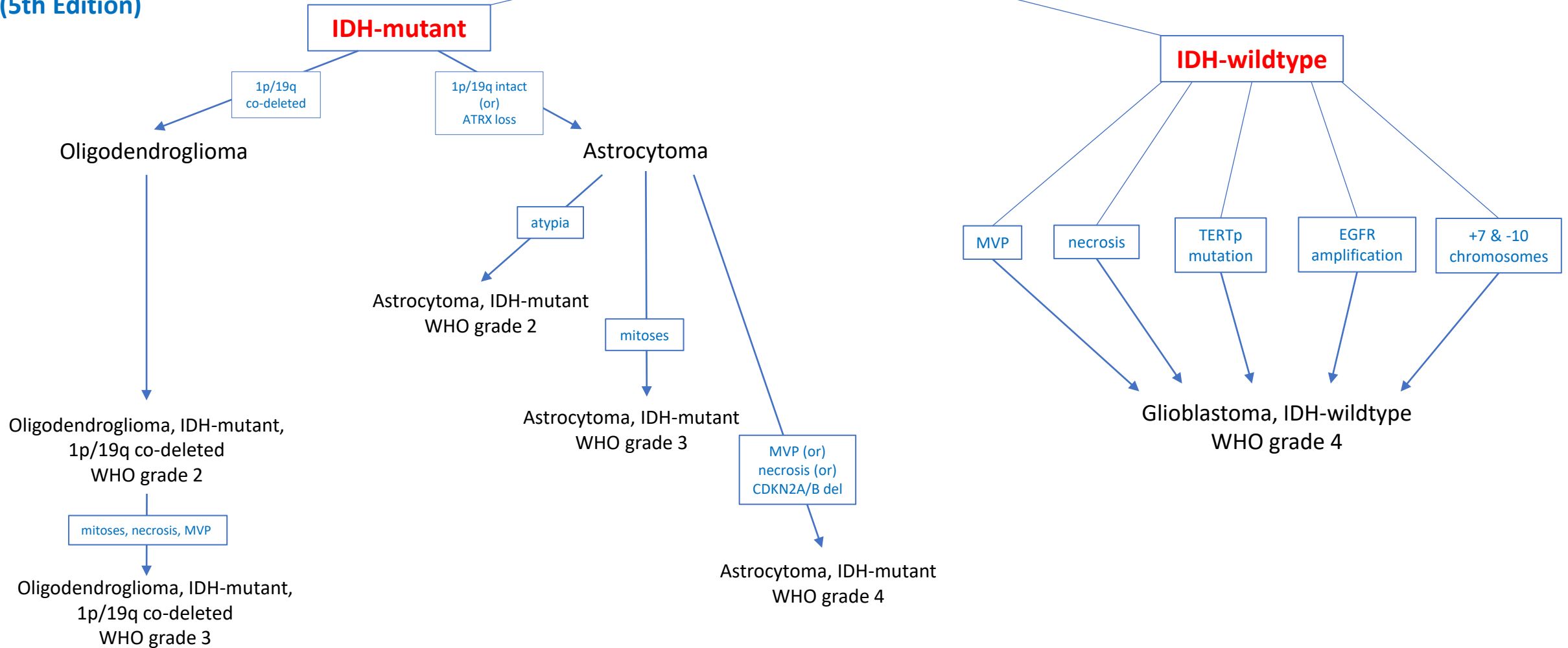
Classic features:
*TERT*p mutation
*MGMT*p methylation

Classic features:
IDH1/2 mutation
 ATRX loss
TP53 mutation
*MGMT*p methylation

Classic features:
*TERT*p mutation
EGFR amplification
PTEN mutation

Adult Infiltrating Gliomas

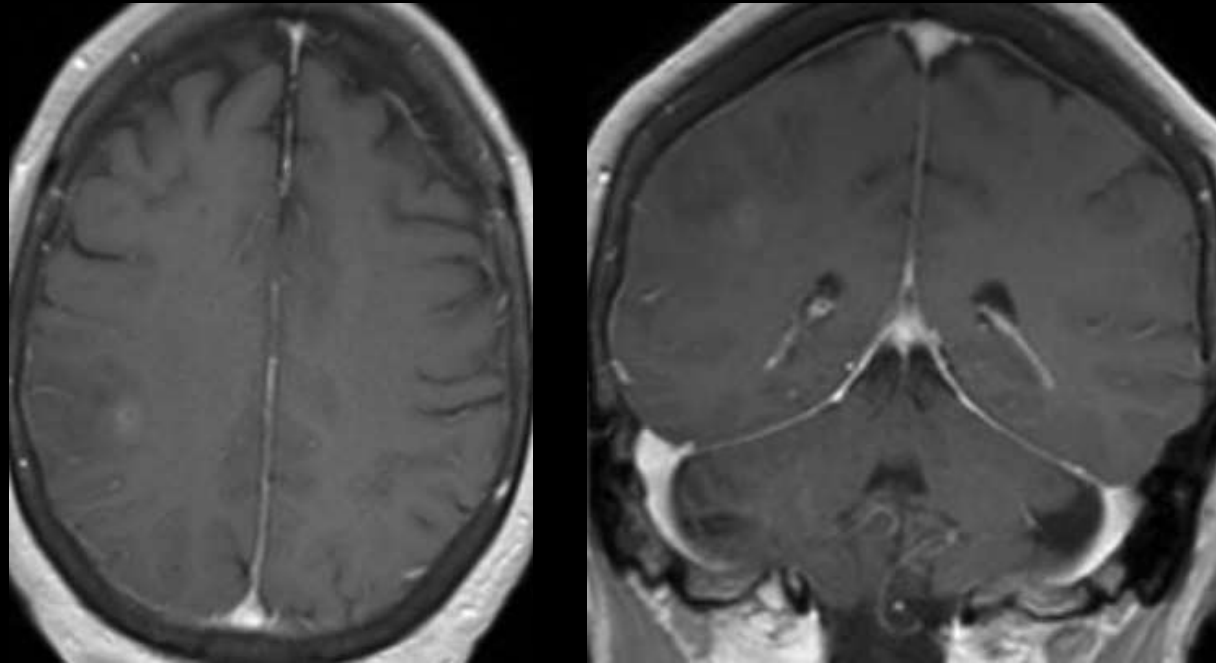
2021 CNS WHO (5th Edition)



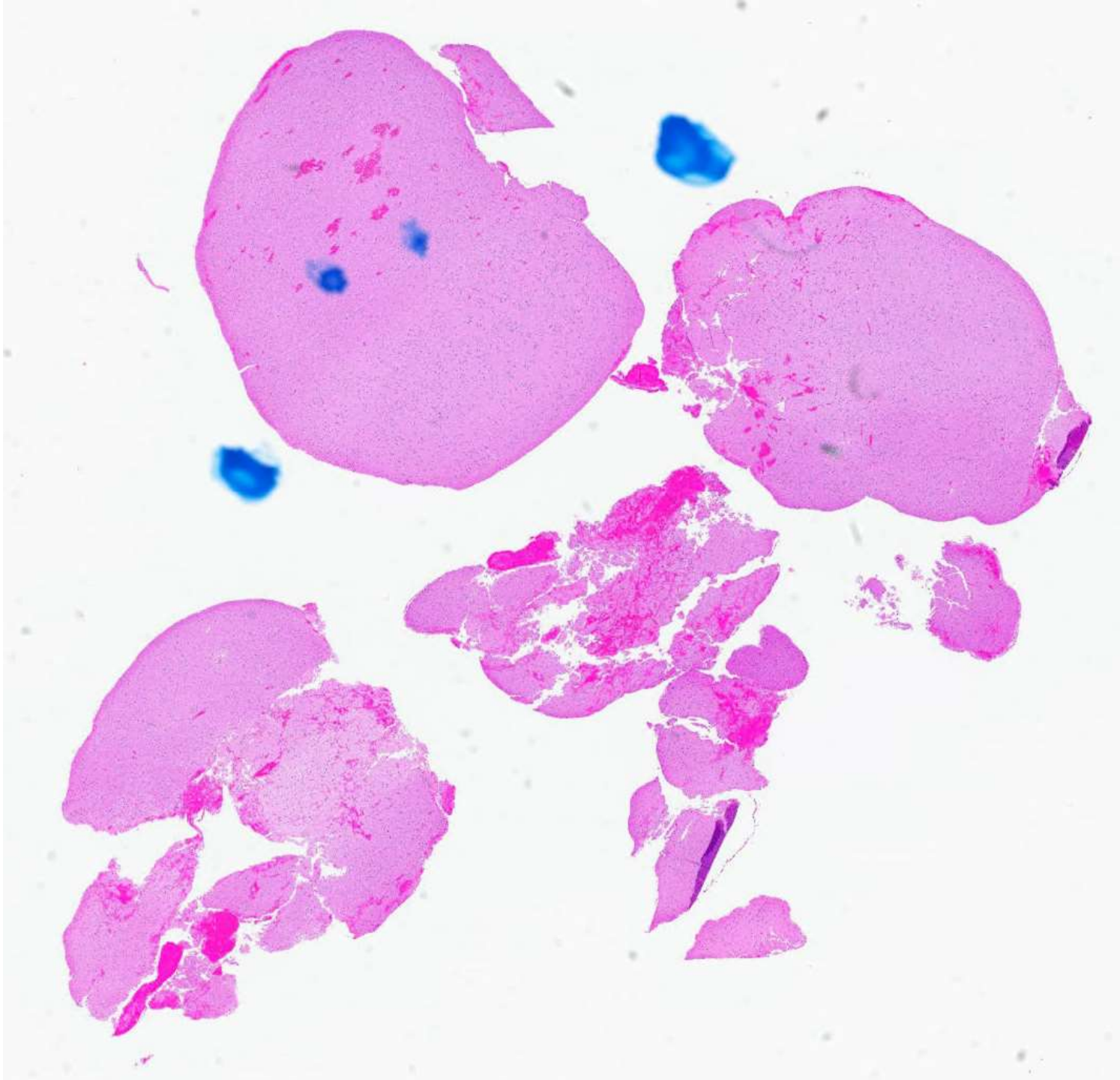
Classic features:
*TERT*p mutation
*MGMT*p methylation

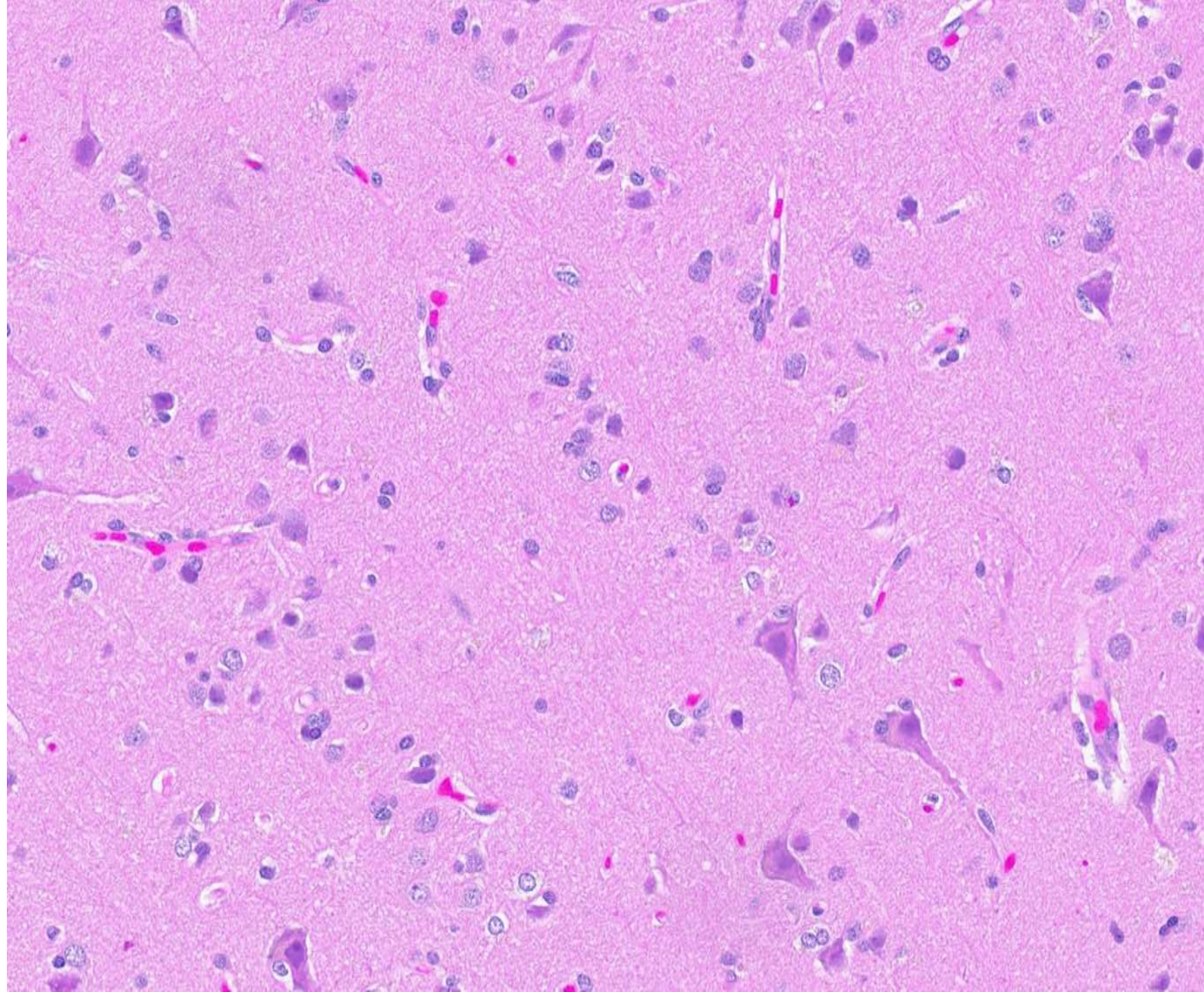


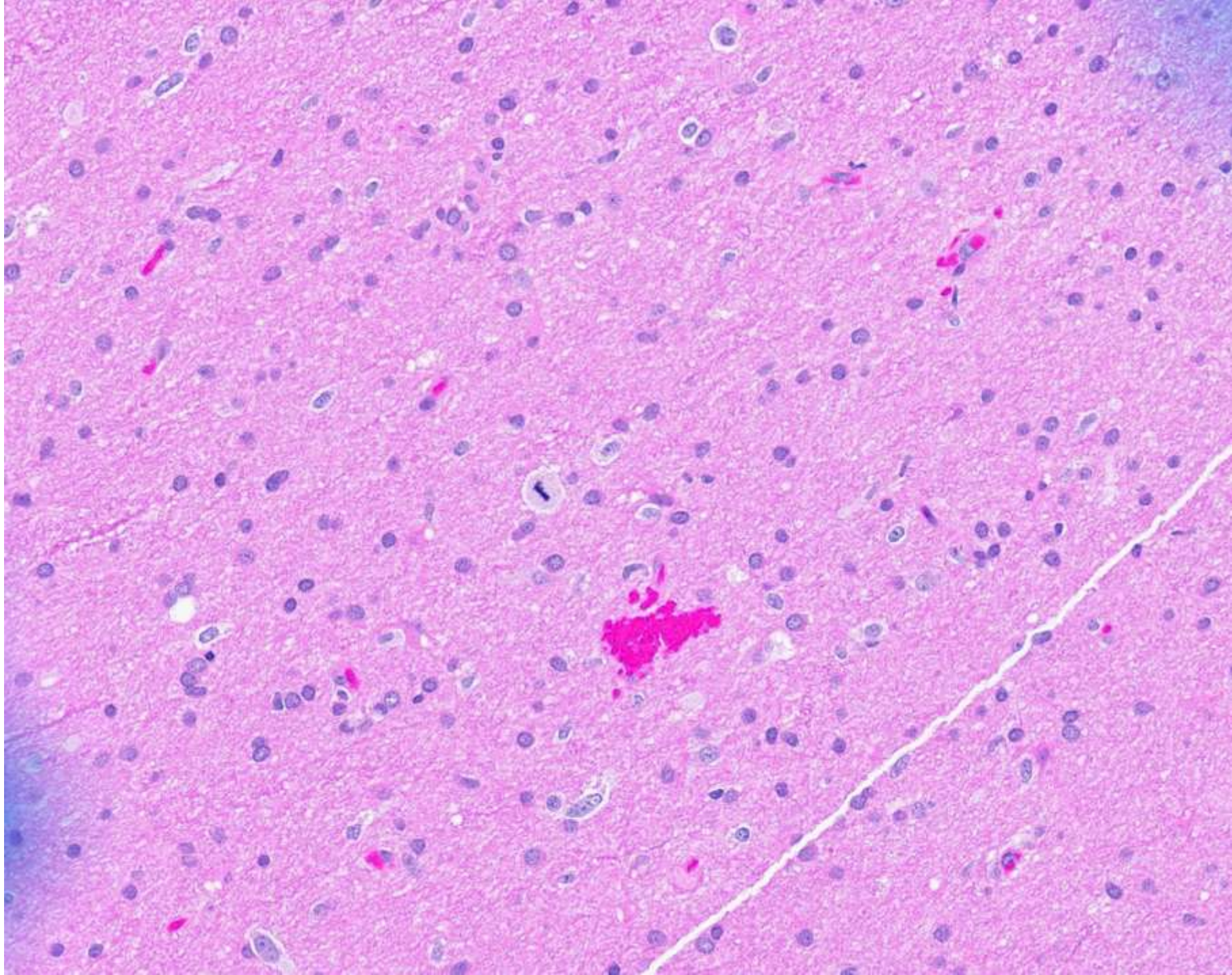
74F admitted to OSH with possible amphetamine overdose and was found to have a brain lesion



T1 +C



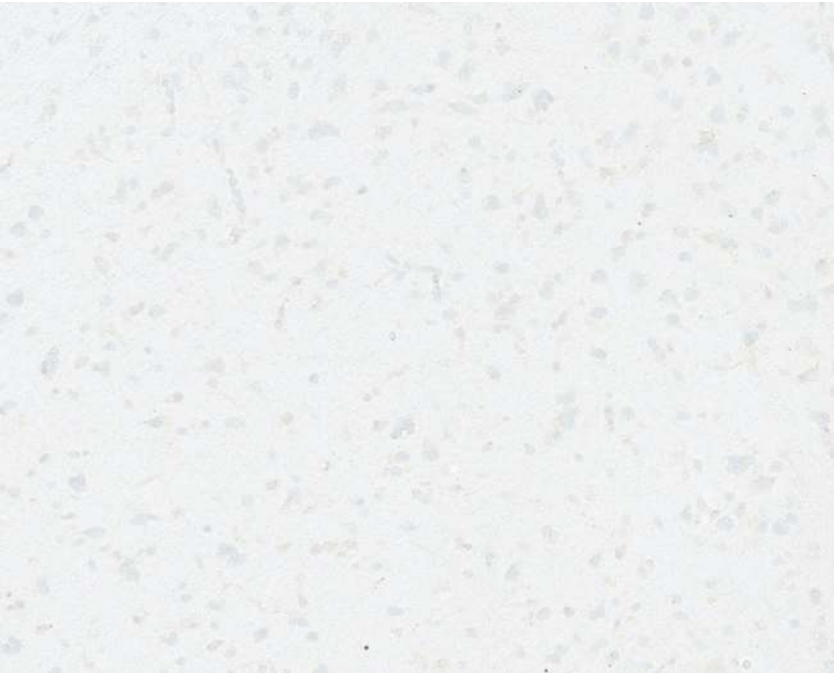




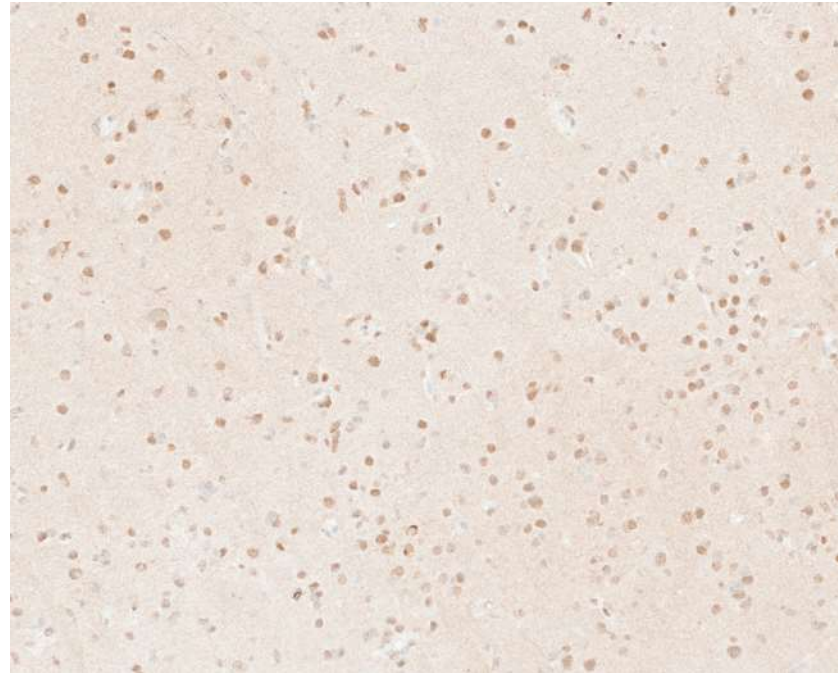
Ki67

CANCER CENTER





IDH1 p.R132H

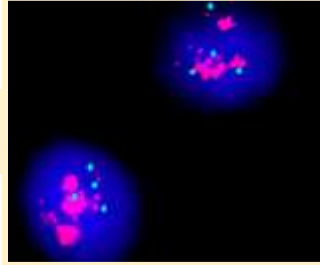


ATRX



p53

Fluorescent In Situ Hybridization (*EGFR*, *PTEN*):



EGFR (7p11.2) FISH Analysis Final Report

AMPLIFICATION of *EGFR* (7p11.2) with five copies of chromosome 7 centromere in 45% of cells.

