

Update on molecular diagnostics in CNS cancers

Tom Jacques

Objectives

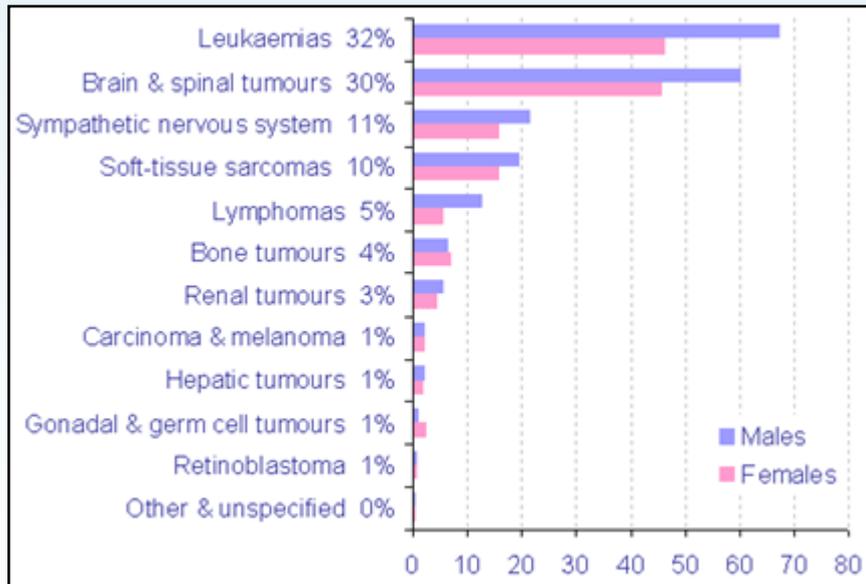
- To describe the current landscape of molecular diagnosis in brain tumours, focussing on:
 - Tests that are currently available
 - Tests that alter clinical management or prognosis
 - The incorporation of the tests in to more traditional pathology

Perspective and biases



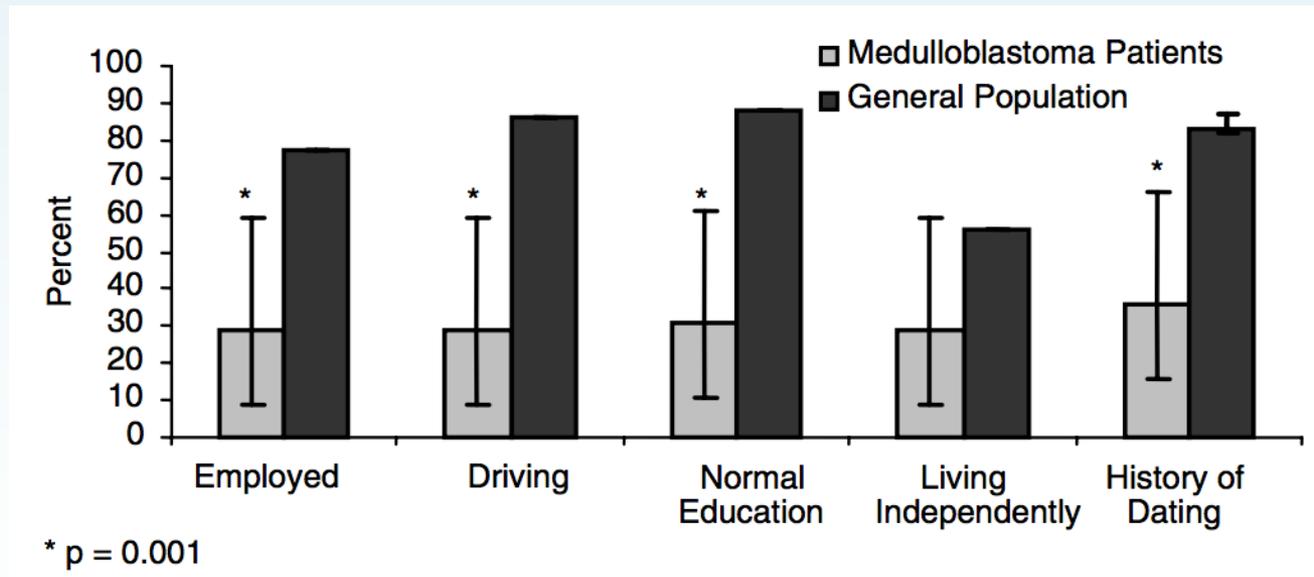
What distinguishes neuropathology for children?

Distinctive clinical features of childhood brain tumours



- CNS tumours are the commonest solid tumours of childhood and the commonest fatal tumours in childhood
- Paediatric brain tumours are often amenable to curative treatment
- ...but significant morbidity may be associated with treatment
- Complete surgical resection is often critical for curative treatment.
- Many patients will be on trials with up-front molecular pathology for stratification

Survival is at the cost of long-term complications

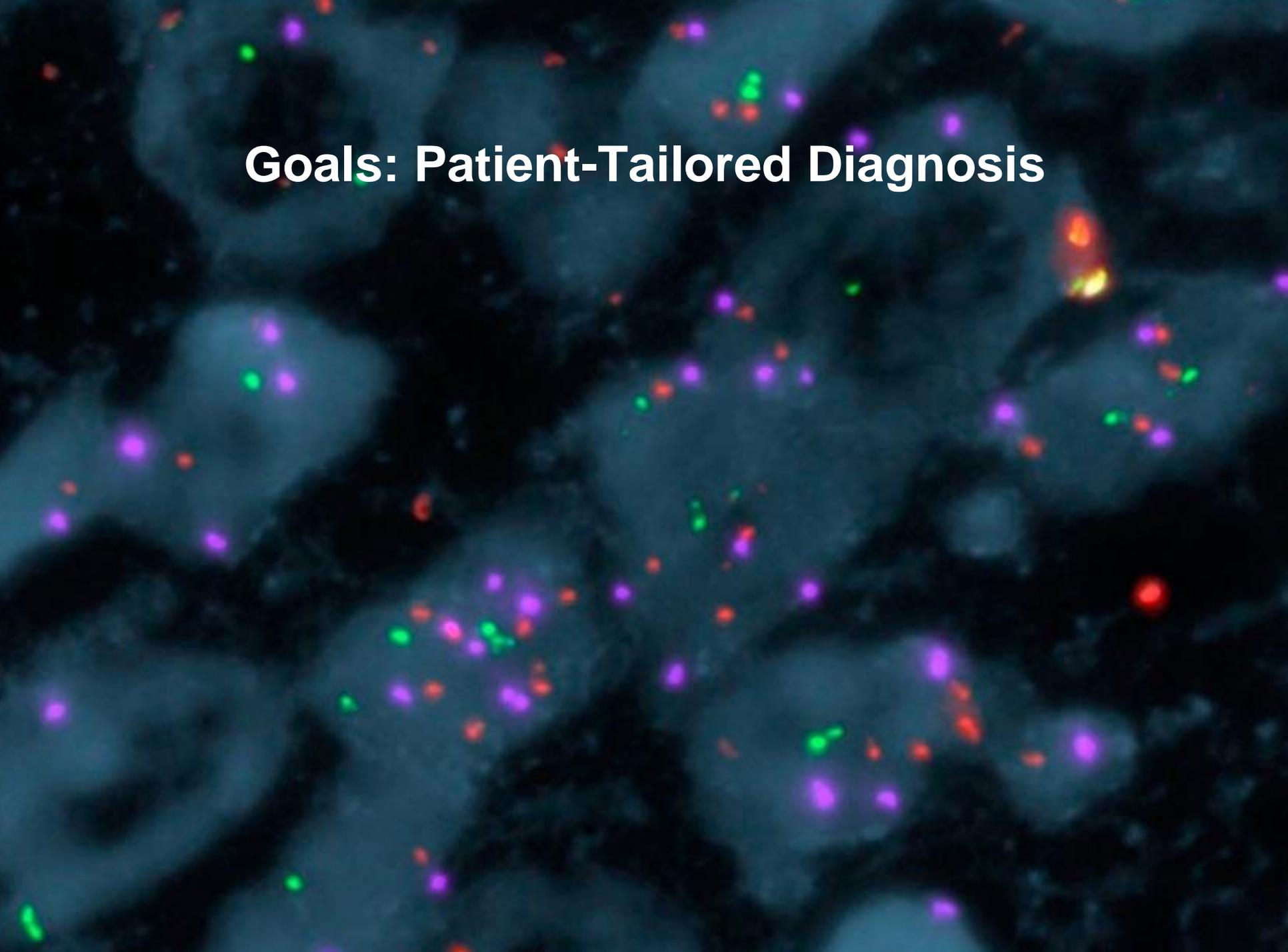


Late effects

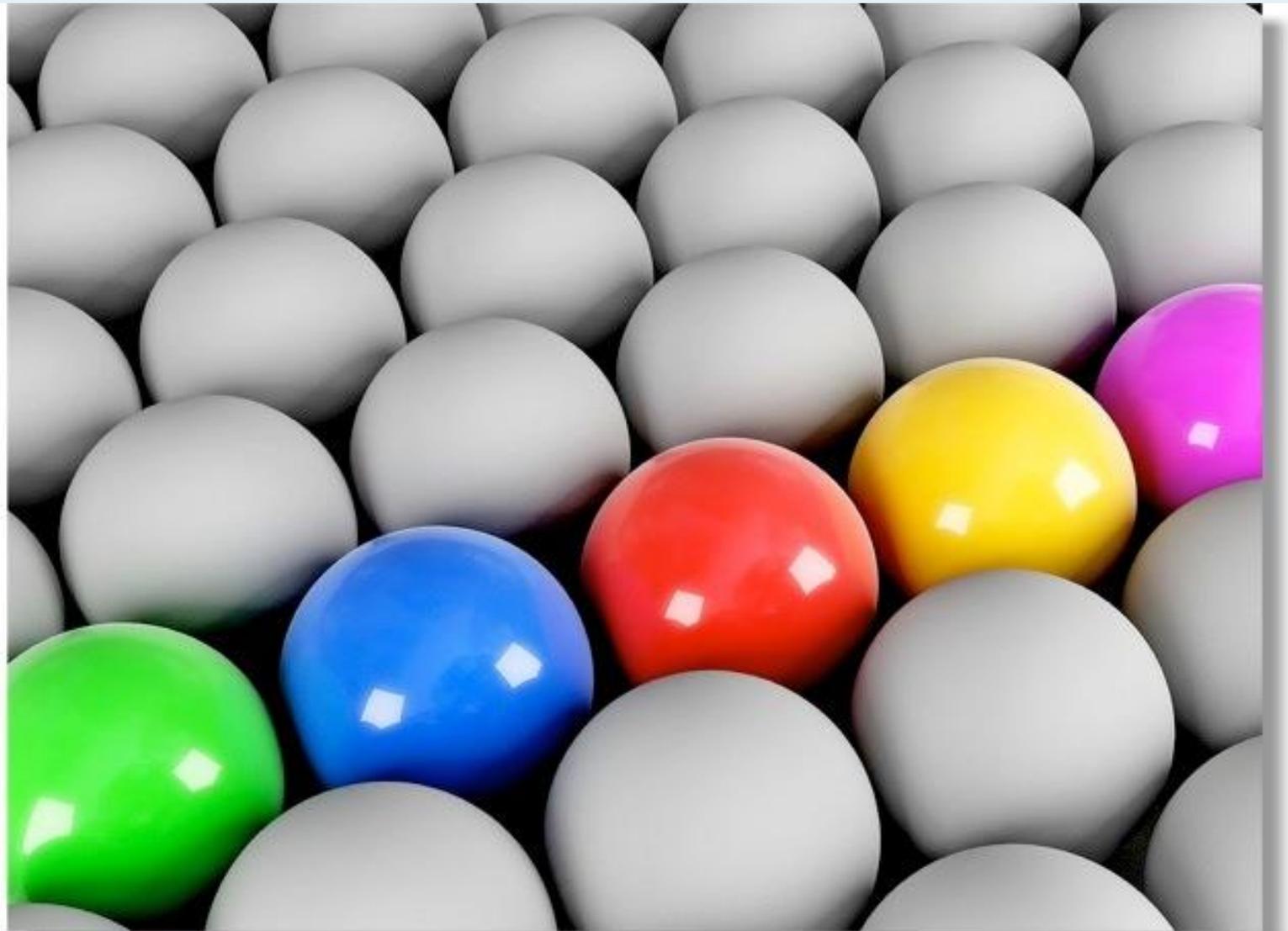


**Long term
cure**

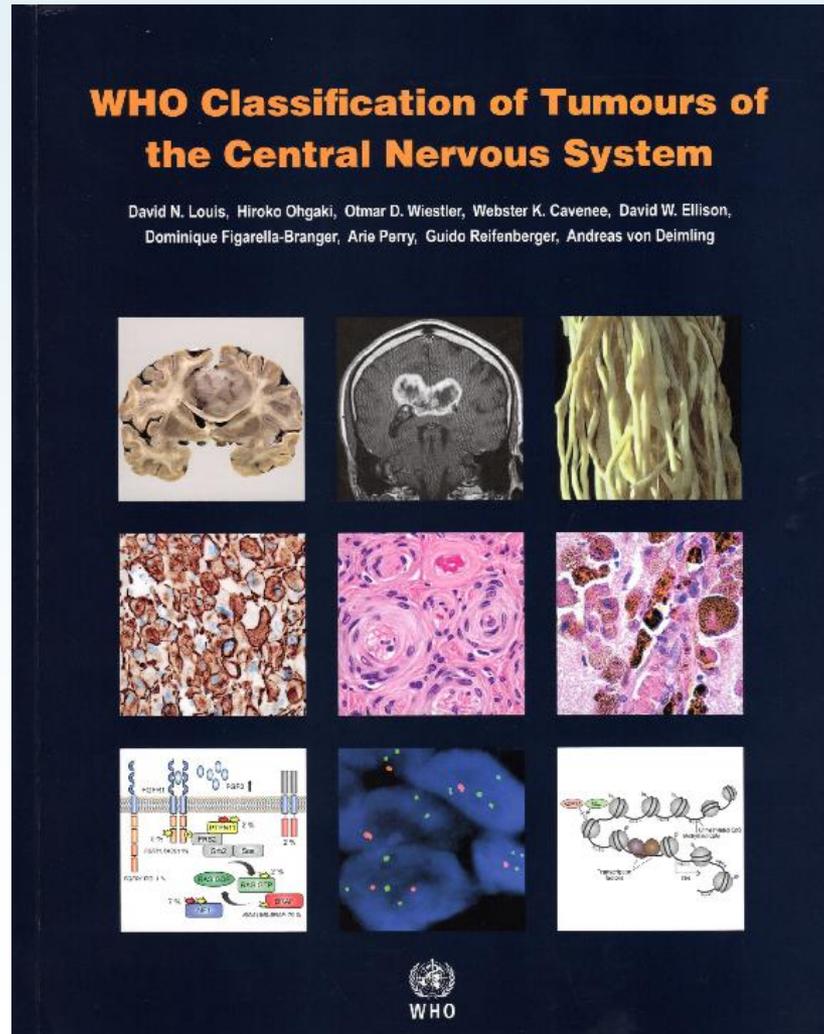
Goals: Patient-Tailored Diagnosis



The challenges of the numbers



The revised WHO classification



For the first time: Molecularly defined tumours

Medulloblastoma, SHH-activated and TP53-mutant

Eberhart C.G.
Giangaspero F.
Ellison D.W.
Haapasalo H.

Pietsch T.
Wiestler O.D.
Pfister S.

Definition

A poorly differentiated embryonal tumour of the cerebellum with evidence of SHH pathway activation and either germline or somatic TP53 mutation

In large series of tumours, SHH-activated medulloblastomas tend to have similar transcriptome, methylome, and micro-RNA profiles. SHH pathway activation in TP53-mutant tumours is associated with amplification of *GLI2*, *MYCN*, or *SHH*. Mutations in *PTCH1*, *SUFU*, and *SMO* are generally absent. Large cell / anaplastic morphology and chromosome 17p loss are also common in SHH-activated and TP53-mutant tumours. Patterns of chromosome shattering known as chromothripsis are often present.

SHH-activated tumours account for approximately 30% of all medulloblastomas and originate from rhombic lip-derived cerebellar granule neuron precursors, the proliferation of which is dependent on SHH signalling activity. SHH-activated and TP53-mutant medulloblastomas are rare and generally found in children aged 4–17 years. Clinical outcomes in patients with SHH-activated and TP53-mutant tumours are very poor.

ICD-O code

9476/3

Grading

Like all medulloblastomas, SHH-activated and TP53-mutant medulloblastoma corresponds histologically to WHO grade IV.

Epidemiology

SEER data from 1973–2007 suggest medulloblastoma incidence rates of 6.0 cases per 1 million children aged 1–9 years and 0.6 cases per 1 million adults aged > 19 years [2382]. SHH-activated medulloblastomas in general show a bimodal age distribution, being most common in infants and young adults, with a male-to-female ratio of approximately 1.5:1 [1804]. In contrast, SHH-activated and TP53-mutant tumours in particular are generally found in children aged 4–17 years [1333]. In one study that included 133 SHH-activated medulloblastomas, 28 patients (21%) had a TP53 mutation, and the median age of these patients was approximately 15 years [2870].

Localization

SHH-activated medulloblastomas were proposed in one study report to predominantly involve the lateral cerebellum, due to their origin from granule neuron precursors [831]. A subsequent study that included 17 SHH-activated medulloblastomas found that although 9

of those tumours were hemispheric, the other 8 were centred in, or significantly involved, the midline vermis [2534]. The localization of SHH-activated tumours may be age-dependent. A third study found that in older children and young adults, SHH-activated medulloblastomas grow predominantly in the rostral cerebellar hemispheres, whereas in infants they more frequently involve the vermis [2716]. Specific data on the localization of SHH-activated and TP53-mutant medulloblastoma are not available.

Imaging

On CT and MRI, medulloblastomas present as solid, intensely contrast-enhancing masses. SHH-activated medulloblastomas are most often identified in the lateral hemispheres, but can also involve midline structures [831,2534]. Oedema was relatively common in one imaging series that included 12 desmoplastic/nodular medulloblastomas and 9 medulloblastomas with extensive nodularity [743]. A nodular, so-called grape-like pattern on MRI often characterizes medulloblastoma with extensive nodularity because of the tumour's distinctive and diffuse nodular architecture [820,1744]. Medulloblastomas involving the peripheral cerebellar hemispheres in adults occasionally present as extra-axial lesions

Medulloblastoma, SHH-activated and *TP53*-mutant

Definition

A poorly differentiated embryonal tumour of the cerebellum with evidence of SHH pathway activation and either germline or somatic TP53 mutation

In large series of tumours, SHH-activated medulloblastomas tend to have similar transcriptome, methylome, and micro-RNA profiles. SHH pathway activation in *TP53*-mutant tumours is associated with

Grading

Like all medulloblastomas, SHH-activated and *TP53*-mutant medulloblastoma corresponds histologically to grade IV.

