

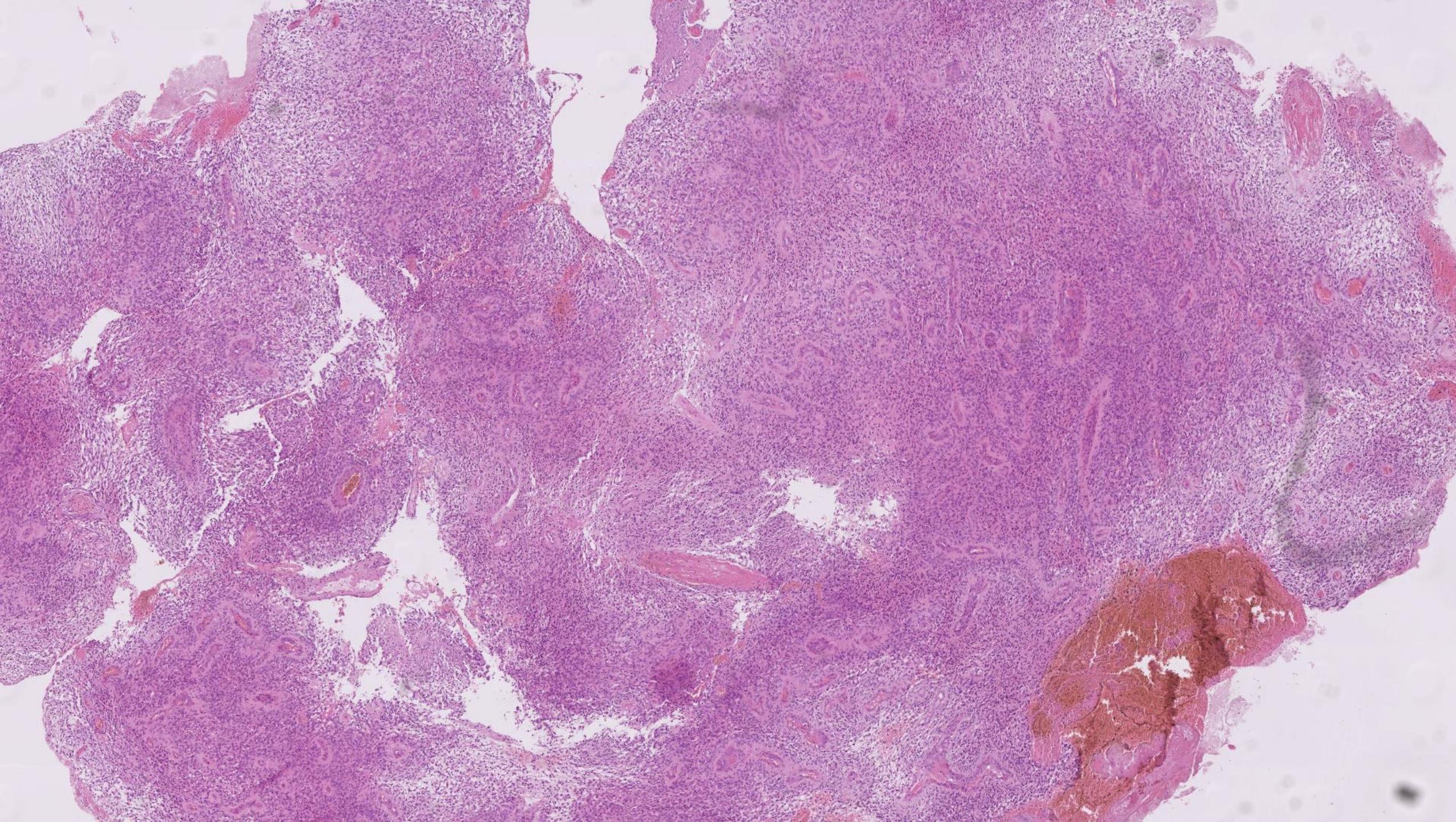
XXX. Martinský bioptický seminár SD-IAP

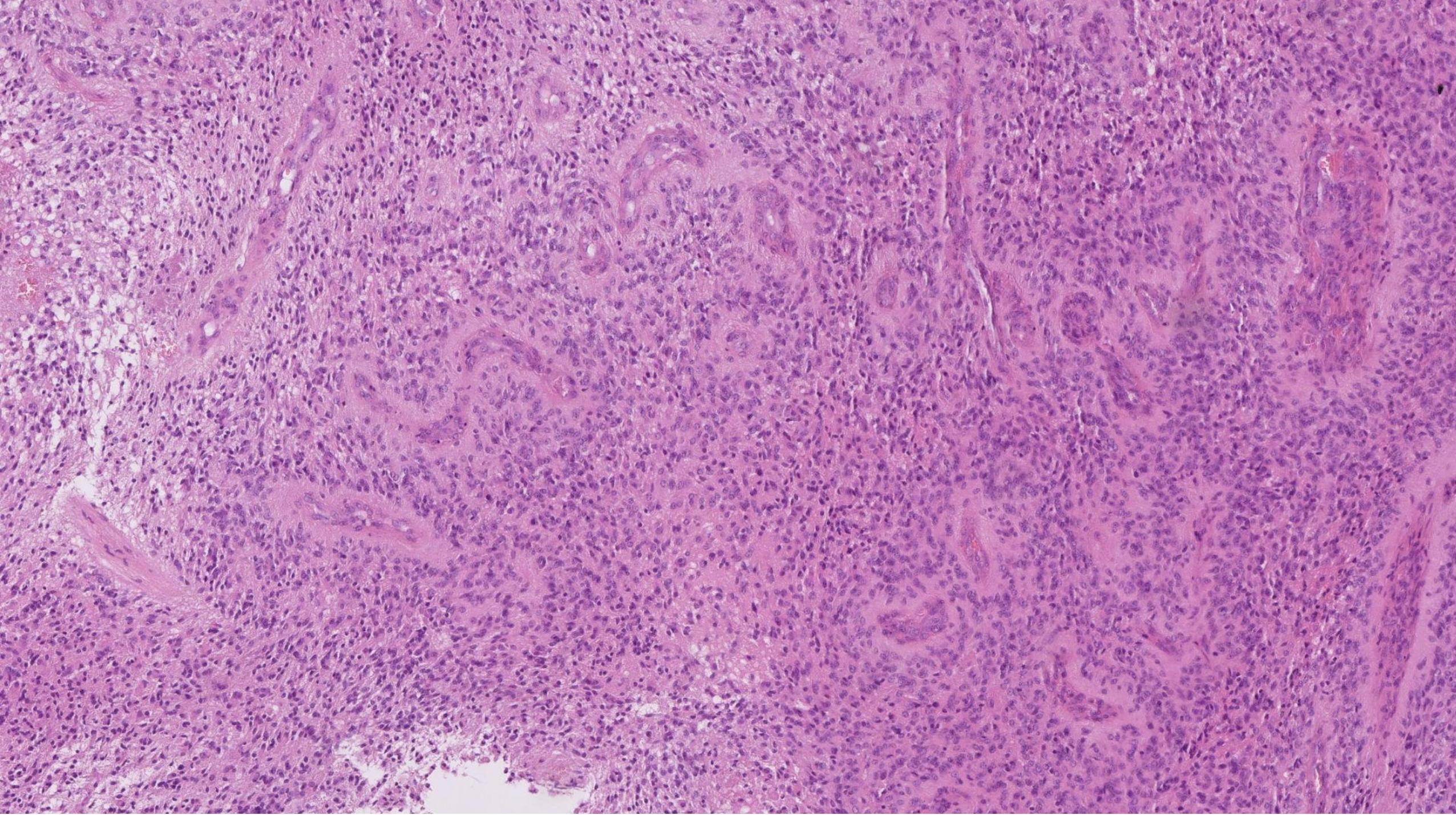
Prípad SD-IAP 854

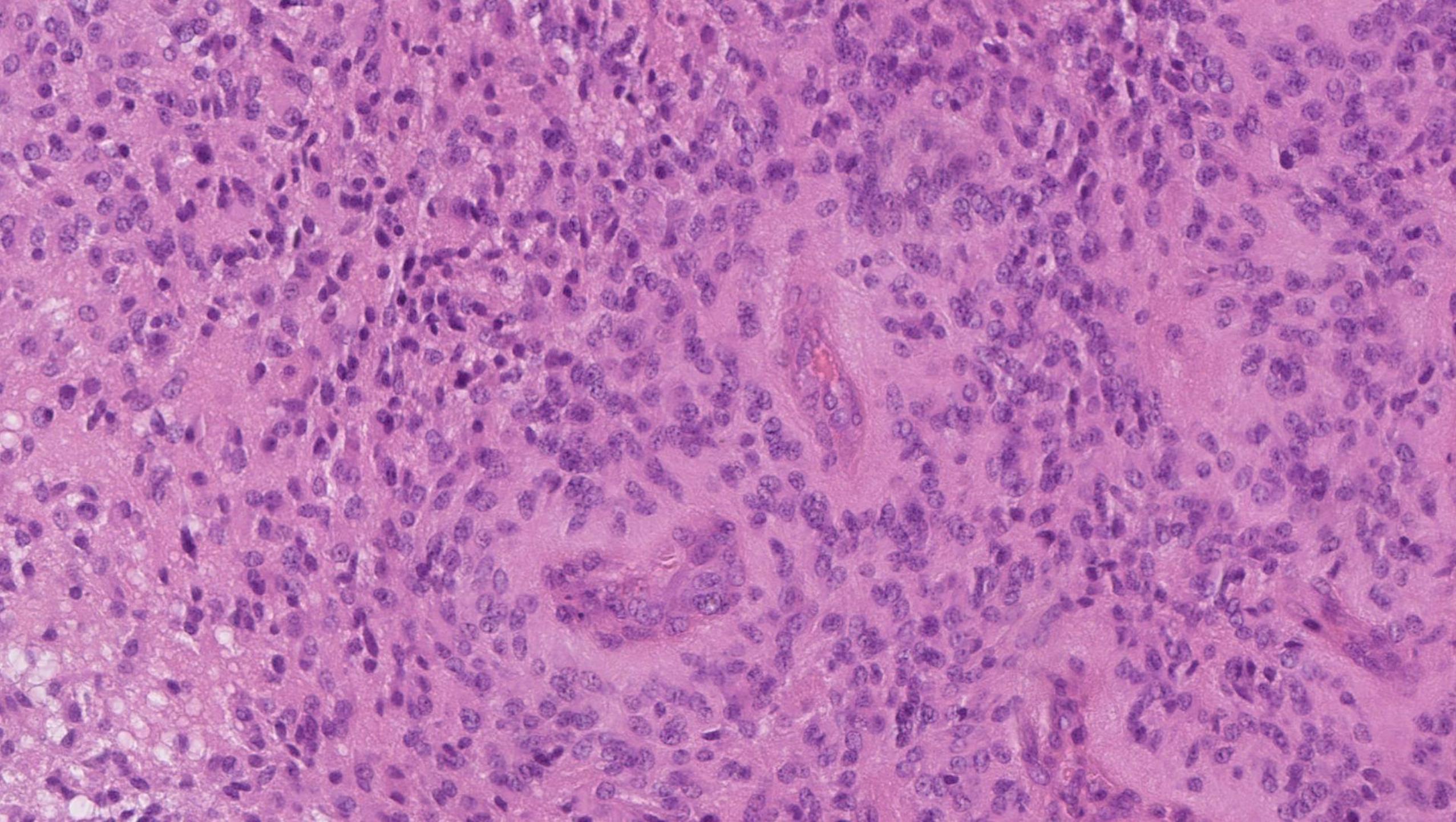
M. Švajdler

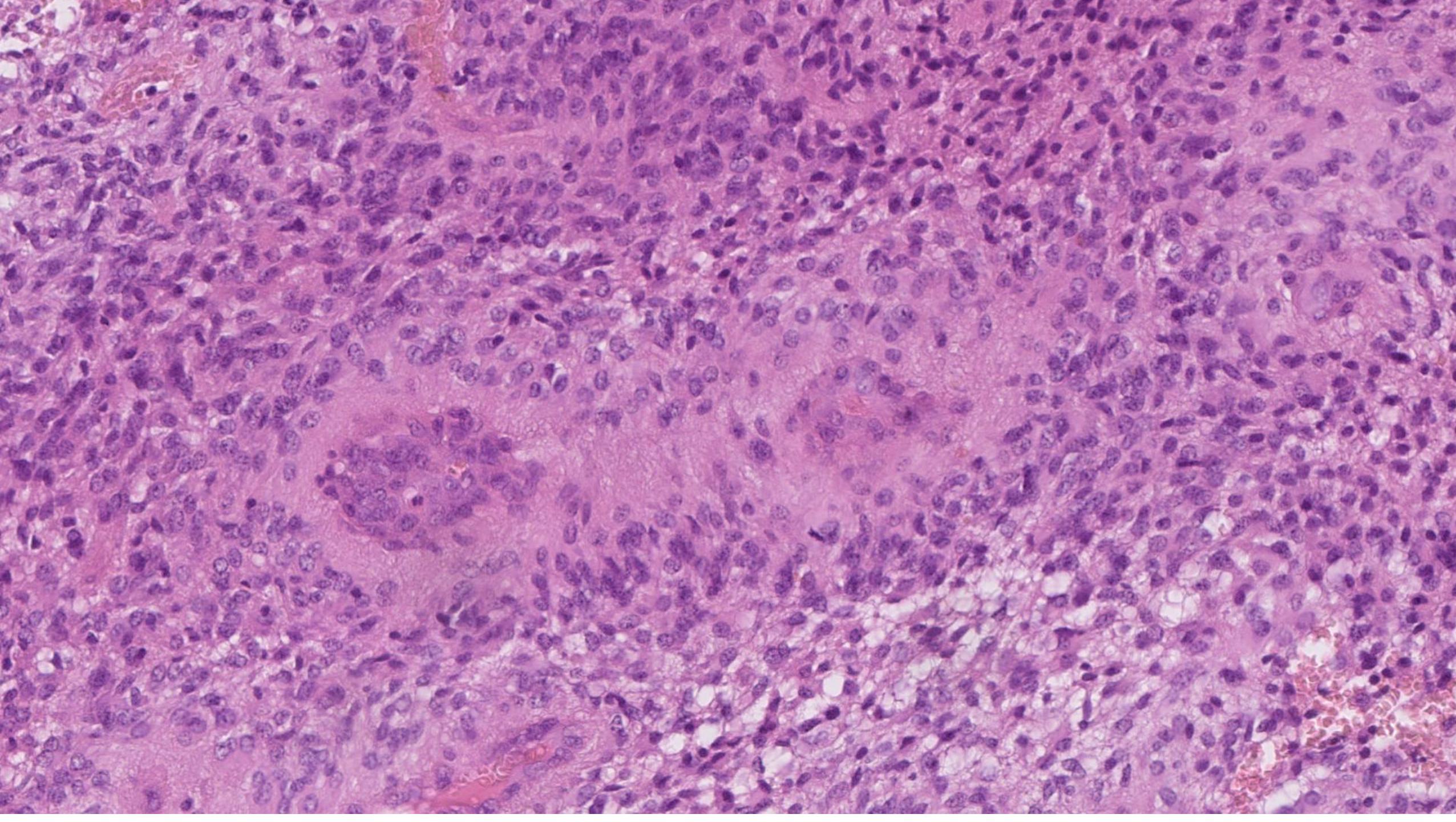


- 11-ročné dievča, tumor mozgu, konzultačné vyšetrenie - **patologická diagnóza ependymóm**
- prípad zaslaný za účelom **molekulovej subklasifikácie pomocou metylačného profilovania**



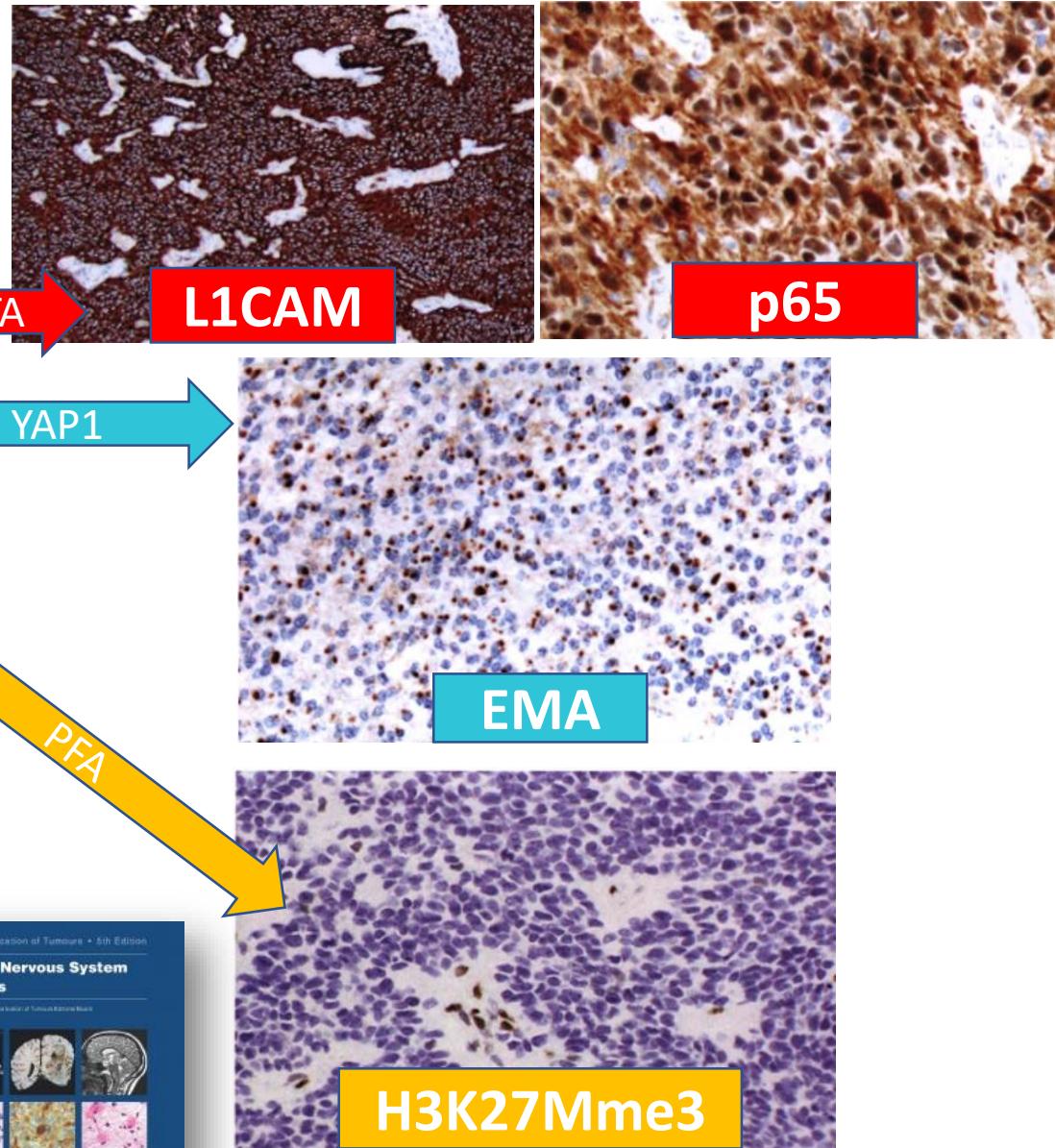
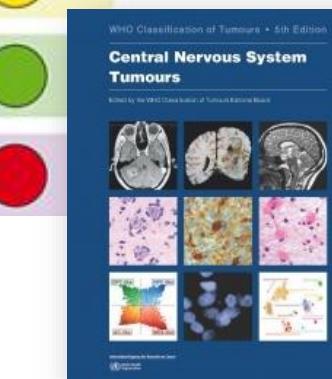