PTEN (10q23.32) FISH Analysis Final Report

ABNORMAL and consistent with monosomy 10 in 54% of cells.

Brain, right parietal lesion, stereotactic biopsy:

Glioblastoma, IDH-wildtype
CNS WHO grade 4

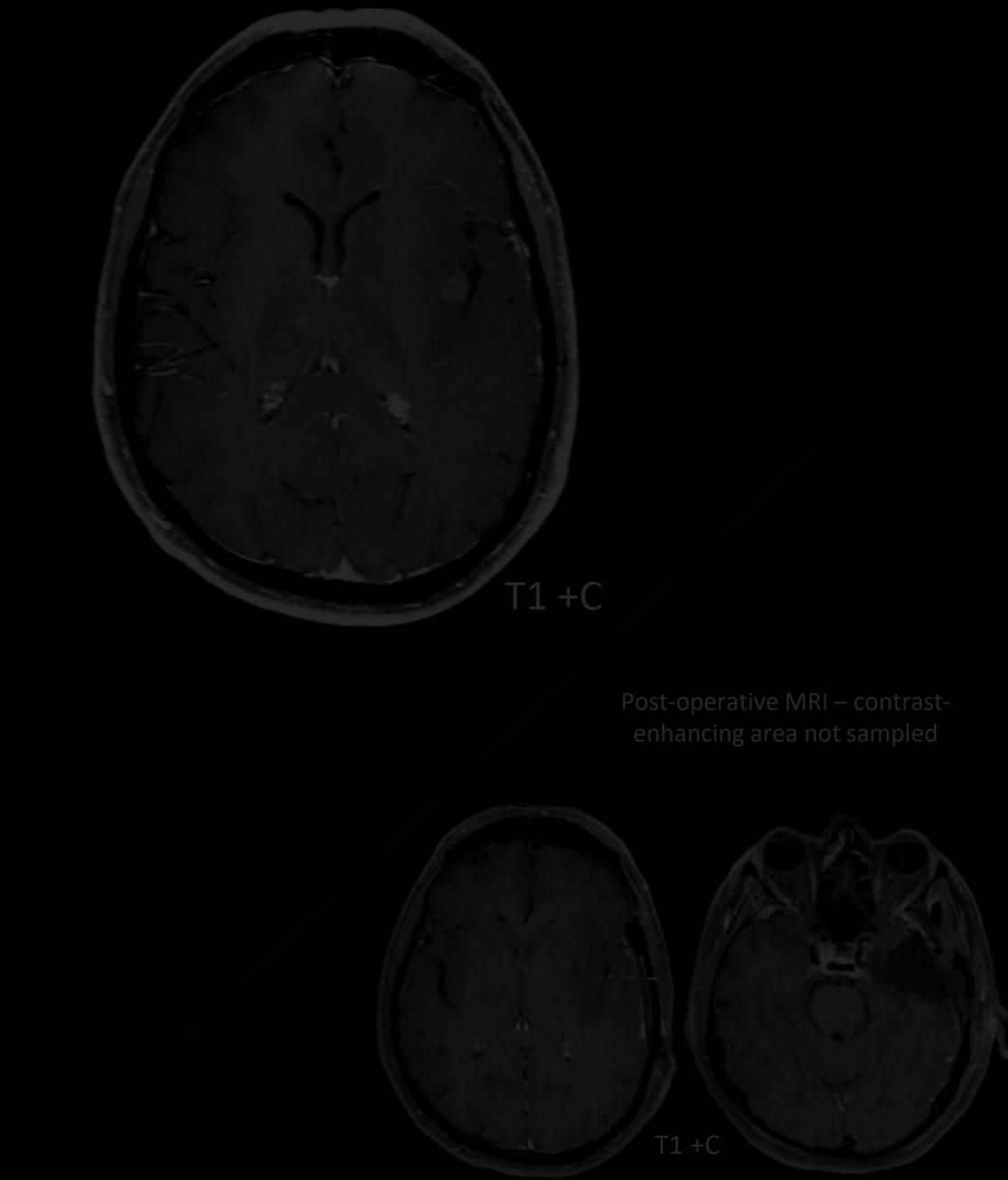
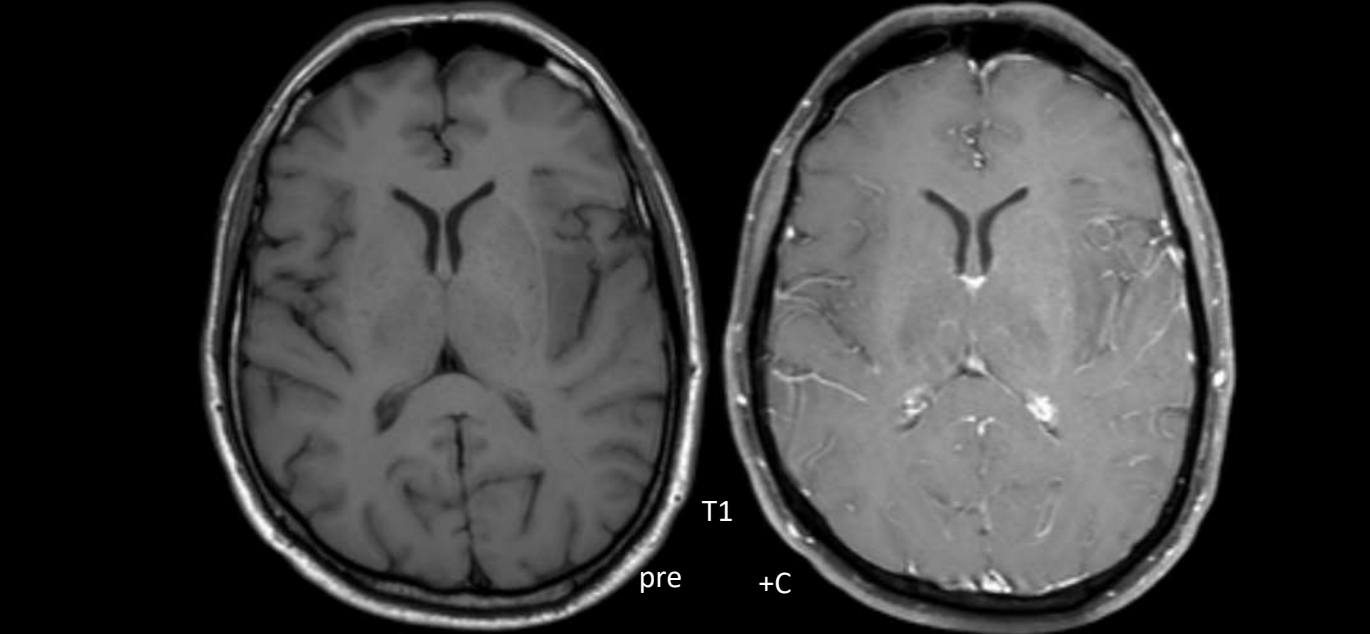
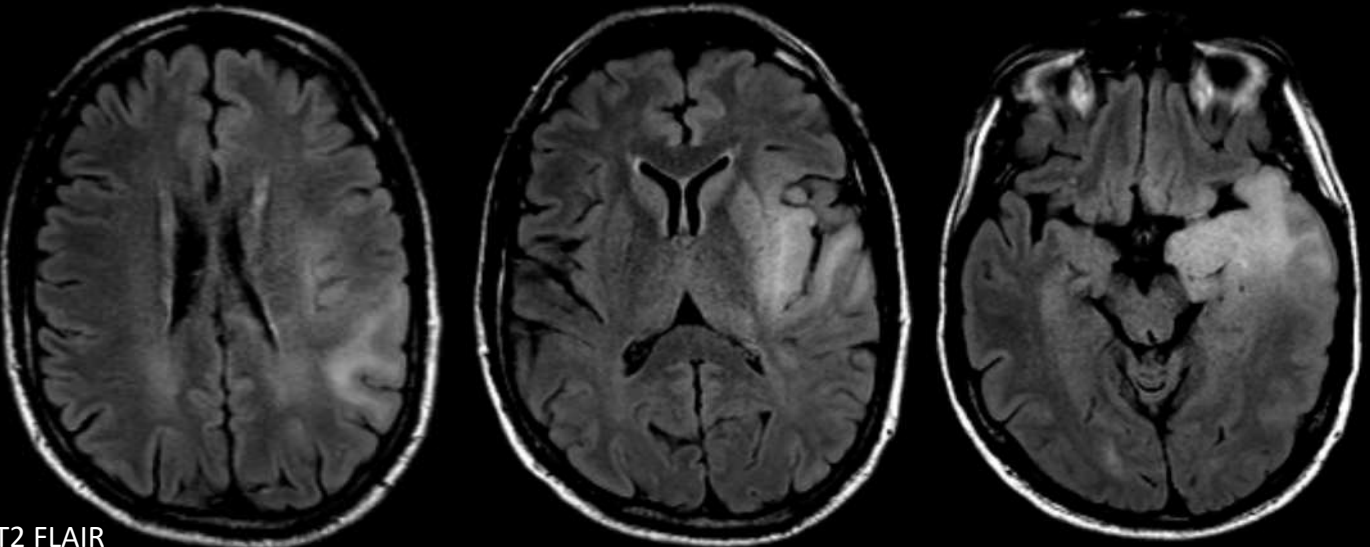
IDH1 (IHC): Negative for p.R132H mutant protein expression

EGFR/Ch7 (FISH): POSITIVE for *EGFR* amplification; POSITIVE for Ch7 gain (45%)

PTEN/Ch10 (FISH): Negative for *PTEN* deletion; POSITIVE for Ch10 monosomy (54%)

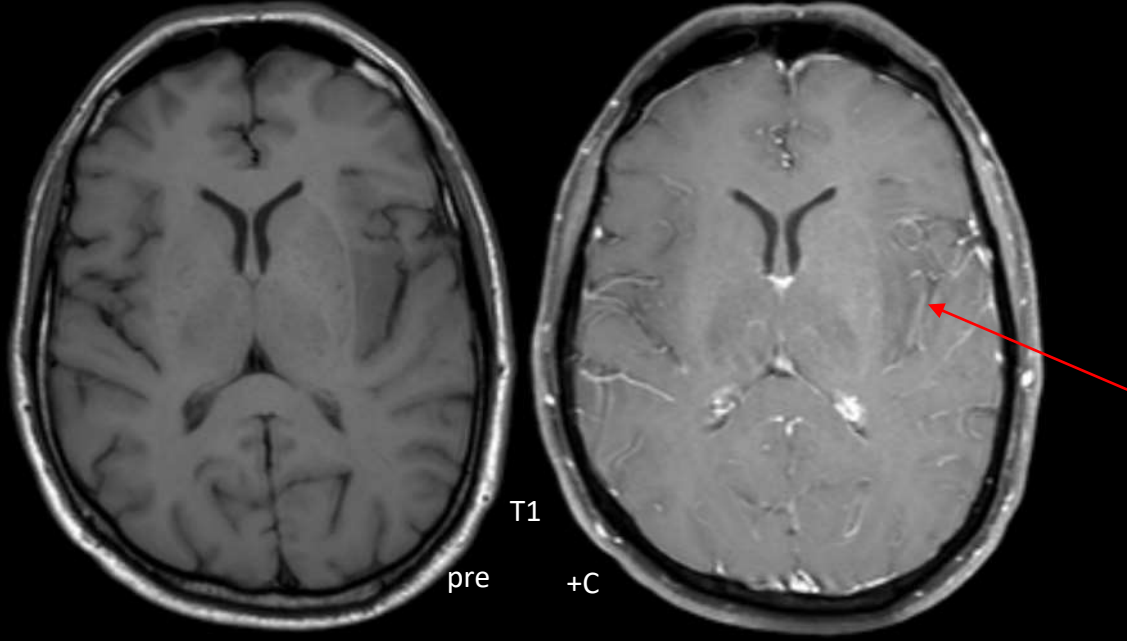
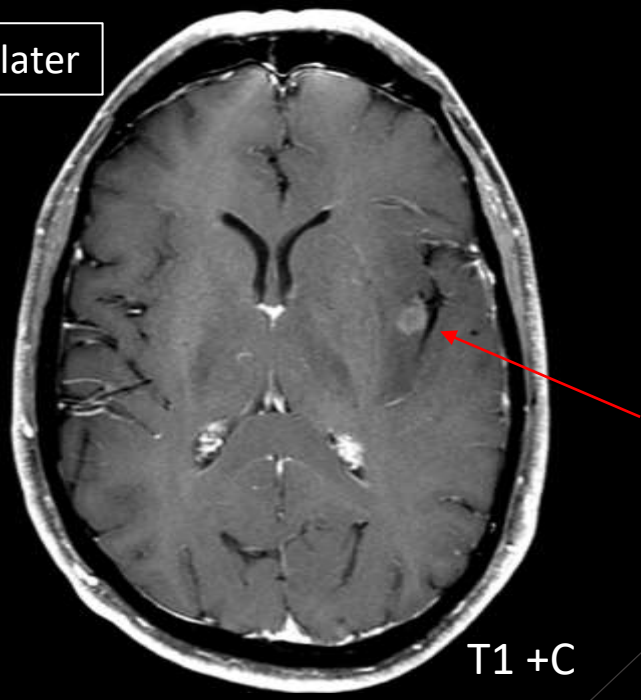
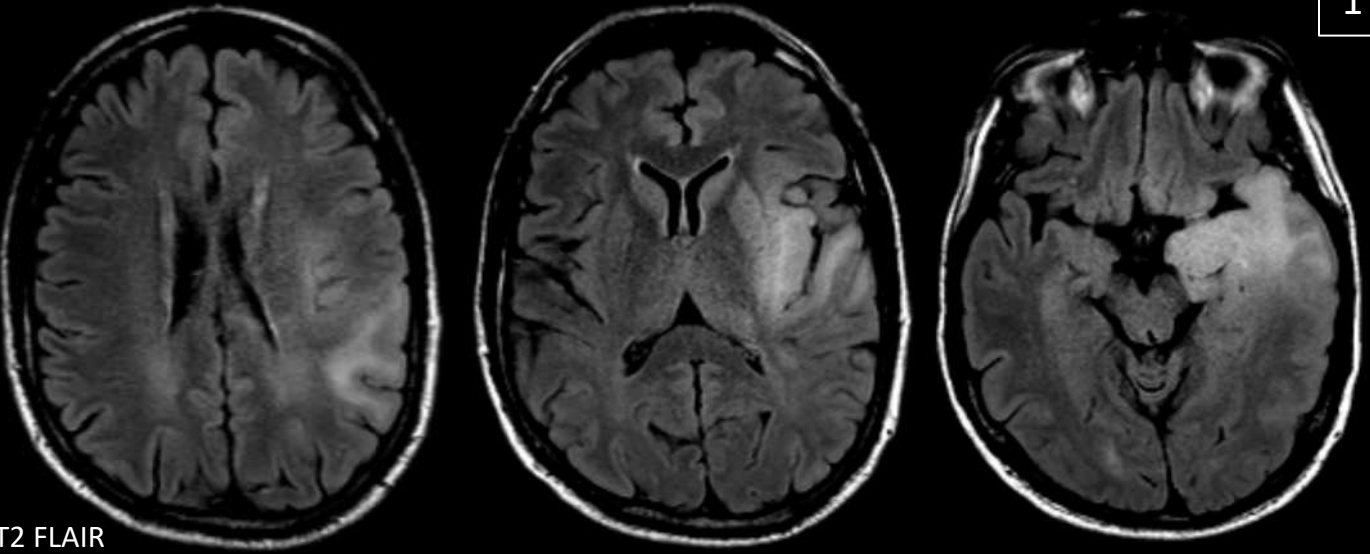
MGMT (methyl-PCR): POSITIVE for promoter hypermethylation

61M who presented with 3-month history of hot flashes, speech difficulties, and forgetfulness