Epidemiology

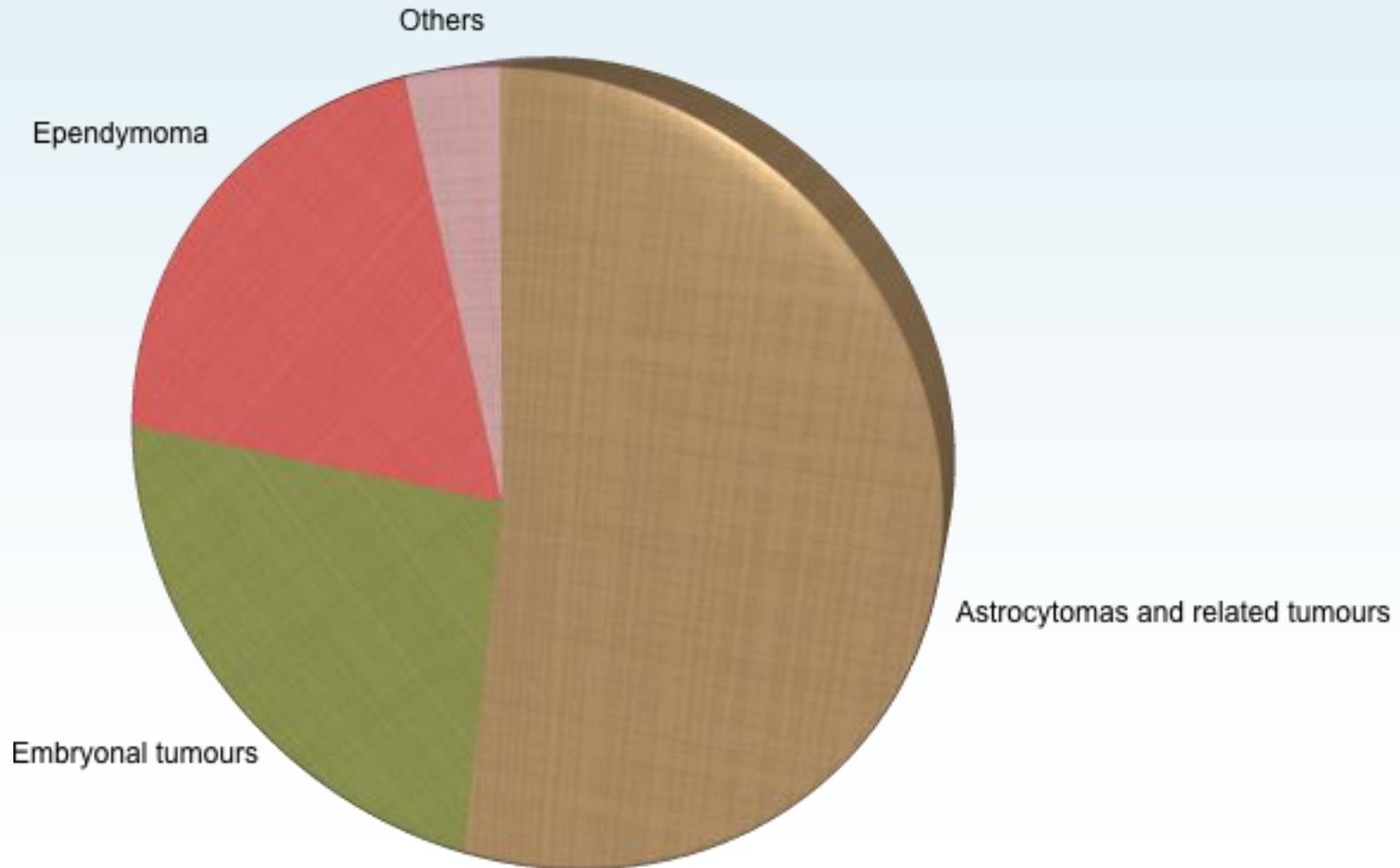
SEER data from 1973–2007 suggests medulloblastoma incidence rates of 6.0 cases per 1 million children aged 1–9 y

INTEGRATED DIAGNOSIS

INTEGRATED DIAGNOSIS: Medulloblastoma, SHH-activated, *TP53*-mutant

- » Histological Diagnosis: Medulloblastoma
- » Histological Grade: IV
- » Molecular Data: SHH activated, *TP53* mutated

There are 3 common brain tumour groups in children



Embryonal Tumours

Terminology

- Embryonal tumours
- Primitive Neuro-ectodermal Tumours (PNET)
 - In essence primary small round blue cell tumours of the brain

Embryonal Tumour classification

- Embryonal Tumour in the cerebellum=Medulloblastoma
- Everything else

CHAPTER 8

Embryonal tumours

Medulloblastoma, genetically defined

Medulloblastoma, WNT-activated

Medulloblastoma, SHH-activated and TP53-mutant

Medulloblastoma, SHH-activated and TP53-wildtype

Medulloblastoma, non-WNT/non-SHH

Medulloblastoma, histologically defined

Medulloblastoma, classic

Desmoplastic/nodular medulloblastoma

Medulloblastoma with extensive nodularity

Large cell / anaplastic medulloblastoma

Embryonal tumour with multilayered rosettes, C19MC-altered

Embryonal tumour with multilayered rosettes, NOS

Medulloepithelioma

CNS neuroblastoma

CNS ganglioneuroblastoma

CNS embryonal tumour, NOS

Atypical teratoid/rhabdoid tumour

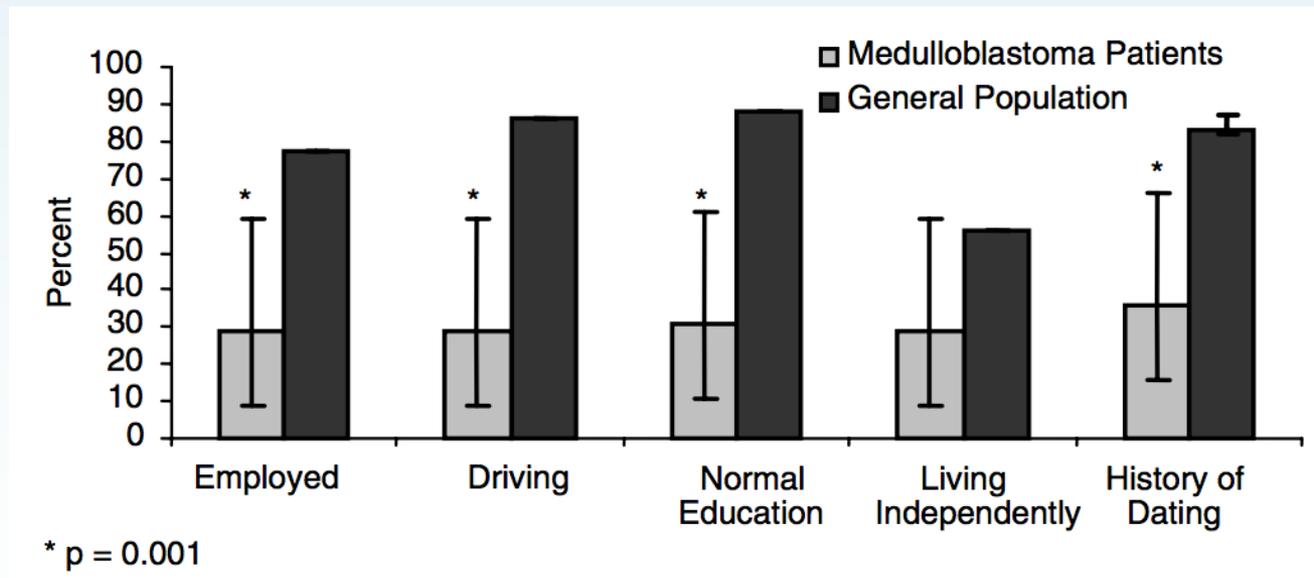
CNS embryonal tumour with rhabdoid features

Medulloblastoma



- Definition-embryonal tumour of the posterior fossa
- Peak age of onset 7 years
- 5-yr survival >70%

Survival is at the cost of long-term complications



Classification-genetics and histology

Medulloblastoma, genetically defined

Medulloblastoma, WNT-activated

Medulloblastoma, SHH-activated and TP53-mutant

Medulloblastoma, SHH-activated and TP53-wildtype

Medulloblastoma, non-WNT/non-SHH

Medulloblastoma, histologically defined

Medulloblastoma, classic

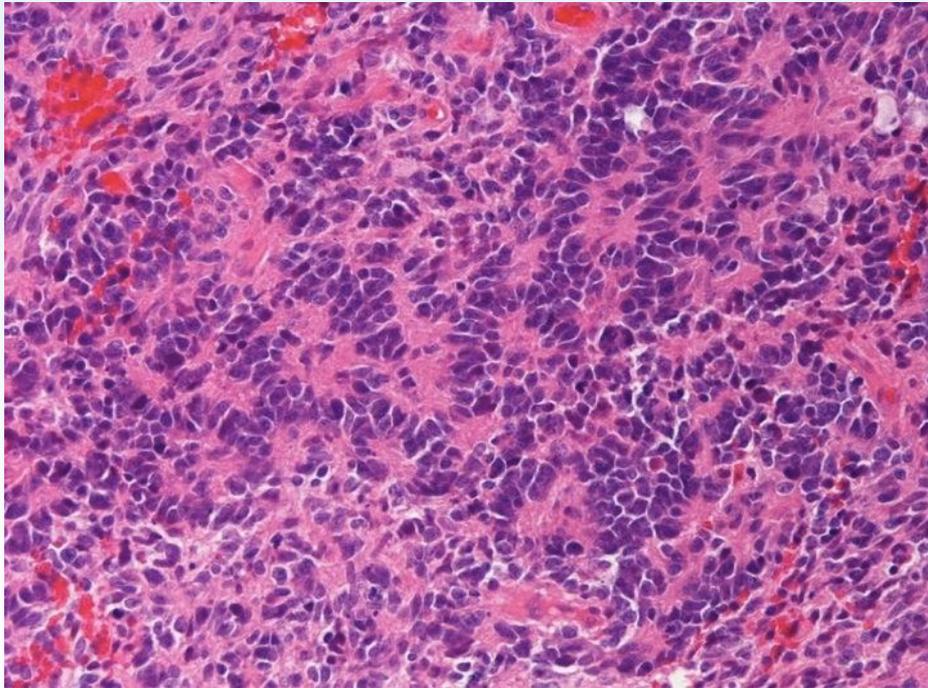
Desmoplastic/nodular medulloblastoma

Medulloblastoma with extensive nodularity

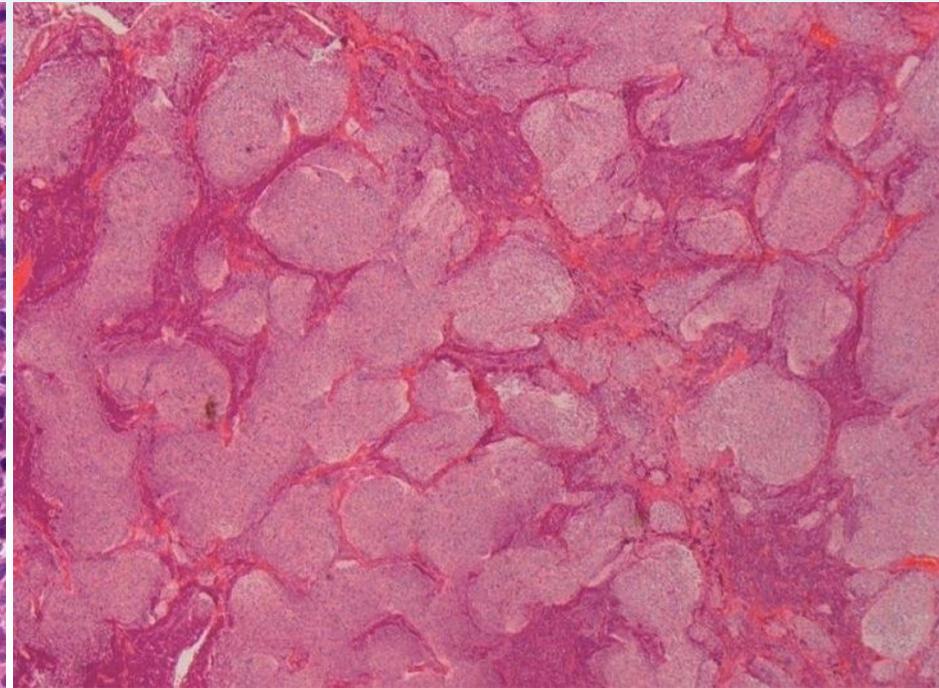
Large cell / anaplastic medulloblastoma

Risk stratification-conventional histology

Medulloblastoma-Architecture

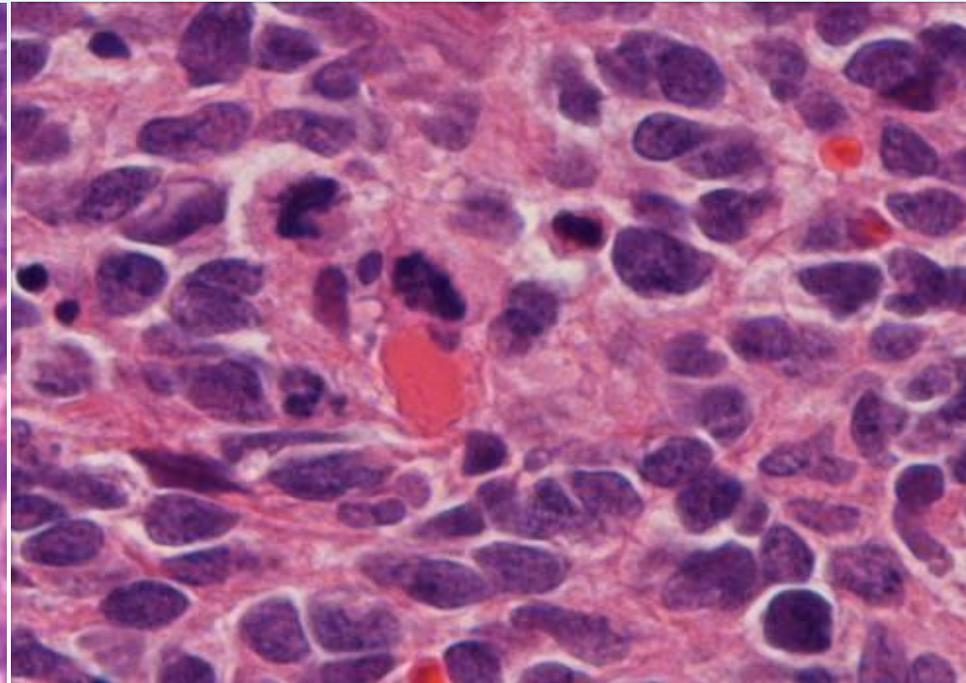
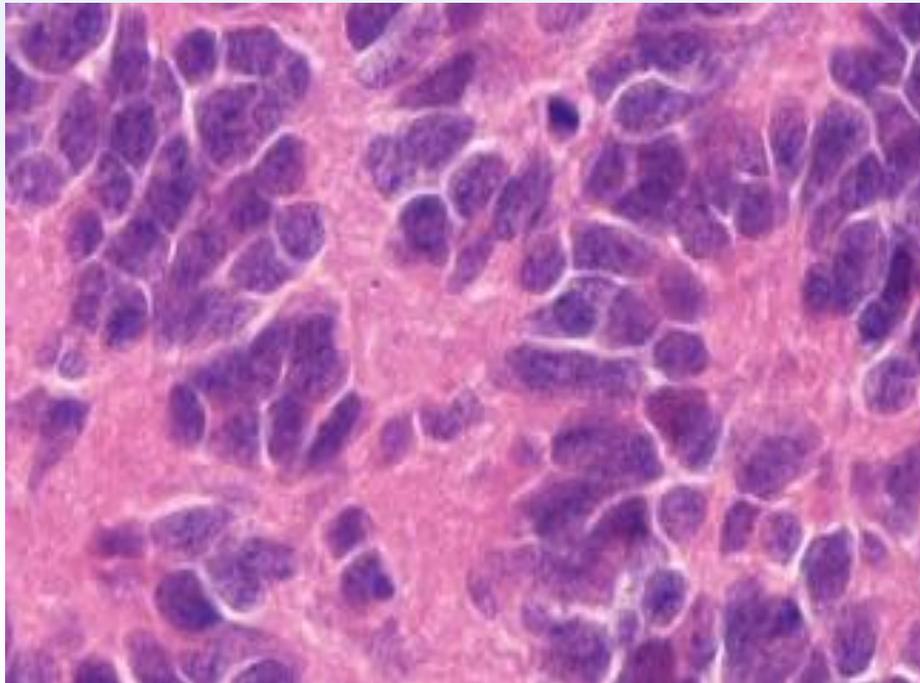