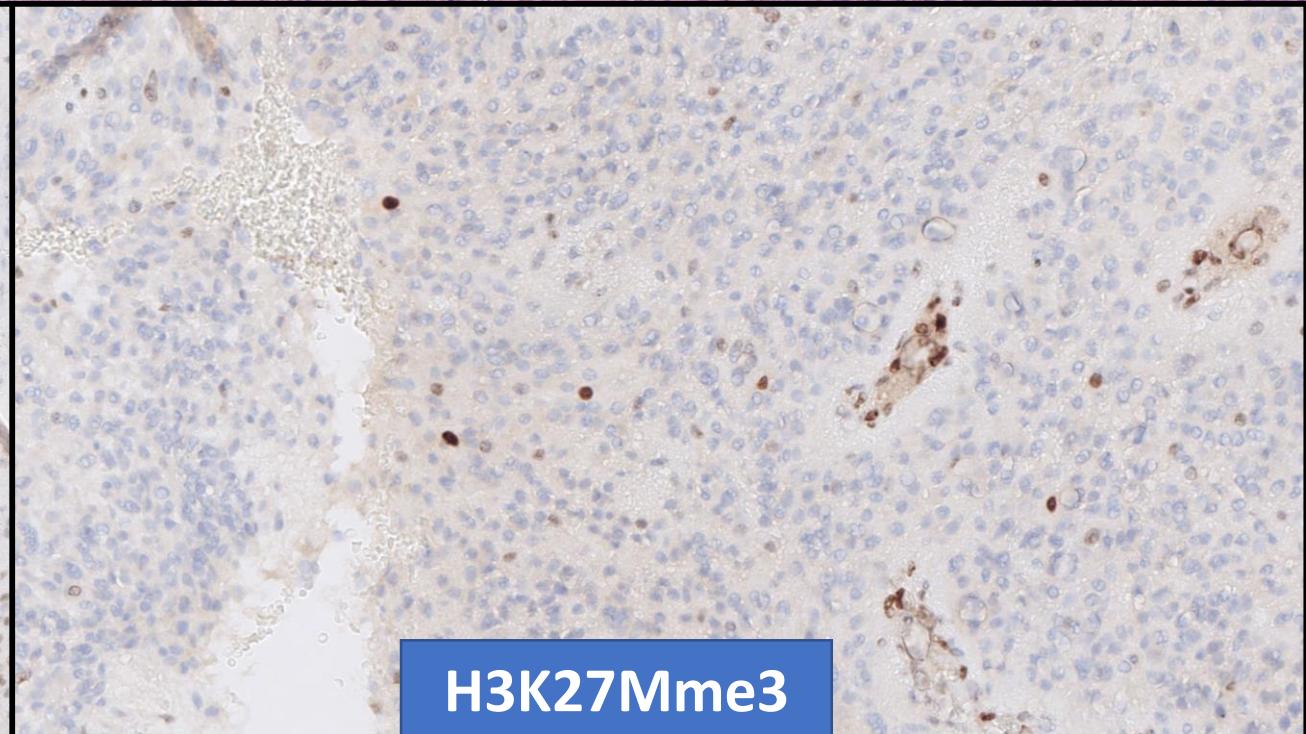
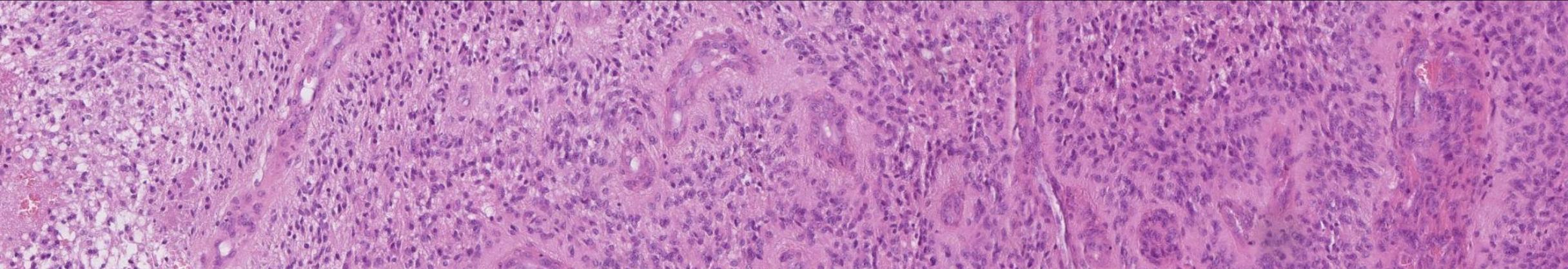


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	Age	Sex	CNS WHO grade	Molecular features	Outcome	
Supratentorial	ST-SE		♂ ♂ ♂ ♀	1	Balanced genome	
	ST-ZFTA		♂ ♂ ♀	2 / 3	ZFTA fusions Chromothripsis CDKN2A and/or CDKN2B loss	ZFTA →
	ST-YAP1		♂ ♀ ♀ ♀	2 / 3	YAP1 fusions	YAP1 →
	PF-SE		♂ ♂ ♂ ♀	1	Balanced genome	
	PFA		♂ ♂ ♀	2 / 3	EZH2 mutations H3 p.K28M (K27M) mutations Chr. 1q gain	PFA →
	PFB		♂ ♀	2 / 3	Chromosomal instability	
	SP-SE		♂ ♀	1	Chr. 6q deletion	
	SP-EP		♂ ♂ ♀	2 / 3	NF2 mutations	
	SP-MP		♂ ♀	2	Chromosomal instability	
	SP-MYCN		♂ ♀		MYCN amplification (Chr. 2p)	



11-ročné dievča, ependymóm



Brain tumor classifier results (12.5)

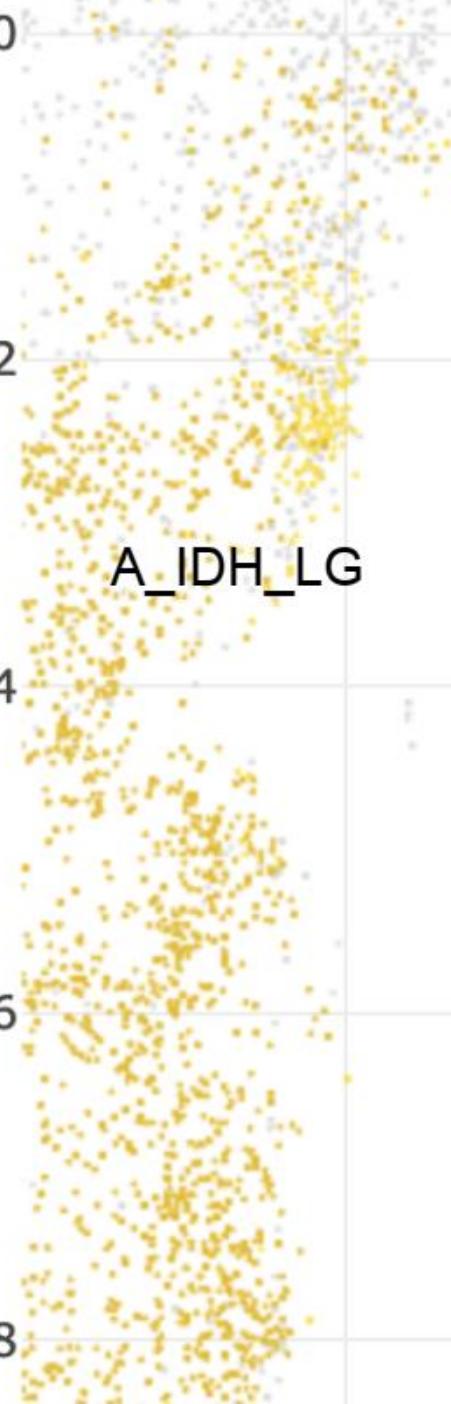
Methylation classes (Highest level >= 0.3, lower levels >= 0.1, all of lowest level)		Calibrated score	Interpretation	
Ependymal Tumours		0.85	no match	✗
	Posterior Fossa Ependymoma Group A	0.78	no match	✗
	Posterior Fossa Ependymoma Group A	0.68	no match	✗
	Mc Posterior Fossa Group A (pfa) Ependymoma Subclass 2b (novel)	0.67	no match	✗
	Mc Posterior Fossa Group A (pfa) Ependymoma Subclass 2c (novel)	0.00	no match	✗
	Mc Posterior Fossa Group A (pfa) Ependymoma Subclass 2a (novel)	0.00	no match	✗

Legend: ✓ Match (score >= 0.9) ✗ No match or low quality
● Match to MC family member (score >= 0.5)

Dx.: ependymóm, morfologicky klasického typu, s expresiou EZHIP a stratou expresie H3K27Mme3 a metylačným profilom konzistentným s ependymómom posterior fossa type A

EPN_PFA_2B

-20



A_IDH_LG

207558600099_R08C01_1



EPN_PFA_2C

-22

-24

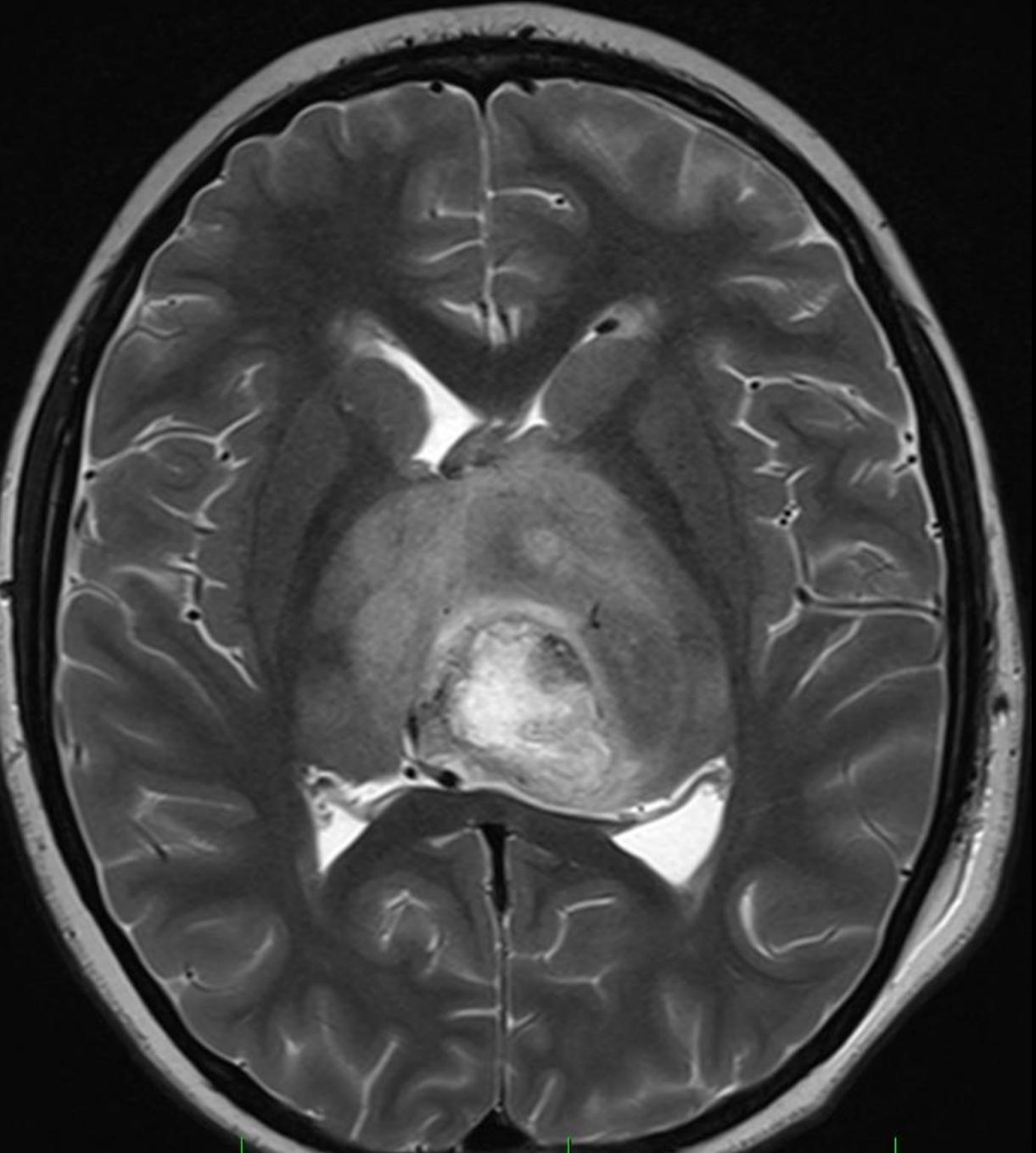
-26

-28

EPN_PFA_1C

EPN_PFA_1B

MB_SHH



Klinicky strednočiarový bitalamický tumor

Žiadny tumor v
zadnej jame

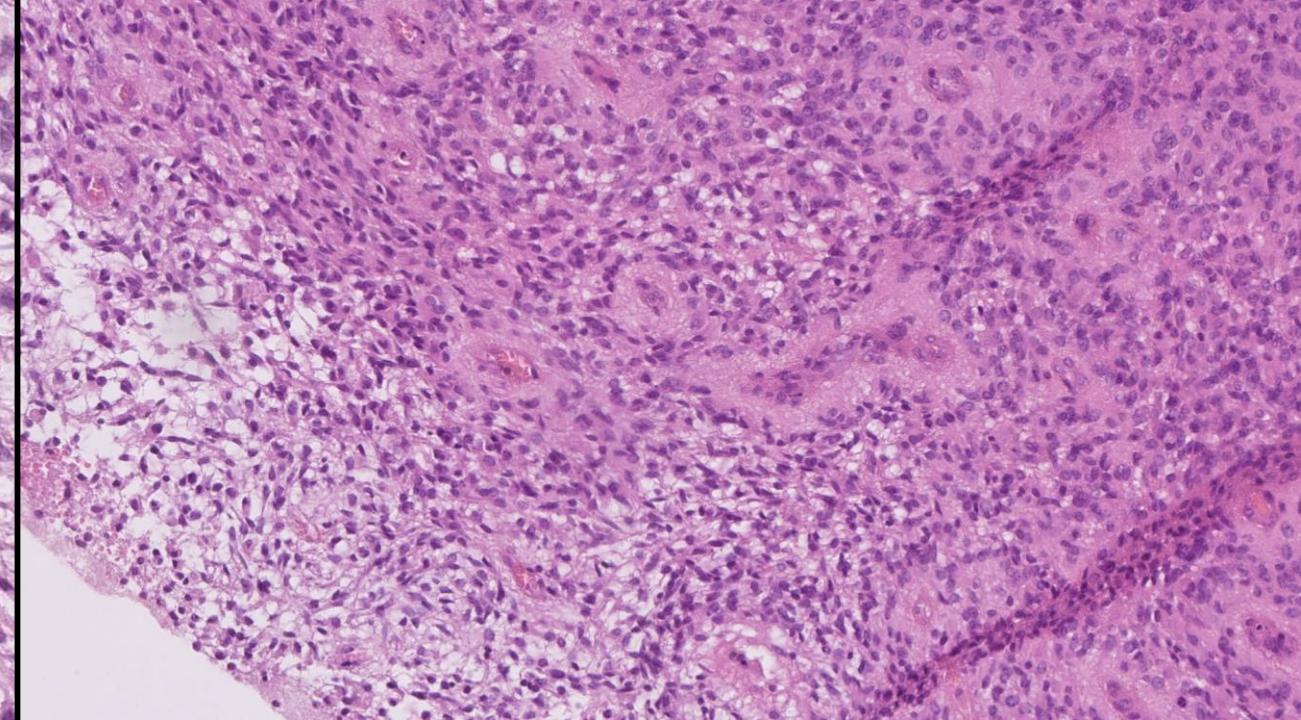
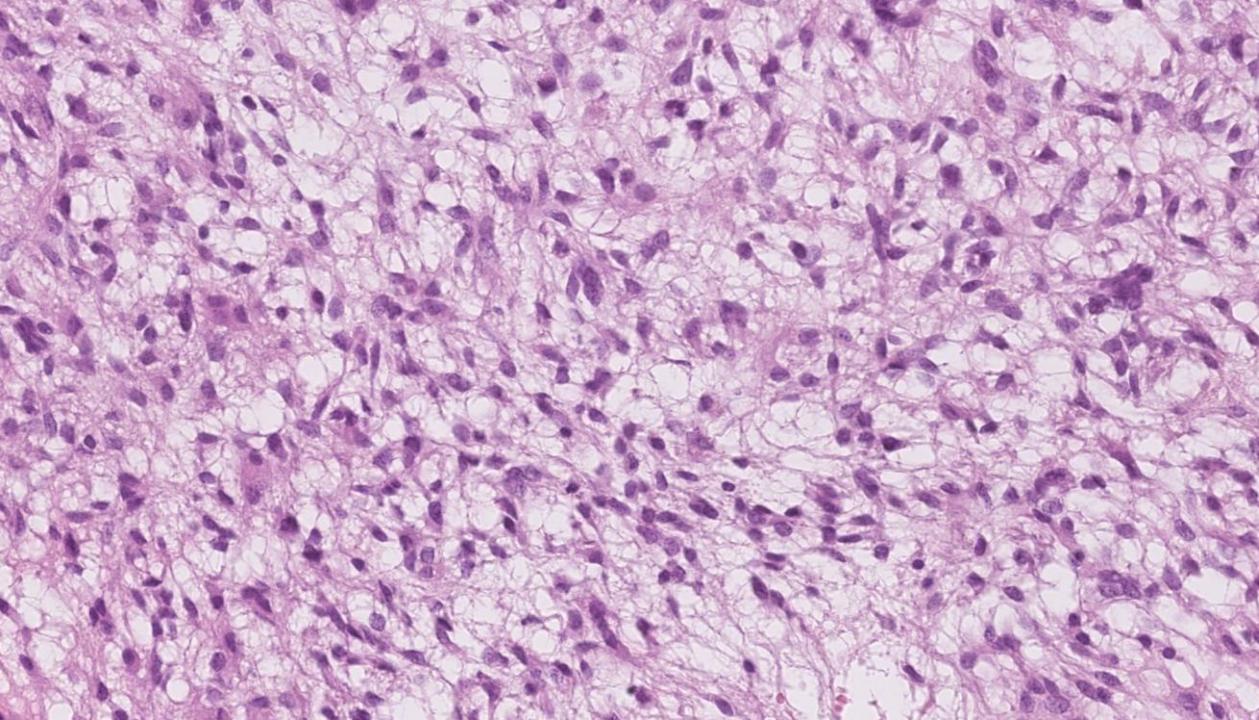
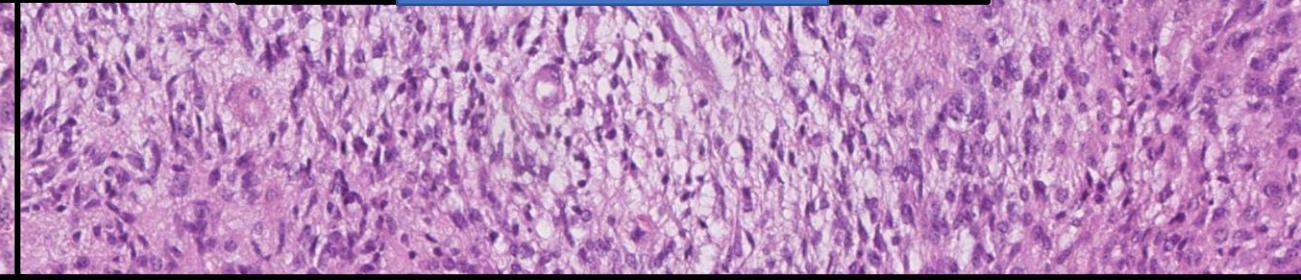
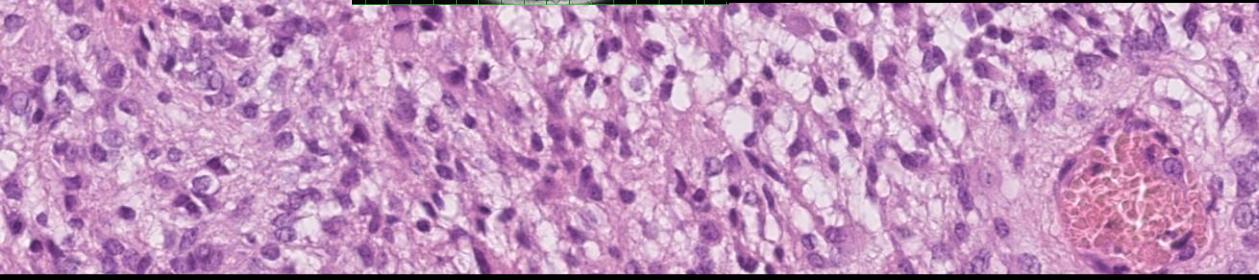
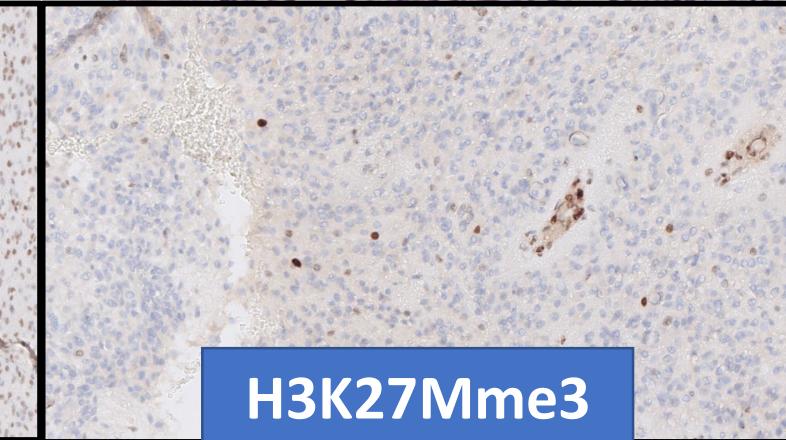
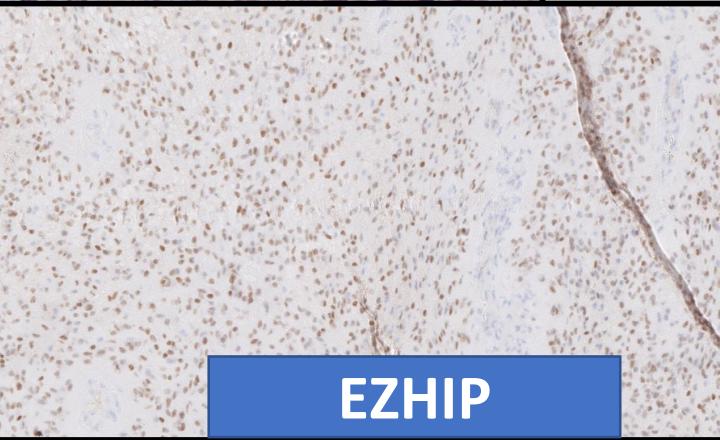
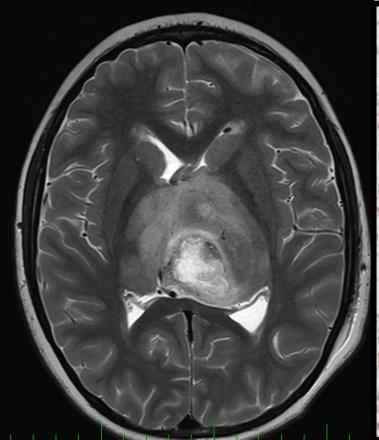