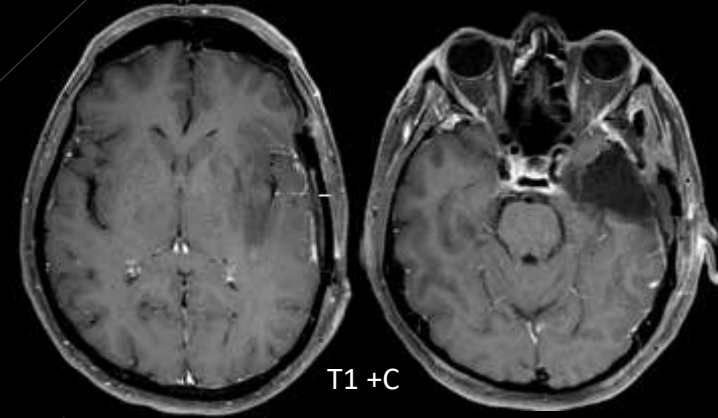


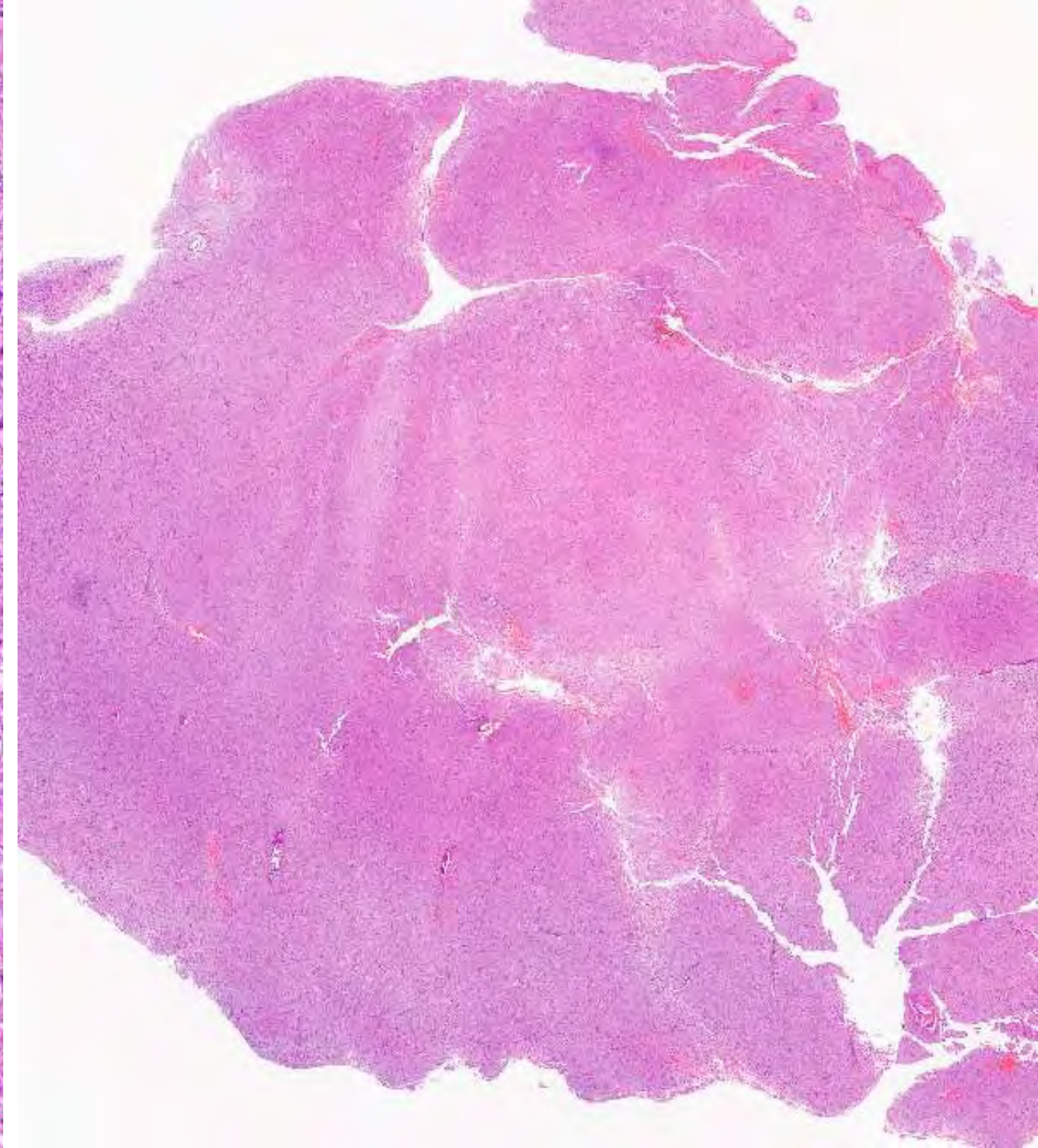
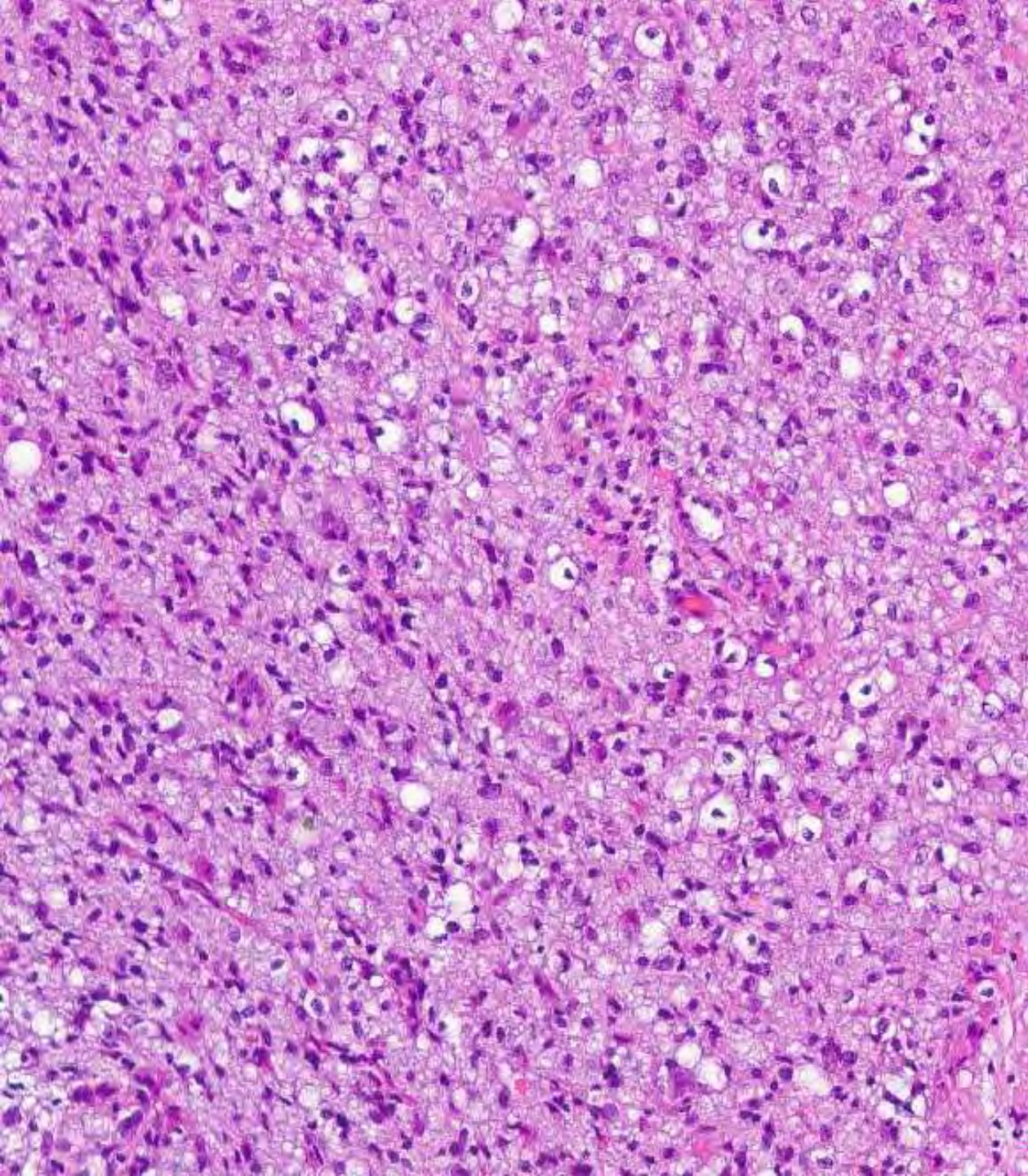
61M who presented with 3-month history of hot flashes, speech difficulties, and forgetfulness

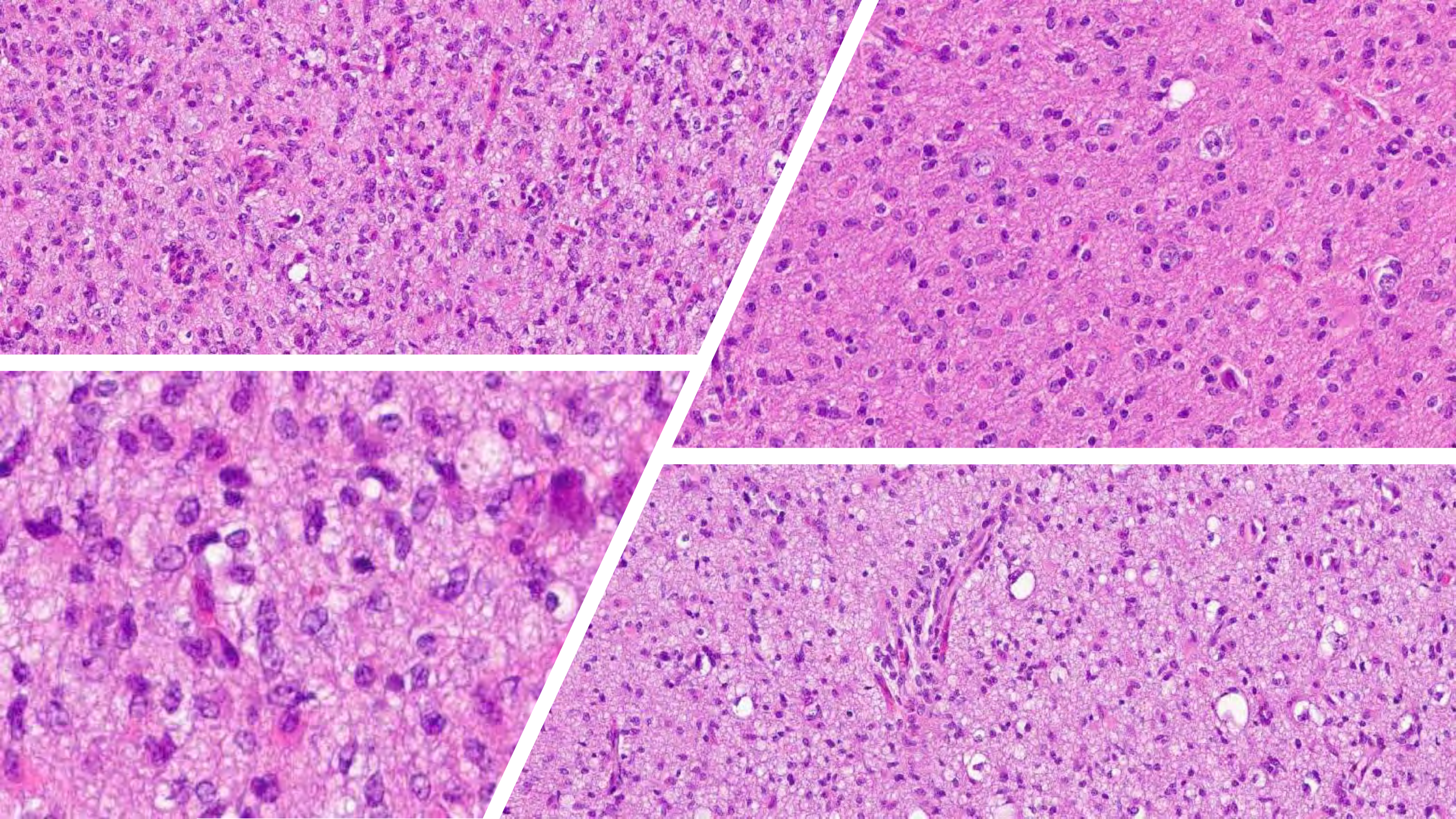
1 month later



Post-operative MRI – contrast-enhancing area not sampled







IDH1 R132H

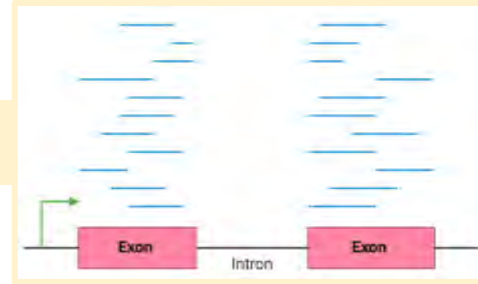
ATRX

p53

Ki67
up to 14%

Mitoses – up to 4 per 10 HPF

Next-Generation Sequencing Panel:



Copy Number Variations

<i>CDK4</i>	Amplification	12q14.1
<i>EGFR</i>	Amplification	7p11.2
<i>MDM4</i>	Amplification	1q32.1

Somatic Mutations

<i>EGFR</i>	p.R108K	(pathogenic)
<i>TERT</i>	c.-124C>T	(pathogenic)

Brain, left temporal lobe, awake craniotomy with resection:

Glioblastoma, IDH-wildtype
CNS WHO grade 4

IDH1/2 (NGS): Negative for mutations

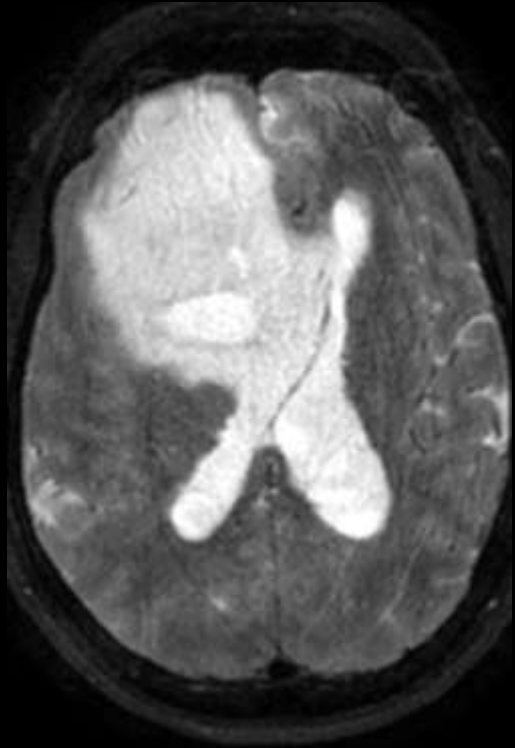
TERT (NGS): POSITIVE for c.-124C>T mutation

EGFR (NGS): POSITIVE for amplification and p.R108K mutation

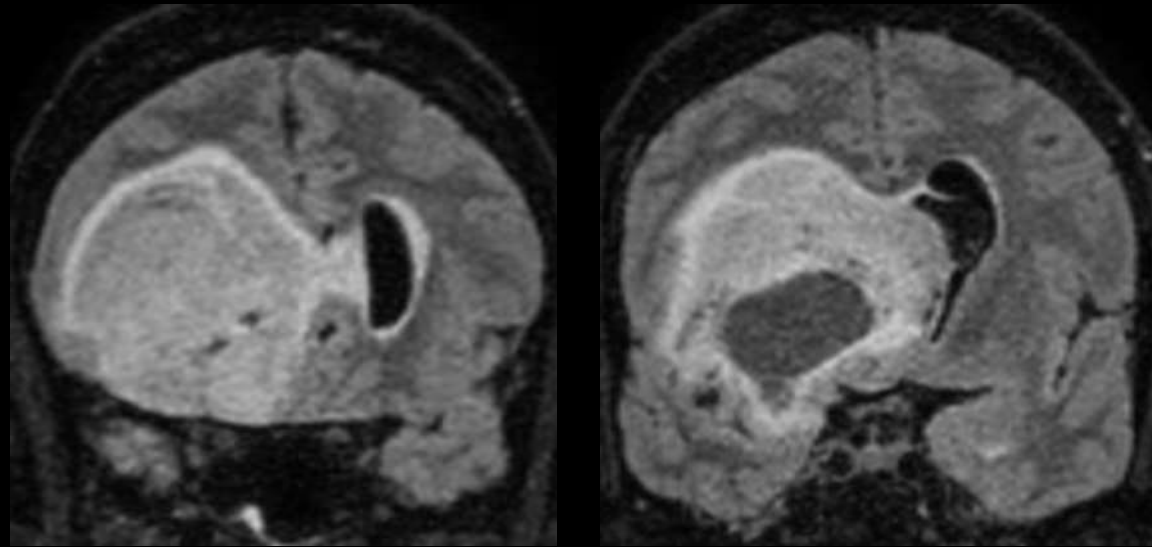
CDK4 (NGS): Positive for amplification

MDM4 (NGS): Positive for amplification

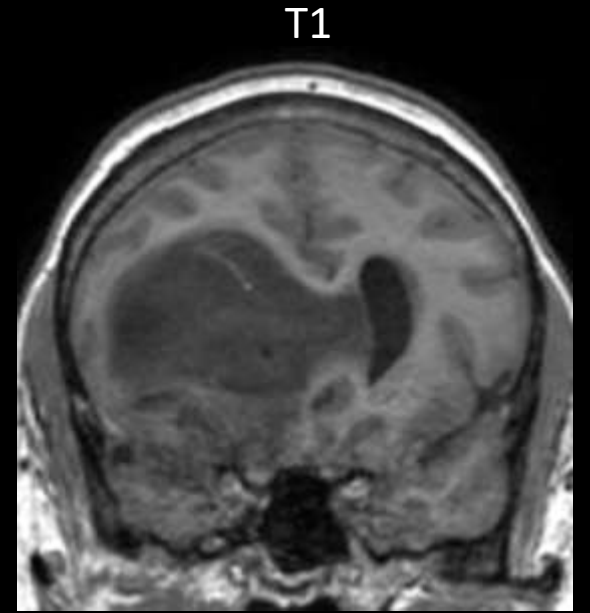
59F who presented with altered mental status over the past few days