Diffuse

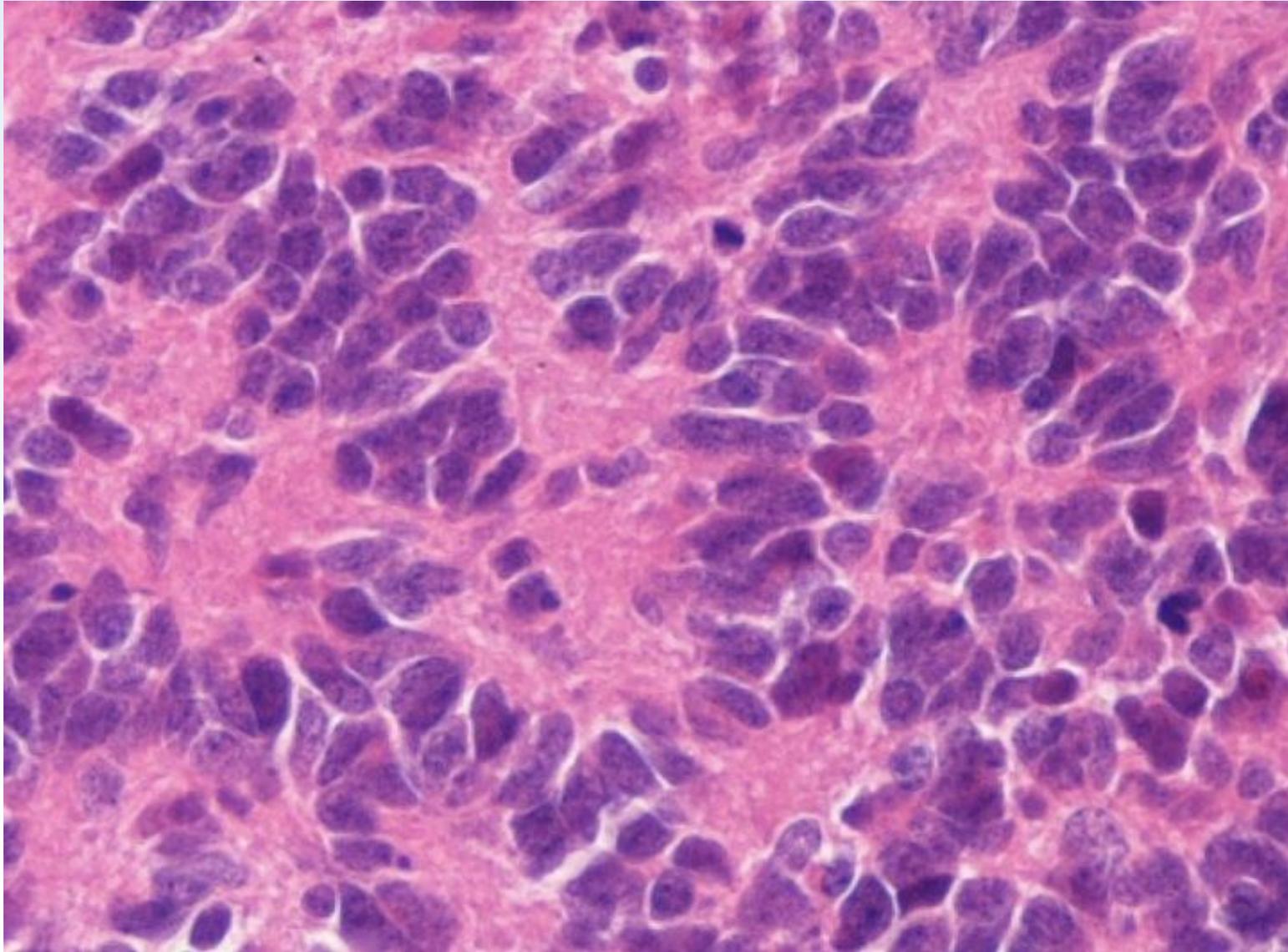


Nodular

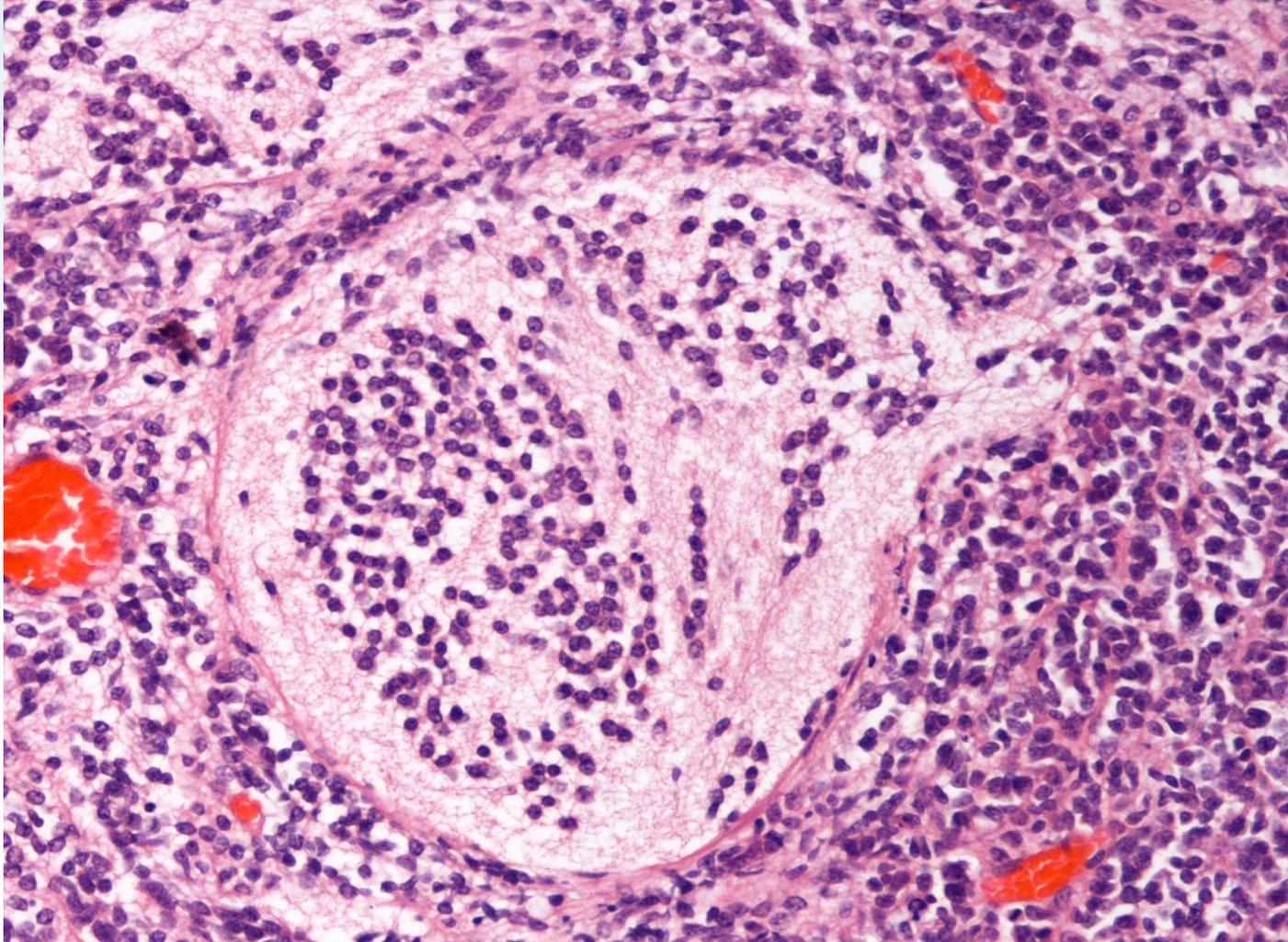
Medulloblastoma-Cytology



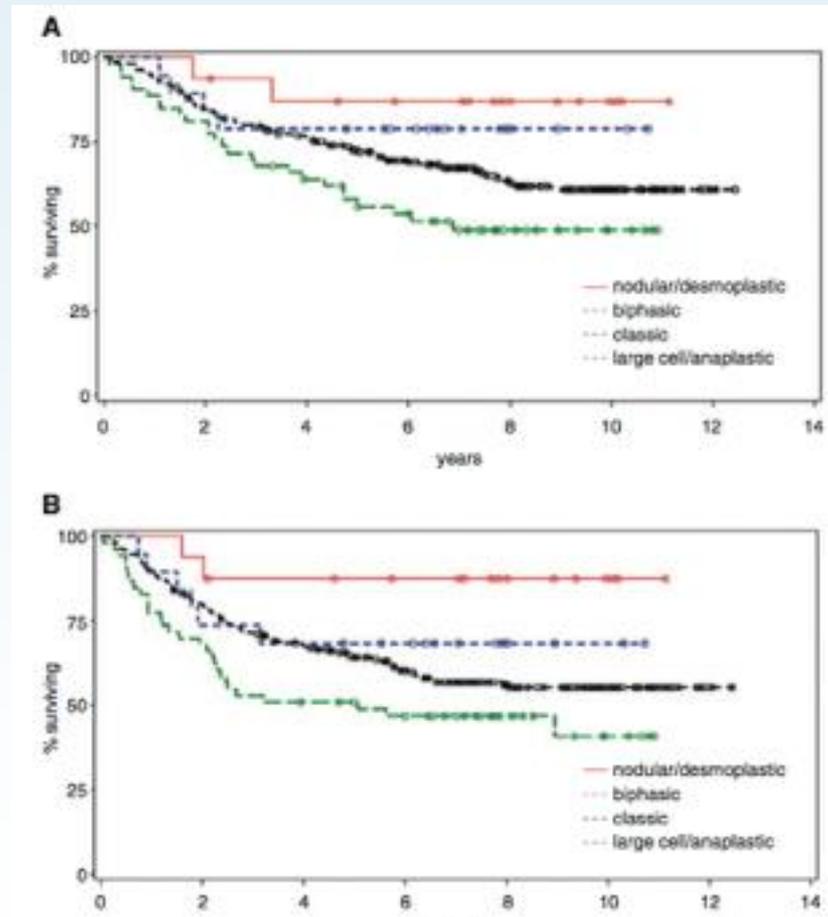
Medulloblastoma-Homer Wright rosettes



Variants-Nodular Medulloblastoma



Nodularity is associated with a better prognosis



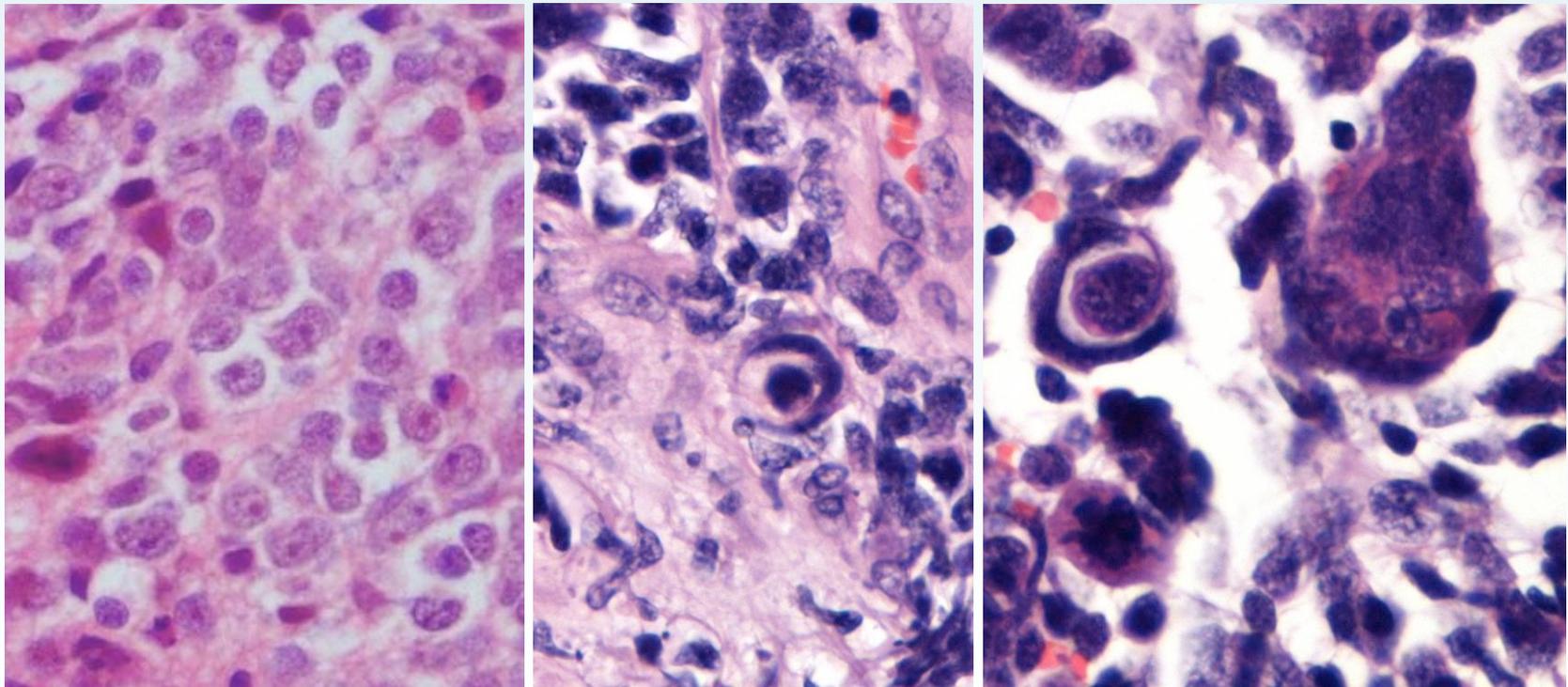
RESEARCH ARTICLE

DOI 10.1111/j.1750-3639.2007.00058.x

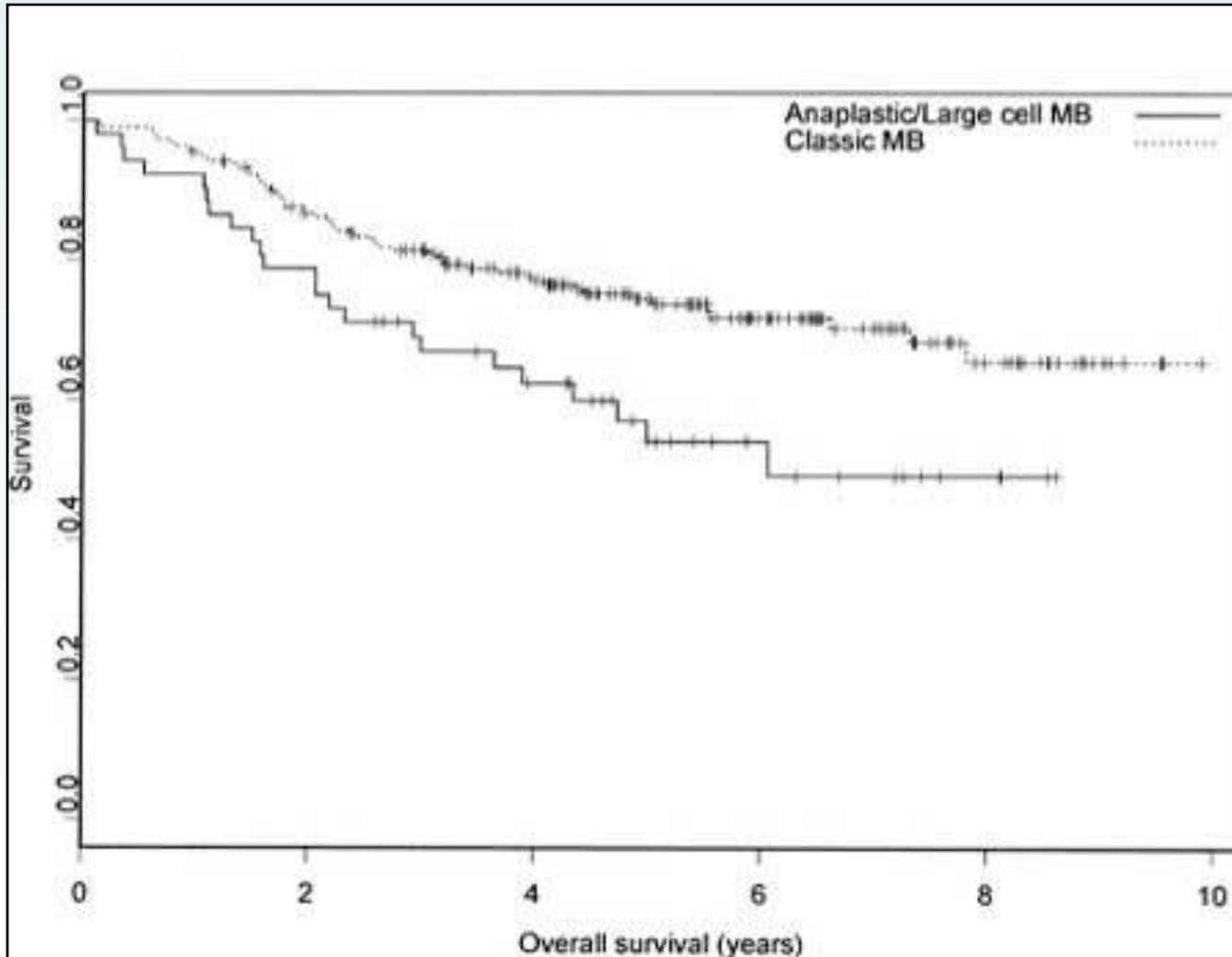
Nodule Formation and Desmoplasia in Medulloblastomas—Defining the Nodular/Desmoplastic Variant and Its Biological Behavior

Charles S. McManamy^{1,2*}, Jane Peam^{1,3*}, Claire L. Weston¹, Zoltan Hanzely¹, James W. Ironside¹, Roger E. Taylor¹, Richard G. Grundy¹, Steven C. Clifford¹, David W. Ellison^{1,2,3,4} on behalf of the Clinical Brain Tumour Group, Children's Cancer and Leukaemia Group (formerly the UK Children's Cancer Study Group), UK

Prognostic Factors: Anaplastic and Large Cell Medulloblastoma



Anaplastic and large cell medulloblastoma carry a worse prognosis



Mc Nanamy *et al.* 2003 JNEN

Risk stratification-molecular subtyping

Molecular Subgroups of Medulloblastoma

CONSENSUS

Cho (2010)
Northcott (2010)
Kool (2008)
Thompson (2006)

WNT

C6
WNT
A
B

SHH

C3
SHH
B
C, D

Group 3

C1/C5
Group C
E
E, A

Group 4

C2/C4
Group D
C/D
A, C

DEMOGRAPHICS

Age Group:
Gender: ♀ ♂



♂♂ : ♀♀

♂♂ : ♀♀

♂♂ : ♀

♂♂ : ♀

CLINICAL FEATURES

Histology
Metastasis
Prognosis

classic, rarely LCA

desmoplastic/nodular,
classic, LCA

classic, LCA

classic, LCA

rarely M+

uncommonly M+

very frequently M+

frequently M+

very good

infants good, others
intermediate

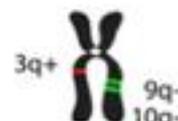
poor

intermediate

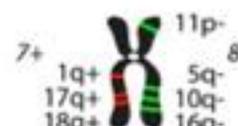
GENETICS



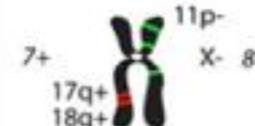
CTNNB1 mutation



PTCH1/SMO/SUFU mutation
GLI2 amplification
MYCN amplification



i17q
MYC amplification



i17q
CDK6 amplification
MYCN amplification

GENE EXPRESSION



WNT signaling

SHH signaling

Photoreceptor/GABAergic

Neuronal/Glutamatergic

MYC+

MYCN+

MYC+++

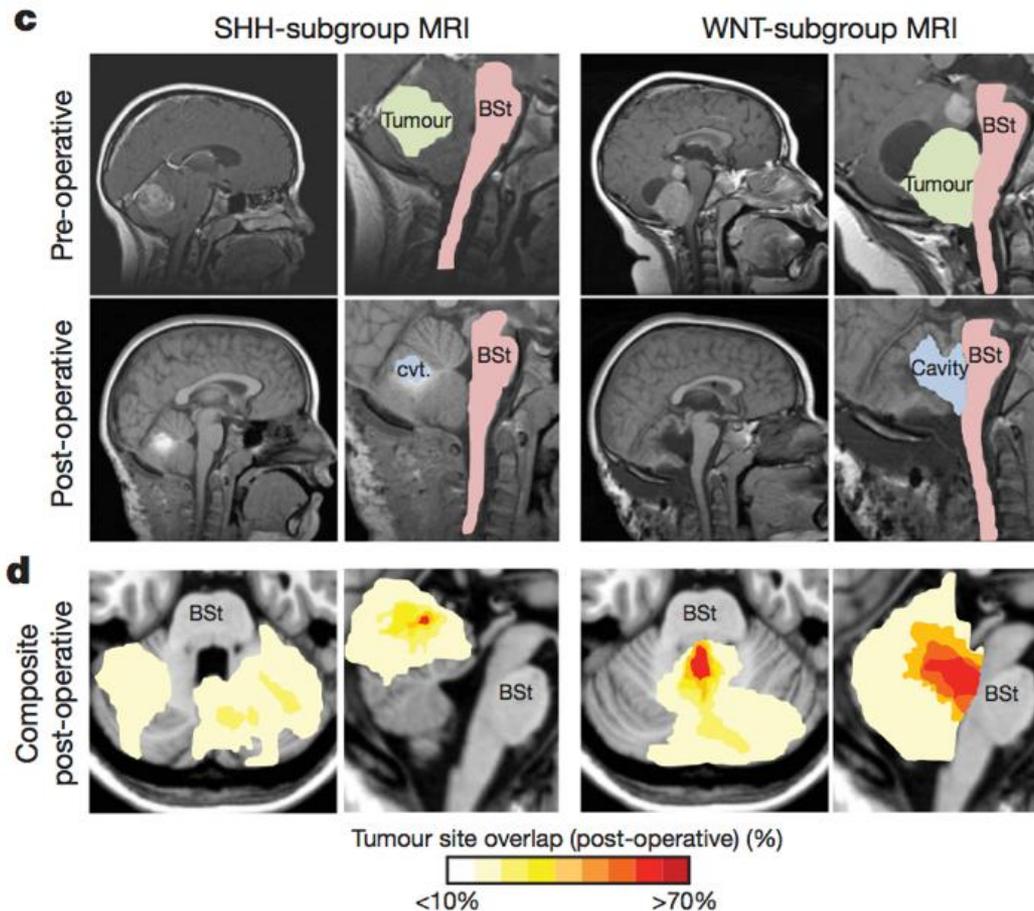
minimal MYC/MYCN

LETTER

doi:10.1038/nature09587

Subtypes of medulloblastoma have distinct developmental origins

Paul Gibson¹, Yiai Tong¹, Giles Robinson^{1,2}, Margaret C. Thompson³, D. Spencer Currie¹, Christopher Eden¹, Tanya A. Kranenburg¹, Twala Hogg¹, Helen Poppleton¹, Julie Martin¹, David Finkelstein³, Stanley Pounds⁴, Aaron Weiss¹⁰, Zoltan Patay⁵, Matthew Scoggins⁵, Robert Ogg⁵, Yanxin Pei¹¹, Zeng-Jie Yang¹¹, Sonja Brun¹¹, Youngsoo Lee⁶, Frederique Zindy⁶, Janet C. Lindsey¹², Makoto M. Taketo¹³, Frederick A. Boop⁷, Robert A. Sanford⁴, Amar Gajjar⁴, Steven C. Clifford¹², Martine F. Roussel⁶, Peter J. McKinnon⁶, David H. Gutmann¹⁴, David W. Ellison⁸, Robert Wechsler-Reya¹¹ & Richard J. Gilbertson^{1,2}



Diagnostic molecular classification

Acta Neuropathol (2011) 121:381–396
DOI 10.1007/s00401-011-0800-8

ORIGINAL PAPER

Medulloblastoma: clinicopathological correlates of SHH, WNT, and non-SHH/WNT molecular subgroups

David W. Ellison · James Dalton · Mehmet Kocak · Sarah Leigh Nicholson · Charles Fraga · Geoff Neale · Anna M. Kenney · Dan J. Brat · Arie Perry · William H. Yong · Roger E. Taylor · Simon Bailey · Steven C. Clifford · Richard J. Gilbertson

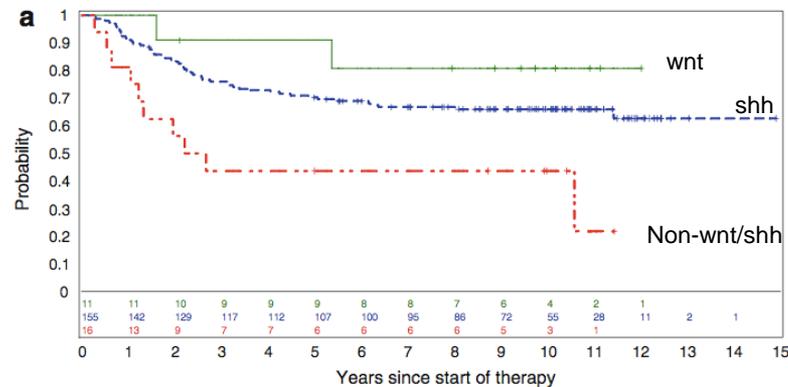
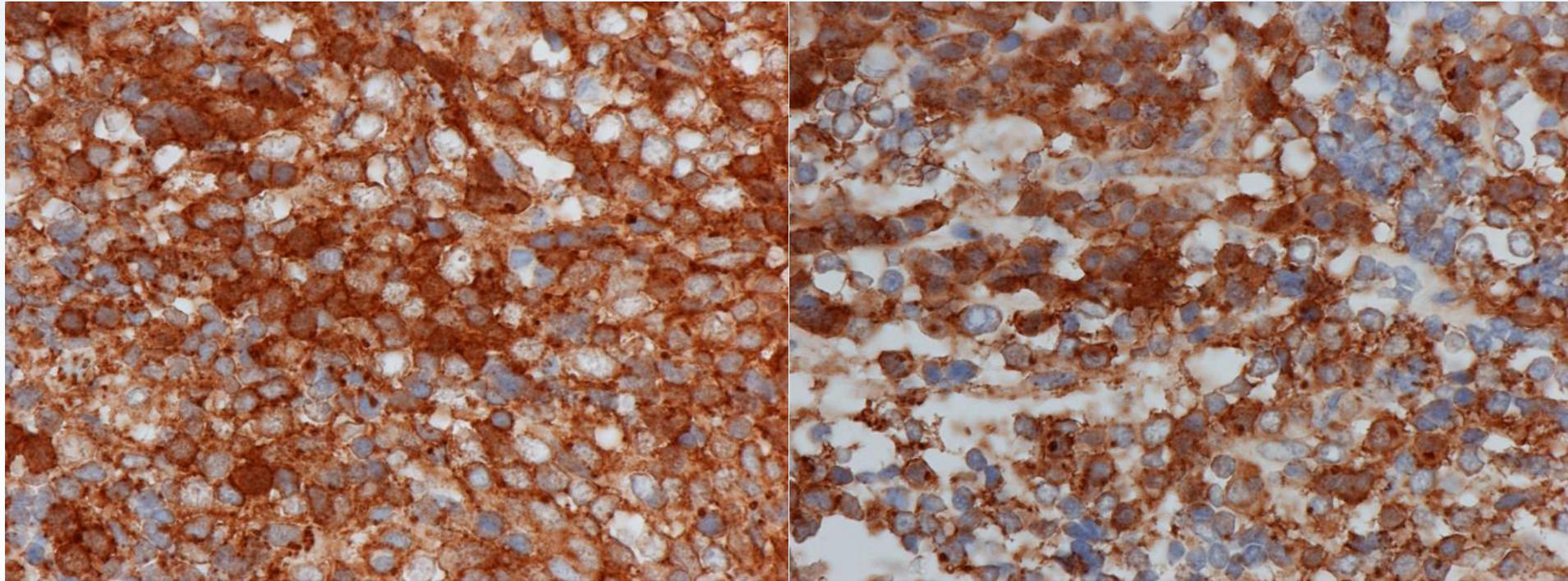