Zdravotnícke zariadenie: Národný ústav detských chorôb

Predmet vyšetrenia
a lokalizácia:

CITO!

1. kápsula tumoru - zadná časť
2. vnútorná časť tumoru - zadná časť (susp nekróza)
3. vnútorná časť tumoru - predná časť (tu robená endohistologia-LGG)

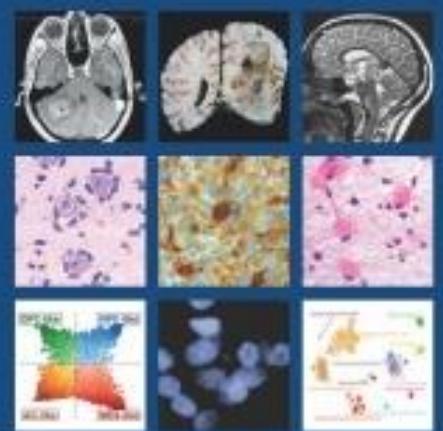


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Central Nervous System

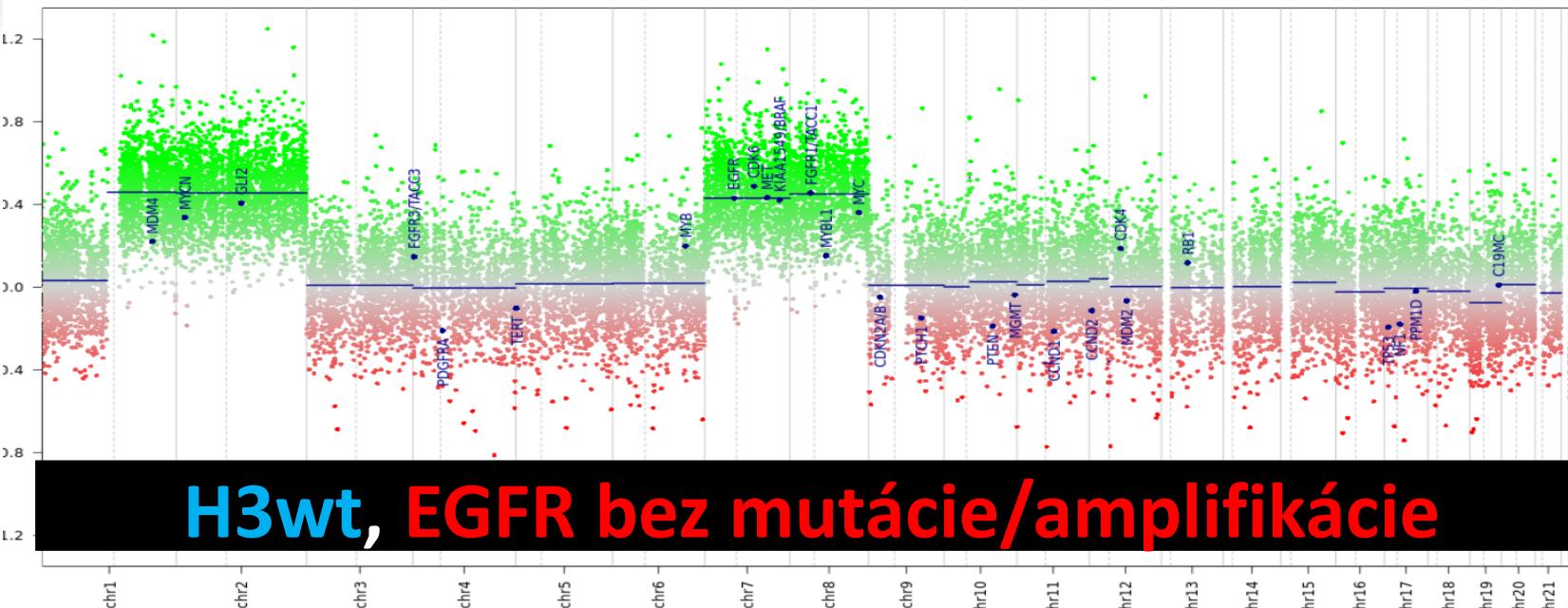
Tumours

Based on the WHO Classification of Tumours, International Agency for Research on Cancer

WHO Classification of Tumours
International Agency for Research on Cancer

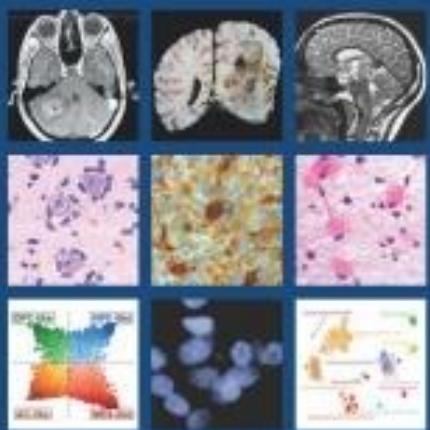
Definition

Diffuse midline glioma, H3 K27-altered, is an infiltrative midline glioma with loss of H3 p.K28me3 (K27me3) and usually either an **H3 c.83A>T p.K28M (K27M) substitution** in one of the histone H3 isoforms, **aberrant overexpression of EZHIP**, or an **EGFR mutation** (CNS WHO grade 4).



PROKAZUJEME klinicky významnou mutaci genu:

- **BCOR c.1550_1551del p.(Asn517ArgfsTer39) AF: 38%; - popísané v DMG aj EPE**
- **BCORL1 c.1825C>T p.(Arg609Ter) AF: 46%;**
- **ACVR1 c.617G>A p.(Arg206His) AF: 27%; - popísané v DMG**
- **LZTR1 c.1321C>T p.(Gln441Ter) AF: 95%.**

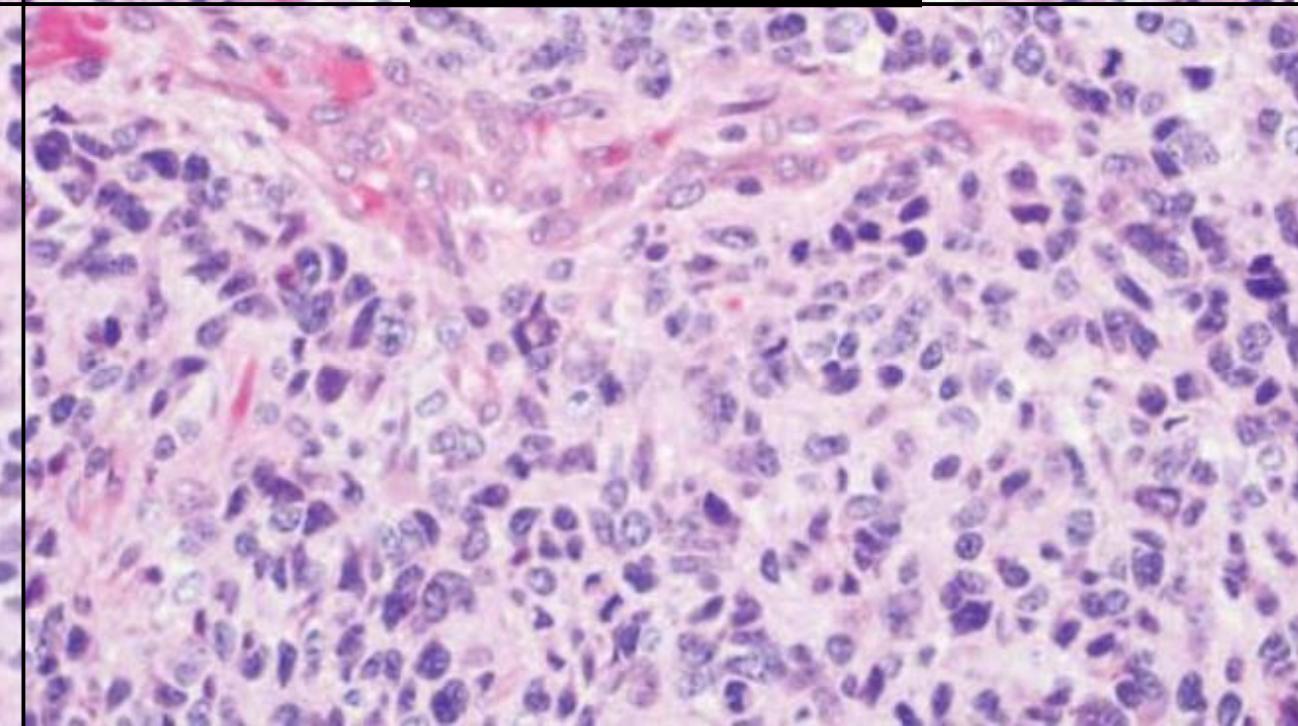
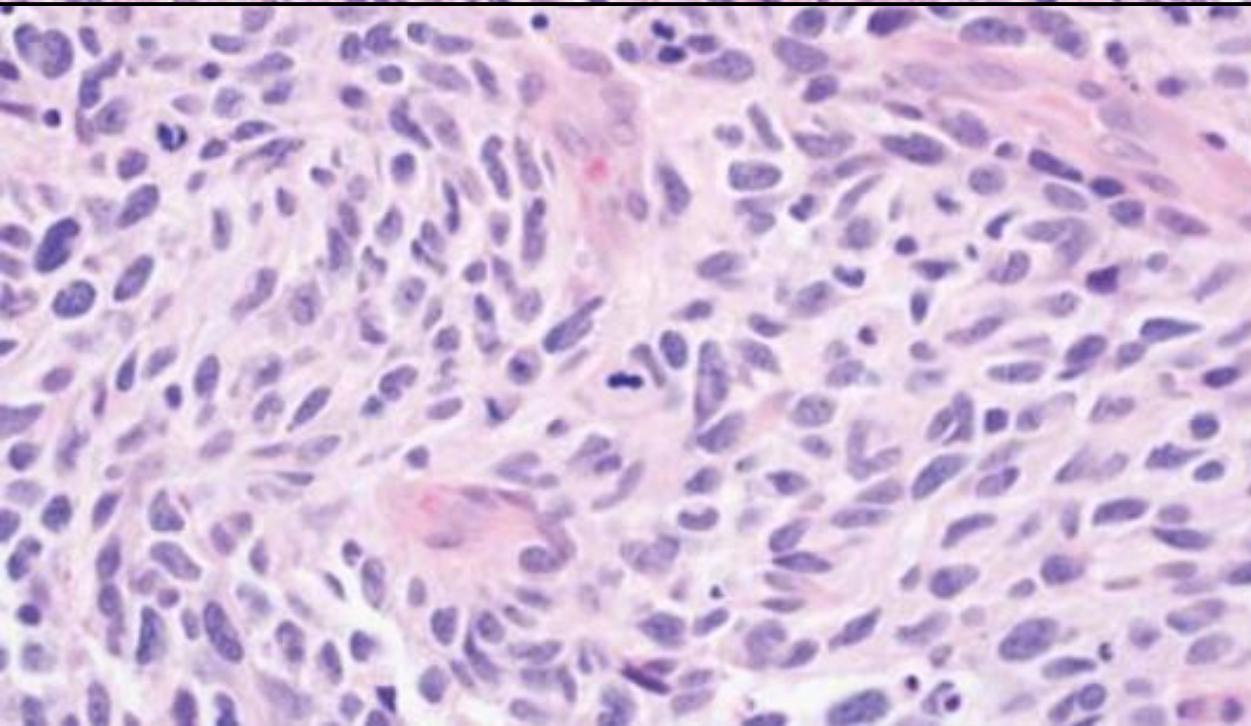
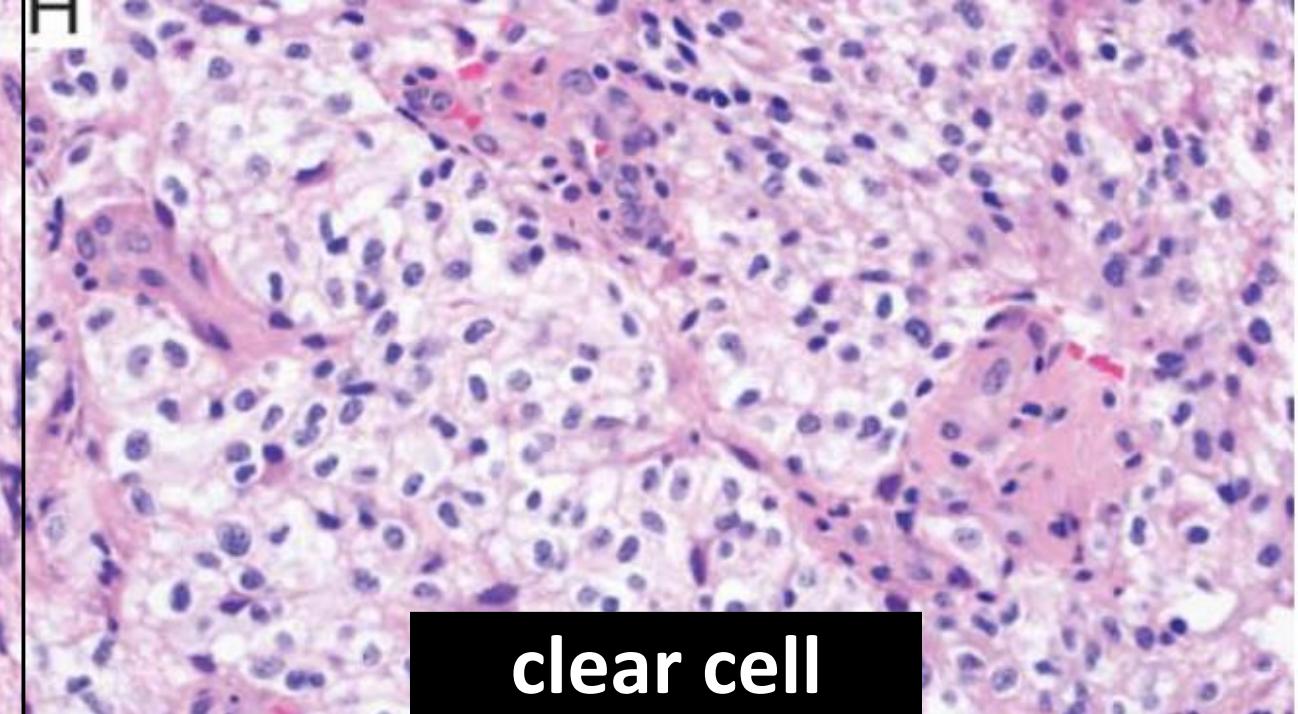
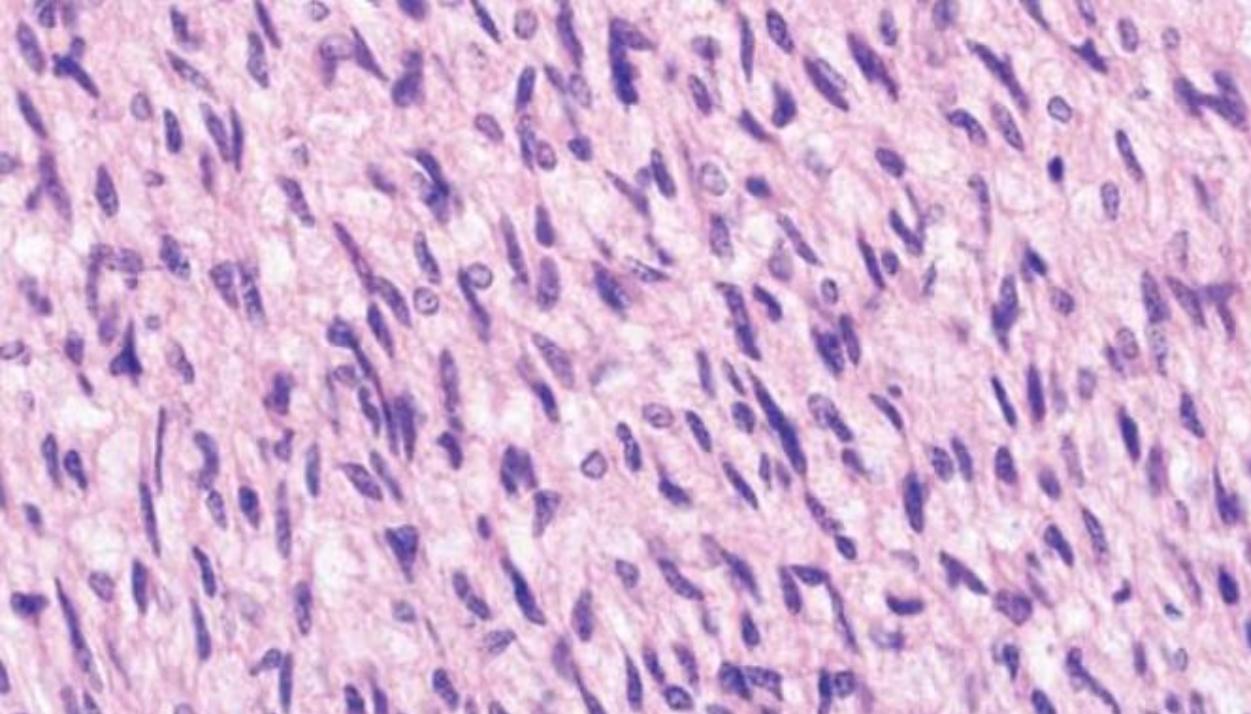


In posterior fossa group A (**PFA) ependymomas**, H3 p.K28 (K27) mutations are extremely rare, but loss of H3 p.K28me3 (K27me3) or EZHIP overexpression occurs; these tumours can be distinguished from **DMGs** on a morphological basis.