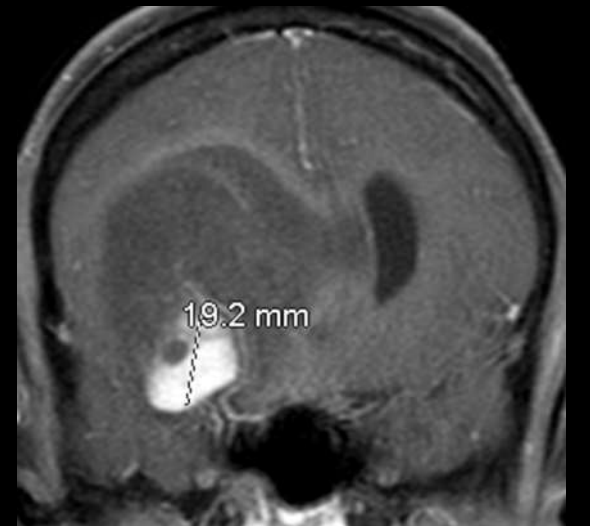
T2



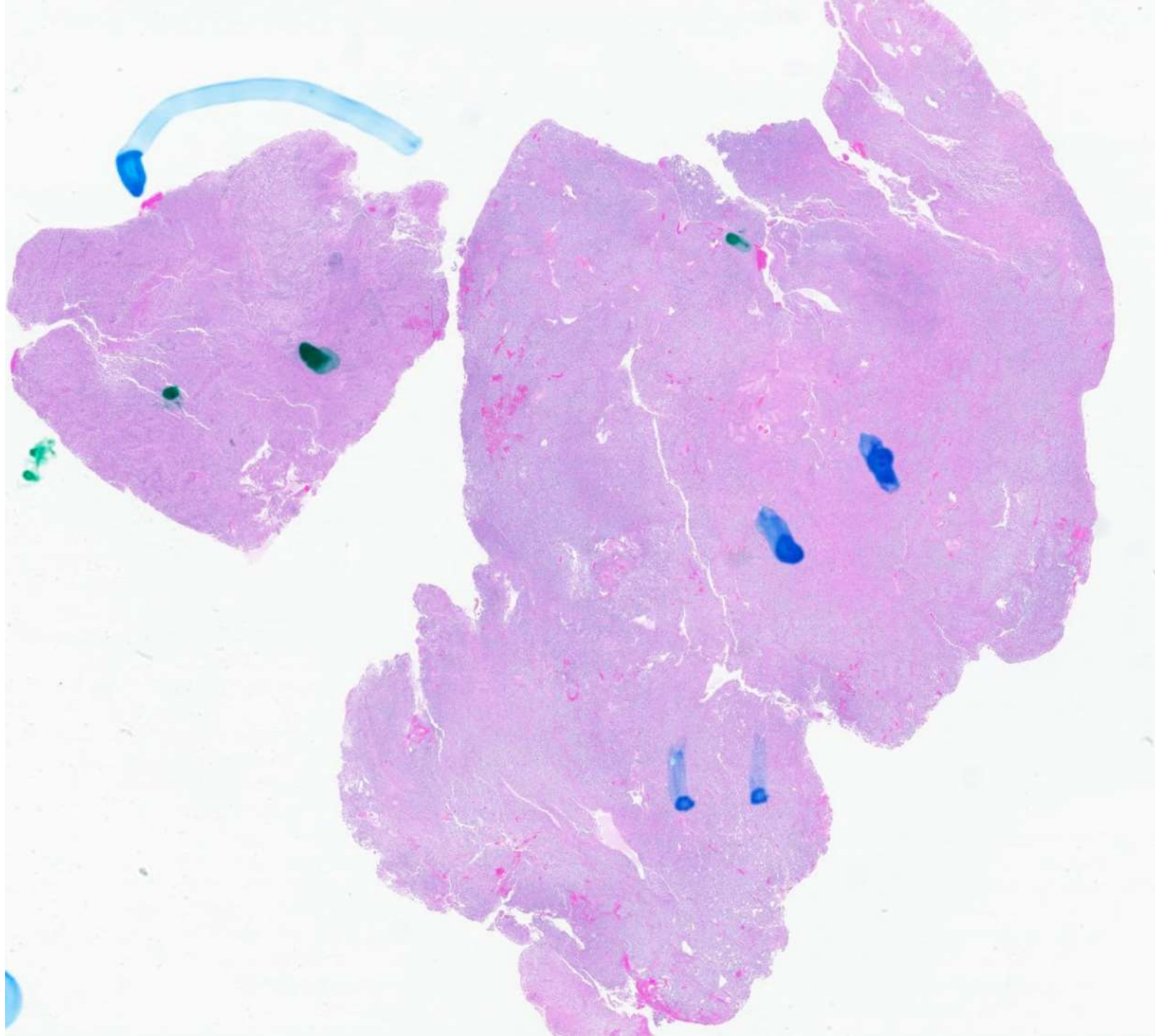
T2 FLAIR

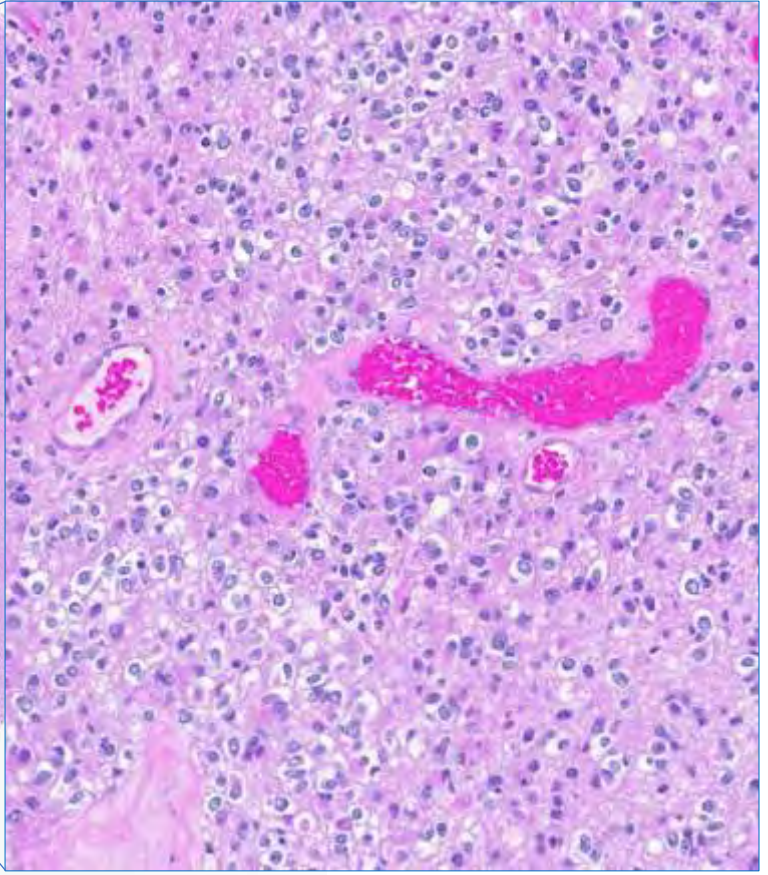
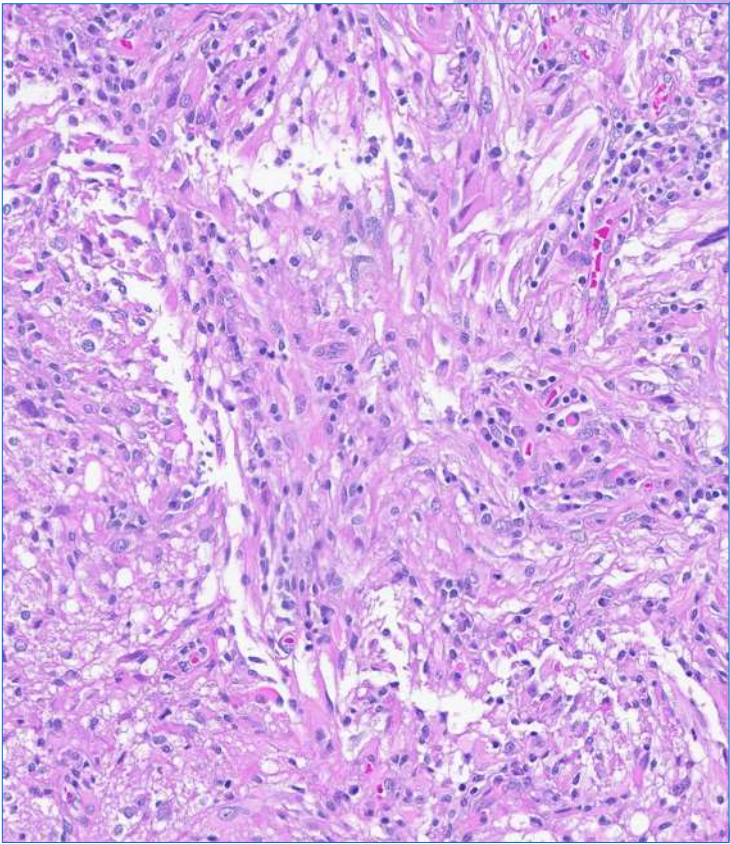


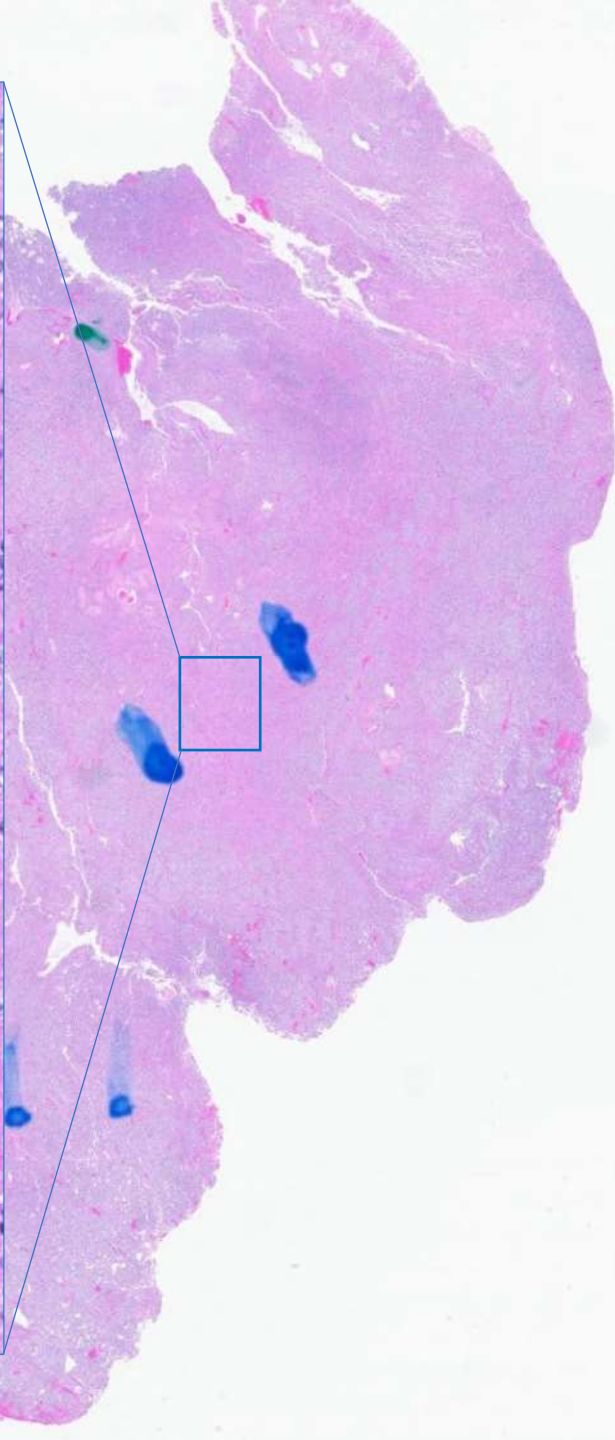
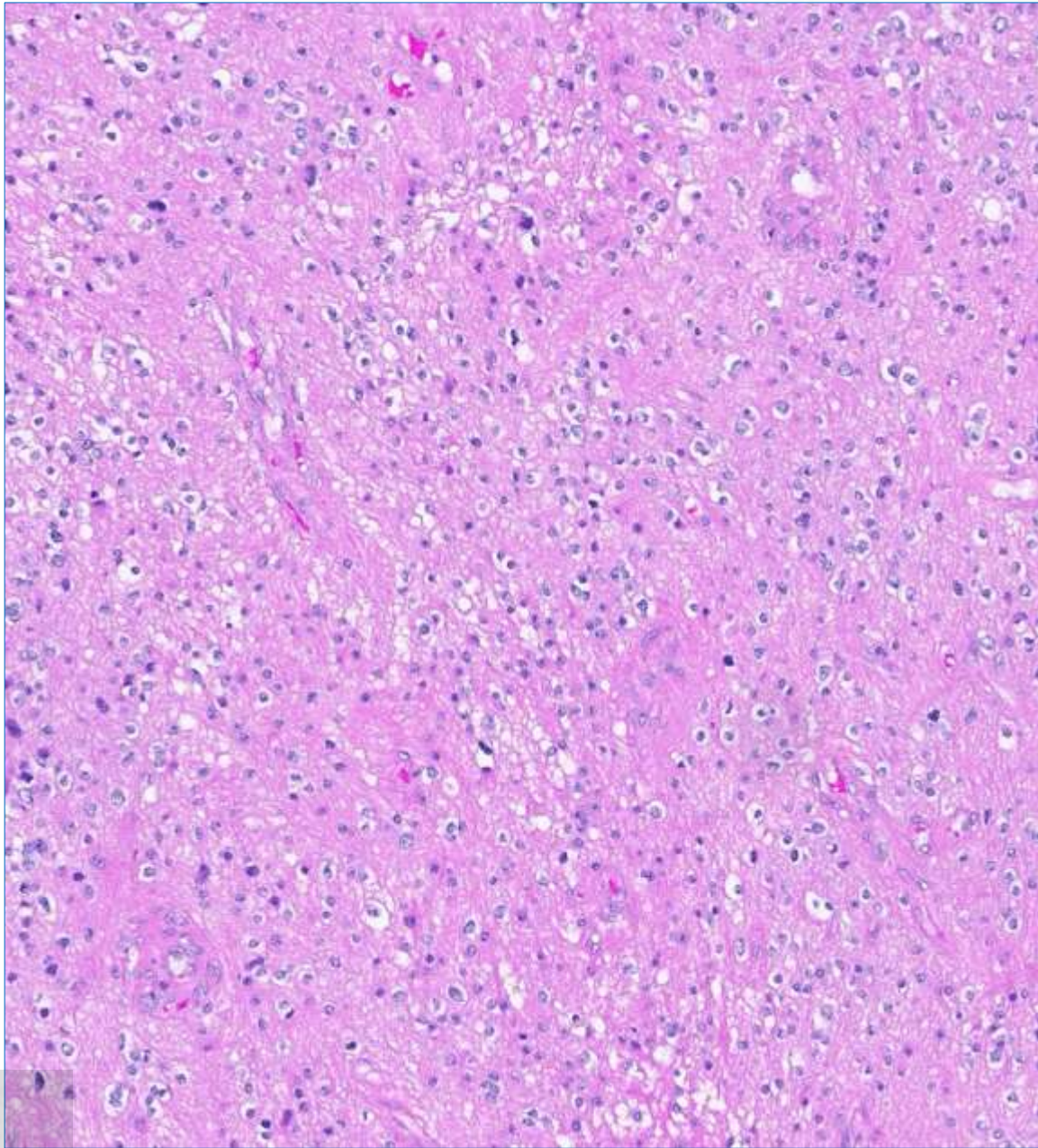
T1