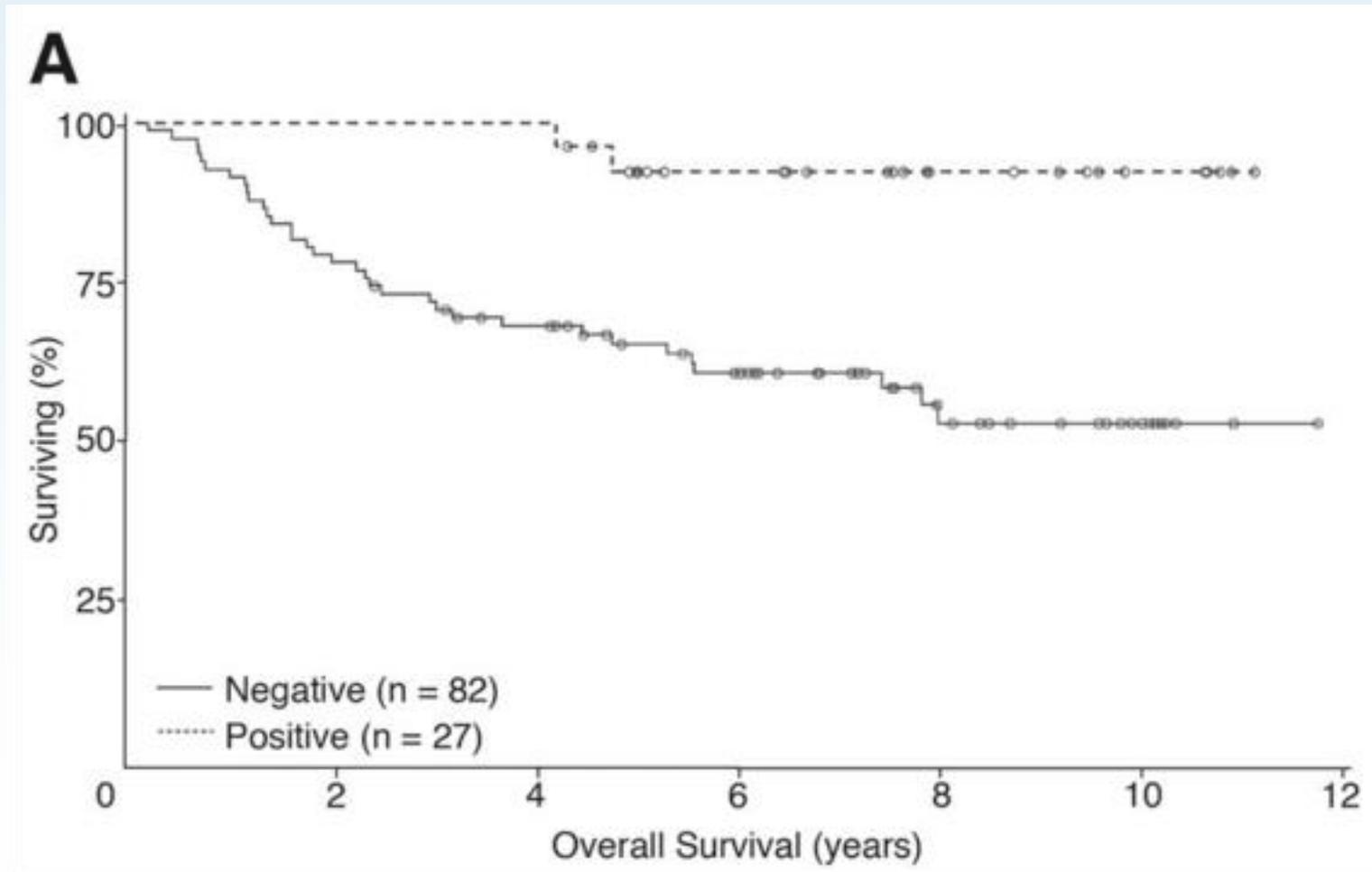
Table 2 Immunophenotypes of SHH, WNT, and non-SHH/WNT molecular subgroups

Molecular group	Immunoreactivity			
	GAB1	β -catenin	Filamin A	YAP1
SHH	Cytoplasmic	Cytoplasmic	Cytoplasmic	Nuclear + cytoplasmic
WNT	Negative	Nuclear + cytoplasmic	Cytoplasmic	Nuclear + cytoplasmic
Non-SHH/WNT	Negative	Cytoplasmic	Negative	Negative

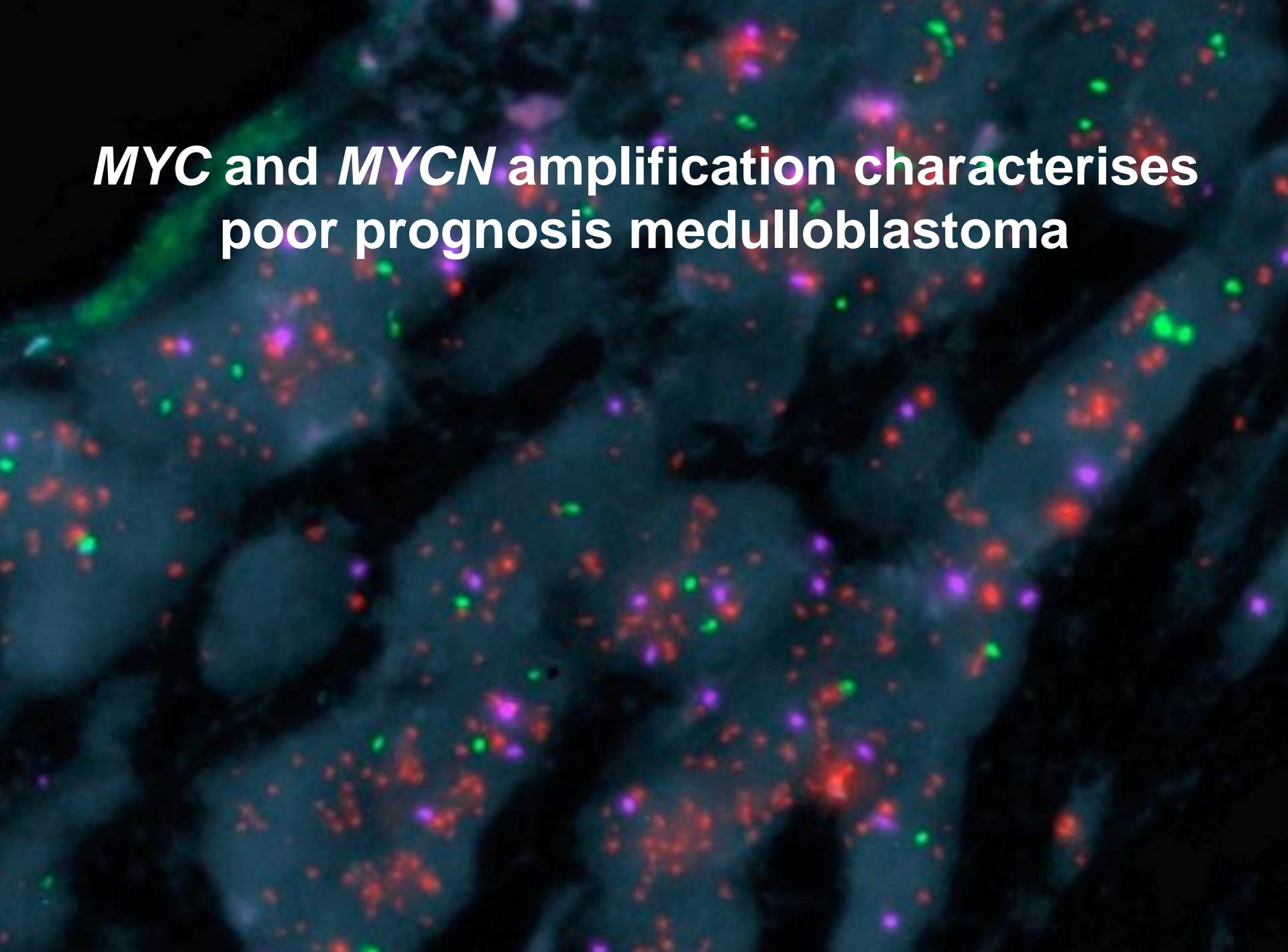
Beta-catenin nuclear staining identifies WNT-subgroup medulloblastoma



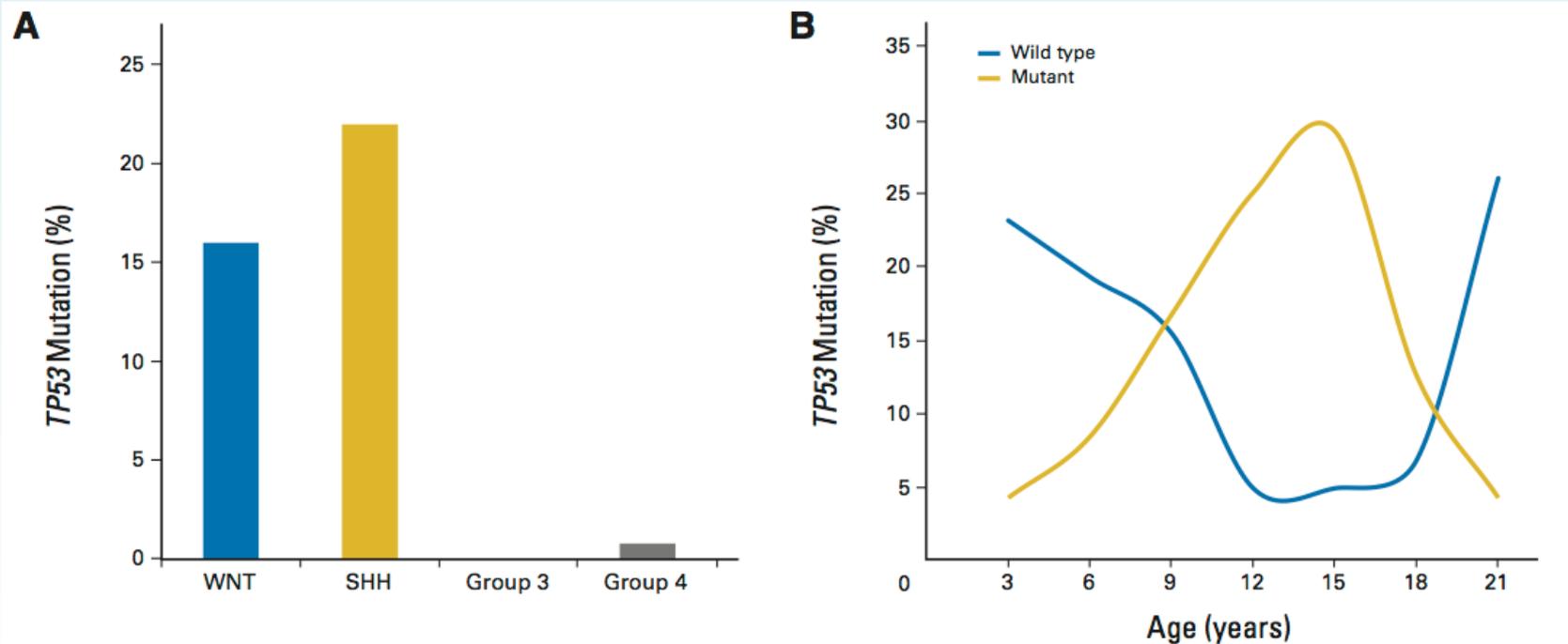
Nuclear beta-catenin expression is associated with an excellent prognosis



***MYC* and *MYCN* amplification characterises
poor prognosis medulloblastoma**



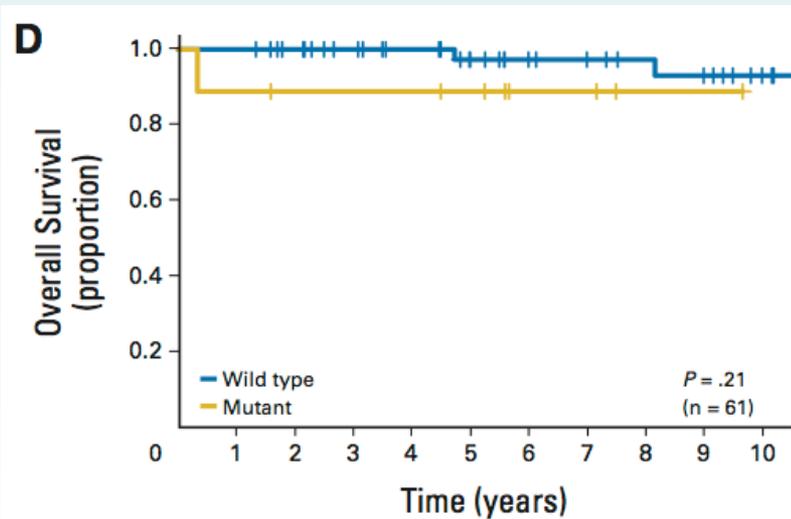
TP53 mutations occur in specific subtypes of medulloblastoma



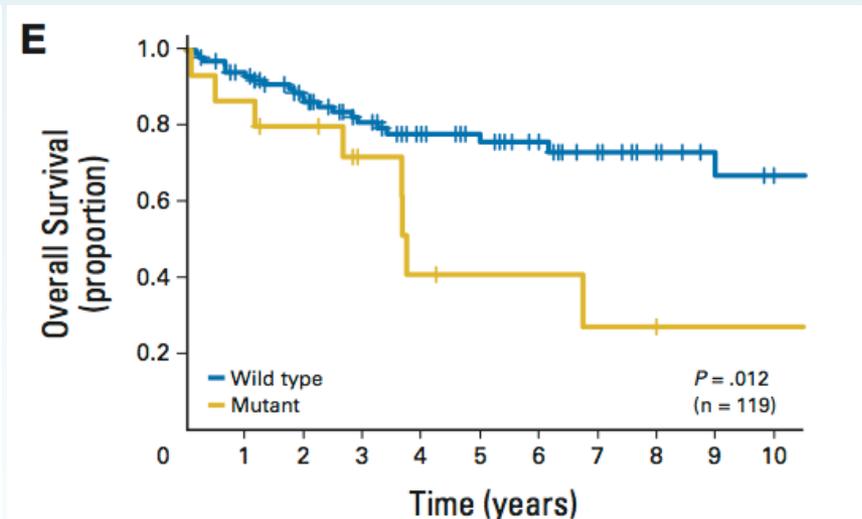
Subgroup-Specific Prognostic Implications of TP53 Mutation in Medulloblastoma

Nataliya Zhukova, Vijay Ramaswamy, Marc Remke, Elke Pfiff, David J.H. Shi, Diana C. Martin, Pedro Castelo-Branco, Berivan Baskin, Peter N. Ray, Eric Bouffet, André O. von Bueren, David T.W. Jones, Paul A. Northcott, Marcel Kool, Dominik Sturm, Trevor J. Fugh, Scott L. Pomeroy, Yoon-Jae Cho, Terence Dutsch, Marco Gessi, Stefan Rutkowski, Lucila Begner, Alois Rieker, Byung-Kyu Cho, Seung-Ki Kim, Kyu-Chang Wang, Charles G. Eberhart, Michelle Ferre-Montange, Maryam Fouladi, Fim J. French, Max Kros, Wiesława A. Czapkowska, Nalin Gupta, William A. Weiss, Peter Hauser, Nada Jabado, Anne Invern, Shin Jung, Toshihiro Kumabe, Boleslaw Lach, Jeffrey R. Leonard, Joshua R. Rubin, Linda M. Liaw, Luca Mastri, Ian F. Pollack, Young Shin Ra, Erwin G. Van Meir, Karol Zitterbart, Ulrich Schaller, Rebecca M. Hill, Janet C. Lindsey, Ed C. Schwalbe, Simon Bulley, David W. Elliott, Cynthia Hawkins, David Malkin, Steven C. Clifford, Andrey Korshunov, Stefan Pfister, Michael D. Taylor, and Uri Tabori

TP53 mutations are a poor prognostic feature in the SHH-subtype of medulloblastoma



WNT-subtype



SHH-subtype

Subgroup-Specific Prognostic Implications of TP53 Mutation in Medulloblastoma

Nataliya Zhukova, Vijay Ramaswamy, Marc Remke, Elke Pfiff, David J.H. Shi, Diana C. Martin, Pedro Castelo-Branco, Berivan Baskin, Peter N. Ray, Eric Bouffet, André O. von Bueren, David T.W. Jones, Paul A. Northcott, Marcel Kool, Dominik Sturm, Trevor J. Fugh, Scott L. Pomeroy, Yoon-Jae Cho, Terence Dutsch, Marco Gessi, Stefan Rutkowski, Lucila Begner, Alois Rickner, Byoung-Kyu Cho, Seung-Ki Kim, Kyu-Chang Wang, Charles G. Eberhart, Michelle Ferre-Montange, Maryam Fouladi, Fim J. French, Max Kros, Wiesława A. Czapkowska, Nalin Gupta, William A. Weiss, Peter Hauser, Nadea Jabado, Anne Invern, Shin Jung, Toshihiro Kamabe, Bolesław Lach, Jeffrey R. Leonard, Joshua R. Rubin, Linda M. Liaw, Luca Mastai, Ian F. Pollack, Young Shin Ra, Erwin G. Van Meir, Karol Zitterbart, Ulrich Schaller, Rebecca M. Hill, Janet C. Lindsay, Ed C. Schwalbe, Simon Bullock, David W. Ellison, Cynthia Hawkins, David Malkin, Steven C. Clifford, Andrey Korshunov, Stefan Pfister, Michael D. Taylor, and Uri Tabori

Risk stratification for medulloblastoma

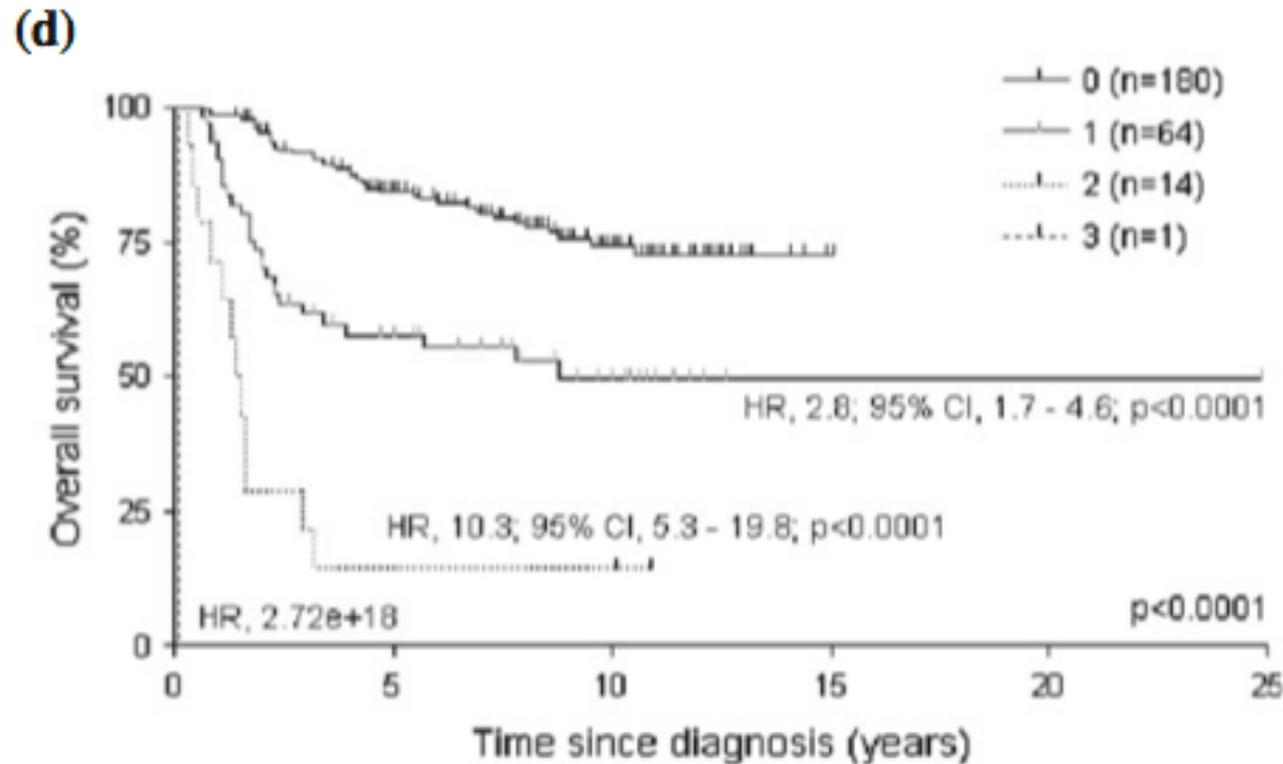
Adverse prognosis

- *MYC* and *MYCN* amplification
- Large Cell/Anaplastic
- *TP53*-mutant, *SHH*-activated