Diffuse Midline Gliomas with Histone H3-K27M Mutation: A Series of 47 Cases Assessing the Spectrum of Morphologic Variation and Associated Genetic Alterations

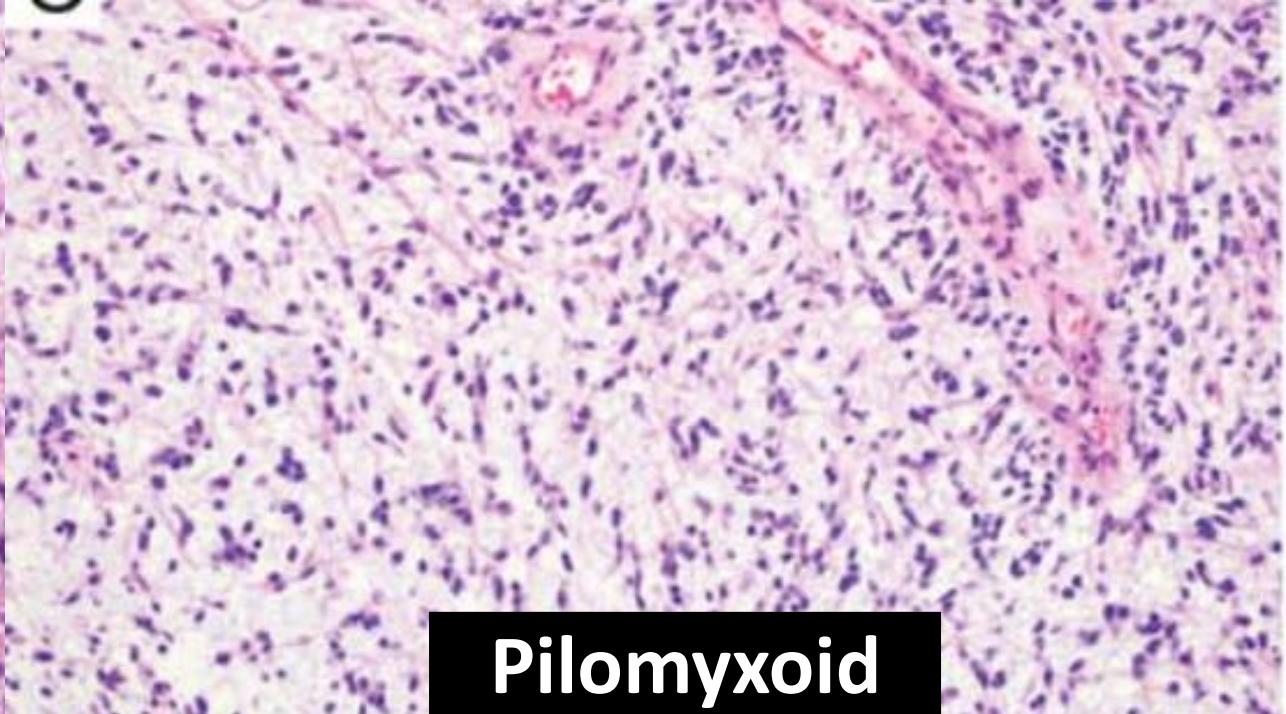
David A. Solomon¹; Matthew D. Wood¹; Tarik Tihan¹; Andrew W. Bollen¹; Nalin Gupta^{2,3};
Joanna J. J. Phillips^{1,2}; Arie Perry^{1,2}

Brain Pathol. 2016 Sep;26(5):569-80.

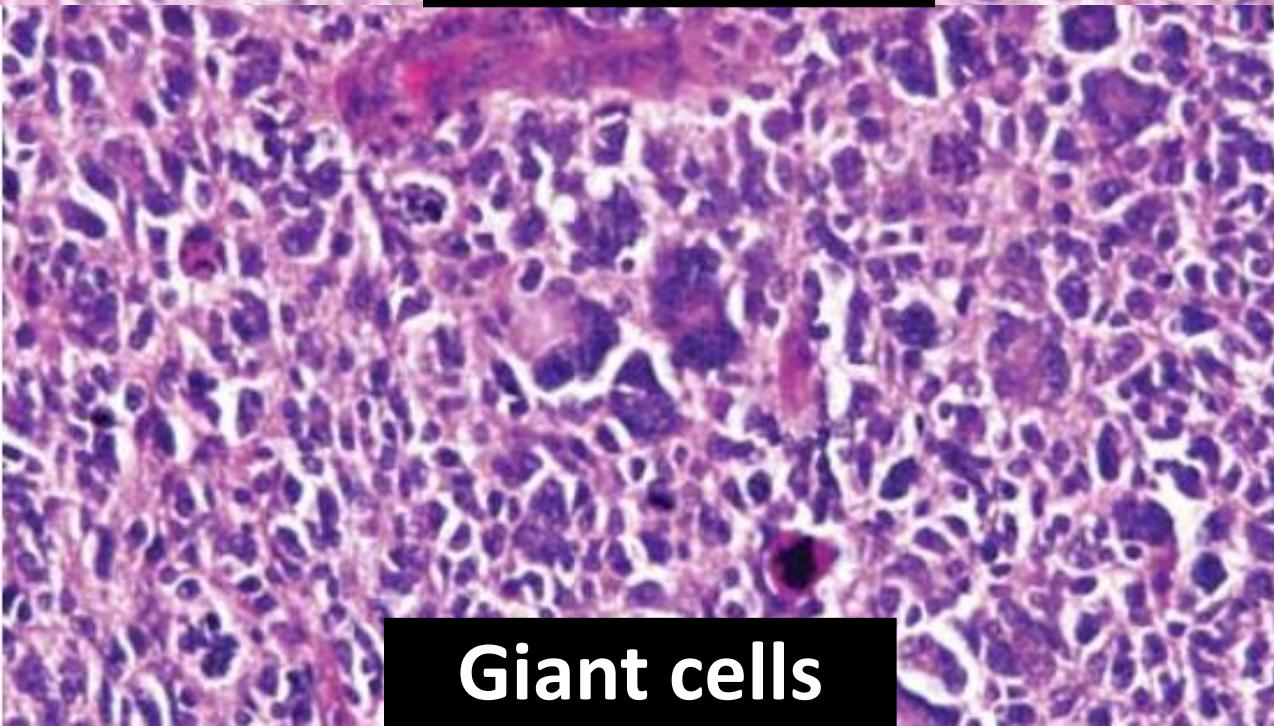




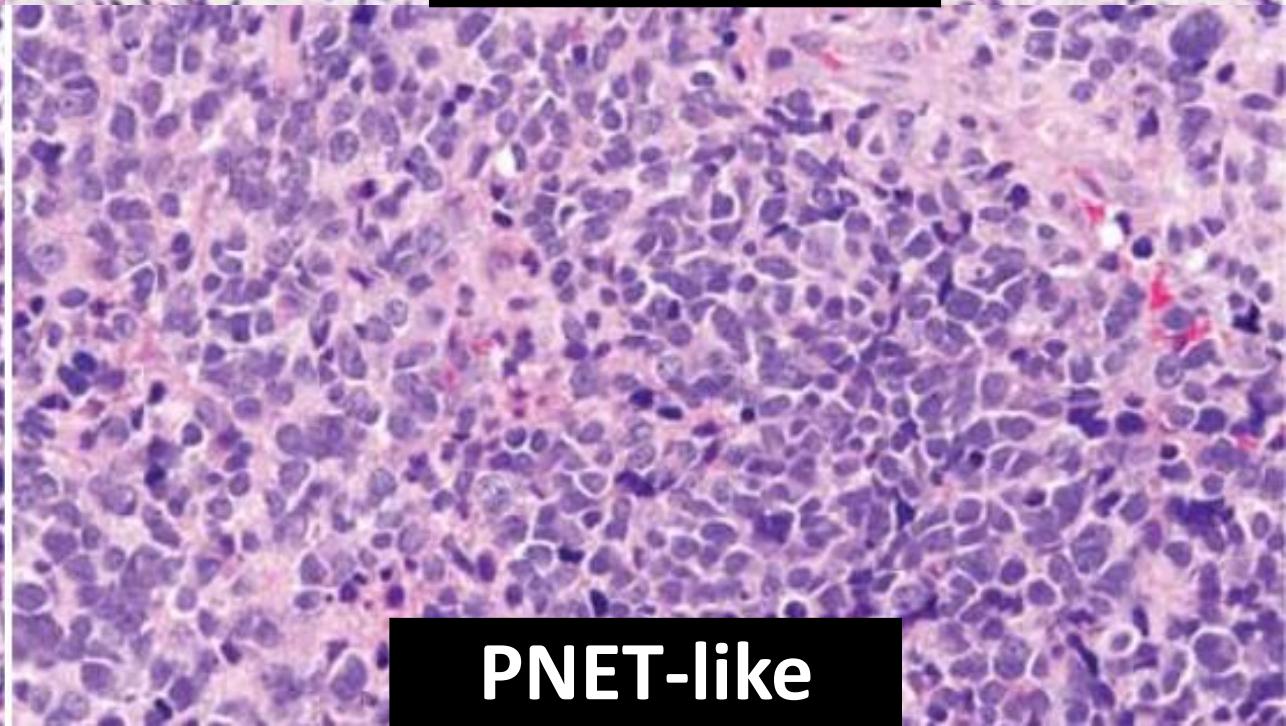
Epithelioid



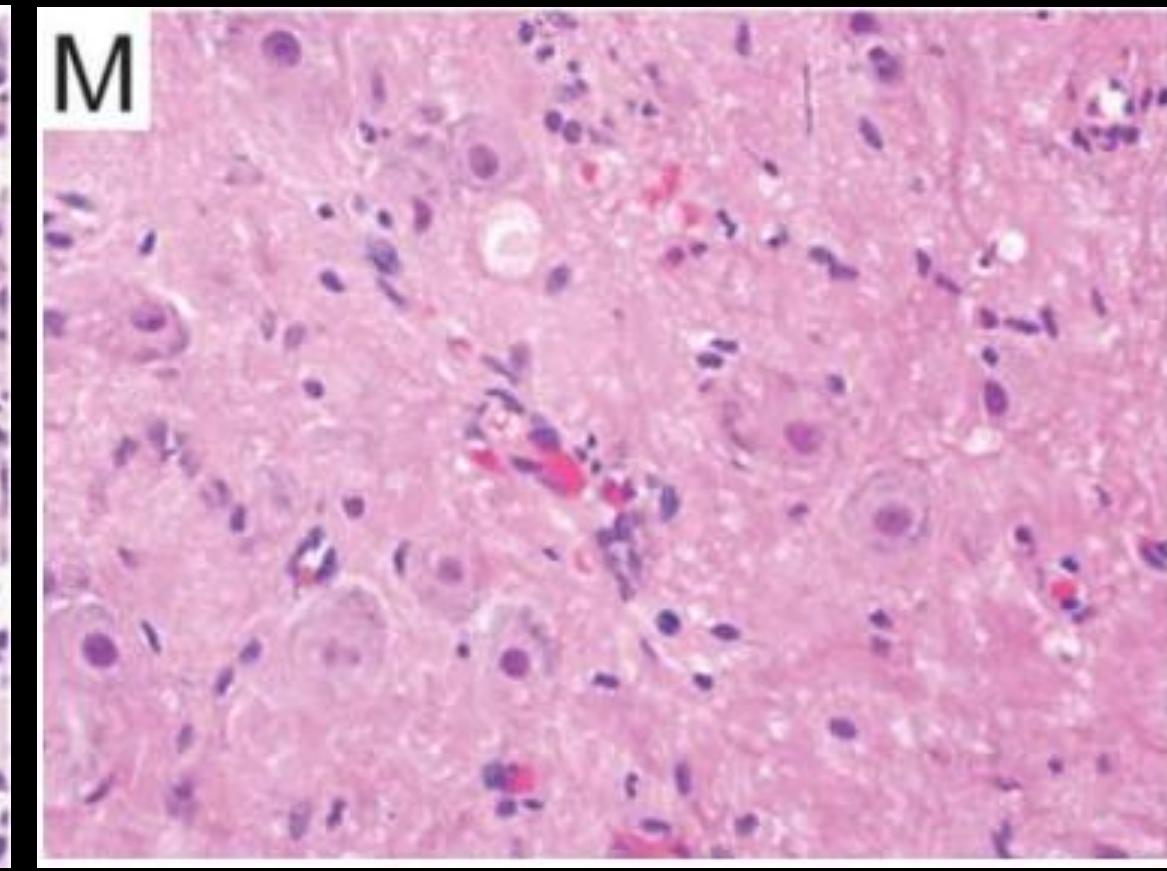
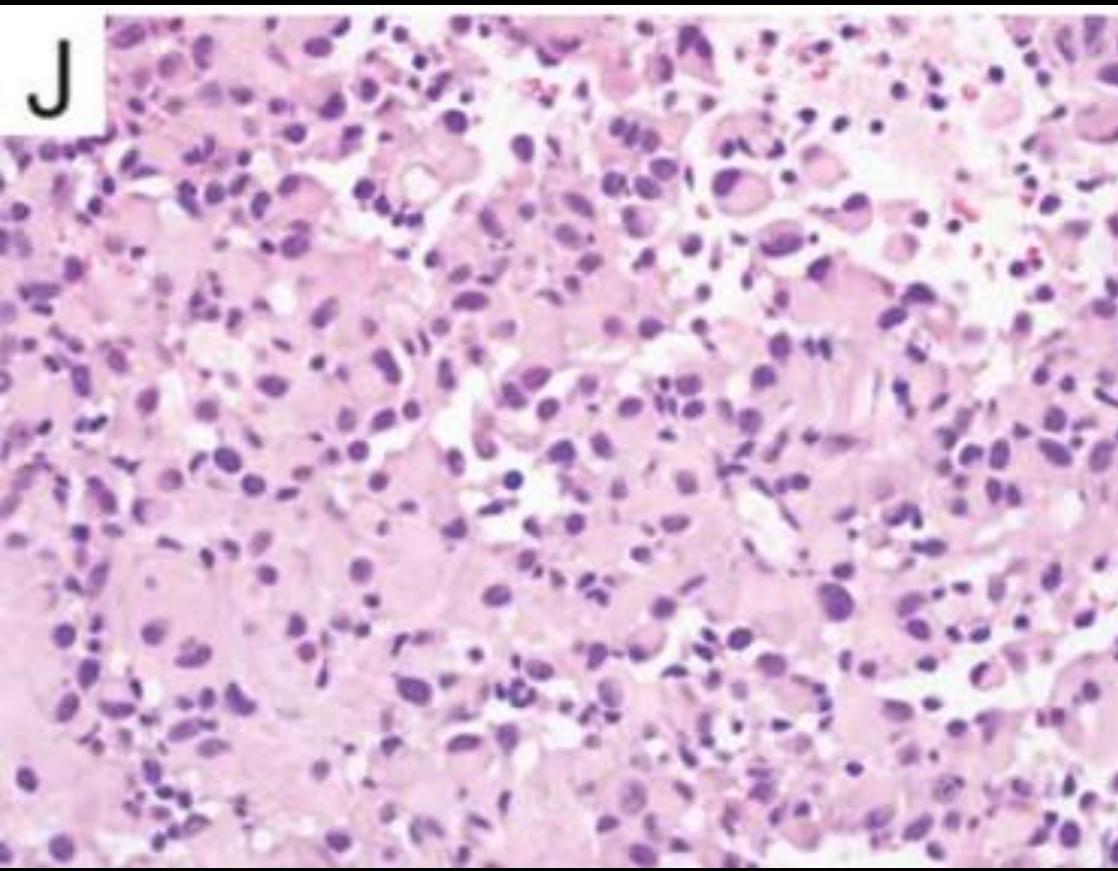
Pilomyxoid