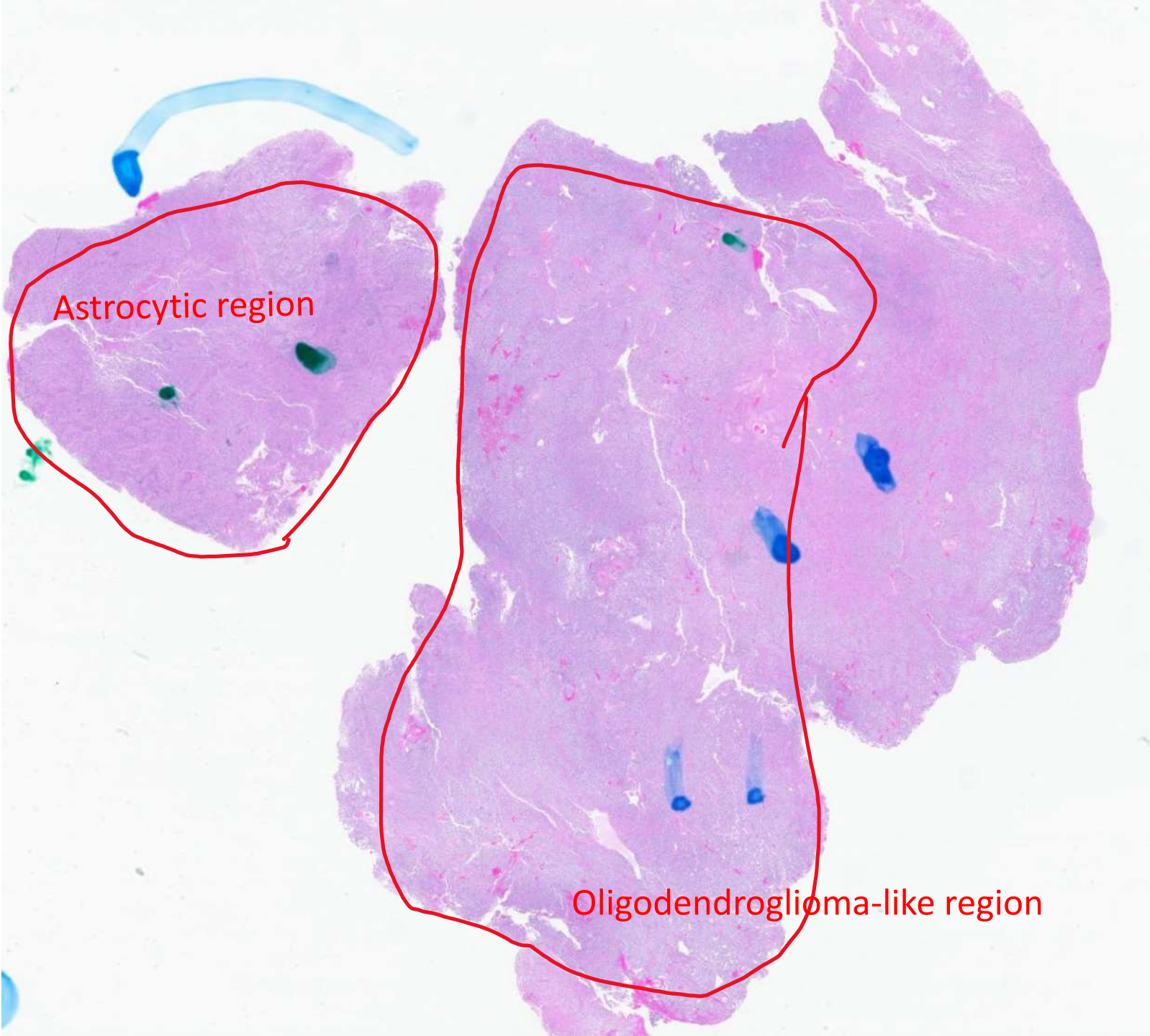


T1 +C



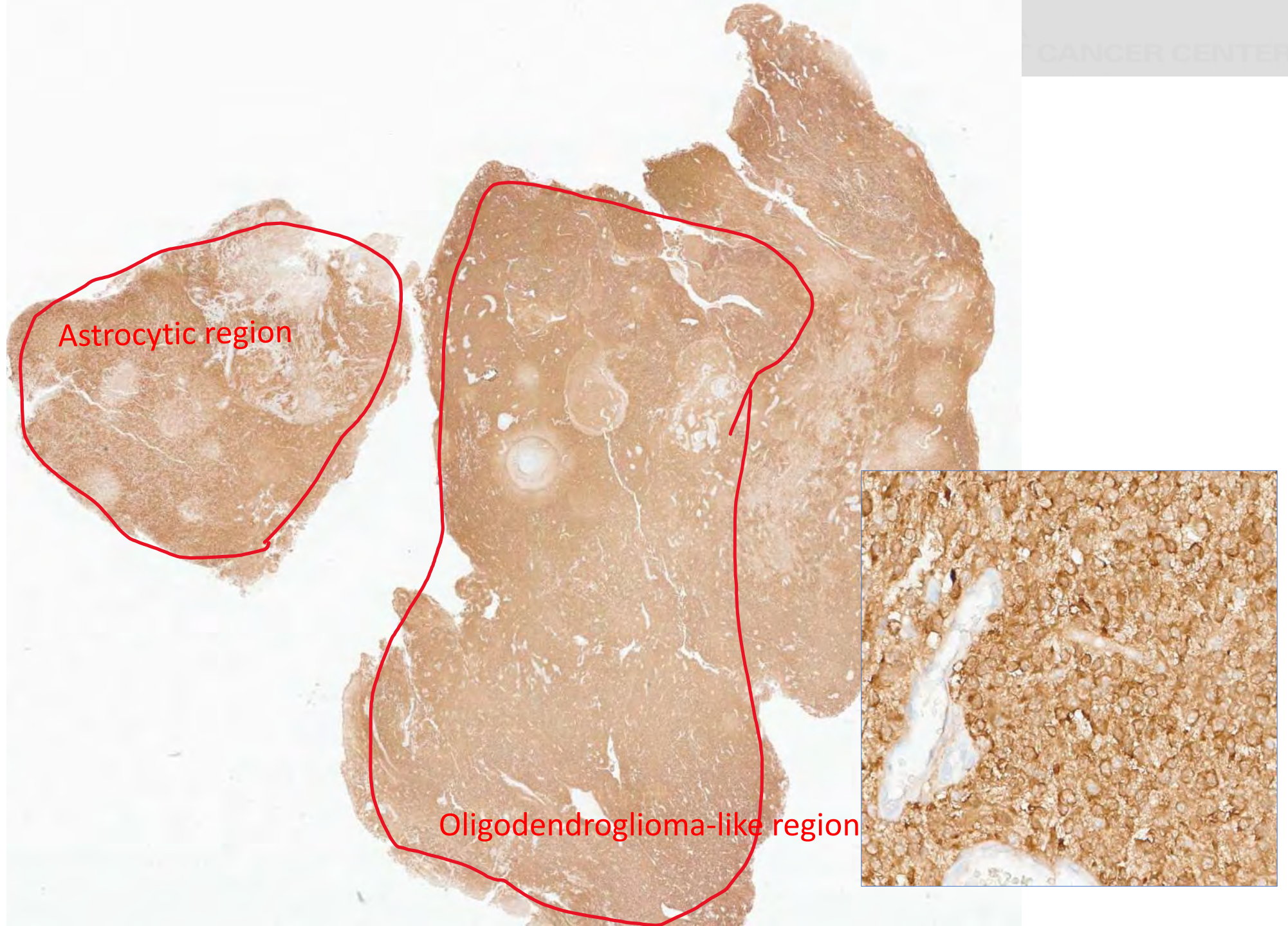






Astrocytic region

Oligodendroglioma-like region



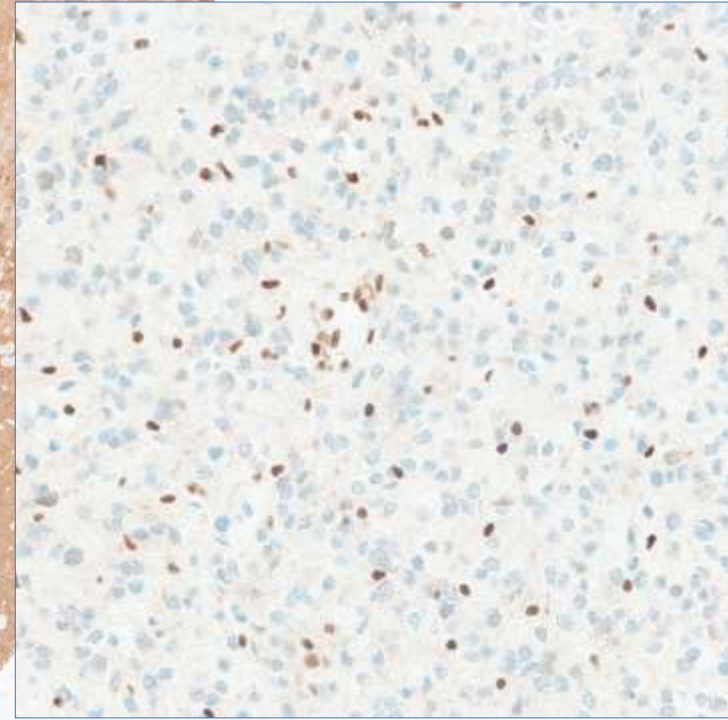
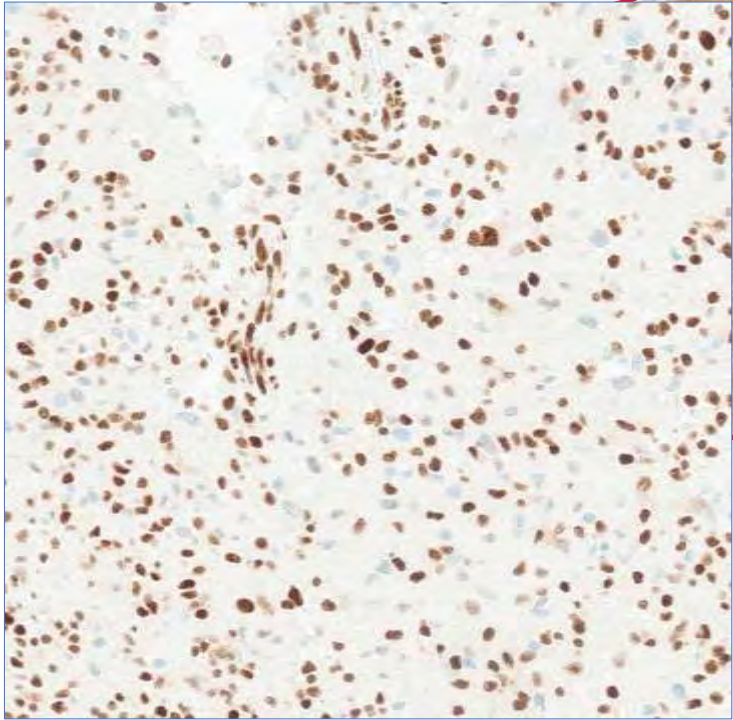
Astrocytic region

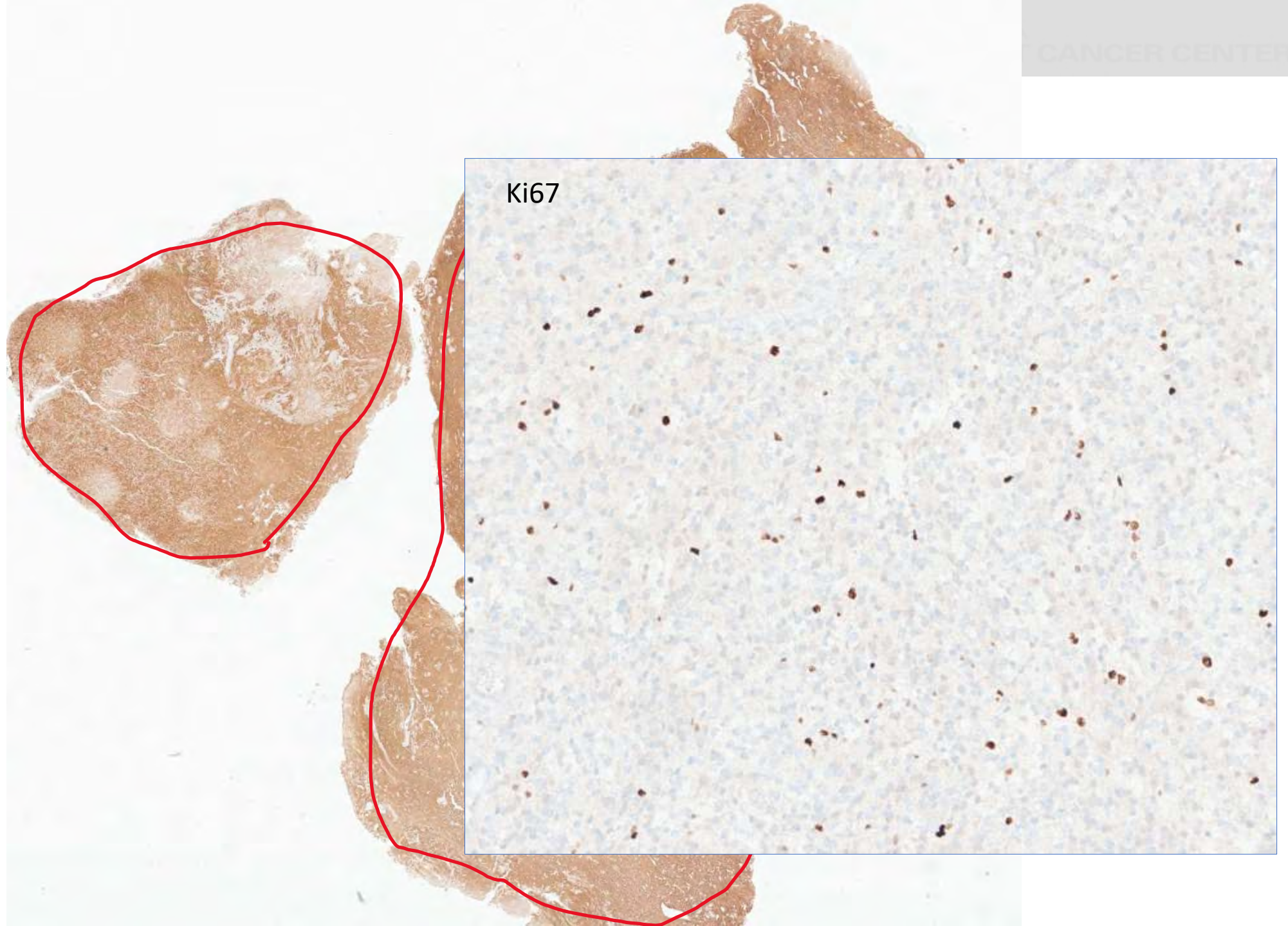
Oligodendroglioma-like region

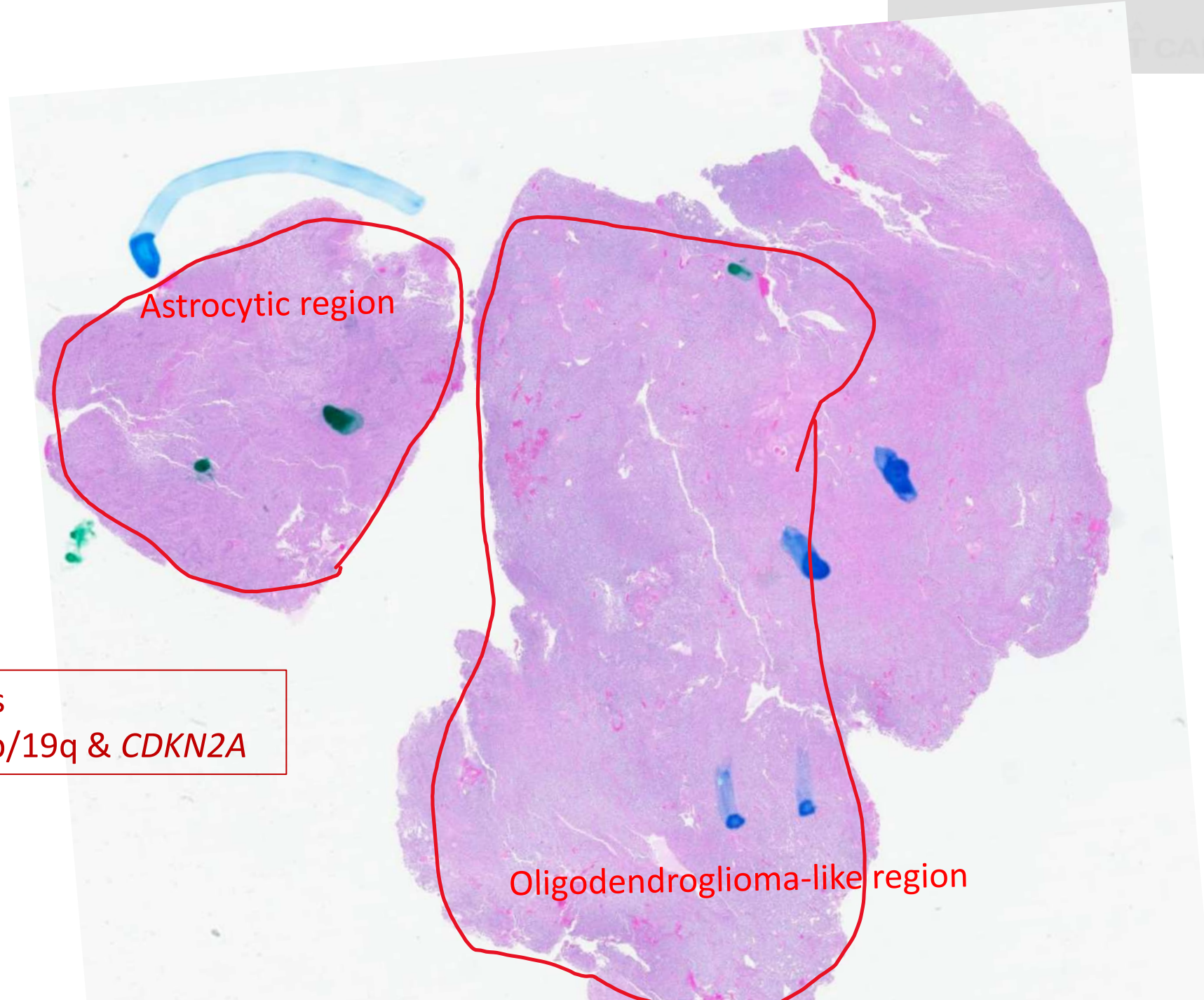
IDH1 p.R132H

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ATRX





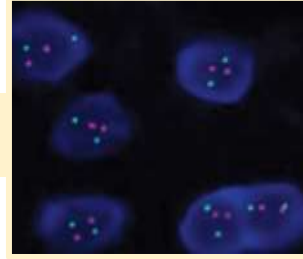


Astrocytic region

Oligodendroglioma-like region

Marked slides
→ FISH for 1p/19q & *CDKN2A*

Fluorescent In Situ Hybridization (*CDKN2A*, 1p/19q):



Astrocytic Morphology Area

POSITIVE for homozygous deletion of the *CDKN2A* (9p21) locus

NEGATIVE for 1p36 deletion

NEGATIVE for 19q13 deletion

Oligo-like Morphology Area

NEGATIVE for homozygous deletion of the *CDKN2A* (9p21) locus

NEGATIVE for 1p36 deletion

NEGATIVE for 19q13 deletion

Brain, right frontotemporal tumor, craniotomy with resection:

Astrocytoma, IDH-mutant
CNS WHO grade 4

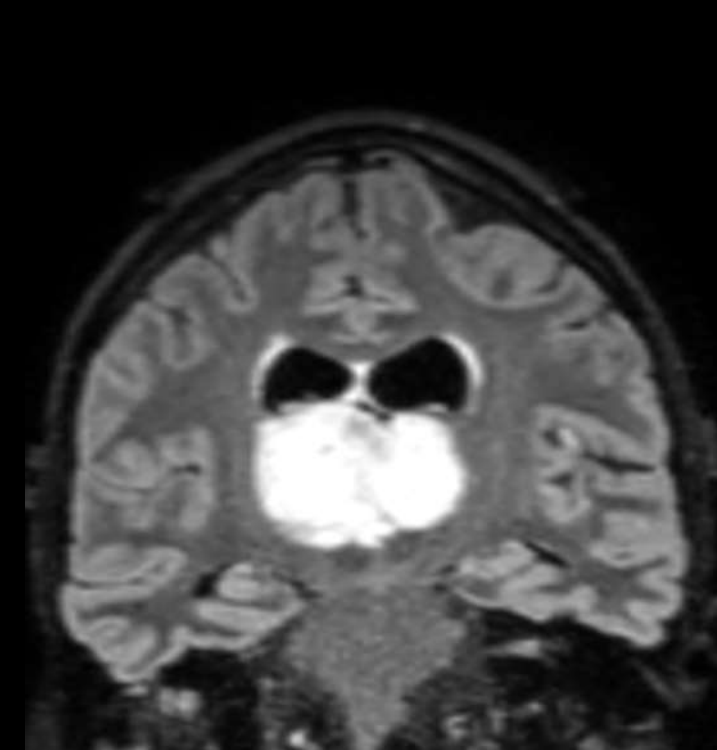
IDH1 (IHC): POSITIVE for p.R132H mutant protein expression