Good prognosis

- Nodular/Desmoplastic
- *WNT*-activated

Poor prognostic factors are additive



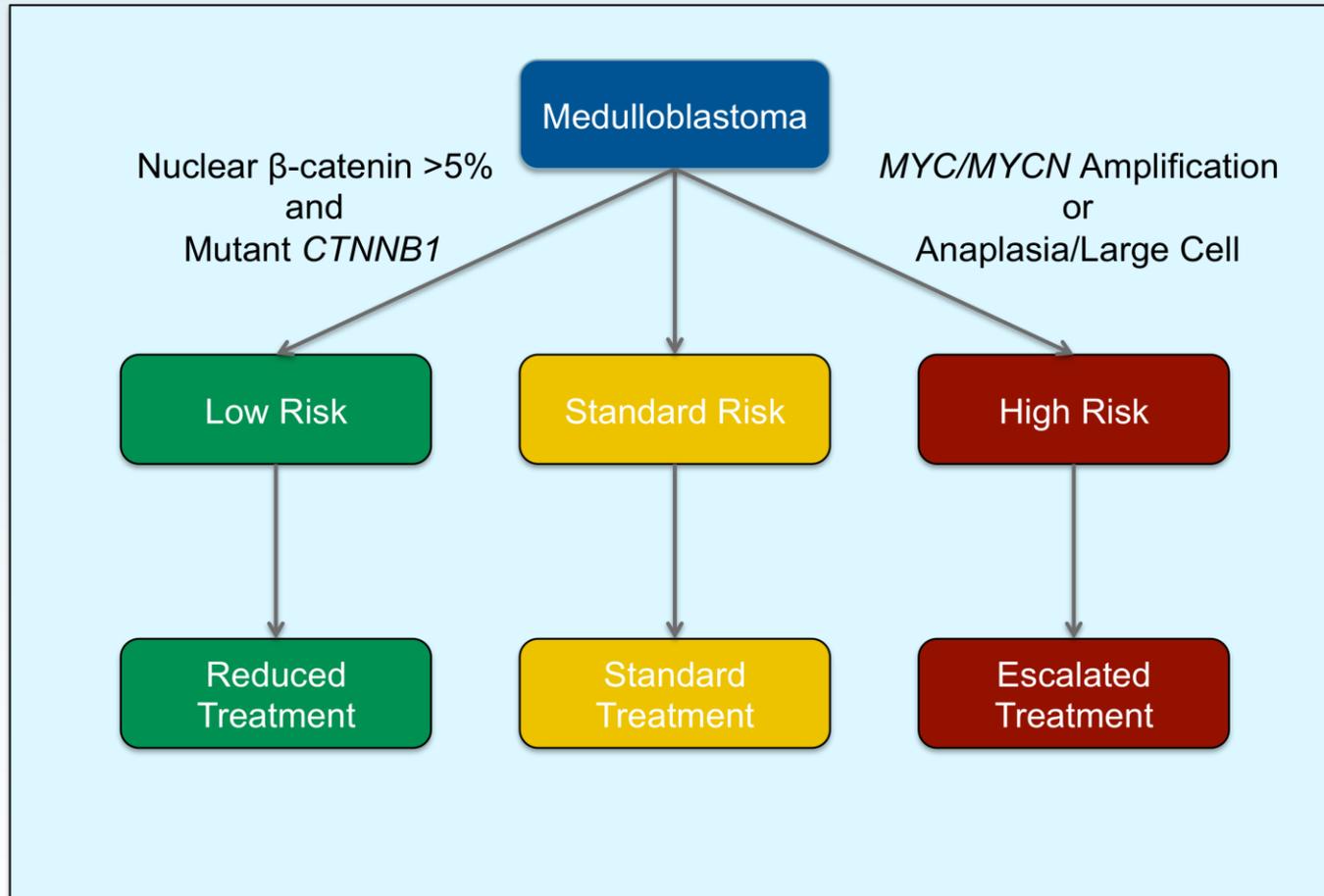
Acta Neuropathol (2012) 123:501–513
 DOI 10.1007/s00401-011-0923-y

ORIGINAL PAPER

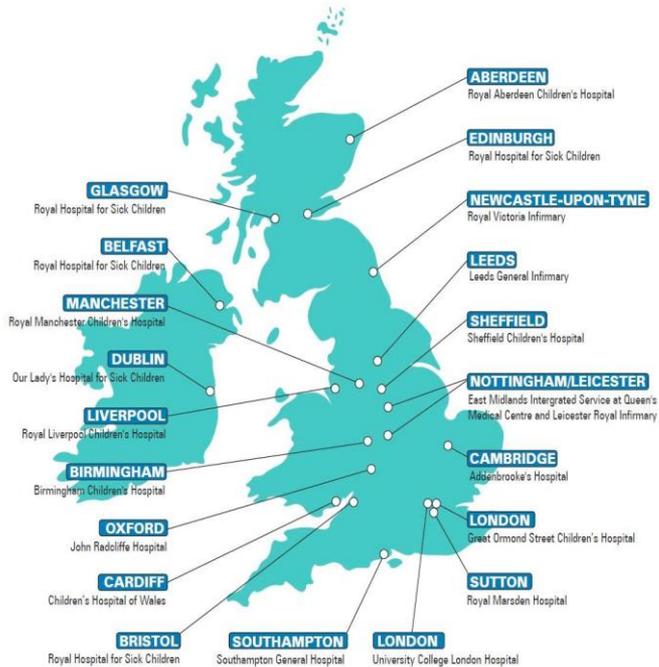
MYC family amplification and clinical risk-factors interact to predict an extremely poor prognosis in childhood medulloblastoma

Sarra L. Ryan · Ed C. Schwalbe · Michael Cole · Yuan Lu · Meryl E. Lusher · Elsham Megahed · Kieran O'Toole · Sarah Leigh Nicholson · Lavinia Bogdan · Miklos Garami · Peter Hammer · Andrey Korshunov · Stefan M. Pfister · Daniel Williamson · Roger E. Taylor · David W. Ellison · Simon Bailey · Steven C. Clifford

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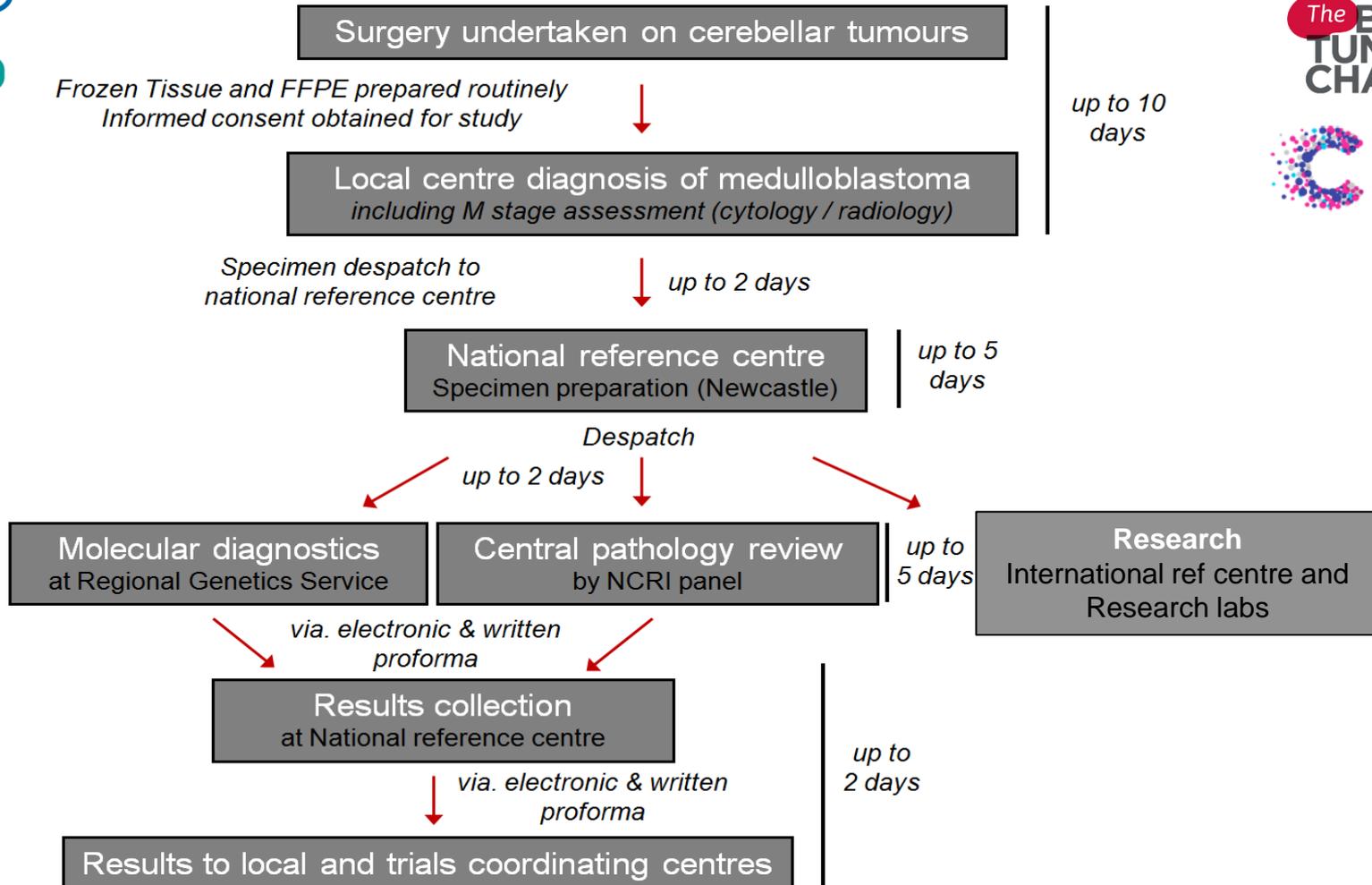
UK feasibility study



22 centres

- Commenced February 2009
- Study information packs circulated
Lead pathologists, oncologists,
centre coordinators
- Frozen tissue
MYC, *MYCN* amplification status
- FFPE tissue:
Central pathology review
WNT (β-catenin) status

Centralised medulloblastoma molecular diagnostics and central pathology review



Does not influence therapy



CHAPTER 8

Embryonal tumours

Medulloblastoma, genetically defined

Medulloblastoma, WNT-activated

Medulloblastoma, SHH-activated and TP53-mutant

Medulloblastoma, SHH-activated and TP53-wildtype

Medulloblastoma, non-WNT/non-SHH

Medulloblastoma, histologically defined

Medulloblastoma, classic

Desmoplastic/nodular medulloblastoma

Medulloblastoma with extensive nodularity

Large cell / anaplastic medulloblastoma

Embryonal tumour with multilayered rosettes, C19MC-altered

Embryonal tumour with multilayered rosettes, NOS

Medulloepithelioma

CNS neuroblastoma

CNS ganglioneuroblastoma

CNS embryonal tumour, NOS

Atypical teratoid/rhabdoid tumour

CNS embryonal tumour with rhabdoid features

Non-Medullo Embryonal Tumours

Embryonal tumour with multilayered rosettes, C19MC-altered

Embryonal tumour with multilayered rosettes, NOS

Medulloepithelioma

CNS neuroblastoma

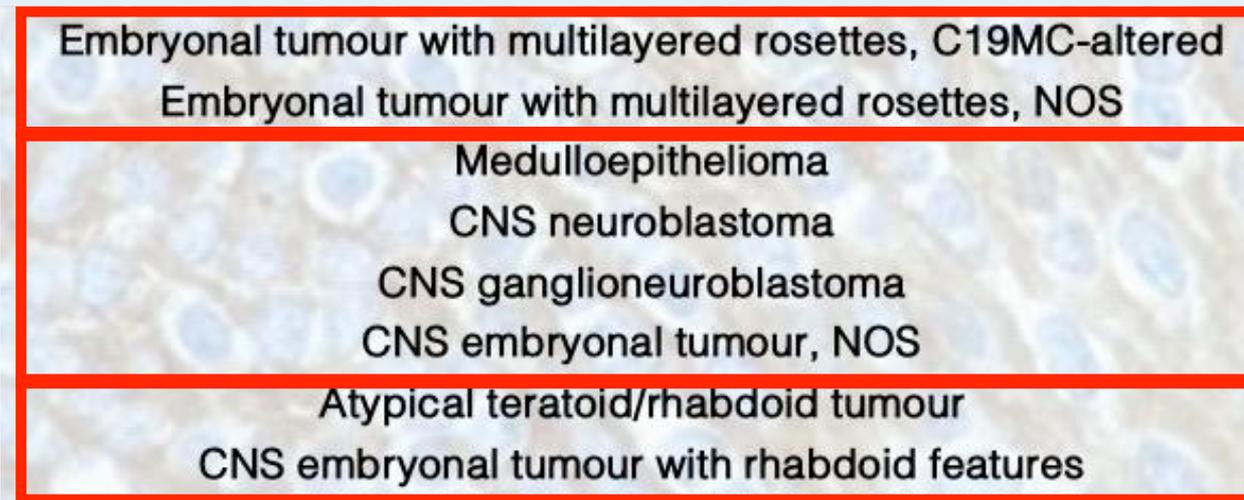
CNS ganglioneuroblastoma

CNS embryonal tumour, NOS

Atypical teratoid/rhabdoid tumour

CNS embryonal tumour with rhabdoid features

Non-Medullo Embryonal Tumours

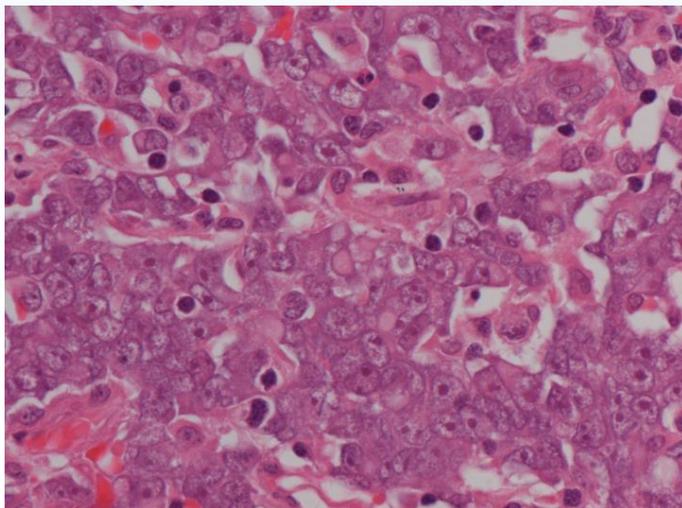
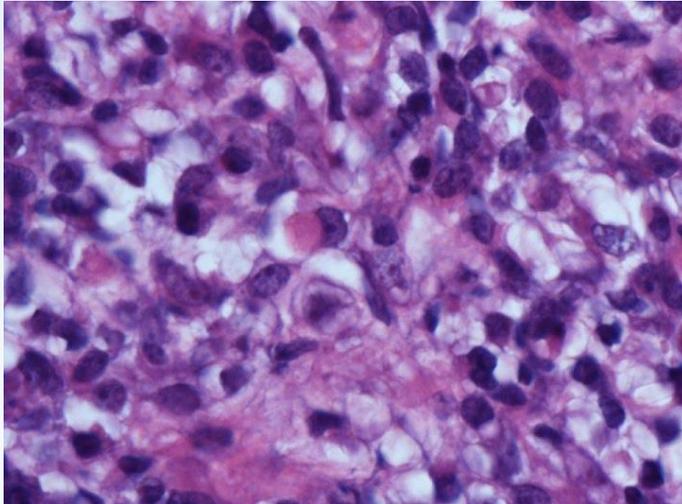


ETMR/ETANTR
 Ependymoblastoma
 Medulloepithelioma

CNS-PNET and variants

ATRT

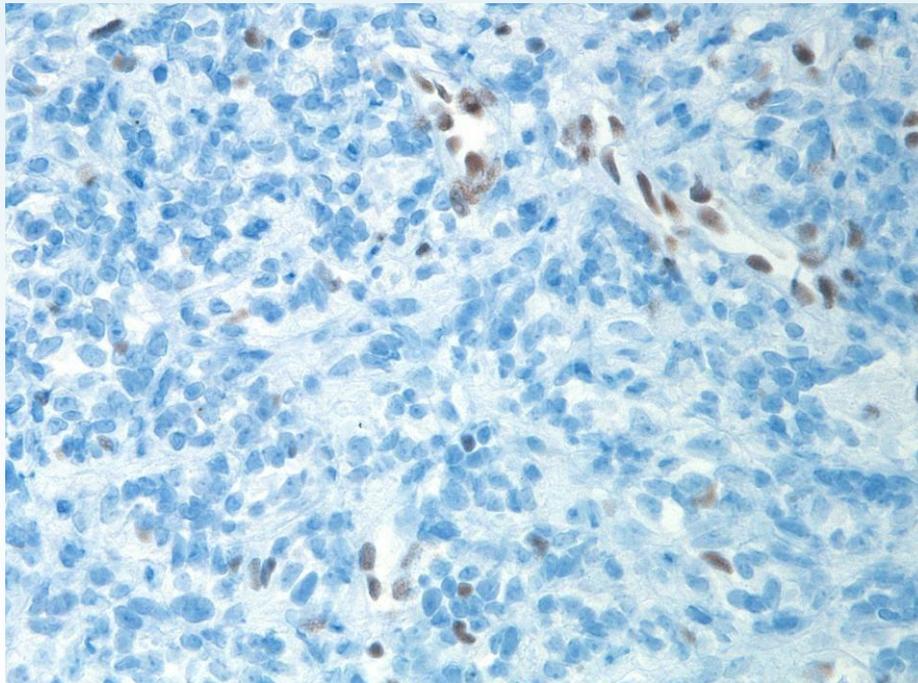
Atypical Teratoid/Rhabdoid Tumour



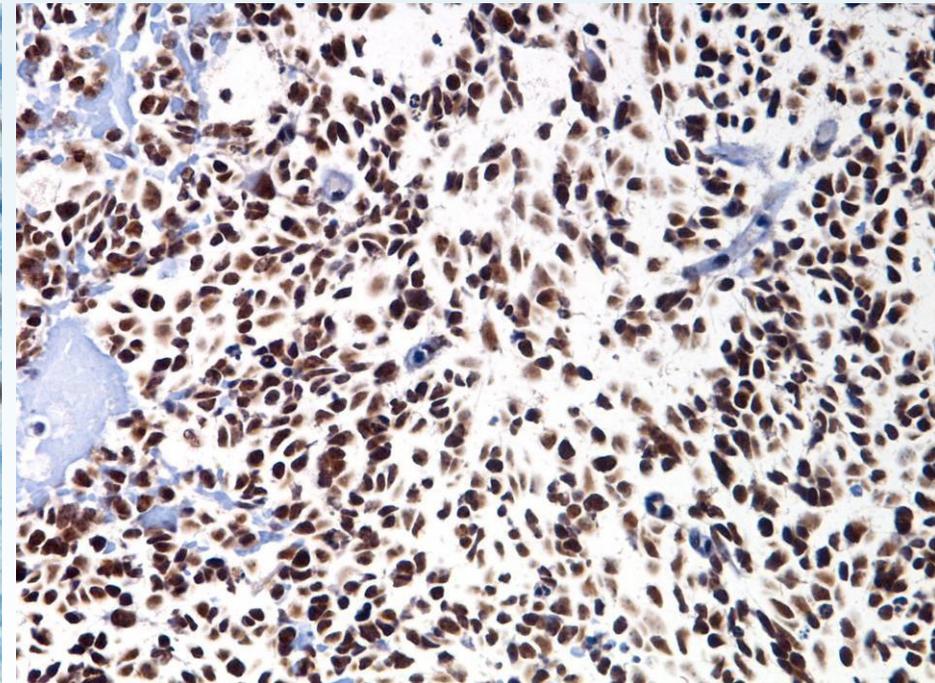
- Age usually <6 years old
- May have typical 'rhabdoid' cytology but usually very unpredictable morphology
- Protean immunohistochemistry
- Associated with mutations in *SMARCB1* (INI1) gene
- Rarely *SMARCA4* (BRG1) mutation
- Median survival typically short
- Germline predisposition