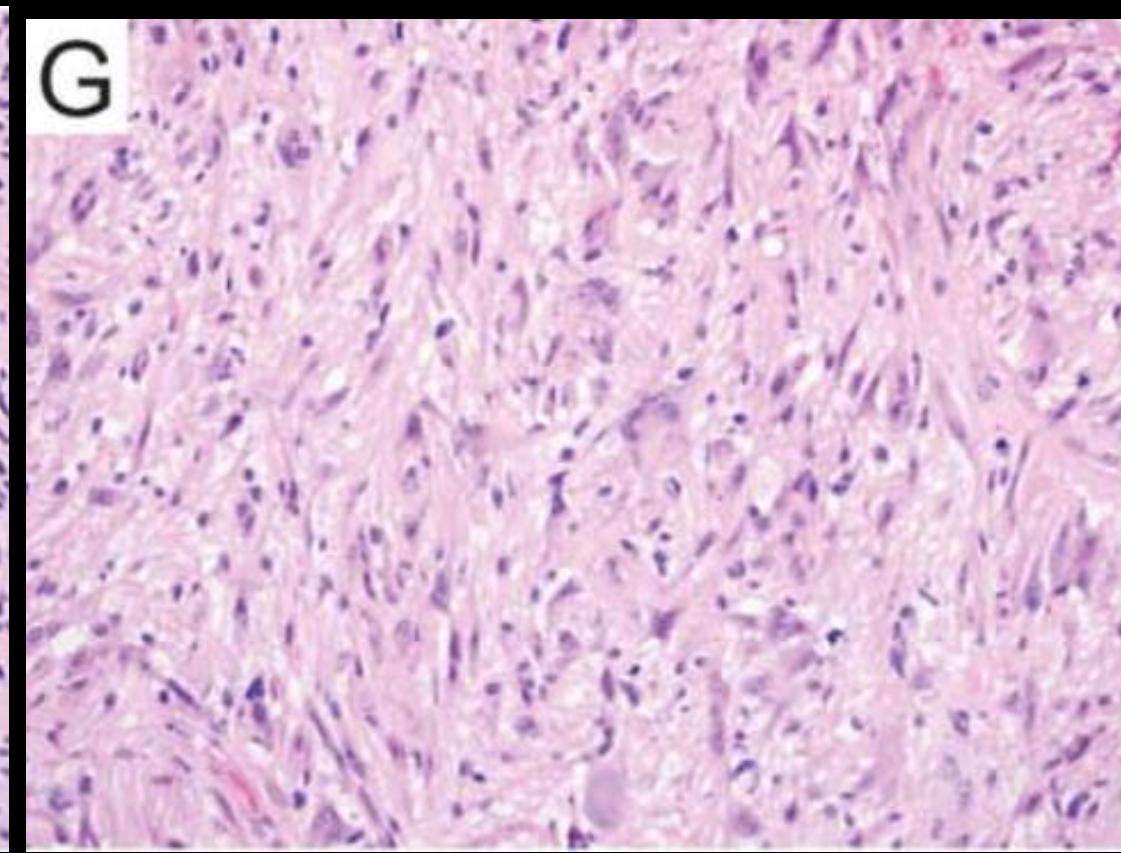
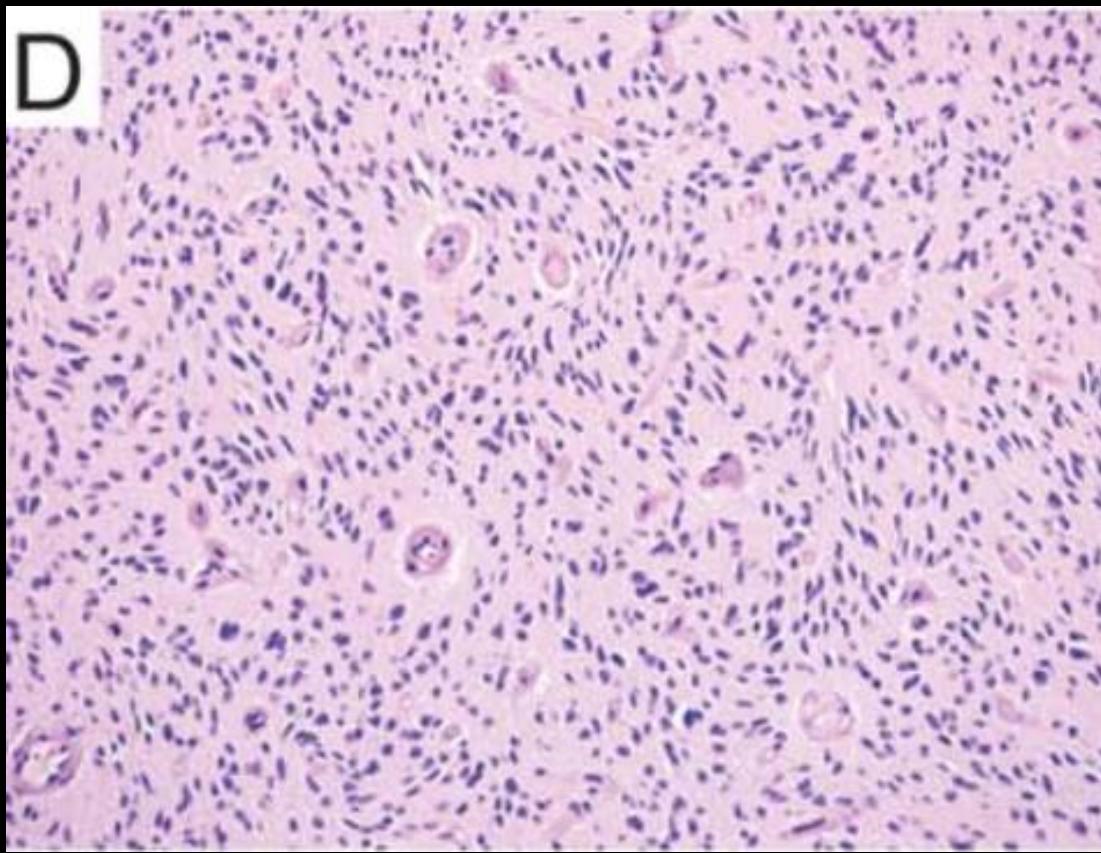


Giant cells

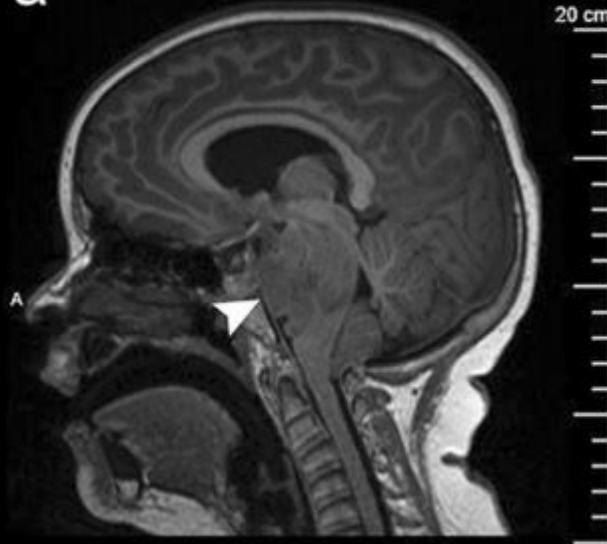


PNET-like





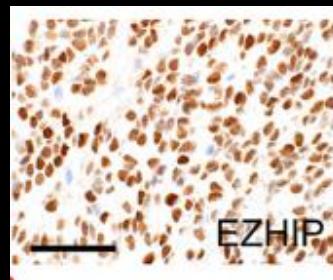
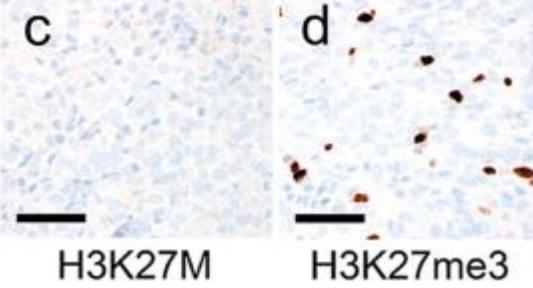
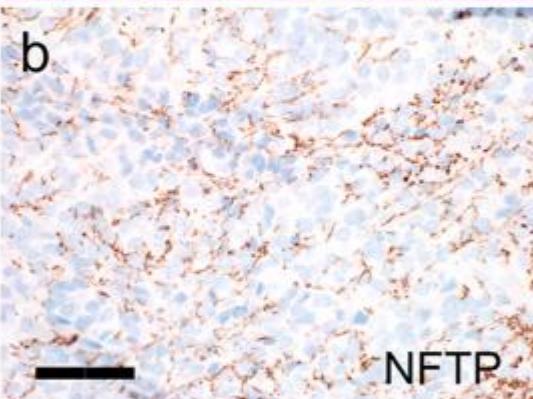
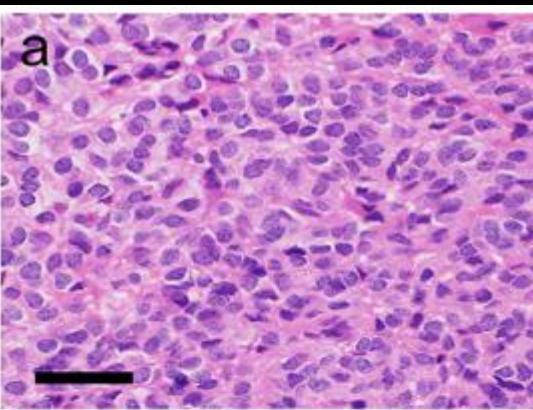
T1-weighted



C



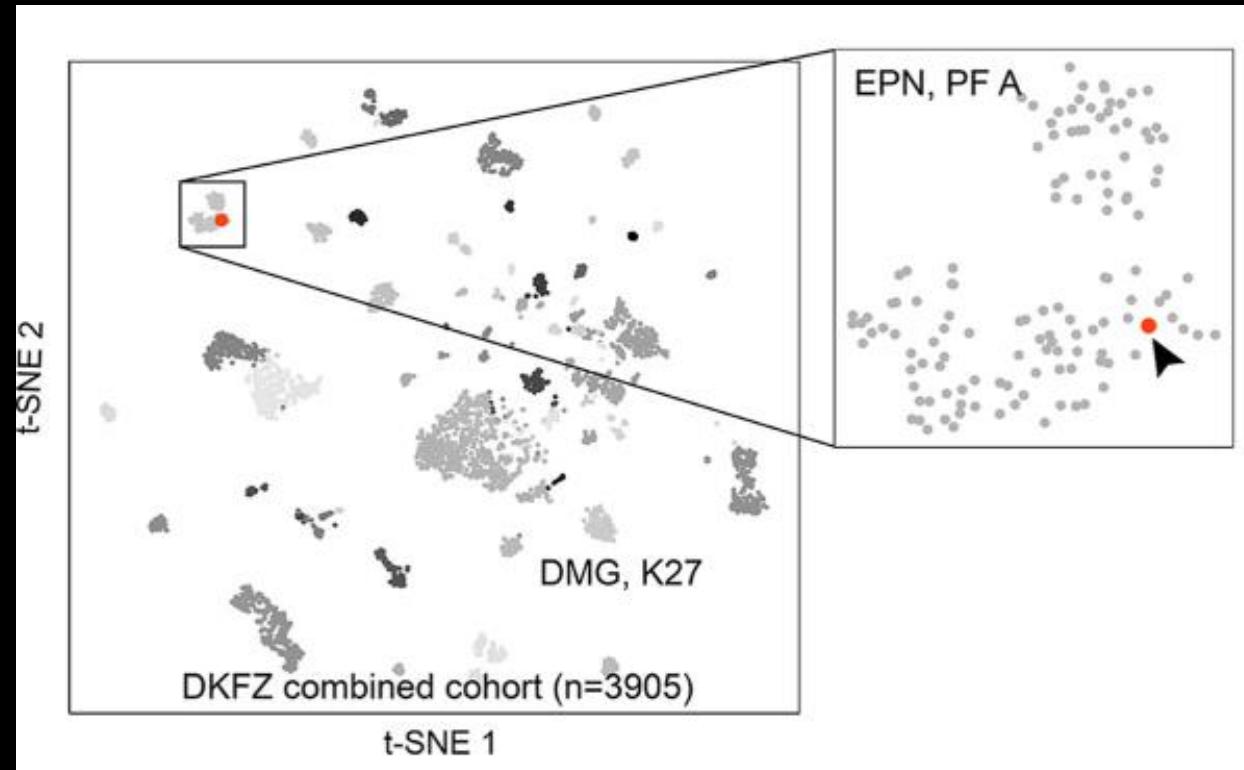
T2-weighted



LETTER TO THE EDITOR

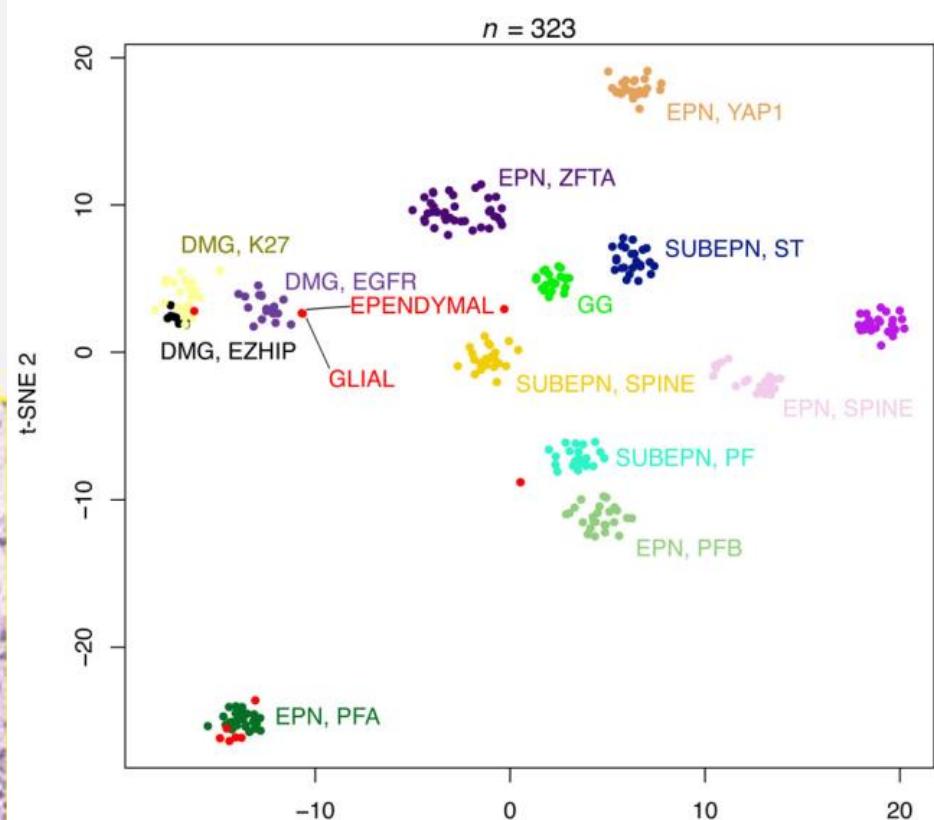
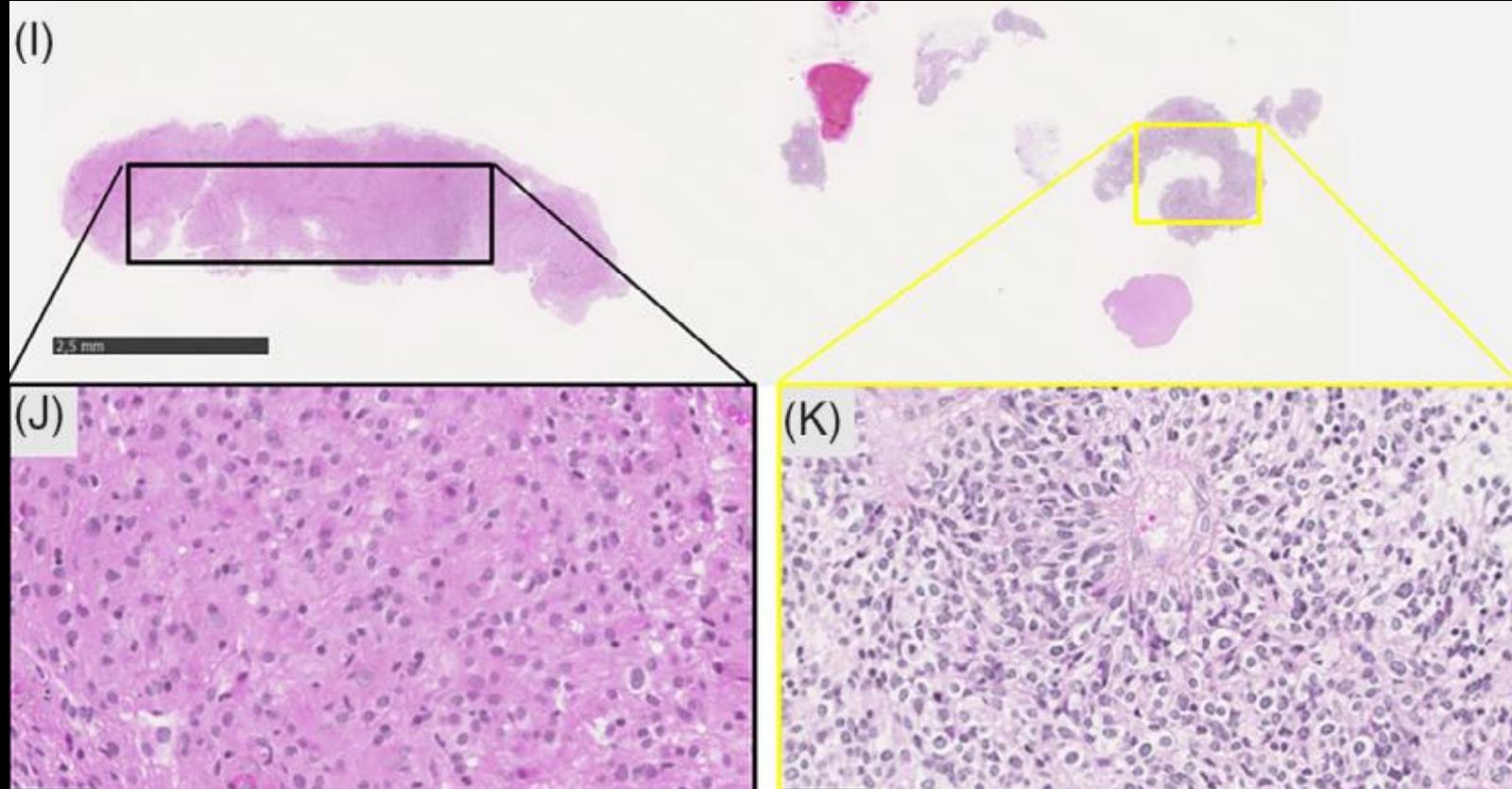
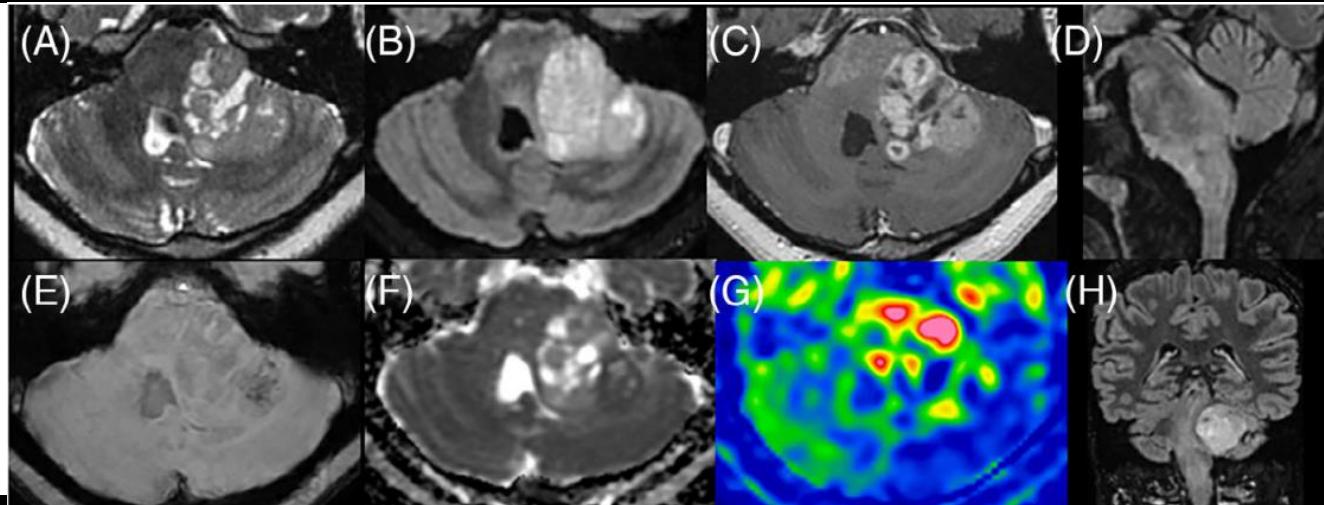
Diffuse intrinsic pontine glioma-like tumor with EZHIP expression and molecular features of PFA ependymoma