ATRX (IHC): LOSS of nuclear expression, regional

CDKN2A (FISH): POSITIVE for homozygous deletion, regional

1p/19q (FISH): Negative for codeletion

29M with diplopia found to have thalamic mass



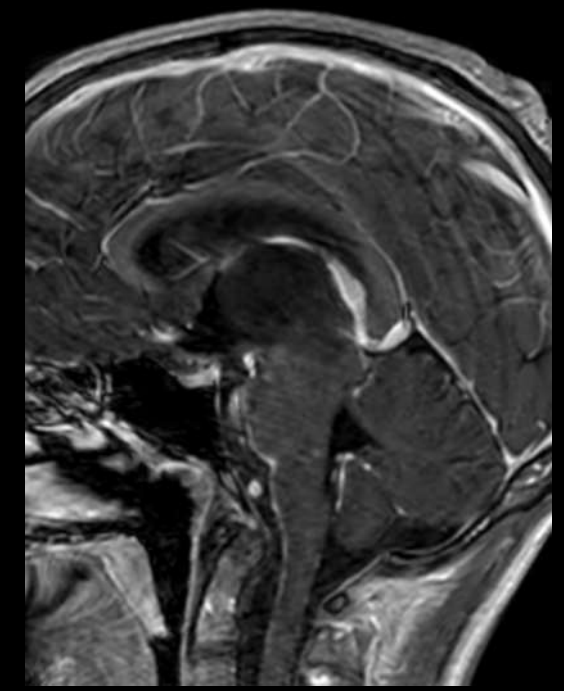
T2 FLAIR



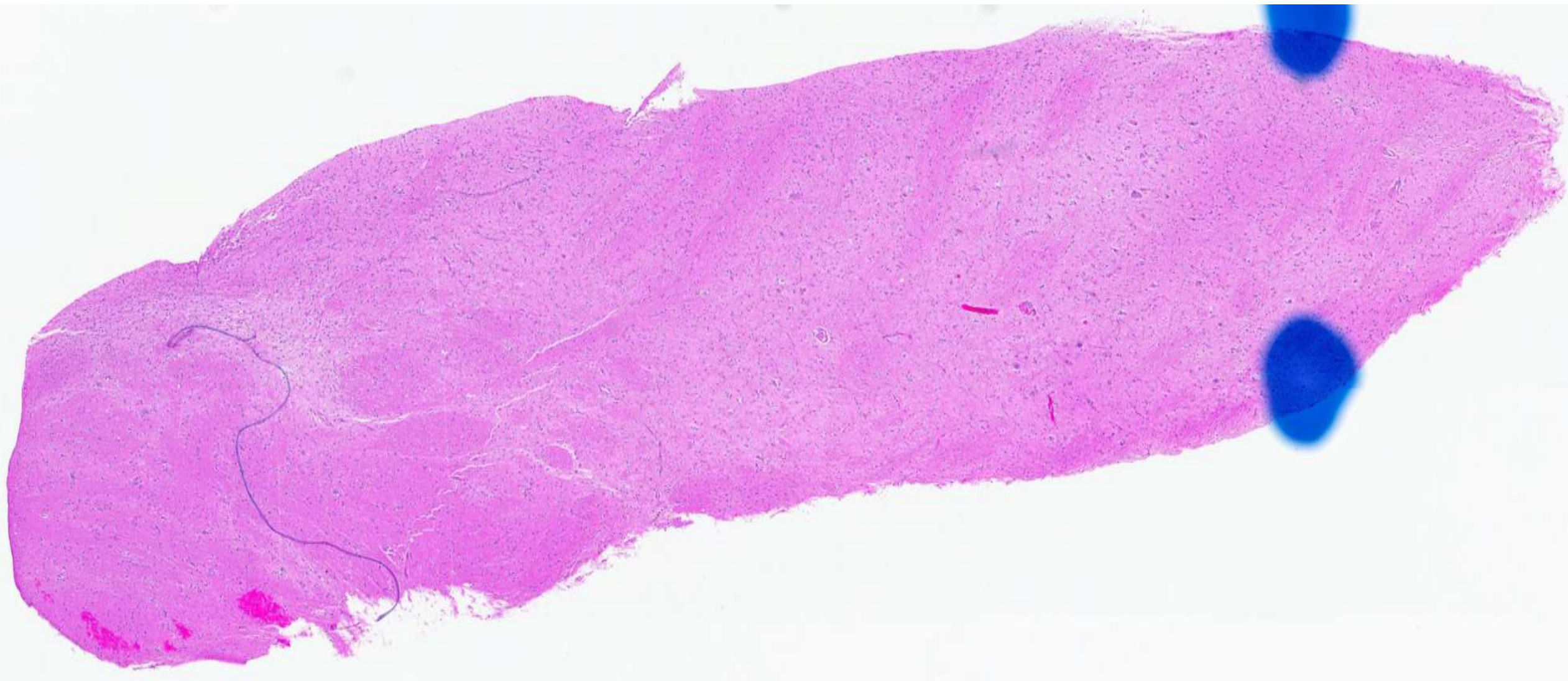
T2

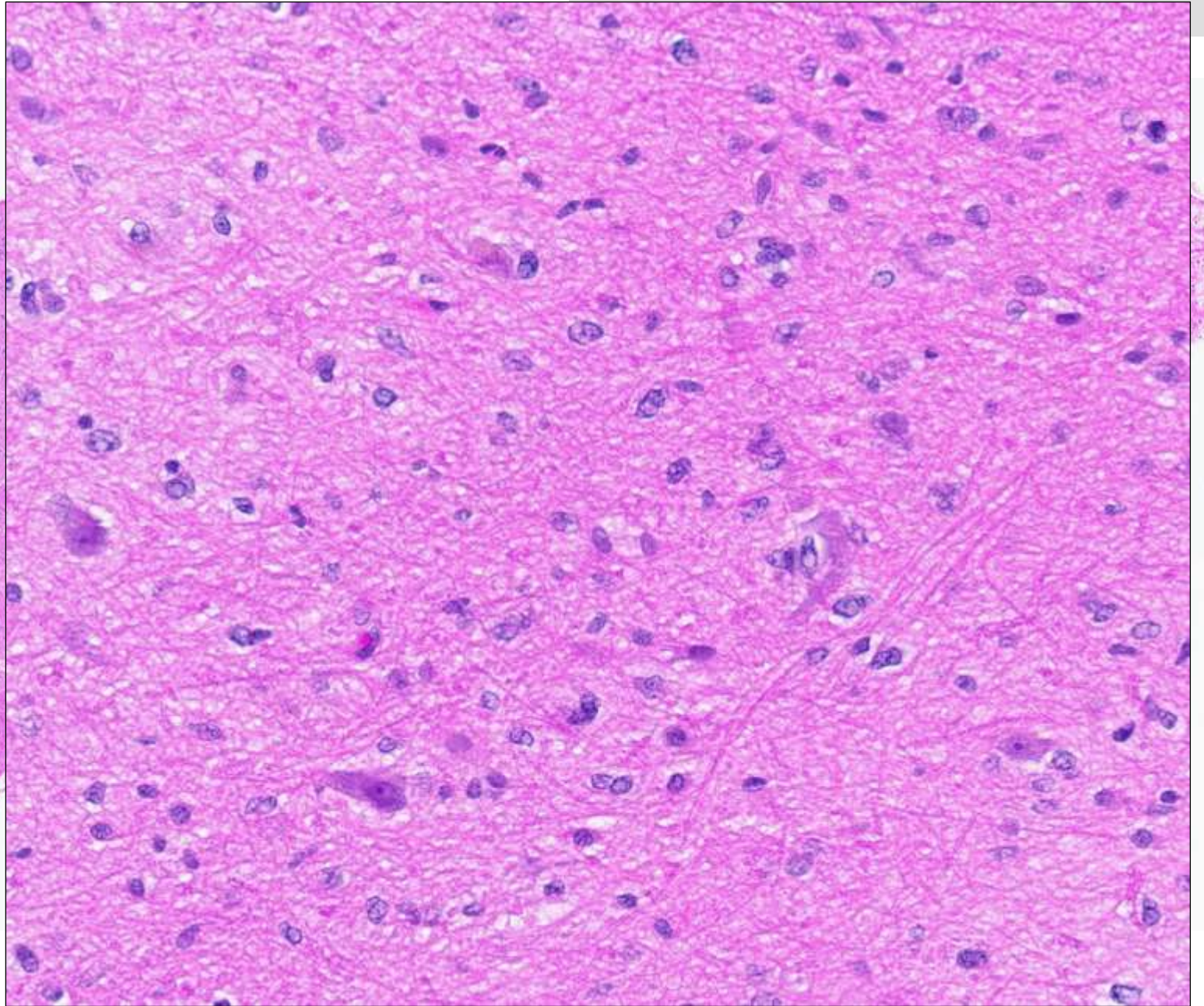
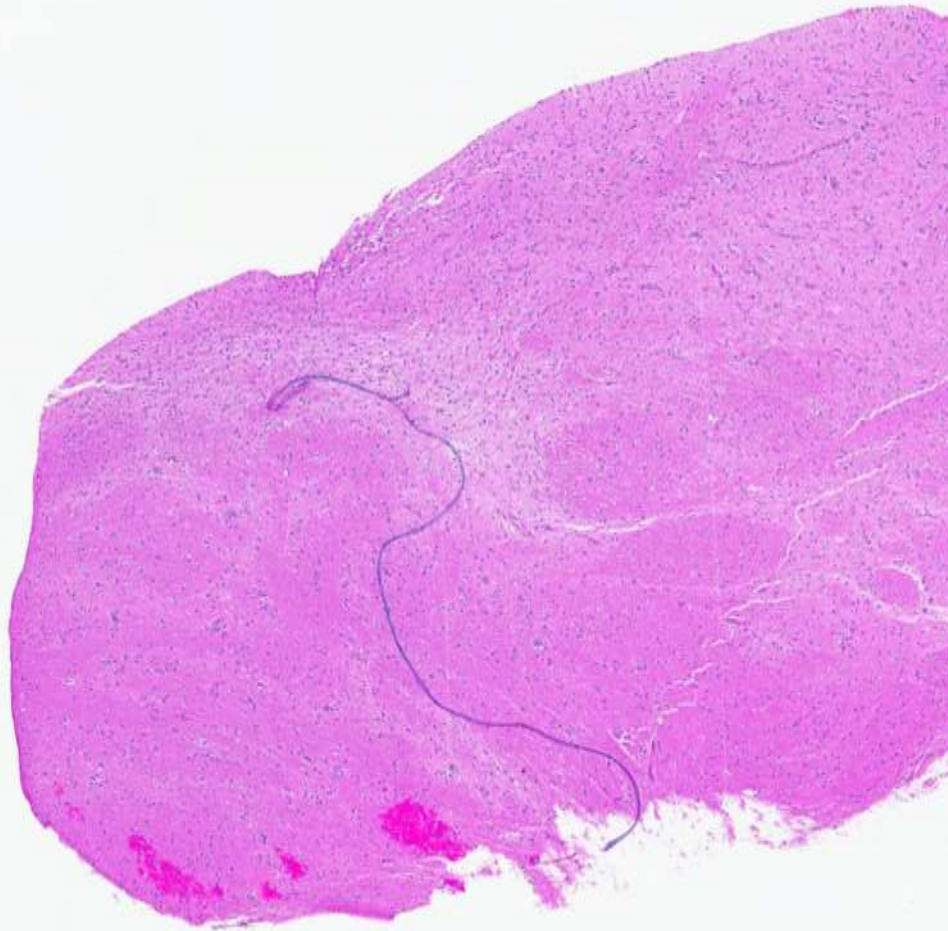


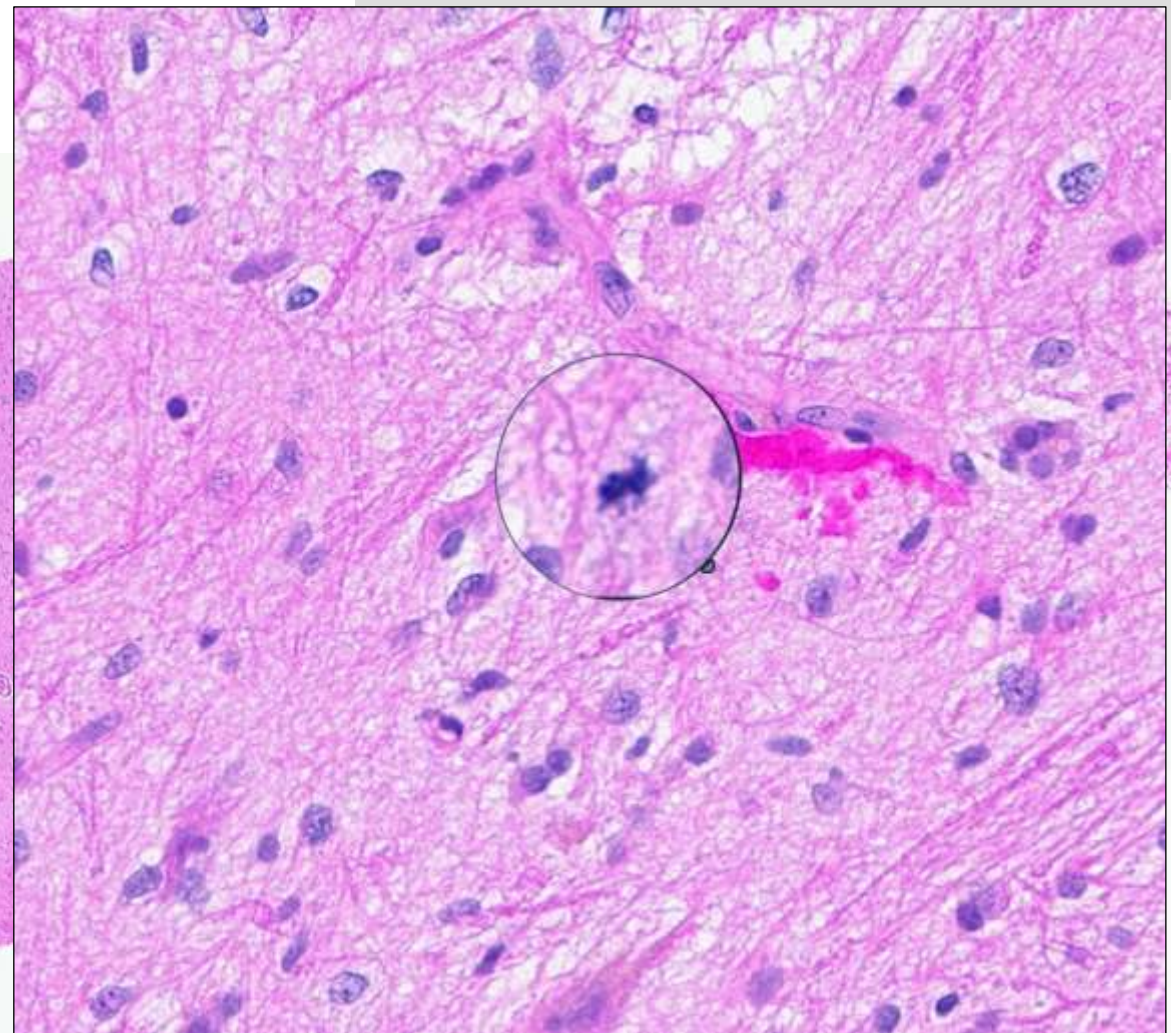
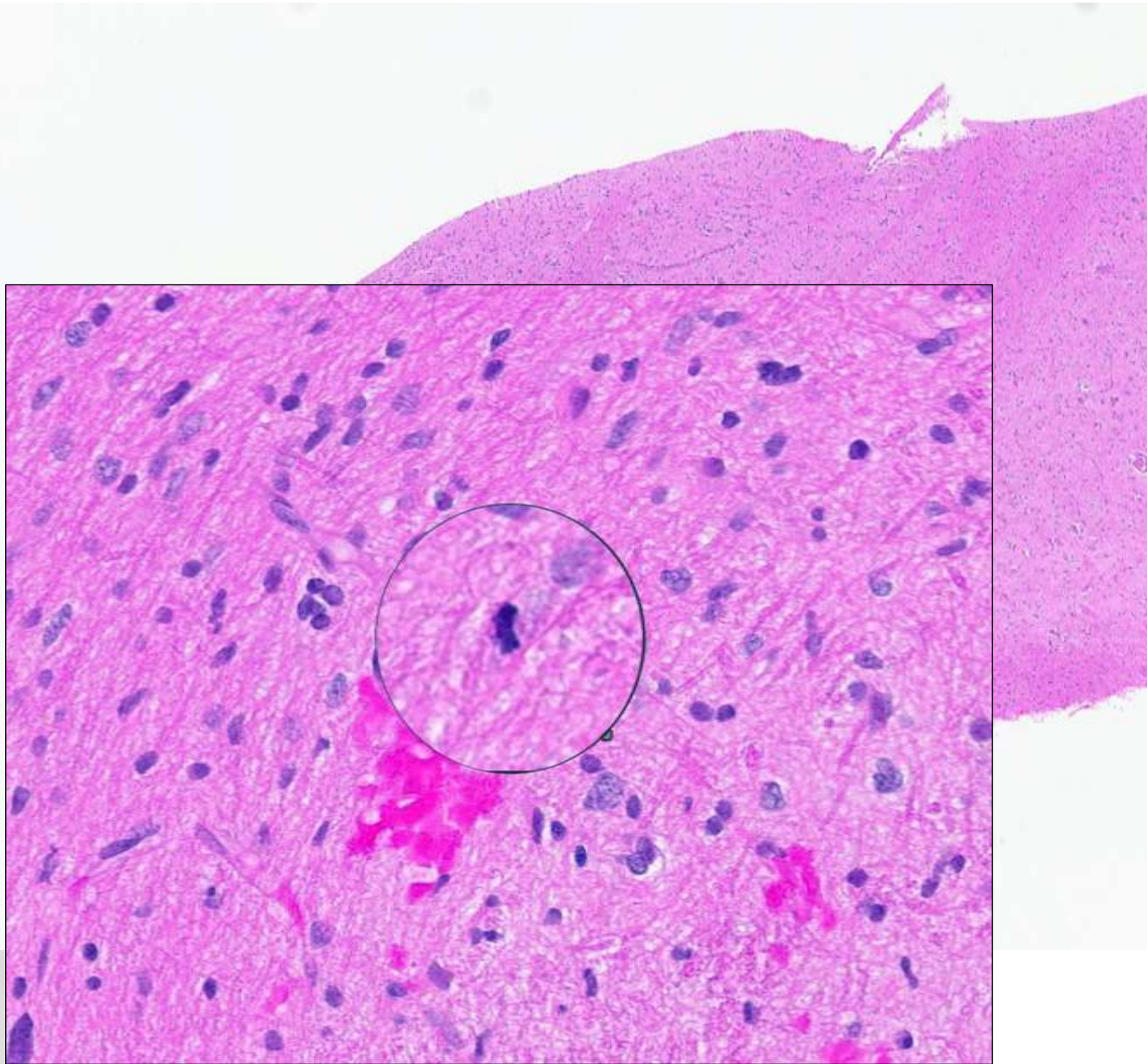
T1