Atypical Teratoid/Rhabdoid Tumour

INI-1

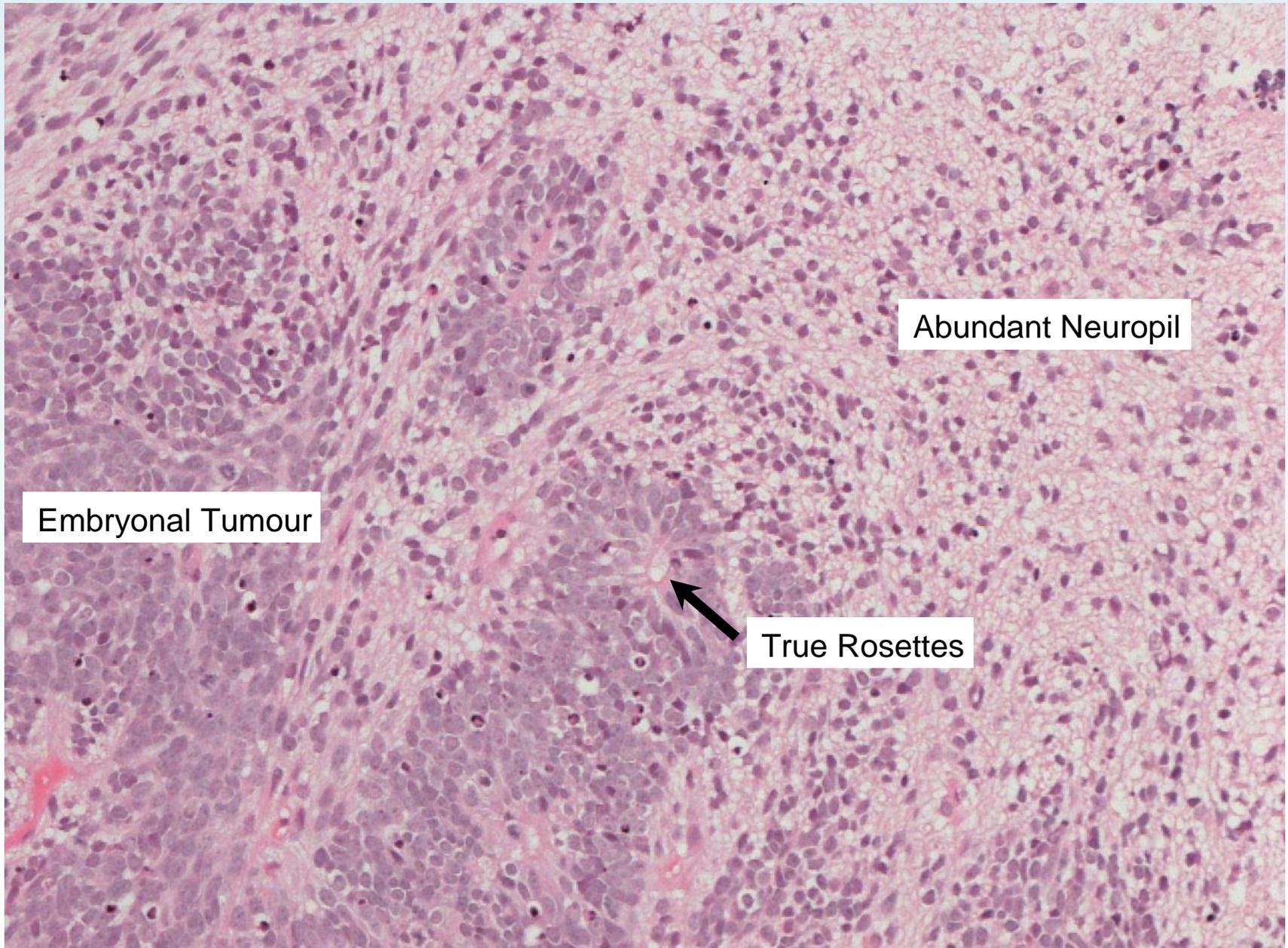


ATRT



Medulloblastoma

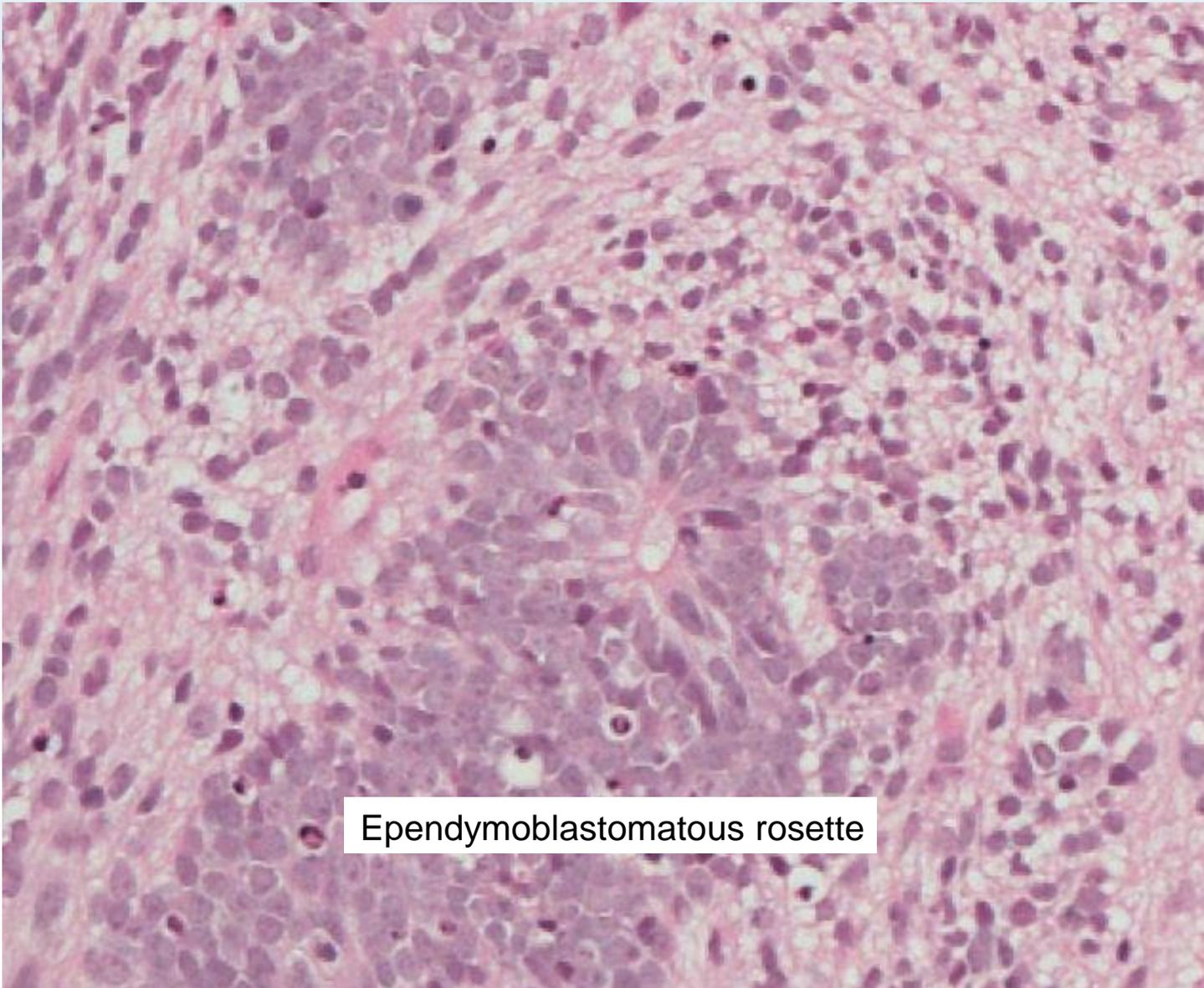
Embryonal Tumour with Multilayered Rosettes (ETMR), C19MC altered



Embryonal Tumour

Abundant Neuropil

True Rosettes

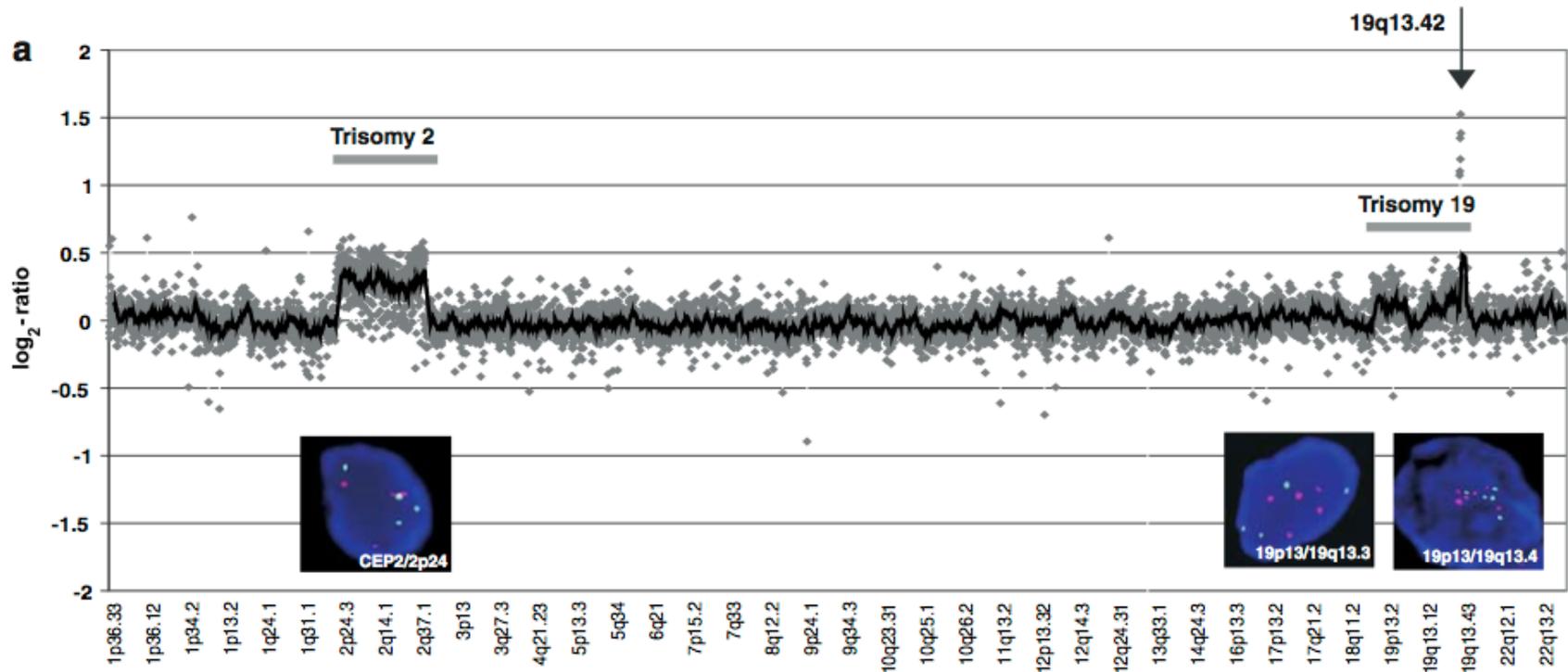


Ependymoblastomatous rosette

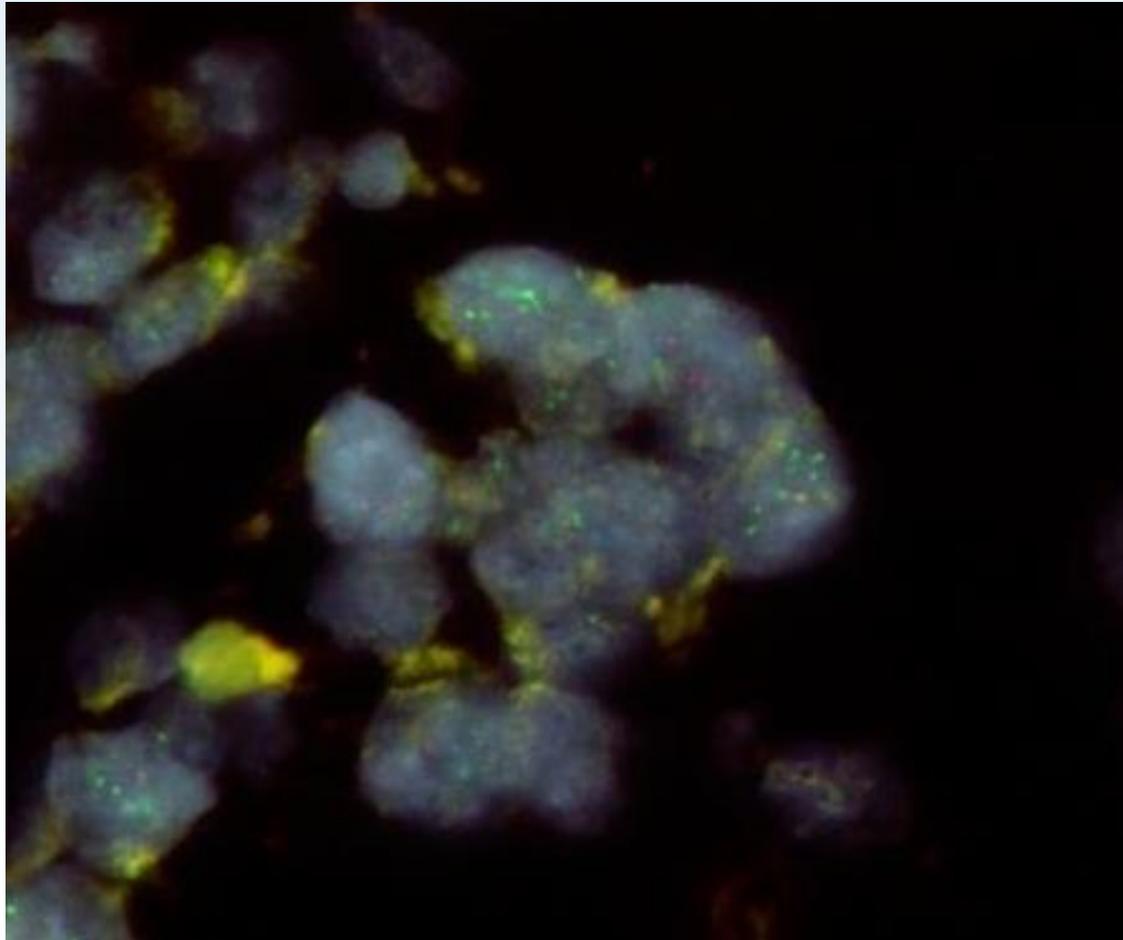
CASE REPORT

Novel genomic amplification targeting the microRNA cluster at 19q13.42 in a pediatric embryonal tumor with abundant neuropil and true rosettes

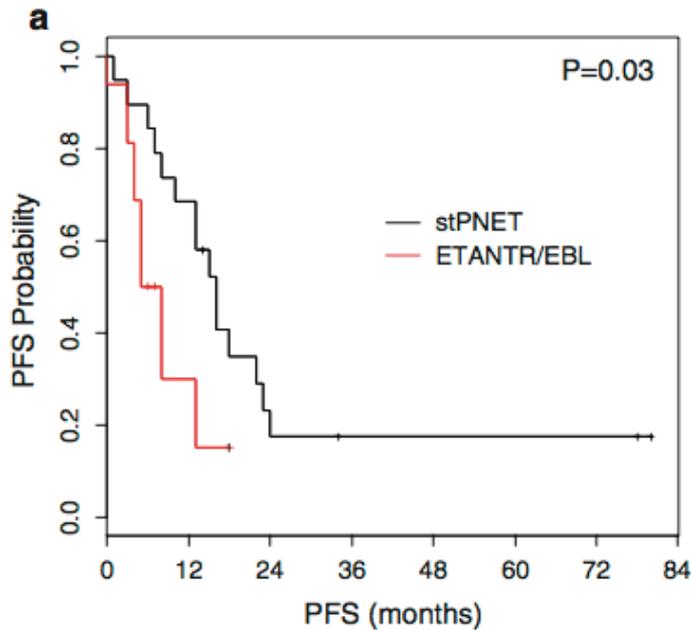
Stefan Pfister · Marc Remke · Mirco Castoldi · Alfa H. C. Bai ·
 Martina U. Muckenthaler · Andreas Kulozik · Andreas von Deimling ·
 Armin Pscherer · Peter Lichter · Andrey Korshunov



FISH for 19q amplification

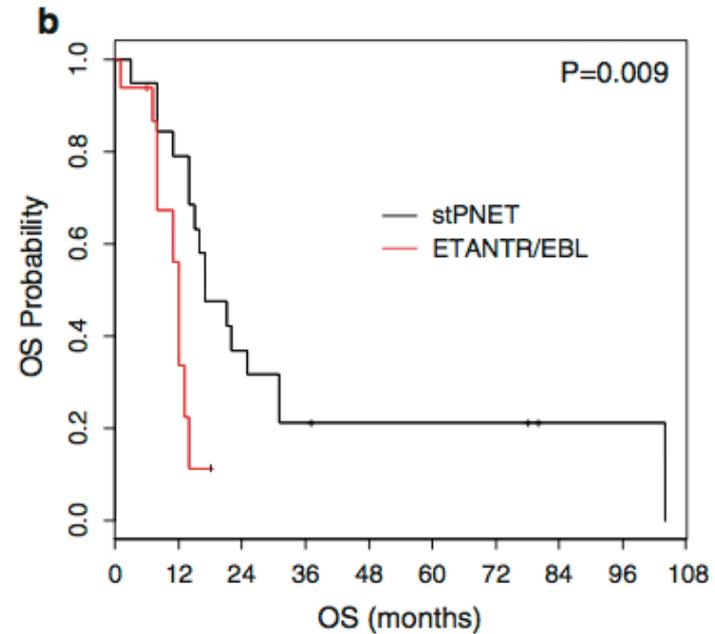


ETMR have a very poor prognosis



Number of patients at risk

19	13	4	2	2	2	2	classic_stPNET
16	2						ETANTR/EBL



Number of patients at risk

19	15	7	4	3	3	3	1	1	classic_stPNET
16	5								ETANTR/EBL

Acta Neuropathol
DOI 10.1007/s00401-010-0688-8

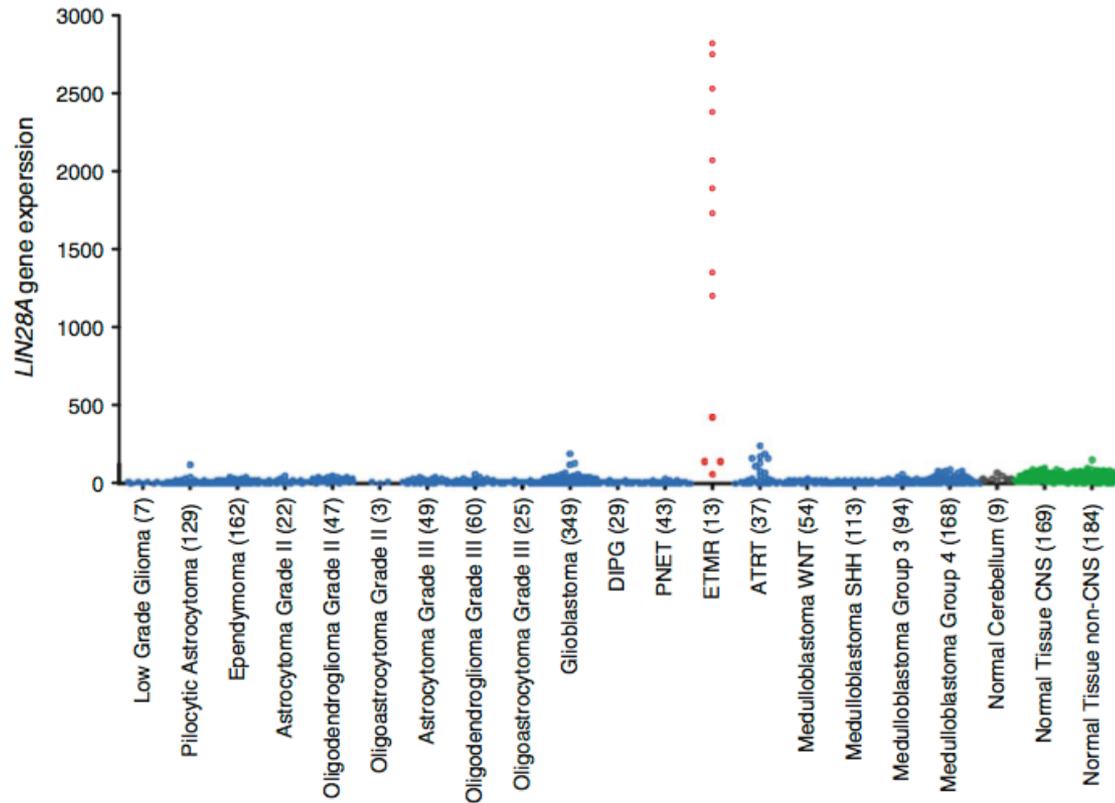
ORIGINAL PAPER

Focal genomic amplification at 19q13.42 comprises a powerful diagnostic marker for embryonal tumors with ependymoblastic rosettes