Pratt et al. Acta Neuropathologica Communications (2020) 8:37



The pontine diffuse midline glioma, *EGFR*-subtype with ependymal features: Yet another face of diffuse midline glioma, H3K27-altered

Arnault Tauziède-Espriat^{1,2}  | Alice Métais^{1,2} | Cassandra Mariet¹ | David Castel³ | Jacques Grill^{3,4} | Raphaël Saffroy⁵ | Lauren Hasty¹ | Volodia Dangouloff-Ros^{6,7} | Nathalie Boddaert^{6,7} | Sandro Benichi⁸ | Fabrice Chrétien¹ | Pascale Varlet^{1,2}

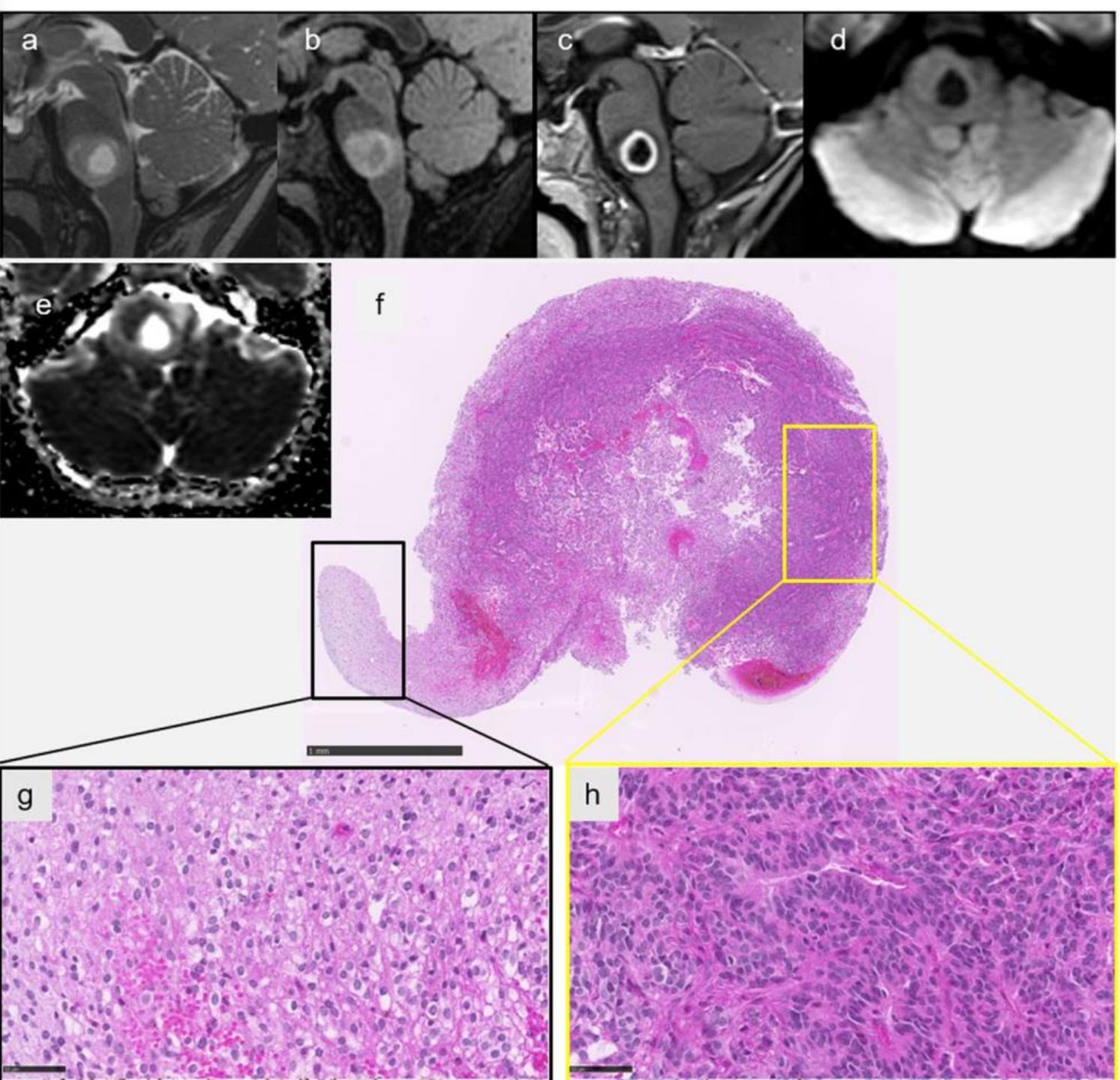


Posterior fossa ependymoma H3 K27-mutant: an integrated radiological and histomolecular tumor analysis

Cassandra Mariet¹, David Castel², Jacques Grill^{1,2}, Raphaël Saffroy³, Volodia Dangouloff-Ros^{4,5}, Nathalie Boddaert^{4,5}, Francisco Llamas-Gutierrez⁶, Céline Chappé⁷, Stéphanie Puget⁸, Lauren Hasty⁹, Fabrice Chrétien⁹, Alice Métais^{9,10}, Pascale Varlet^{9,10} and Arnault Tauziède-Espriat^{9,10*}

Acta Neuropathol Commun. 2022 Sep 14;10(1):137

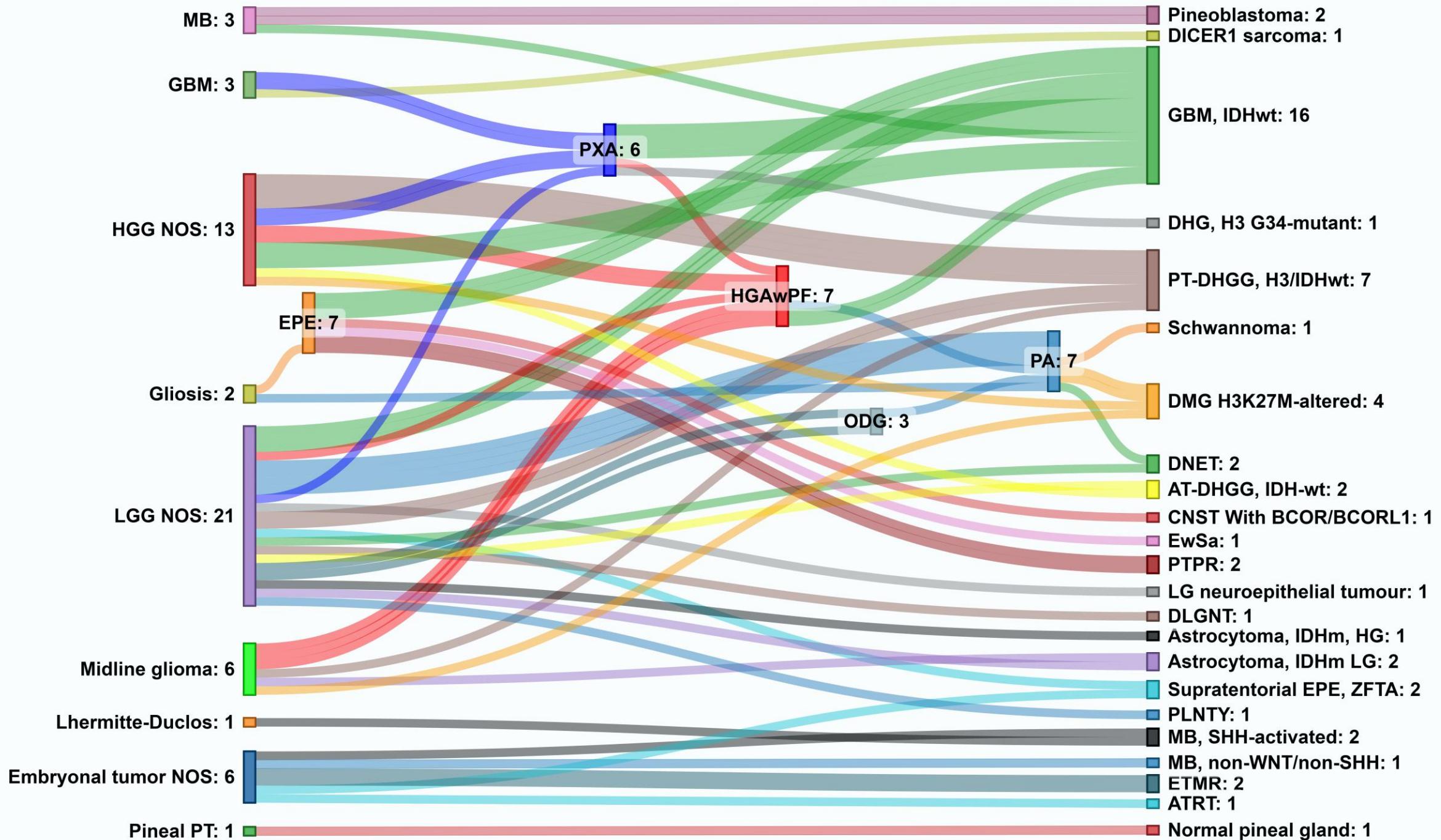
- Seven tumors clustered within EPN_PFA by DNA-methylation analysis
- Among the two remaining cases, one was reclassified as a DMG and the last was unclassified
- The diagnosis of EPN_PFA must include tumor location, growth pattern, Olig2 expression, and DNA-methylation profiling before it can be differentiated from DMG, H3 K27-altered



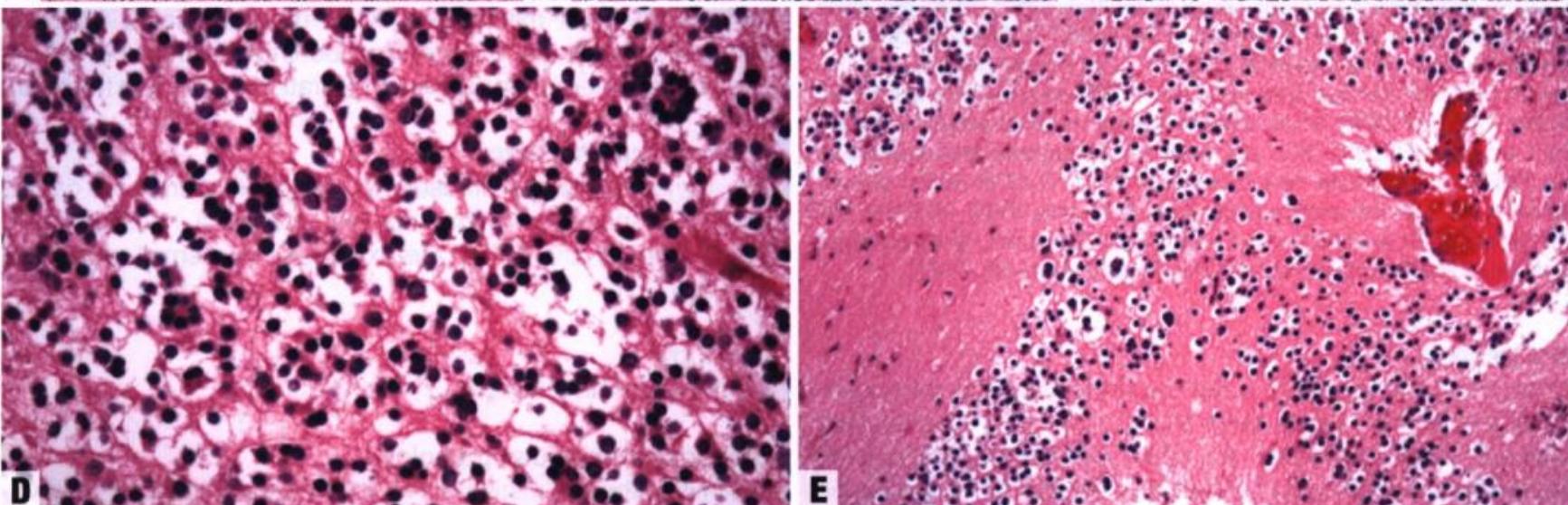
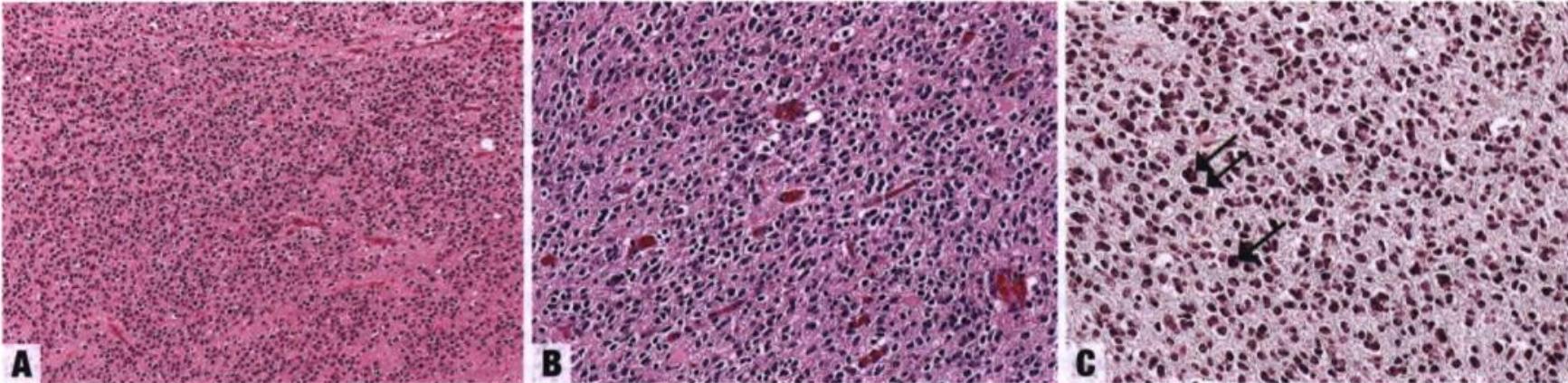
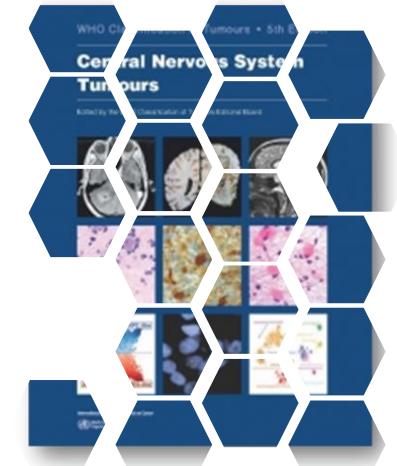
Naše skúsenosti s metylačným profilovaním za 2,5 roka: 216 nádorov CNS

206 analyzovateľných, metylóm užitočný v 78.2% prípadov

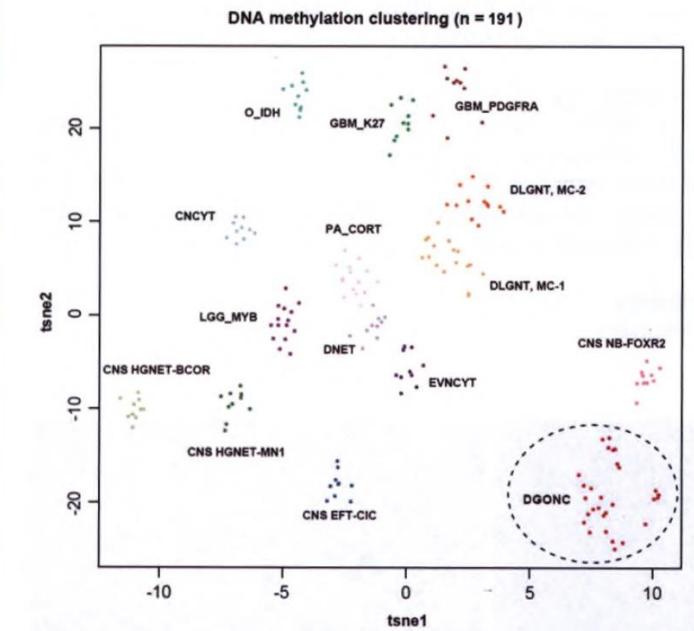
Diagnóza pomocou metylómu/významná zmena diagnózy	N=58 (28.2%)	Including 3 cases with CS ≥ 0.3 and < 0.5, and MC supporting final diagnosis
Vyriešenie diferenciálnej diagnózy	N=20 (9.7%)	Including 1 case with CS ≥ 0.3 and < 0.5, and MC supporting final diagnosis
Potvrdenie preferovanej diagnózy	N=29 (14.1%)	including 4 cases with CS ≥ 0.3 and < 0.5, and MC supporting final diagnosis
Úspešná molekulová subklasifikácia (MB, EPE)	N=54 (26.2%)	41 medulloblastomas, 10 ependymomas, 2 meningiomas, 1 AT/RT
Nediagnostický / nejednoznačný/neprínosný výsledok	N=44 (21,3%)	no CS greater 0.3 (N=6), CS ≥ 3 and < 0.5 and no other supportive findings (N=19), 9 cases with CS ≥ 0.5 < 0.84 and 10 cases with CS ≥ 0.84
Zavádzajúci výsledok	N=1 (0.5%)	