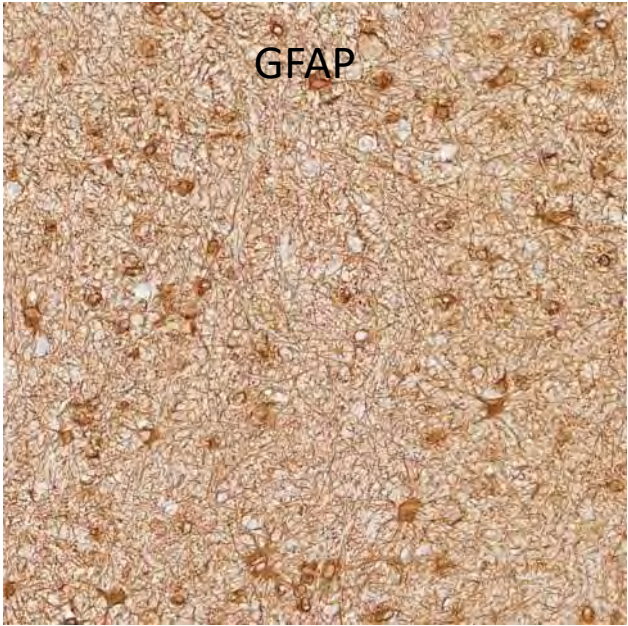
T1 +C



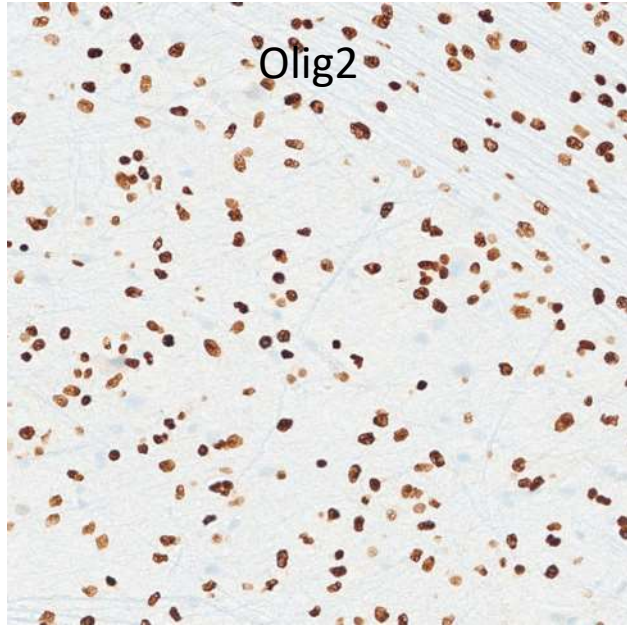




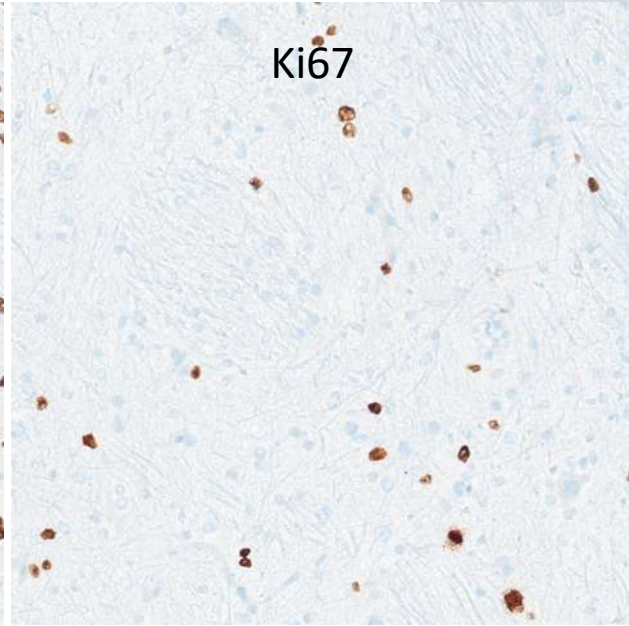
GFAP



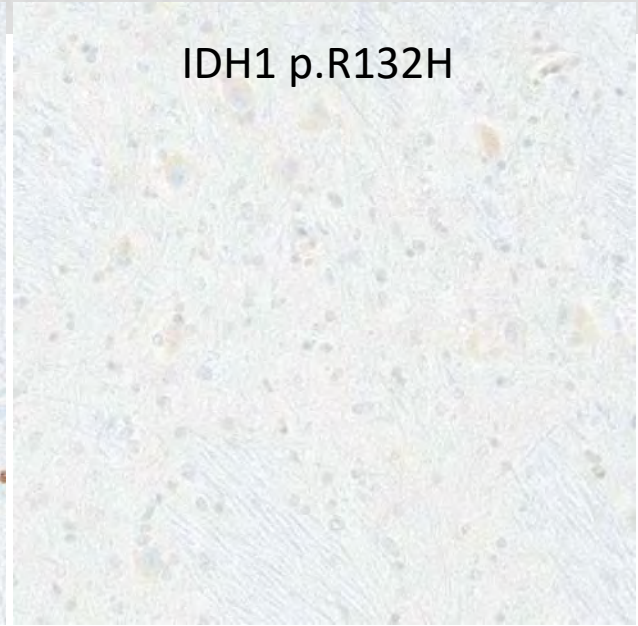
Olig2



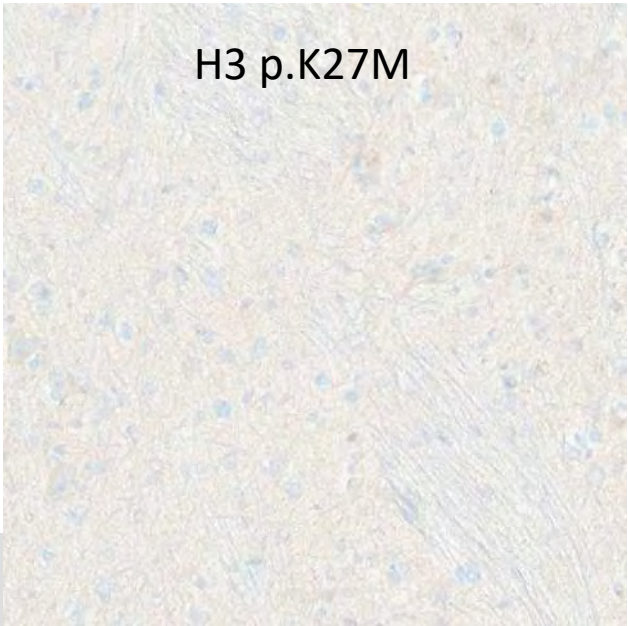
Ki67



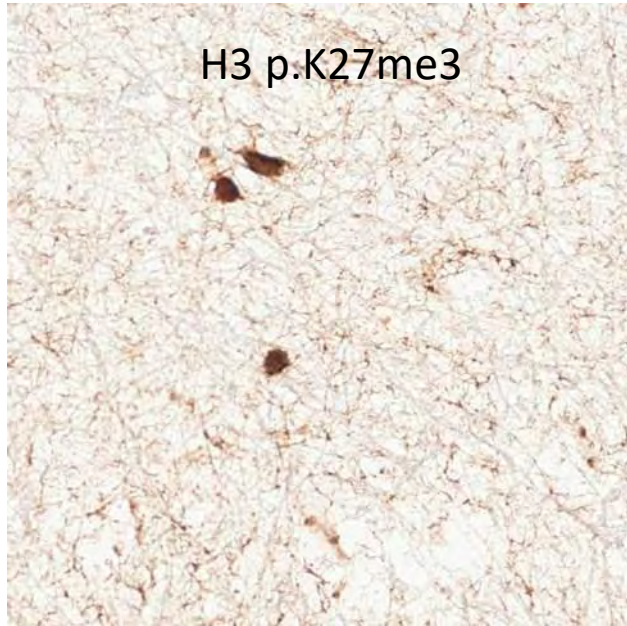
IDH1 p.R132H



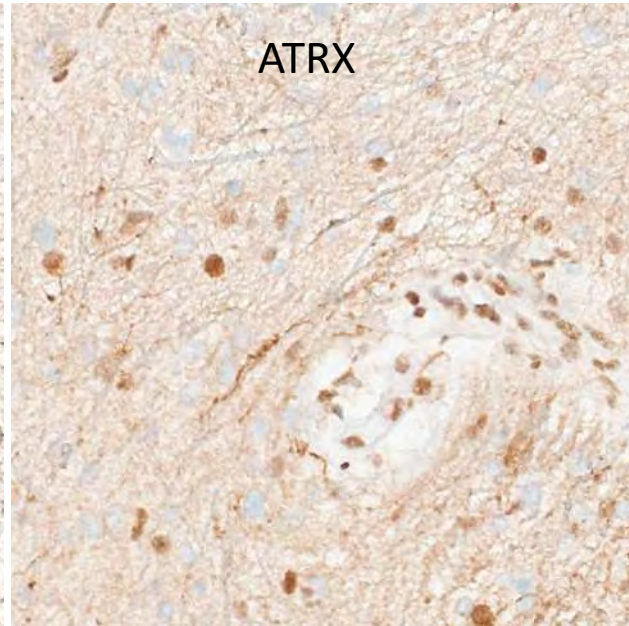
H3 p.K27M



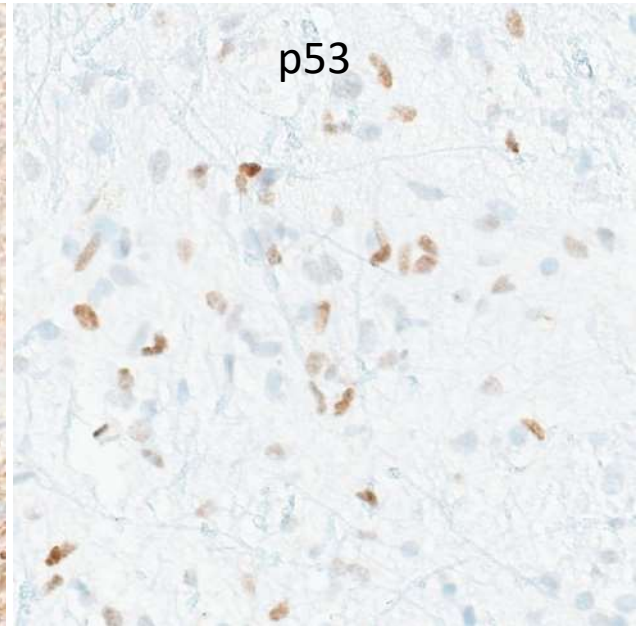
H3 p.K27me3



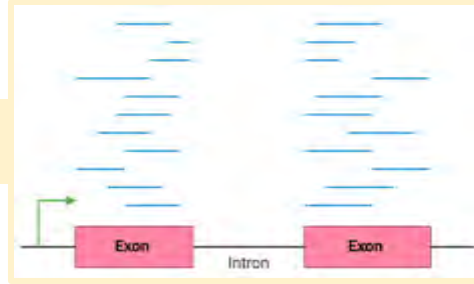
ATRX



p53



Next-Generation Sequencing Panel:



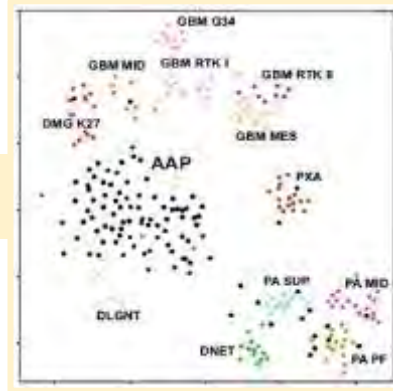
Copy Number Variations

CDKN2A Homozygous deletion
CDKN2B Homozygous deletion
NF1 Homozygous deletion

Somatic Mutations

<i>SETD2</i>	p.E1478*	21% (pathogenic)
<i>SETD2</i>	p.L1804S	25% (likely pathogenic)
<i>ATRX</i>	c.595-2A>G	50% (likely pathogenic)

DNA Methylation-Based Tumor Classification:



Methylation Class Name	High-grade astrocytoma with piloid features
Methylation Class Confidence Score	High confidence
Methylation Class Description:	The “methylation class high-grade astrocytoma with piloid features” is mainly comprised of tumors with the histological diagnosis of anaplastic pilocytic astrocytoma or less commonly glioblastoma. The tumors mostly occur in the posterior fossa and rarely in the diencephalic/thalamic region; median age is 40 years (age range 24 to 75). Deletions of CDKN2A/B are very frequent (>70%). BRAF duplications are also observed in a fraction of cases. Around 50% of cases show an immunohistochemical loss of ATRX. Loss of chromosome 19q (total or partial) is observed in over 50% of cases.

Brain, thalamic tumor, stereotactic biopsy:

High-grade astrocytoma with piloid features (see comment)

Diagnosis comment:

The biopsy show an infiltrating astrocytoma with scattered mitotic figures and a high Ki67 labeling index (10-15%), morphologically consistent with a high-grade astrocytoma. NGS reveals a mutation in ATRX and homozygous deletions in CDKN2A/B and NF1. Methylation profiling reveals consensus match to High-grade astrocytoma with piloid features (HGAP) with a high confidence score. HGAP is a new tumor type in the 2021 WHO classification system with limited data available. A definitive WHO grade has not been established, although current data suggest a clinical behavior roughly corresponding to CNS WHO grade 3.



Anaplastic astrocytoma with piloid features, a novel molecular class of IDH wildtype glioma with recurrent MAPK pathway, CDKN2A/B and ATRX alterations

Annekathrin Reinhardt^{1,2} · Damian Stichel^{1,2} · Daniel Schrimpf^{1,2} · Felix Sahm^{1,2} · Andrey Korshunov^{1,2} · David E. Reuss^{1,2} · Christian Koelsche^{1,2} · Kristin Huang^{1,2} · Annika K. Wefers^{1,2} · Volker Hovestadt^{3,4} · Martin Sill^{4,48} · Dorothee Gramatzki²⁹ · Joerg Felsberg⁹ · Guido Reifenberger^{9,30} · Arend Koch⁷ · Ulrich-W. Thomale³⁵ · Albert Becker⁸ · Volkmar H. Hans¹⁰ · Marco Prinz^{11,47} · Ori Staszewski¹¹ · Till Acker¹² · Hildegard Dohmen¹² · Christian Hartmann¹³ · Wolf Mueller¹⁴ · Muin S. A. Tuffaha³⁶ · Werner Paulus¹⁵ · Katharina Heß¹⁵ · Benjamin Brokinkel¹⁵ · Jens Schittenhelm¹⁶ · Camelia-Maria Monoranu¹⁷ · Almuth Friederike Kessler³⁷ · Mario Loehr³⁷ · Rolf Buslei^{18,19} · Martina Deckert²⁰ · Christian Mawrin²¹ · Patricia Kohlhof²² · Ekkehard Hewer²³ · Adriana Olar^{24,25,26} · Fausto J. Rodriguez²⁷ · Caterina Giannini²⁸ · Amulya A. NageswaraRao²⁸ · Uri Tabori^{38,39,40,41} · Nuno Miguel Nunes^{40,41} · Michael Weller²⁹ · Ute Pohl³¹ · Zane Jaunmuktane³² · Sebastian Brandner³² · Andreas Unterberg⁴² · Daniel Hänggi⁴³ · Michael Platten^{44,45} · Stefan M. Pfister^{4,5,6,48} · Wolfgang Wick^{33,4} · Christel Herold-Mende³⁴ · David T. W. Jones^{4,48,49} · Andreas von Deimling^{1,2,4} · David Capper^{1,2,46,50}

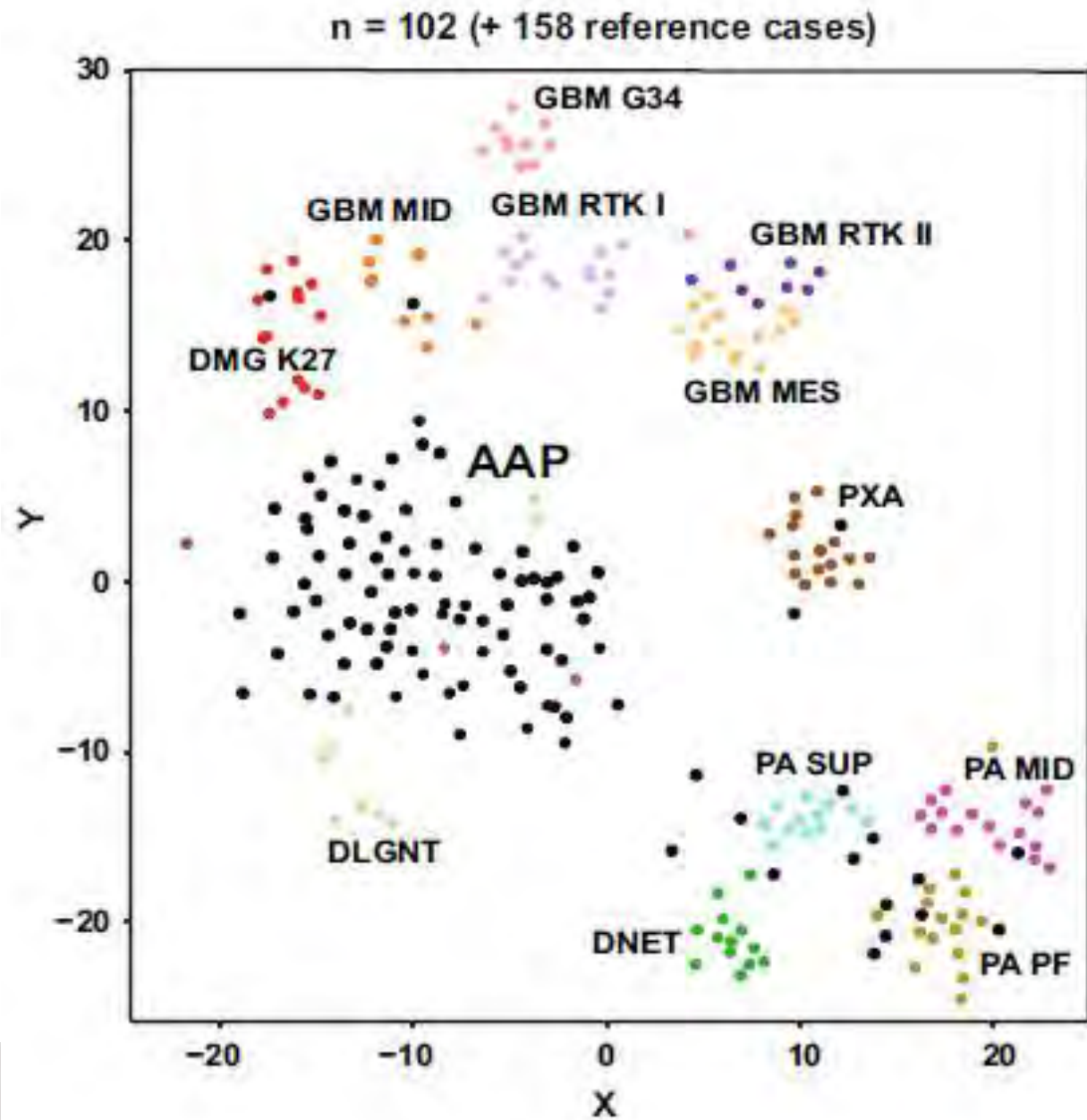


Fig. 1 t-SNE analysis of 102 cases with histological features compatible with the diagnosis of anaplastic pilocytic astrocytoma, indicated in black, and 158 reference cases of established glioma methylation classes, indicated in different colors. Tumors of the same class are depicted in one color. For this analysis, the 20,000 most variably methylated CPG sites were used. 83 of the histologically