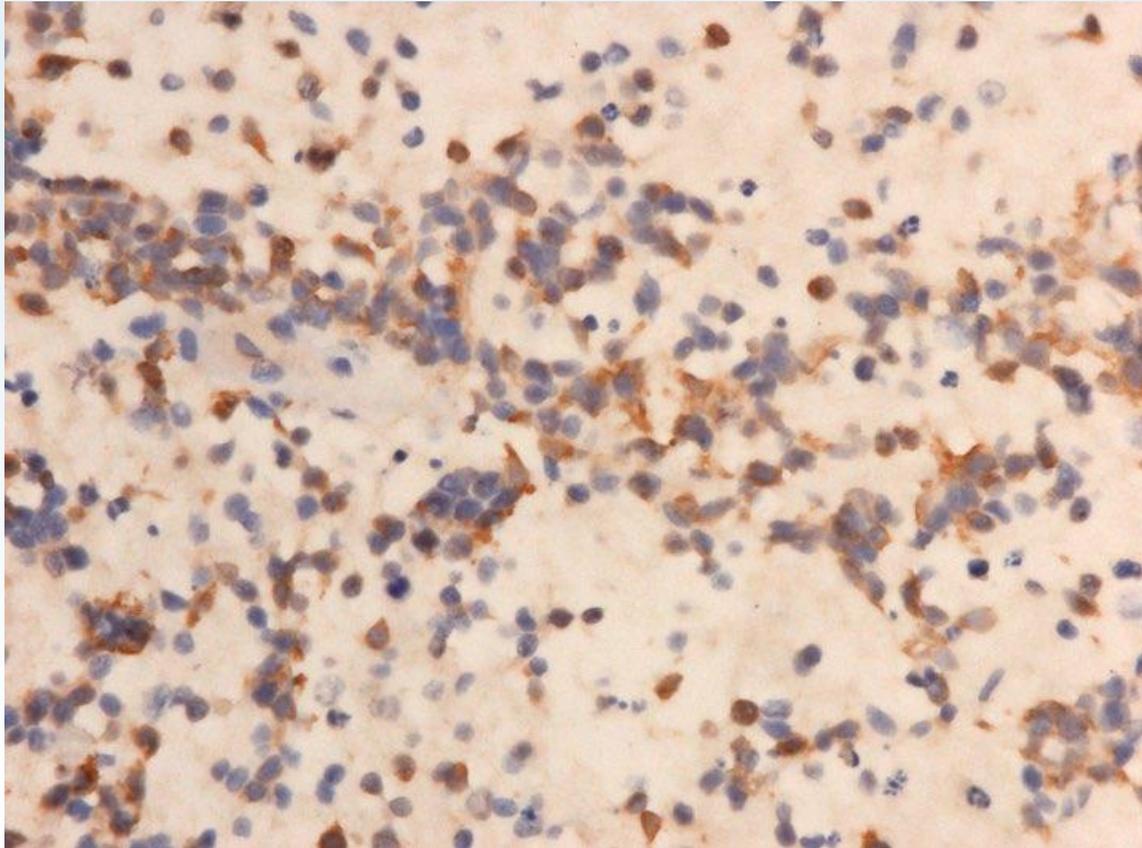
Andrey Korshunov · Marc Renke · Marco Gessi · Marina Ryzhova · Thomas Hielscher · Hendrik Witt · Vivienne Tobias · Anna Maria Buccolero · Jacopo Sardi · Marina Paola Gardiman · Jose Bonnin · Bernd Scheithauer · Andreas E. Kaloupek · Olaf Witt · Sverre Mørk · Andreas von Deimling · Omar D. Wiestler · Felice Giangaspero · Marc Rosenblum · Torsten Pietsch · Peter Lichter · Stefan M. Pfister

LIN28A immunoreactivity is a potent diagnostic marker of embryonal tumor with multilayered rosettes (ETMR)

Andrey Korshunov · Marina Ryzhova · David T. W. Jones · Paul A. Northcott · Peter van Sluis · Richard Volckmann · Jan Koster · Rogier Versteeg · Cynthia Cowdrey · Arie Perry · Daniel Picard · Marc Rosenblum · Felice Giangaspero · Eleonora Aronica · Ulrich Schüller · Martin Hasselblatt · V. Peter Collins · Andreas von Deimling · Peter Lichter · Annie Huang · Stefan M. Pfister · Marcel Kool

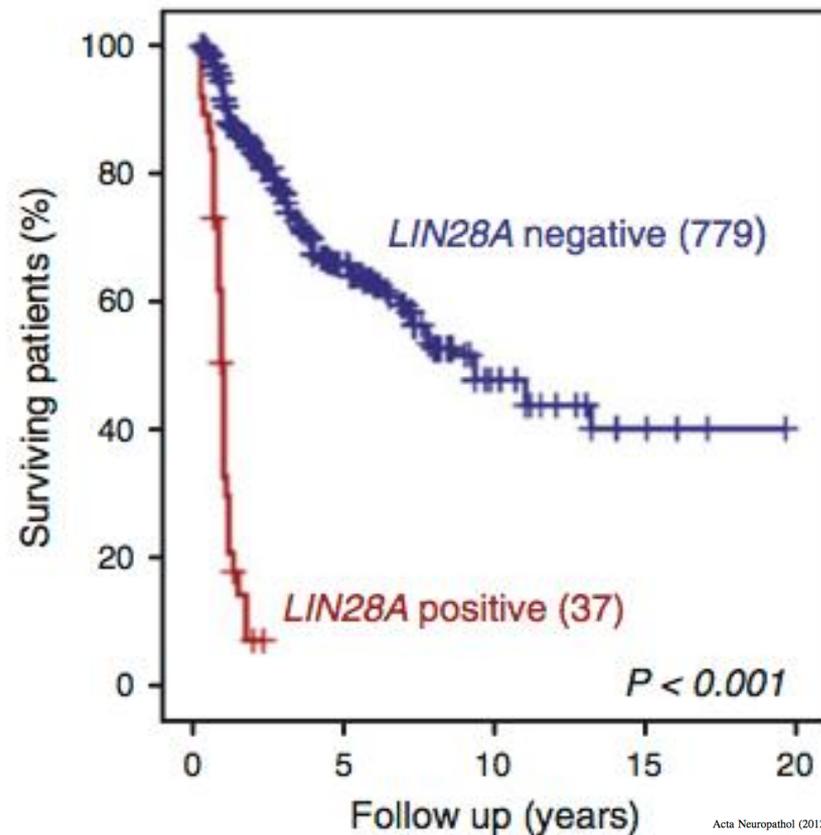


ETMR: Immunohistochemistry



LIN28a

LIN28A tumours have a poor prognosis



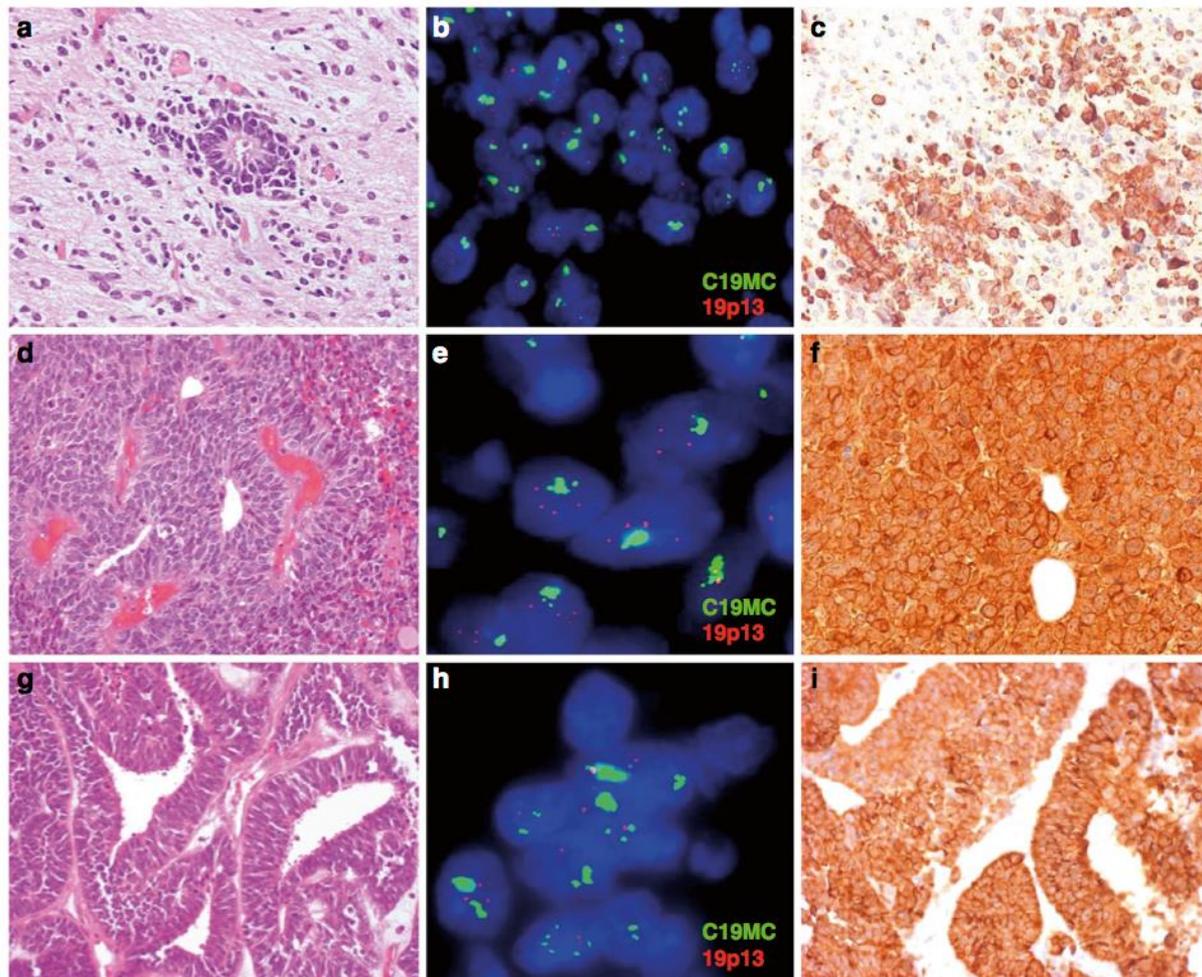
Acta Neuropathol (2012) 124:875–881
DOI 10.1007/s00401-012-1068-3

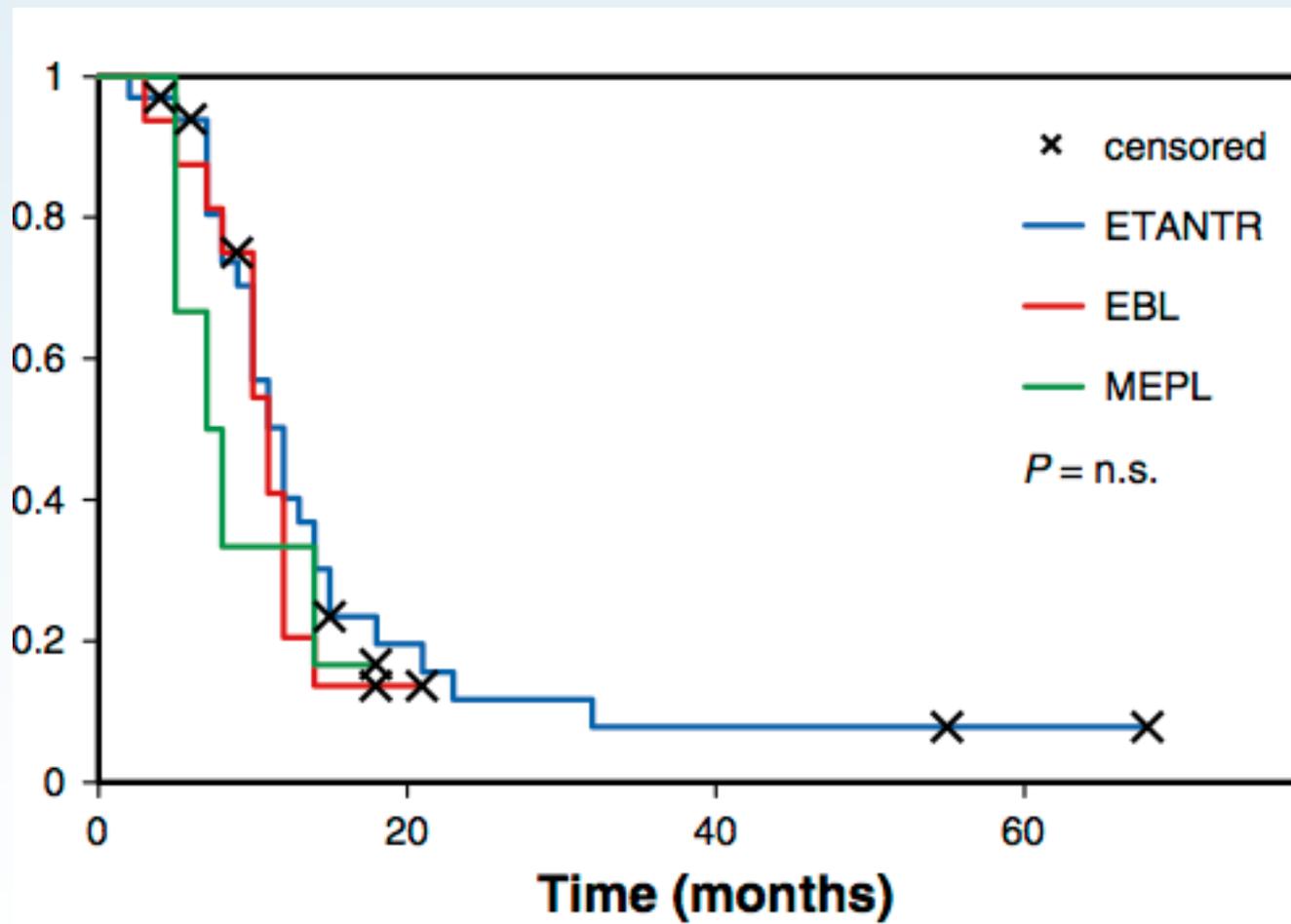
ORIGINAL PAPER

LIN28A immunoreactivity is a potent diagnostic marker of embryonal tumor with multilayered rosettes (ETMR)

Andrey Korshunov · Marina Ryzhova · David T. W. Jones · Paul A. Northcott · Peter van Sluis · Richard Volckmann · Jan Koster · Rogier Versteeg · Cynthia Cowdrey · Arie Perry · Daniel Picard · Marc Rosenblum · Felice Giangaspero · Eleonora Aronica · Ulrich Schüller · Martin Hasselblatt · V. Peter Collins · Andreas von Deimling · Peter Lichter · Annie Huang · Stefan M. Pfister · Marcel Kool

Embryonal tumor with abundant neuropil and true rosettes (ETANTR), ependymoblastoma, and medulloepithelioma share molecular similarity and comprise a single clinicopathological entity





Fusion of *TTYH1* with the C19MC microRNA cluster drives expression of a brain-specific *DNMT3B* isoform in the embryonal brain tumor ETMR

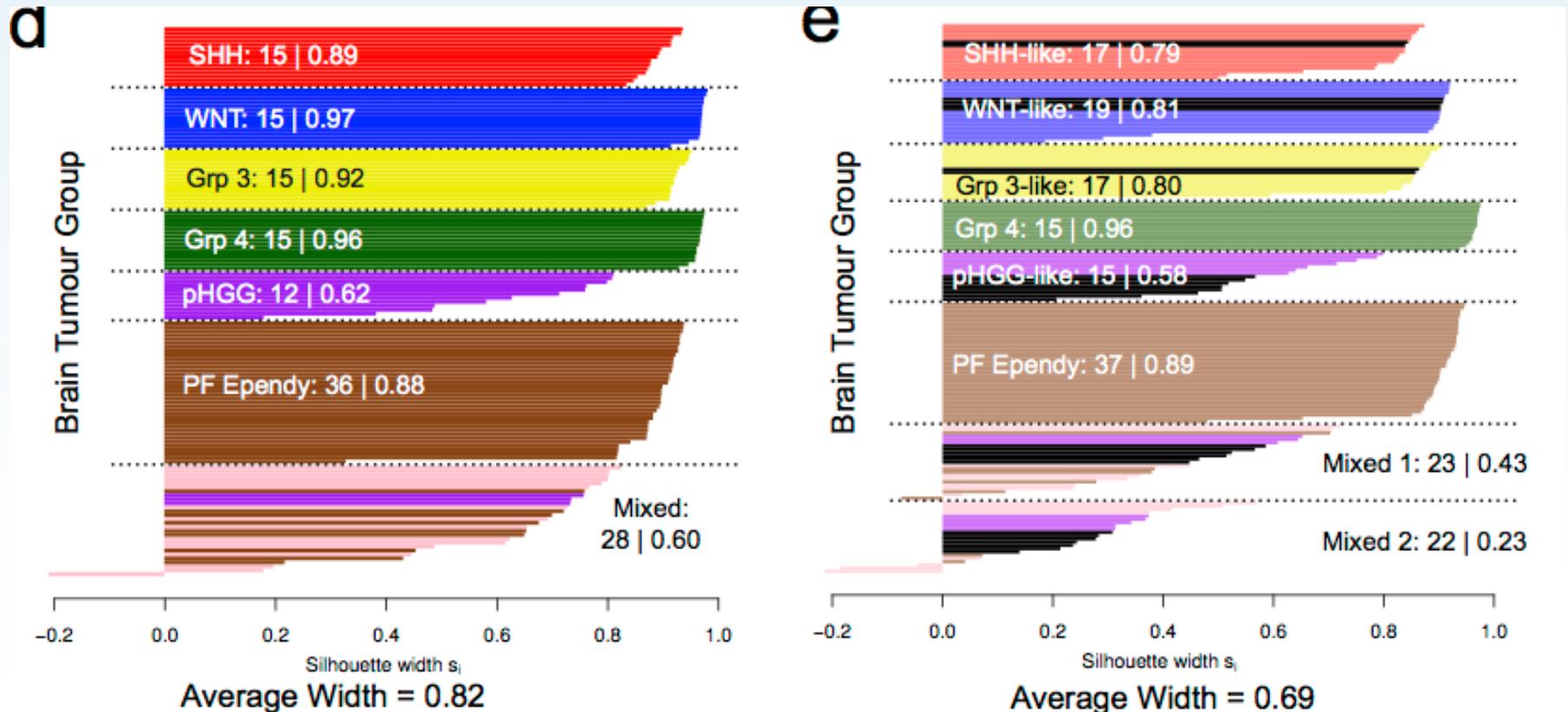
Claudia L Kleinman^{1,2,17}, Noha Gerges^{2,17}, Simon Papillon-Cavanagh¹, Patrick Sin-Chan³, Albena Pramatarova¹, Dong-Anh Khuong Quang², Véronique Adoue¹, Stephan Busche¹, Maxime Caron¹, Haig Djambazian¹, Amandine Bemmo¹, Adam M Fontebasso⁴, Tara Spence³, Jeremy Schwartzentruber¹, Steffen Albrecht⁵, Peter Hauser⁶, Miklos Garami⁶, Almos Klekner⁷, Laszlo Bognar⁷, Jose-Luis Montes⁸, Alfredo Staffa¹, Alexandre Montpetit¹, Pierre Berube¹, Magdalena Zakrzewska⁹, Krzysztof Zakrzewski¹⁰, Pawel P Liberski⁹, Zhifeng Dong¹¹, Peter M Siegel¹¹, Thomas Duchaine¹², Christian Perotti¹³, Adam Fleming¹⁴, Damien Faury¹⁴, Marc Remke¹⁵, Marco Gallo¹⁵, Peter Dirks¹⁵, Michael D Taylor¹⁵, Robert Sladek^{1,2}, Tomi Pastinen¹, Jennifer A Chan¹³, Annie Huang^{3,16,18}, Jacek Majewski^{1,2,18} & Nada Jabado^{2,4,18}

What other tumour are there?