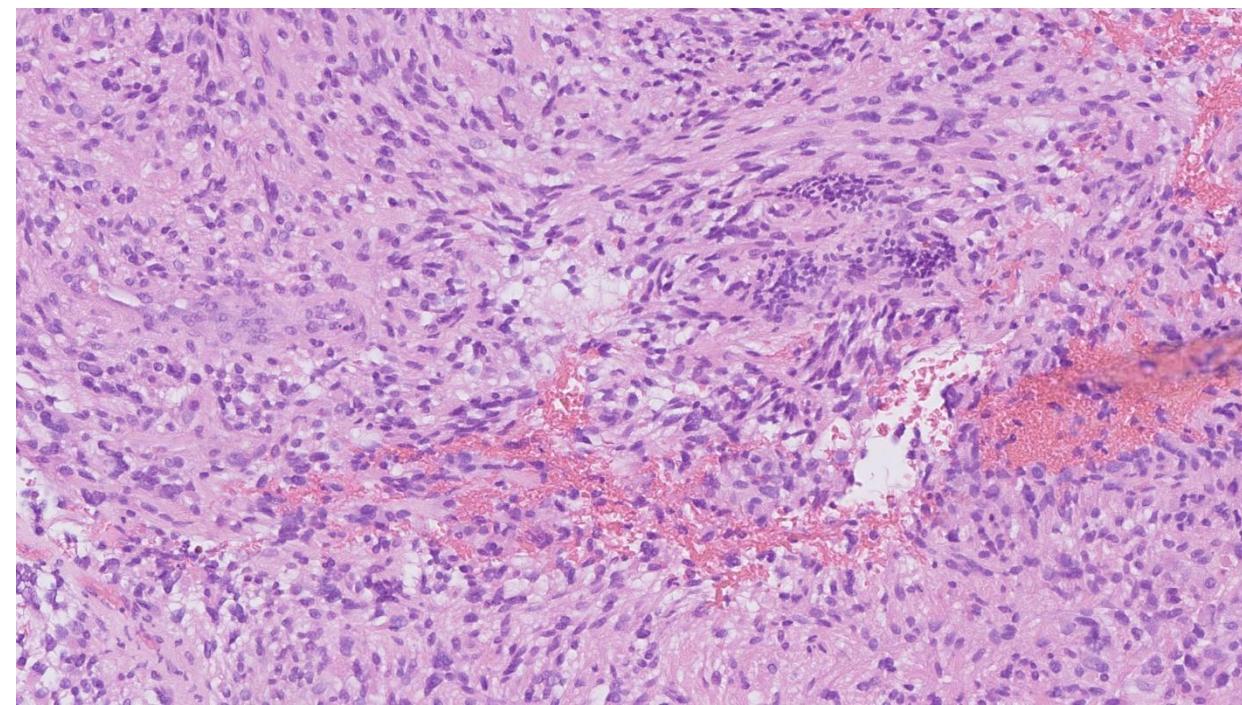
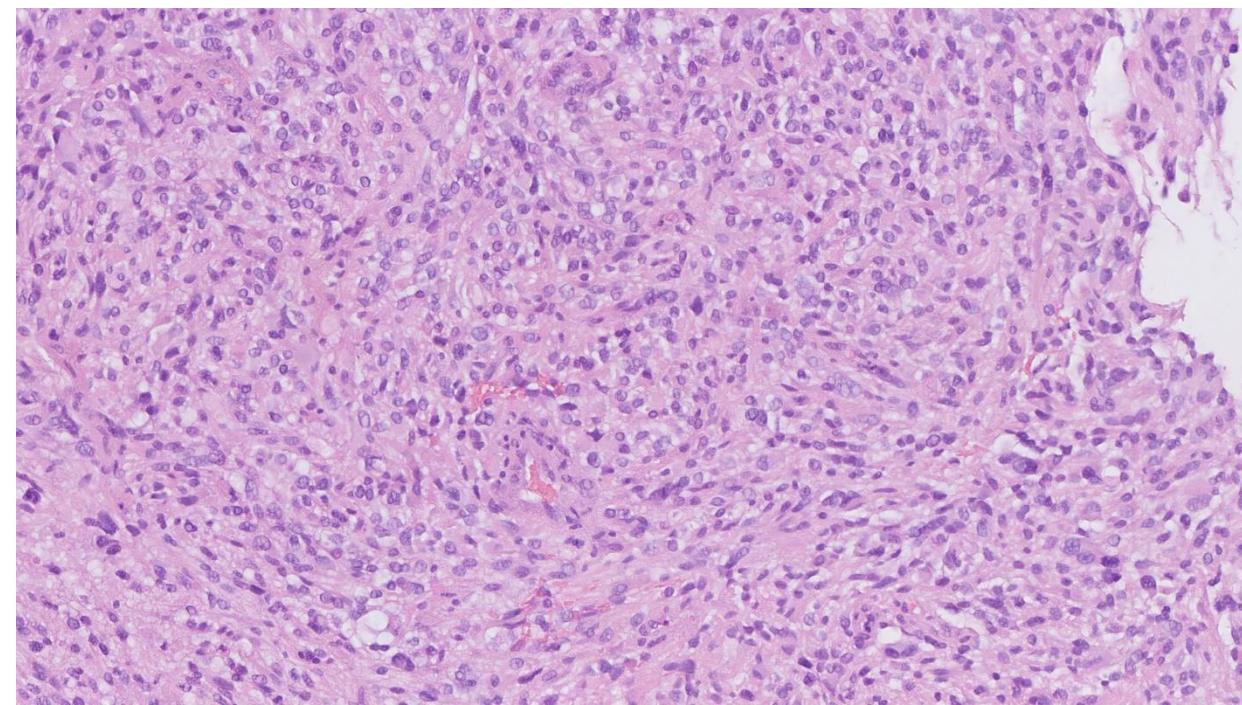
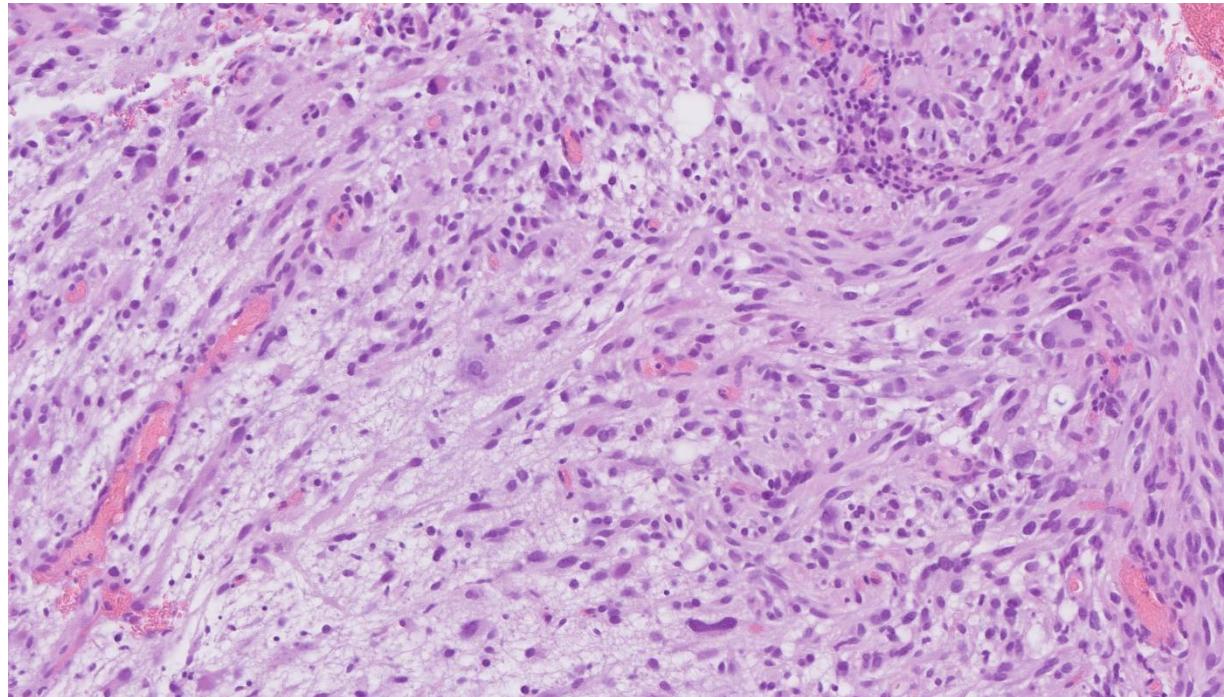
Diffuse glioneuronal tumor with ODG-like features and nuclear clusters(DGONC)



monozómia 14
žiadne ďalšie alterácie



**82-ročný muž,, konzult. Slovinsko,
tumor ponsu a cerebella (171148/22)**



**MC: High grade astrocytoma with piloid
features**

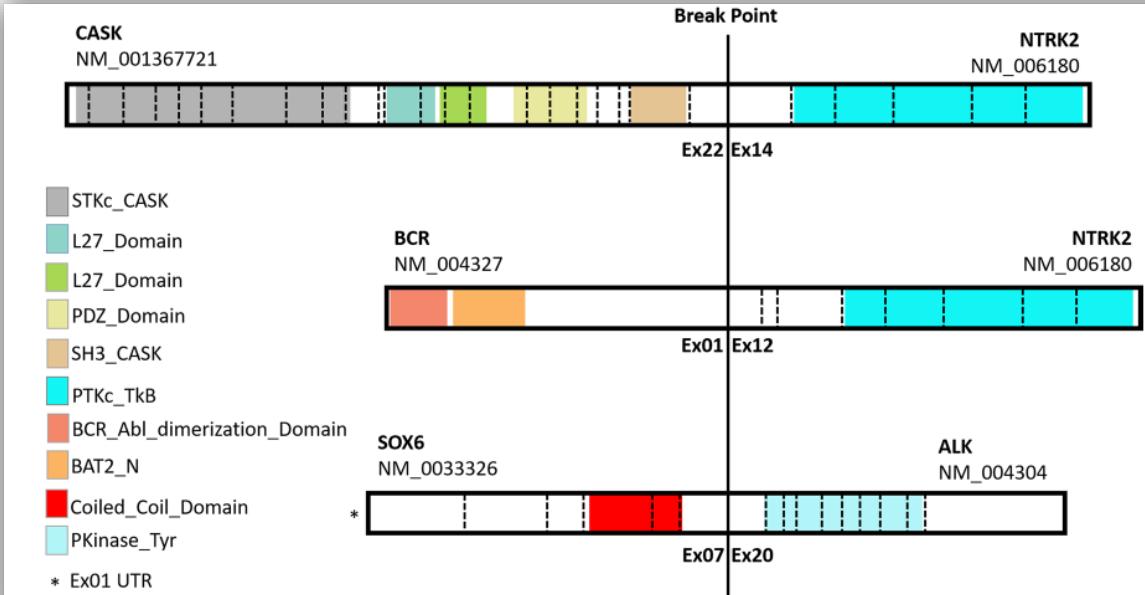
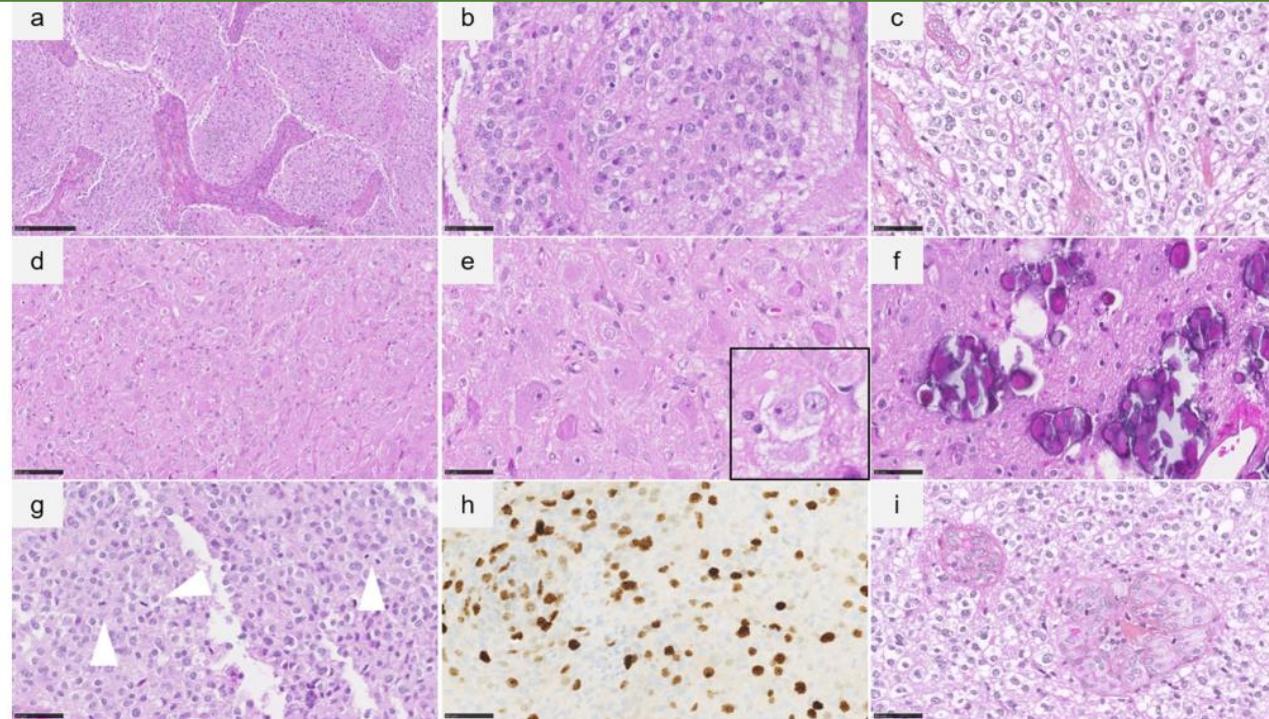
CORRESPONDENCE



Clinicopathological and molecular characterization of three cases classified by DNA-methylation profiling as “Glioneuronal Tumors, NOS, Subtype A”

Arnault Tauziède-Espriat^{1,2} · Volodia-Dangouloff-Ros^{3,4} · Dominique Figarella-Branger⁵ ·
Emmanuelle Uro-Coste^{6,7,8} · Yvan Nicaise⁶ · Nicolas André^{9,10} · Didier Scavarda¹¹ · Benoît Testud¹² · Nadine Girard¹² ·
Audrey Rousseau^{13,14} · Laetitia Basset^{13,14} · Guillaume Chotard¹⁵ · Vincent Jecko¹⁶ · François le Loarer^{17,18} ·
Isabelle Hostein^{17,18} · Marie-Christine Machet¹⁹ · Matthias Tallegas¹⁹ · Antoine Listrat²⁰ · Lauren Hasty¹ ·
Alice Métais^{1,2} · Fabrice Chrétien¹ · Nathalie Boddaert^{3,4} · Pascale Varlet^{1,2} on behalf of the RENOCLIP-LOC

not included in WHO 2021



Tumor detského veku, potenciálne agresívny,
ale prognóza aktuálne neistá

Identifikateľný iba metylačným profilovaním
fúzie NTRK a ALK

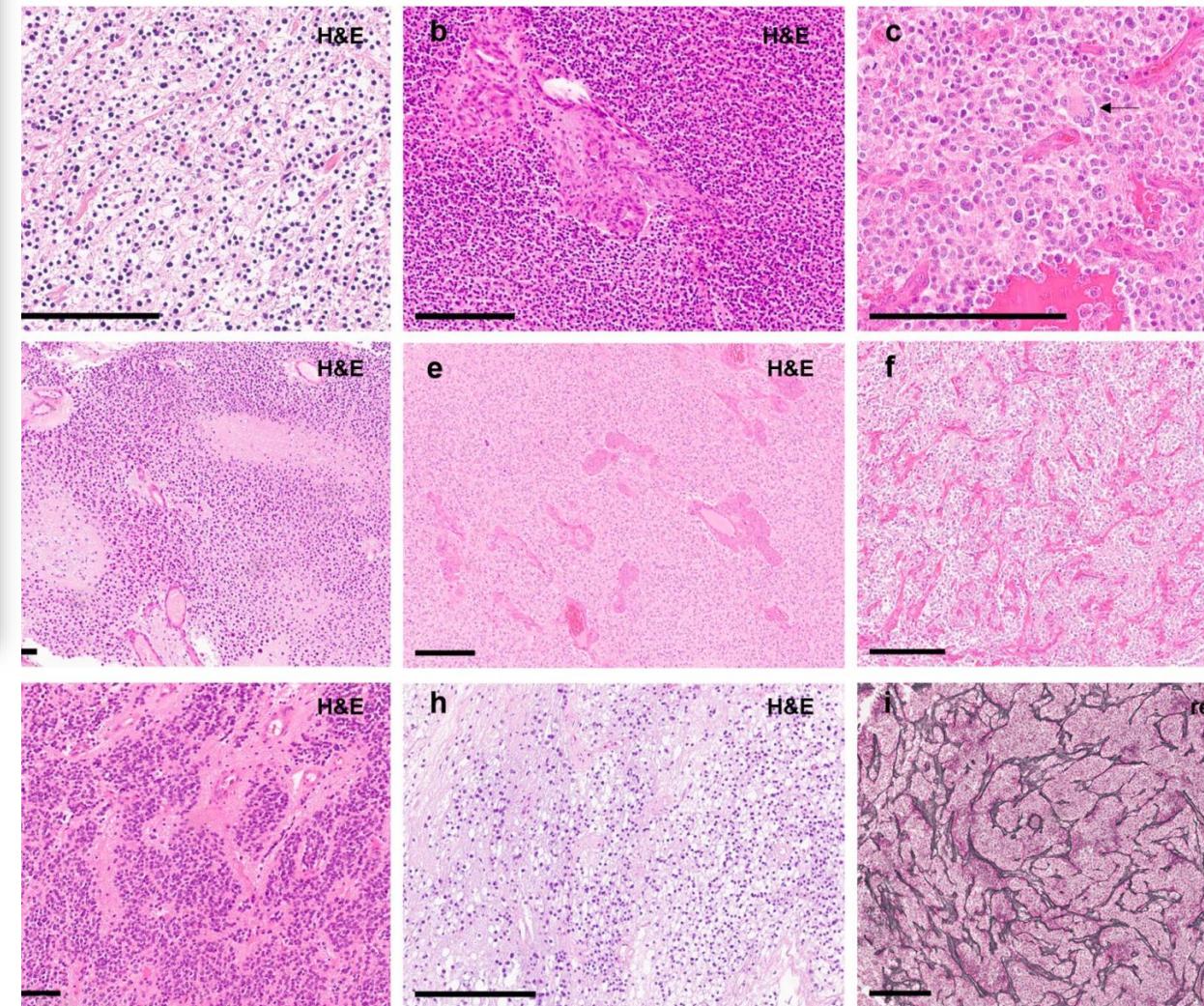


Glioneuronal tumor with ATRX alteration, kinase fusion and anaplastic features (GTAKA): a molecularly distinct brain tumor type with recurrent NTRK gene fusions

Henri Bogumil^{1,2} · Martin Sill^{3,4} · Daniel Schrimpf^{1,2} · Britta Ismer^{3,5,6} · Christina Blume^{1,2} · Ramin Rahmazade^{1,2} · Felix Hinz^{1,2} · Asan Cherkezov^{1,2} · Rouzbeh Banan^{1,2} · Dennis Friedel^{1,2} · David E. Reuss^{1,2} · Florian Selt^{3,7,8,9} · Jonas Ecker^{3,7,8,9} · Till Milde^{3,7,8,9} · Kristian W. Pajtler^{3,4,8} · Jens Schittenhelm^{10,11,12} · Jürgen Hench¹³ · Stephan Frank¹³ · Henning B. Boldt^{14,15} · Bjarne Winther Kristensen^{14,15,16,17} · David Scheie¹⁸ · Linea C. Melchior¹⁸ · Viola Olesen¹⁹ · Astrid Sehested²⁰ · Daniel R. Boué²¹ · Zied Abdullaev²² · Laveniya Satgunaseelan²³ · Ina Kurth²⁴ · Annekatrin Seidlitz^{9,25,26,27,28,29,30,31} · Christine L. White^{32,33,34} · Ho-Keung Ng^{35,36} · Zhi-Feng Shi^{36,37} · Christine Haberler³⁸ · Martina Deckert³⁹ · Marco Timmer⁴⁰ · Roland Goldbrunner⁴⁰ · Arnault Tauziède-Espriat^{41,42} · Pascale Varlet^{41,42} · Sebastian Brandner^{43,44} · Sanda Alexandrescu⁴⁵ · Matija Snuderl⁴⁶ · Kenneth Aldape²² · Andrey Korshunov^{1,2,3} · Olaf Witt^{3,7,8,9} · Christel Herold-Mende⁴⁷ · Andreas Unterberg⁴⁸ · Wolfgang Wick^{49,50} · Stefan M. Pfister^{3,4,8,9} · Andreas von Deimling^{1,2} · David T. W. Jones^{3,5} · Felix Sahm^{1,2,3,9} · Philipp Sievers^{1,2}

not included in WHO 2021

Supratentoriálny tumor, vek 4 – 76 rokov
(median 19 rokov), potenciálne agresívny
tumor, rekurencia priemerne po ~ 23
mesiacoch