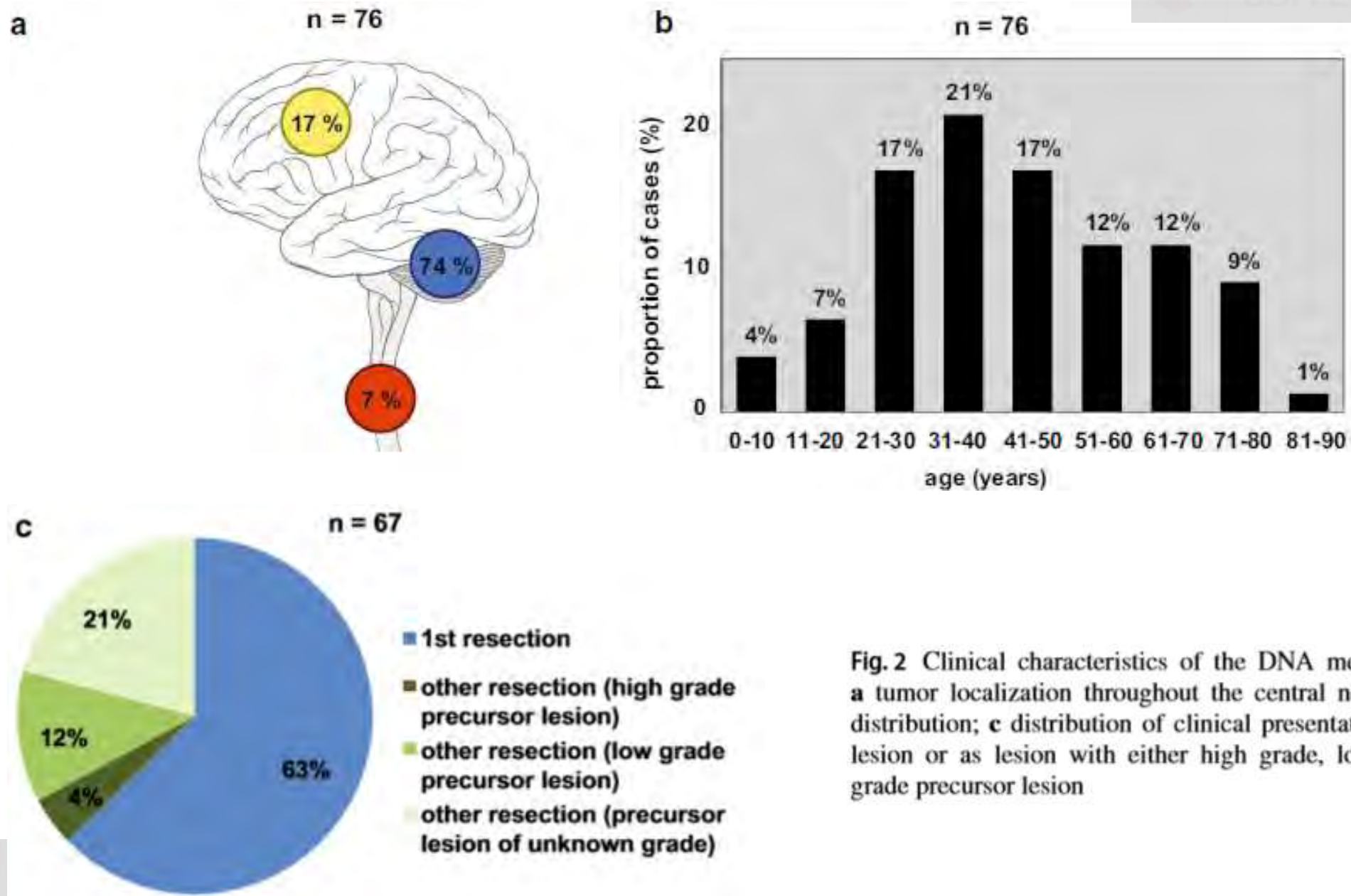


Fig. 2 Clinical characteristics of the DNA methylation class AAP: **a** tumor localization throughout the central nervous system; **b** age distribution; **c** distribution of clinical presentation as either primary lesion or as lesion with either high grade, low-grade or unknown grade precursor lesion

n = 83

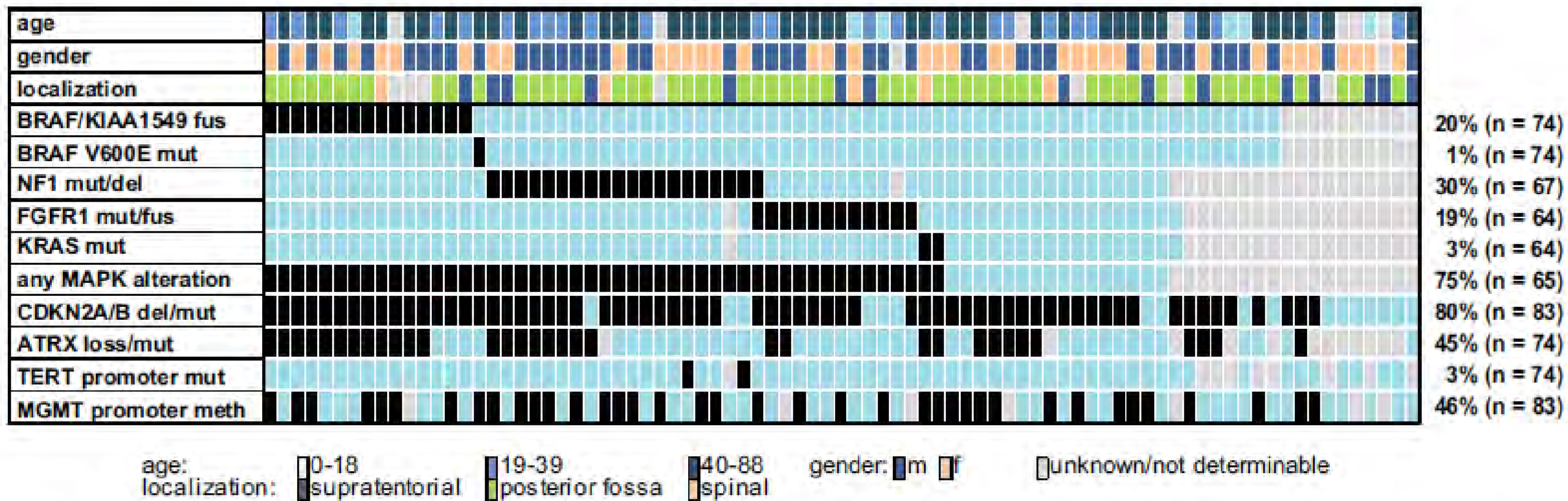


Fig. 6 Summary of molecular alterations and patient characteristics for the methylation class anaplastic astrocytoma with piloid features. For molecular alterations, black fields indicate the presence and light blue fields the absence of the respective alteration. Grey fields indi-

cate that the lesion was not tested for the specific alteration or that the respective parameter was not determinable. *mut* mutation, *del* deletion, *fus* fusion, *meth* methylated

Meningioma, CNS WHO grade 1-3

WHO grade 1

Meningothelial
Fibrous (fibroblastic)
Transitional (mixed)
Psammomatous
Angiomatous
Secretory
Lymphoplasmacyte-rich
Metaplastic
Rhabdoid (?)

Recurrence risk: 7-25%

WHO grade 2

Chordoid
Clear cell

Atypical meningioma

Brain invasion
(or)
 ≥ 2.5 mitosis/mm²
(or)
At least 3 of the following:
increased cellularity
small cells with high N:C
prominent nucleoli
sheeting
spontaneous necrosis

29-52%

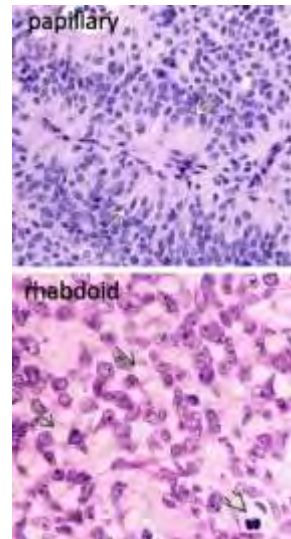
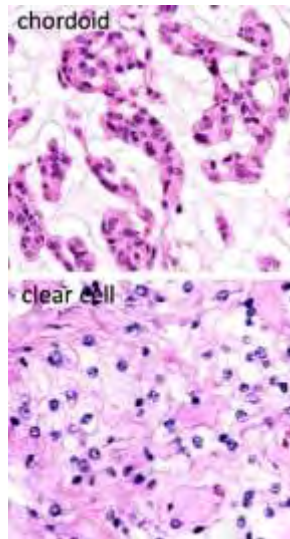
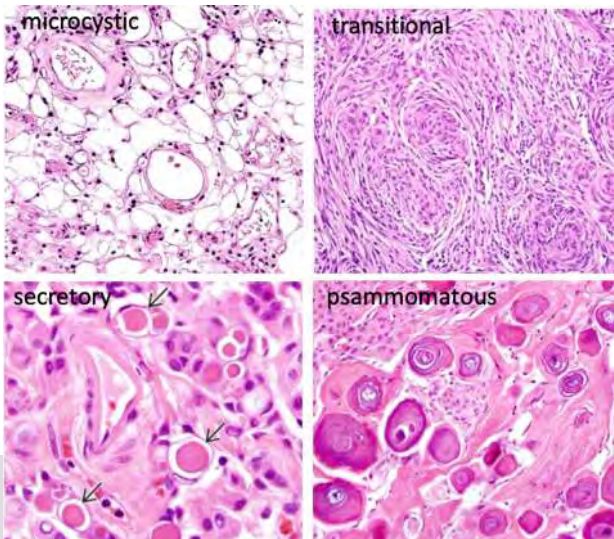
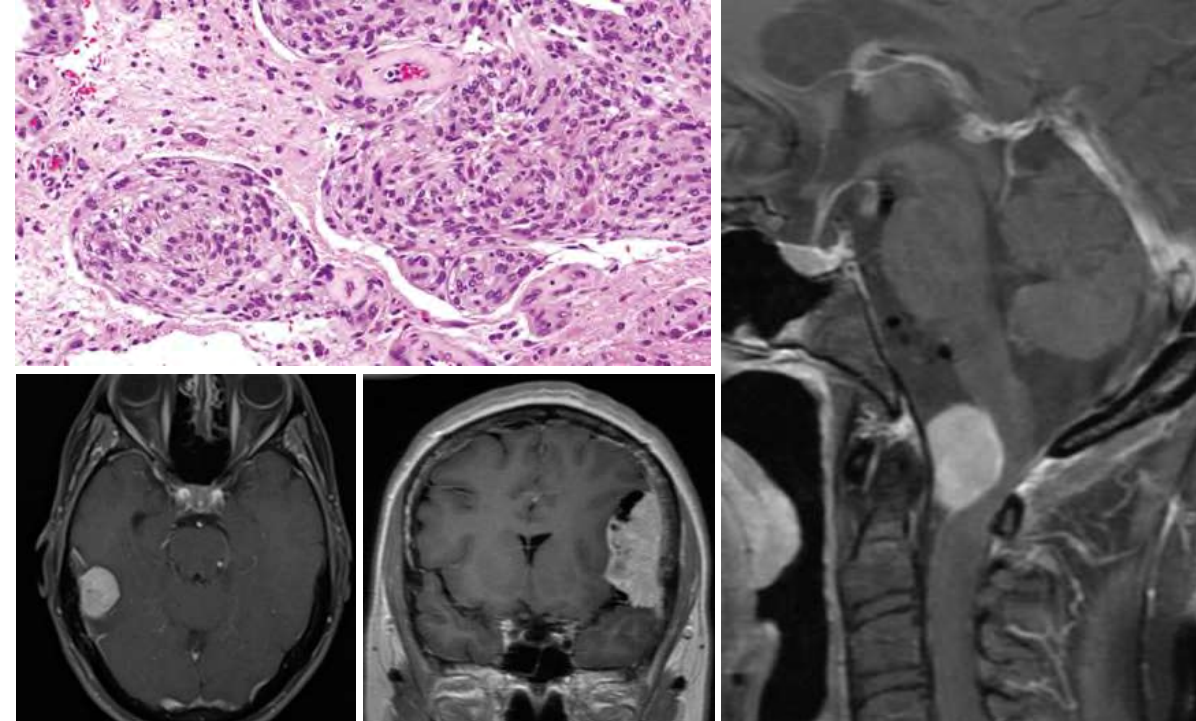
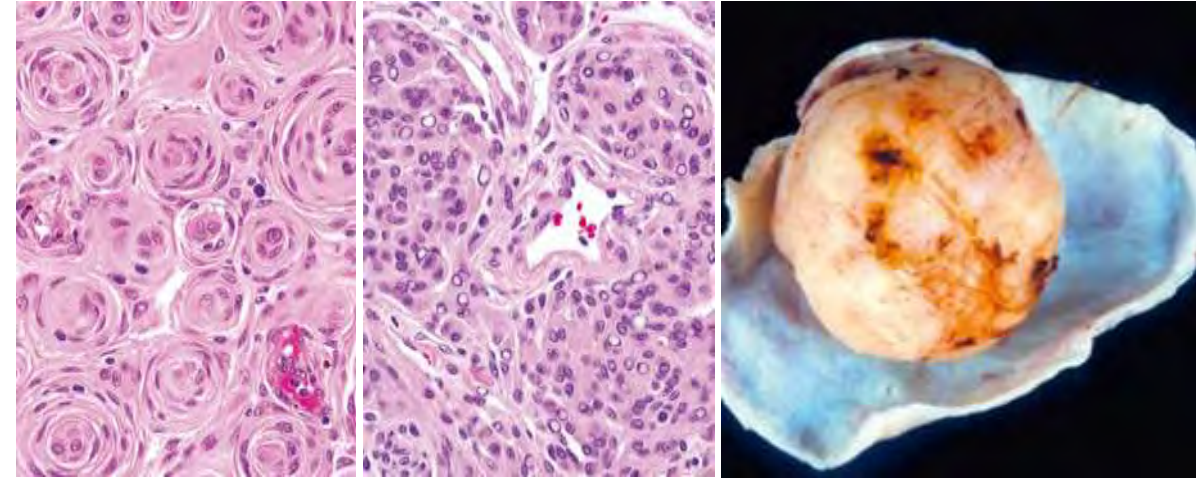
WHO grade 3

Papillary (?)
Rhabdoid (?)

Anaplastic meningioma

Overtly malignant (resembling carcinoma, melanoma, high-grade sarcoma)
(and/or)
 ≥ 12.5 mitosis/mm²
(and/or)
Molecular features:
TERT p mutation
CDKN2A/B loss

50-94%



Additional proposed gliomas supported by tumor methylation profiling since WHO CNS 2021 publication:

High-grade glioma with pleomorphic and pseudopapillary features (HPAP)

Neuroepithelial tumor with *PATZ1* fusion

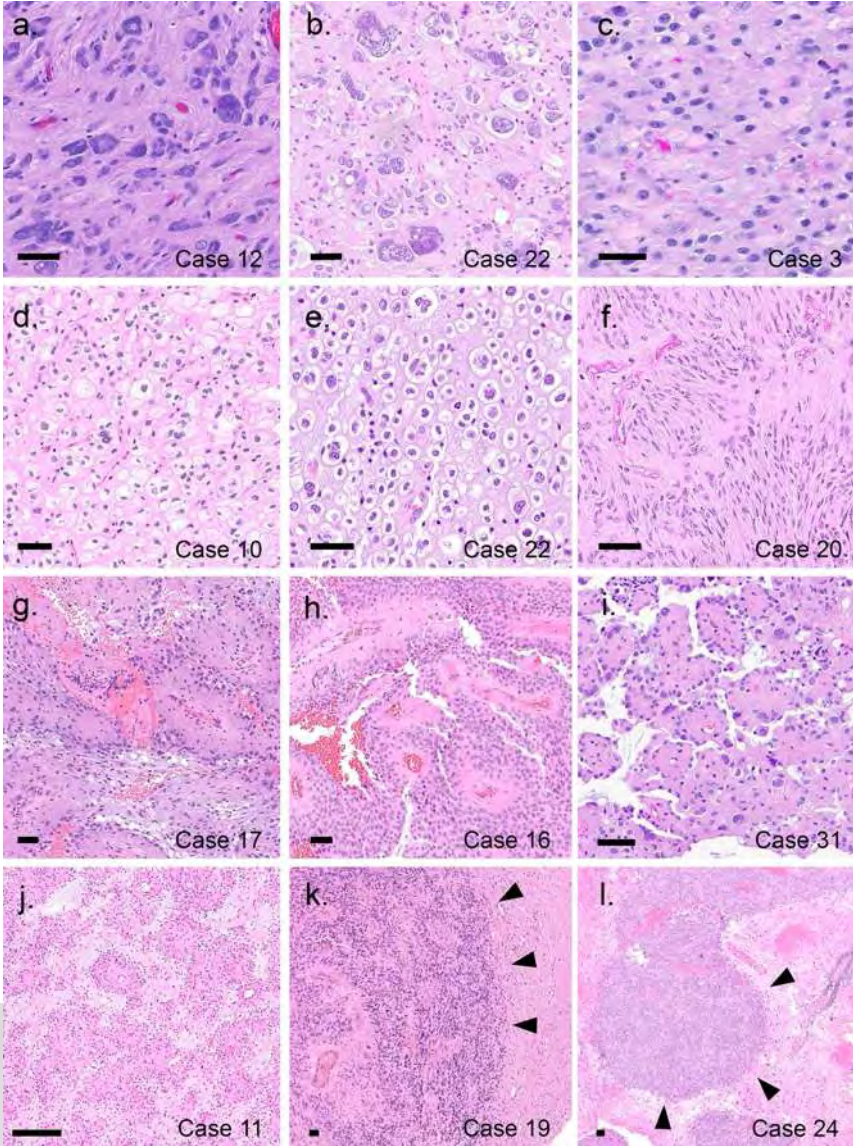
Glial tumor with *BCOR* fusion

Additional proposed gliomas supported by tumor methylation profiling since WHO CNS 2021 publication:

High-grade glioma with pleomorphic and pseudopapillary features (HPAP)

Neuroepithelial tumor with *PATZ1* fusion

Glial tumor with *BCOR* fusion

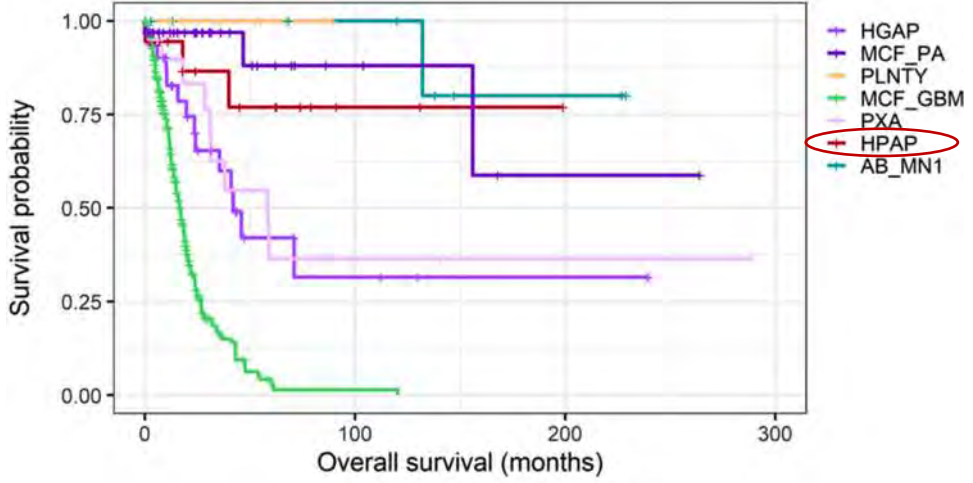
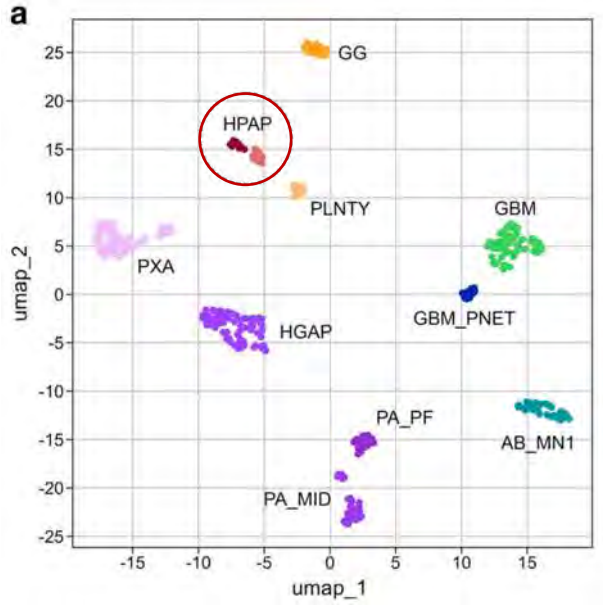


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ORIGINAL PAPER

High-grade glioma with pleomorphic and pseudopapillary features (HPAP): a proposed type of circumscribed glioma in adults harboring frequent *TP53* mutations and recurrent monosomy 13

Drew Pratt¹ · Zied Abdullaev¹ · Antonios Papanicolau-Sengos¹ · Courtney Ketchum¹ · Pavalan Panneer Selvam¹





Thank you!

Division of Neuropathology



Jie Chen, MD, PhD



Sahara Cathcart, MD