Identifikateľný iba metylačným profilovaním; alterácie ATRX, fúzie NTRK, FGFR1, EGFR, BRAF, MET; homozygotná delécia CDKN2A/B (50 %)

ORIGINAL PAPER

Amplification of the PLAG-family genes—*PLAGL1* and *PLAGL2*—is a key feature of the novel tumor type *CNS embryonal tumor with PLAGL amplification*

Michaela-Kristina Keck^{1,2} · Martin Sill^{1,3} · Andrea Wittmann^{1,2} · Piyush Joshi¹ · Damian Stichel^{4,5} · Pengbo Beck^{1,3,6} · Konstantin Okonechnikow^{1,3} · Philipp Sievers^{4,5} · Annika K. Wefers⁷ · Federico Roncaroli⁸ · Shivaram Avula⁹ · Martin G. McCabe¹⁰ · James T. Hayden¹¹ · Pieter Wesseling^{12,13} · Ingrid Øra¹⁴ · Monica Nistér¹⁵ · Mariëtte E. G. Kranendonk¹² · Bastiaan B. J. Tops¹² · Michal Zapotocky^{16,17} · Josef Zamecník¹⁸ · Alexandre Vasiljevic¹⁹ · Tanguy Fenouil¹⁹ · David Meyronet¹⁹ · Katja von Hoff²⁰ · Ulrich Schüller^{7,21,22} · Hugues Loiseau²³ · Dominique Figarella-Branger²⁴ · Christof M. Kramm²⁵ · Dominik Sturm^{1,2,26} · David Scheie²⁷ · Tuomas Rauramaa²⁸ · Jouni Pesola²⁹ · Johannes Gojo³⁰ · Christine Haberler³¹ · Sebastian Brandner^{32,33} · Tom Jacques³⁴ · Alexandra Sexton Oates³⁵ · Richard Saffery³⁵ · Ewa Koscielniak³⁶ · Suzanne J. Baker³⁷ · Stephen Yip³⁸ · Matija Snuderl³⁹ · Nasir Ud Din⁴⁰ · David Samuel⁴¹ · Kathrin Schramm^{1,2} · Mirjam Blattner-Johnson^{1,2} · Florian Selt^{1,42,43} · Jonas Ecker^{1,42,43} · Till Milde^{1,42,43} · Andreas von Deimling^{4,5} · Andrey Korshunov^{1,4,5} · Arie Perry⁴⁴ · Stefan M. Pfister^{1,3,26} · Felix Sahm^{1,4,5} · David A. Solomon⁴⁴ · David T. W. Jones^{1,2} 

not included in WHO 2021 najlepšie identifikovateľný metylačným profilovaním

Acta Neuropathologica
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CORRESPONDENCE

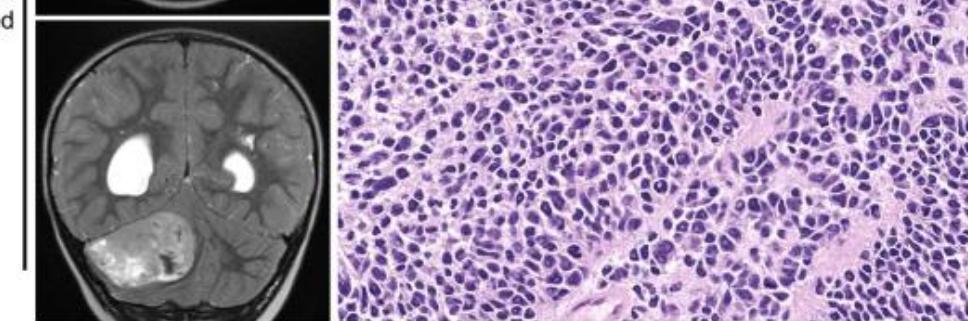
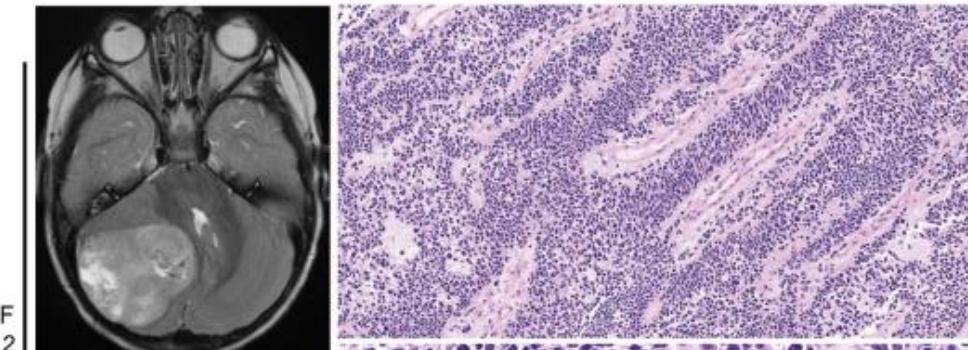


PLAG1 fusions extend the spectrum of PLAG(L)-altered CNS tumors

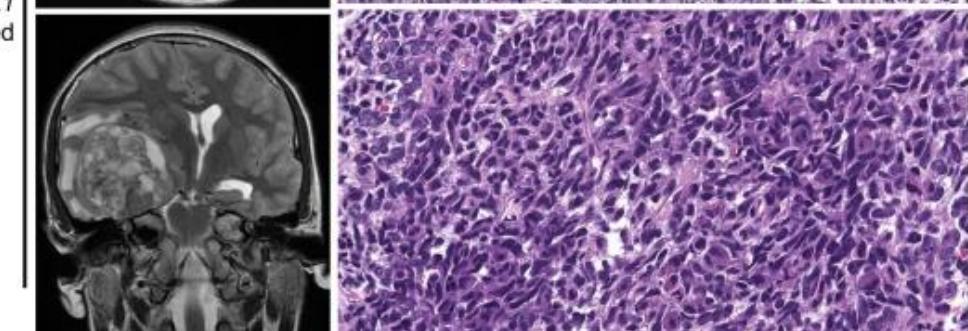
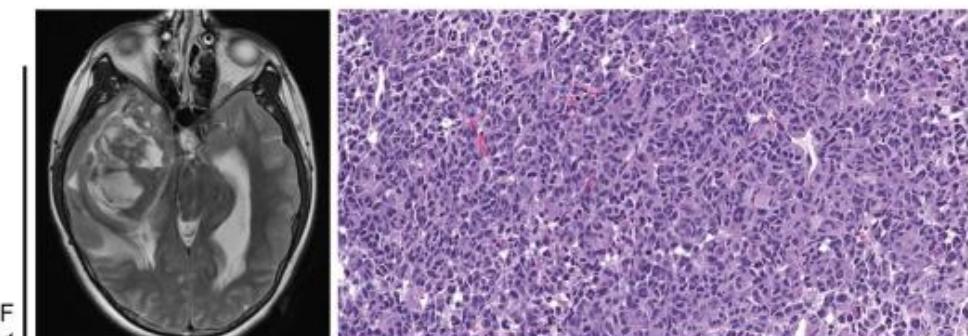
Arnaud Tauziède-Espriat¹ · Aurore Siegfried^{2,3,4} · Yvan Nicaise³ · Delphine Dghayem⁵ · Anne Laprie⁶ · Vincent Lubrano⁷ · Pomone Richard⁸ · Guillaume Gauchotte⁹ · Joséphine Malczuk¹⁰ · Olivier Klein¹⁰ · Lauren Hasty¹ · Alice Métais^{1,11} · Fabrice Chrétien¹ · Volodia Dangouloff-Ros¹² · Nathalie Boddaert¹² · Felix Sahm^{13,14} · Philipp Sievers^{13,14} · Pascale Varlet^{1,11} · Emmanuelle Uro-Coste^{2,3,4} on behalf of the RENOCLIP-LOC

variable GFAP, MAP2, SYP, desmin
PLAGL1/2 amplification, rarely fusion

(a)



(b)

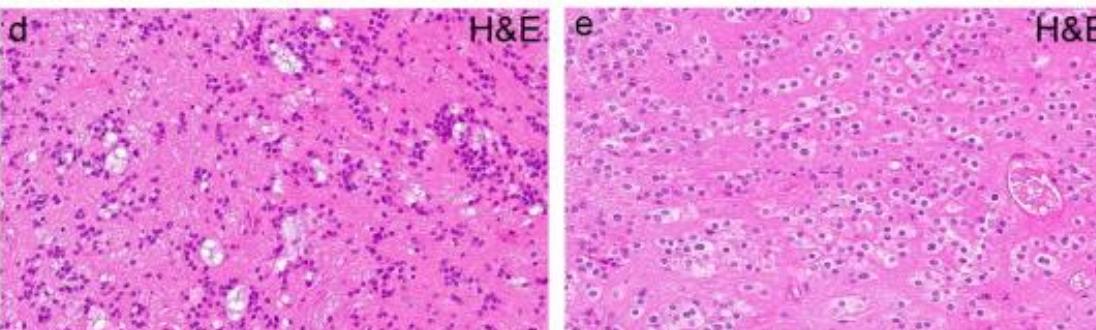




Recurrent fusions in *PLAGL1* define a distinct subset of pediatric-type supratentorial neuroepithelial tumors

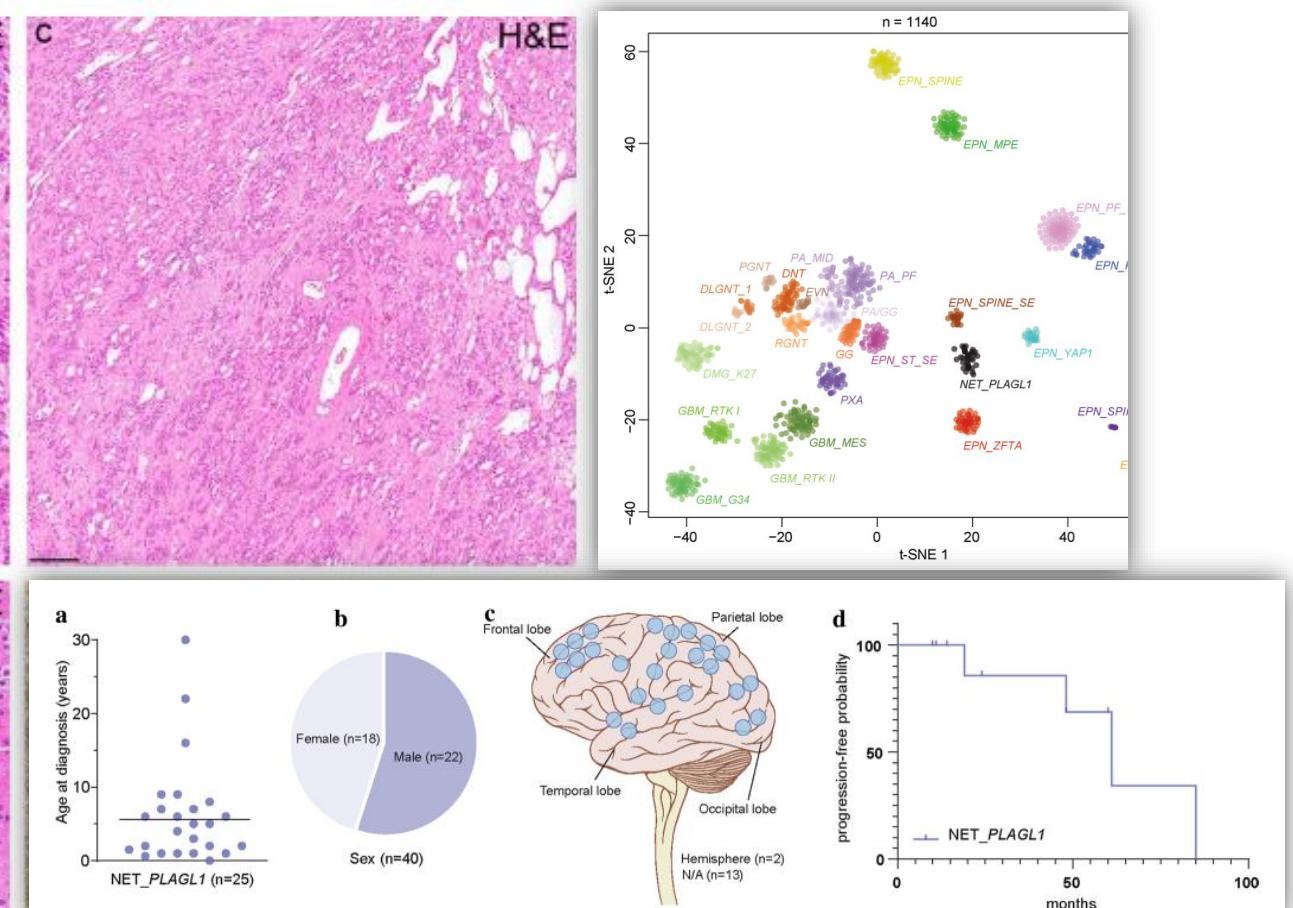
Philipp Sievers^{1,2} · Sophie C. Henneken^{3,4} · Christina Blume^{1,2,5} · Martin Sill^{3,4} · Daniel Schrimpf^{1,2} ·
 Damian Stichel^{1,2} · Konstantin Okonechnikov^{3,4} · David E. Reuss^{1,2} · Julia Benzel^{3,4} · Kendra K. Maaß^{3,4,6} ·
 Marcel Kool^{3,4,7} · Dominik Sturm^{3,6,8} · Tuyu Zheng^{3,4,9} · David R. Ghasemi^{3,4} · Patricia Kohlhof-Meinecke¹⁰ ·
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 Stephanie Bunkowski²⁵ · Christine Stadelmann²⁵ · Ulrich Schüller^{26,27,28} · Wolf C. Mueller²⁹ · Hildegard Dohmen³⁰ ·
 Till Acker³⁰ · Patrick N. Harter^{31,32,33,34} · Christian Mawrin³⁵ · Rudi Beschorner³⁶ · Sebastian Brandner^{37,38} ·
 Matija Snuderl³⁹ · Zied Abdullaev⁴⁰ · Kenneth Aldape⁴⁰ · Mark R. Gilbert⁴¹ · Terri S. Armstrong⁴¹ · David W. Ellison⁴² ·
 David Capper^{20,21} · Koichi Ichimura⁴³ · Guido Reifenberger^{44,45} · Richard G. Grundy⁴⁶ · Nada Jabado^{47,48,49} ·
 Lenka Krskova^{50,51} · Michal Zapotocky^{50,52} · Ales Vicha^{50,52} · Pascale Varlet¹⁷ · Pieter Wesseling^{7,53} ·
 Stefan Rutkowski²⁷ · Andrey Korshunov^{1,2,3} · Wolfgang Wick^{54,55} · Stefan M. Pfister^{3,4,6} · David T. W. Jones^{3,8} ·
 Andreas von Deimling^{1,2} · Kristian W. Pajtler^{3,4,6} · Felix Sahm^{1,2,3,15}

not included in WHO 2021



Tumor prevažne detského veku, pravdepodobne s ependymálnou diferenciáciou, median PFS = 35 mesiacov

Identifikovateľný iba metylačným profilovaním

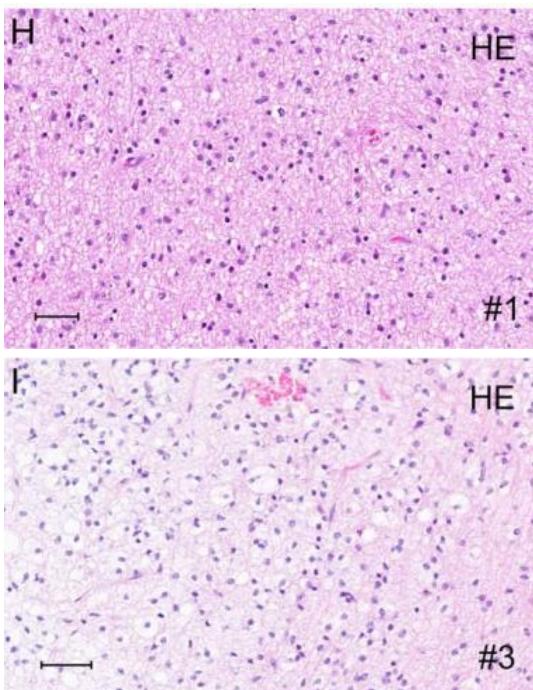
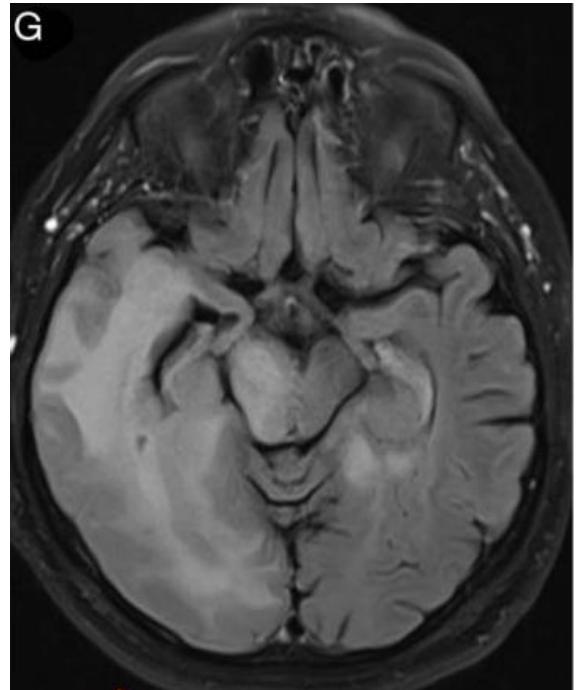


A Novel Type of IDH-wildtype Glioma Characterized by Gliomatosis Cerebri-like Growth Pattern, TERT Promoter Mutation, and Distinct Epigenetic Profile

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Shokufe Nazari-Dehkordi, MD,¶ Julia Onken, MD,# Peter Vajkoczy, MD,#
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David Capper, MD,†||| David Kaul, MD,*** Christian Thomas, MD,§§ and
Leonille Schweizer, MD*†|||††††††††

Am J Surg Pathol 2023;47:1364–1375

not included in WHO 2021



Radiologicky často obraz podobný gliomatosis cerebri, morfologicky difúzny glióm s astrocytárnou morfológiou, bez nekrózy alebo mikrovaskulárnej proliferácie.

Prognóza signifikantne lepšia ako pri glioblastóme, IDHwt, podobná IDH-mutovaným astrocytómom

Identifikateľný iba metylačným profilovaním mutácie promótora TERT, PIK3R1, p53



Zdroj: Vysoká škola života.

Katedra mikrobiológie a virológ