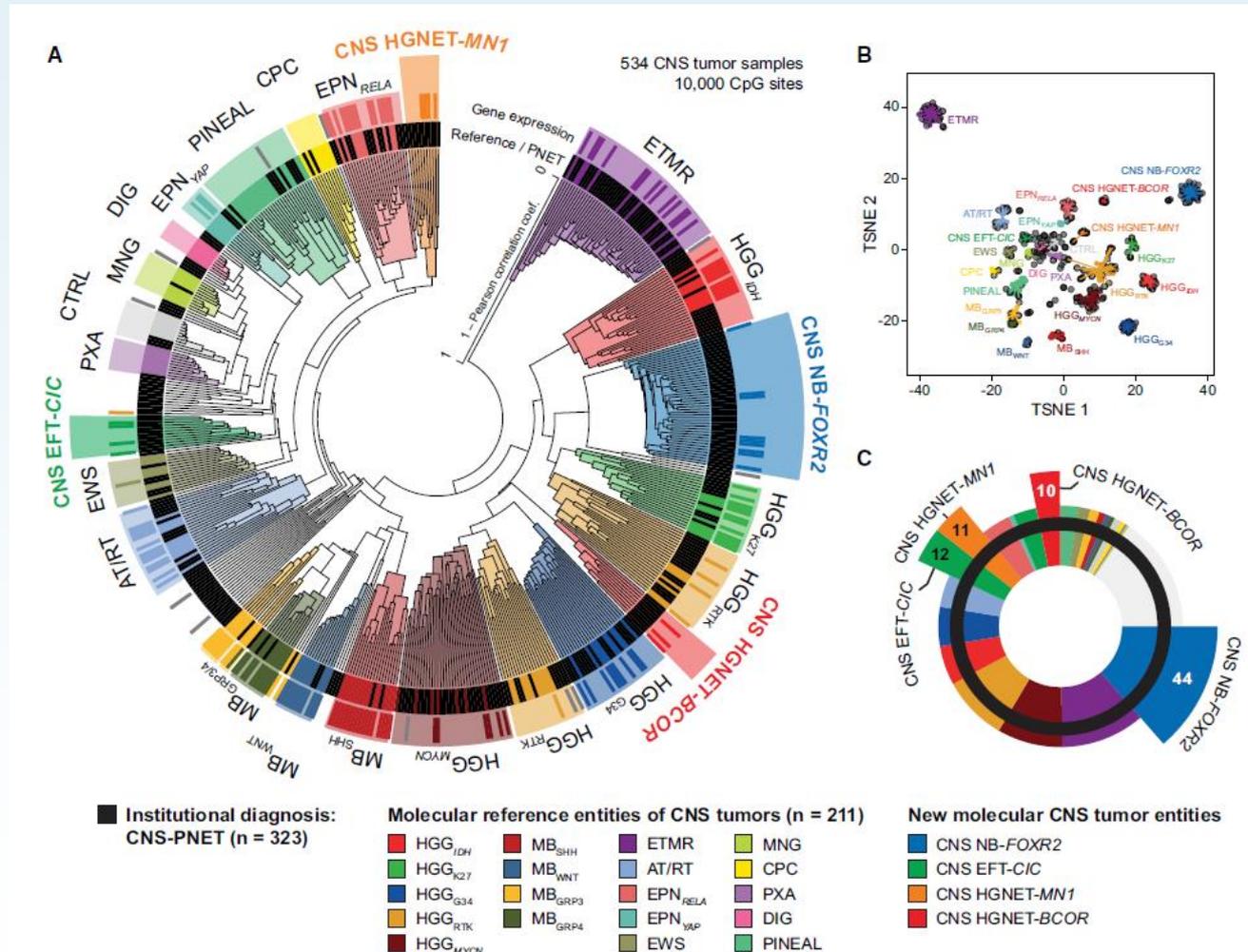
Histologically defined central nervous system primitive neuro-ectodermal tumours (CNS-PNETs) display heterogeneous DNA methylation profiles and show relationships to other paediatric brain tumour types

Ed. C. Schwalbe · James T. Hayden · Hazel A. Rogers · Suzanne Miller · Janet C. Lindsey · Rebecca M. Hill · Sarah-Leigh Nicholson · John-Paul Kilday · Martyna Adamowicz-Brice · Lisa Storer · Thomas S. Jacques · Keith Robson · Jim Lowe · Daniel Williamson · Richard G. Grundy · Simon Bailey · Steven C. Clifford

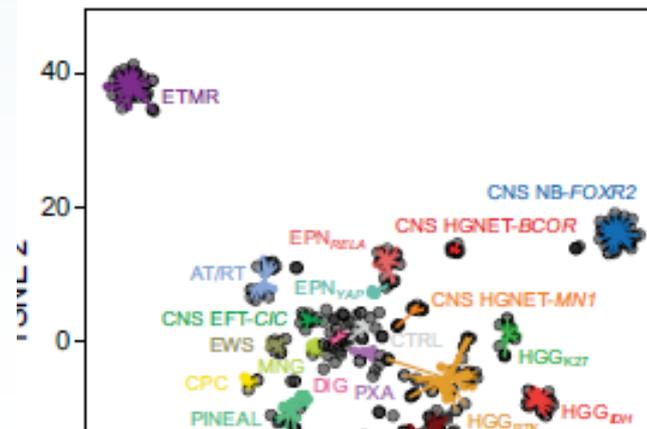
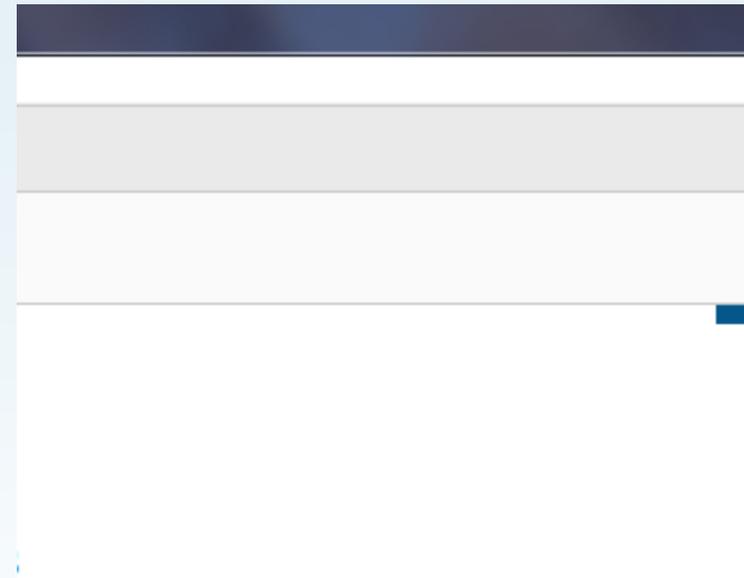
Received: 29 October 2013 / Accepted: 1 November 2013 / Published online: 9 November 2013
 © Springer-Verlag Berlin Heidelberg 2013

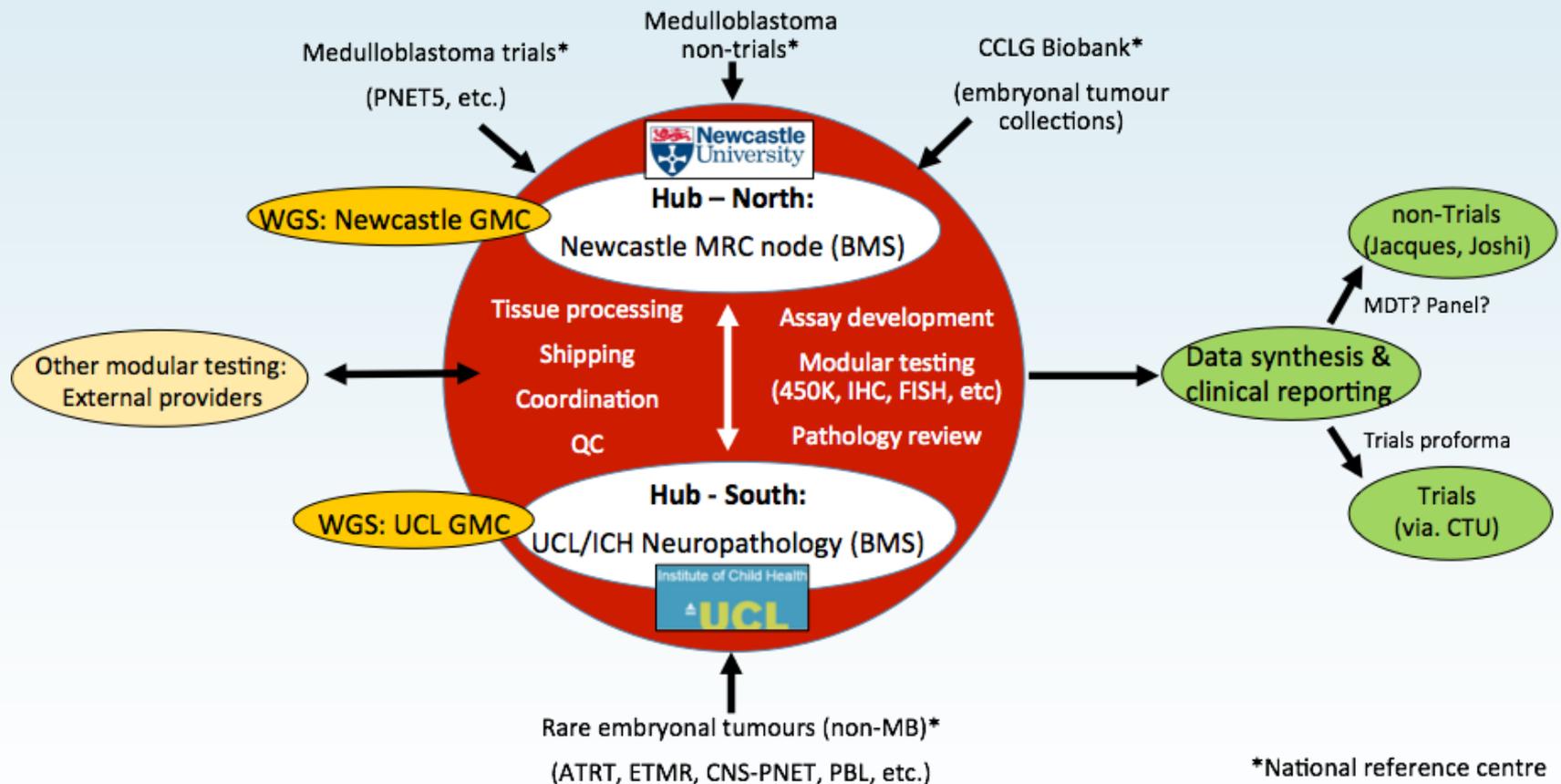


Novel tumour types



Novel tumour types





Astrocytomas in childhood

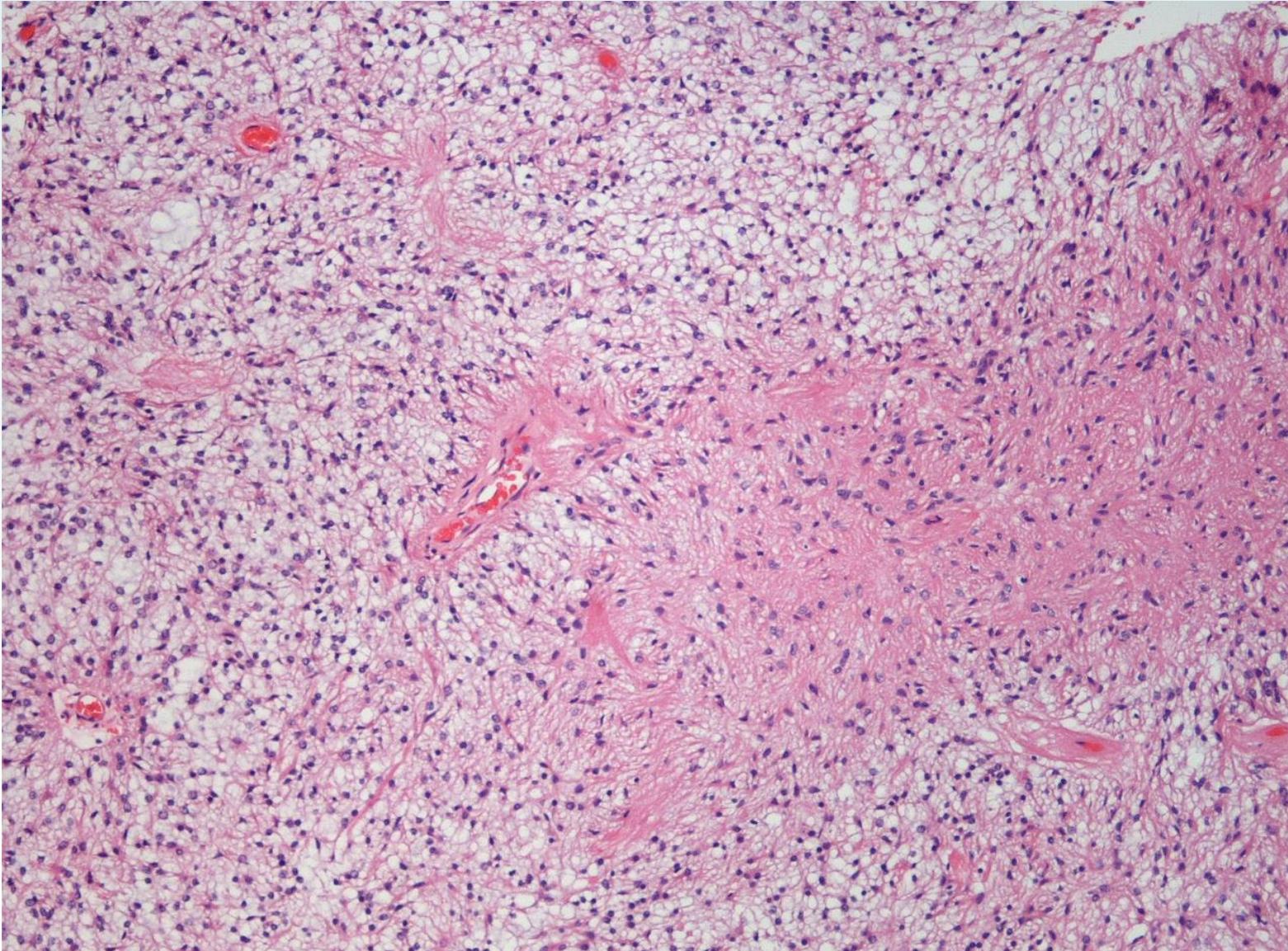
Gliomas

Diffuse astrocytic and oligodendroglial tumours

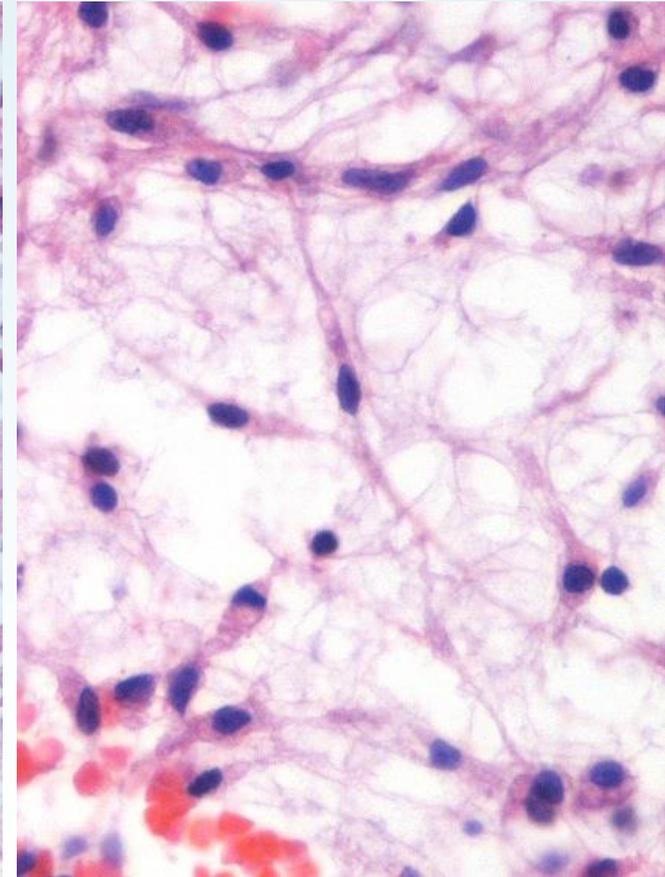
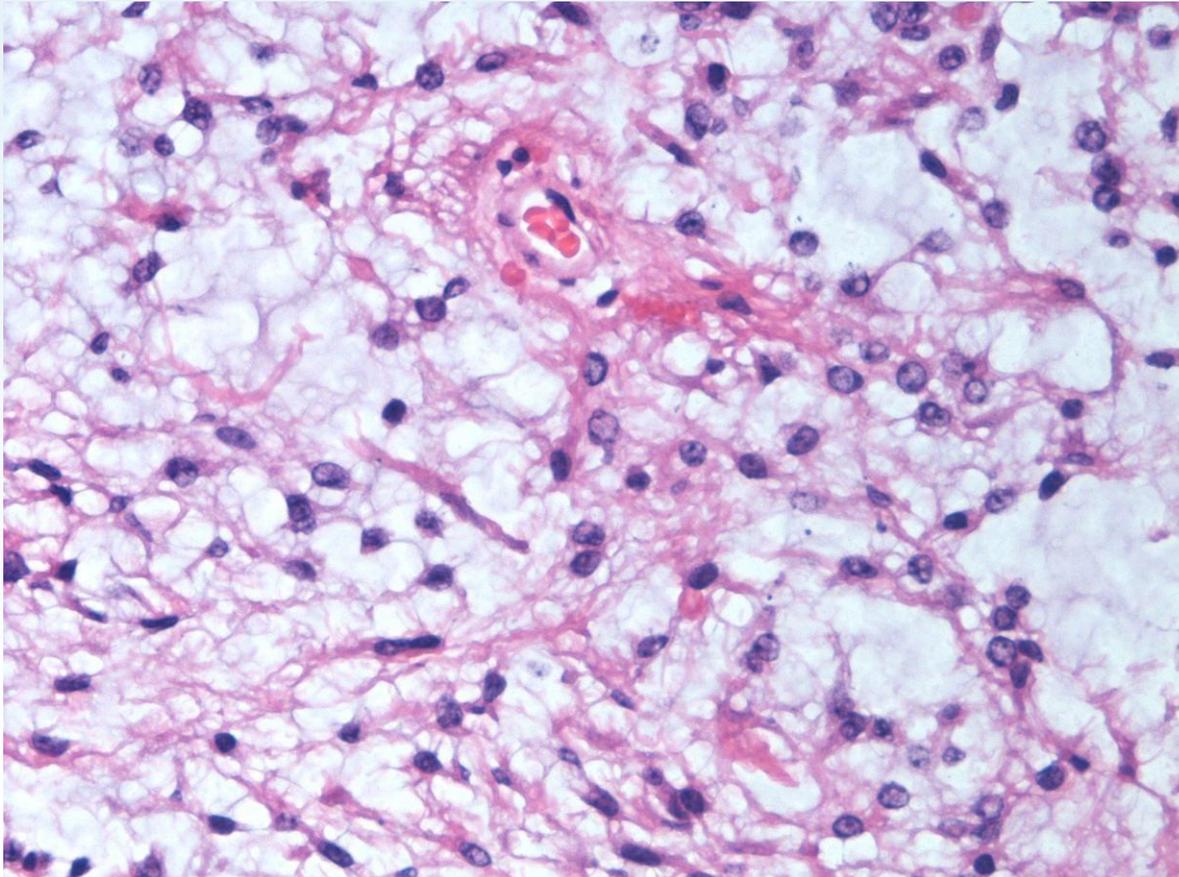
Other astrocytic tumours

- Pilocytic astrocytoma
- Pilomyxoid astrocytoma
- SEGA
- PXA
- Anaplastic PXA

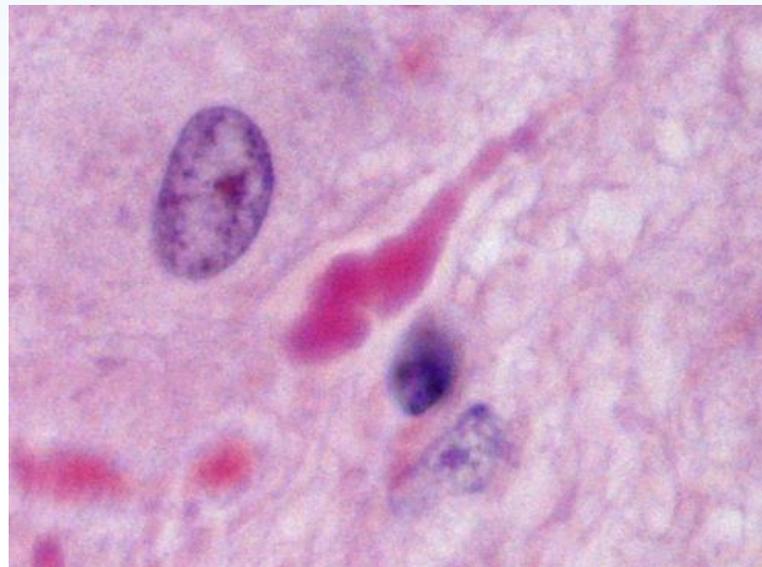
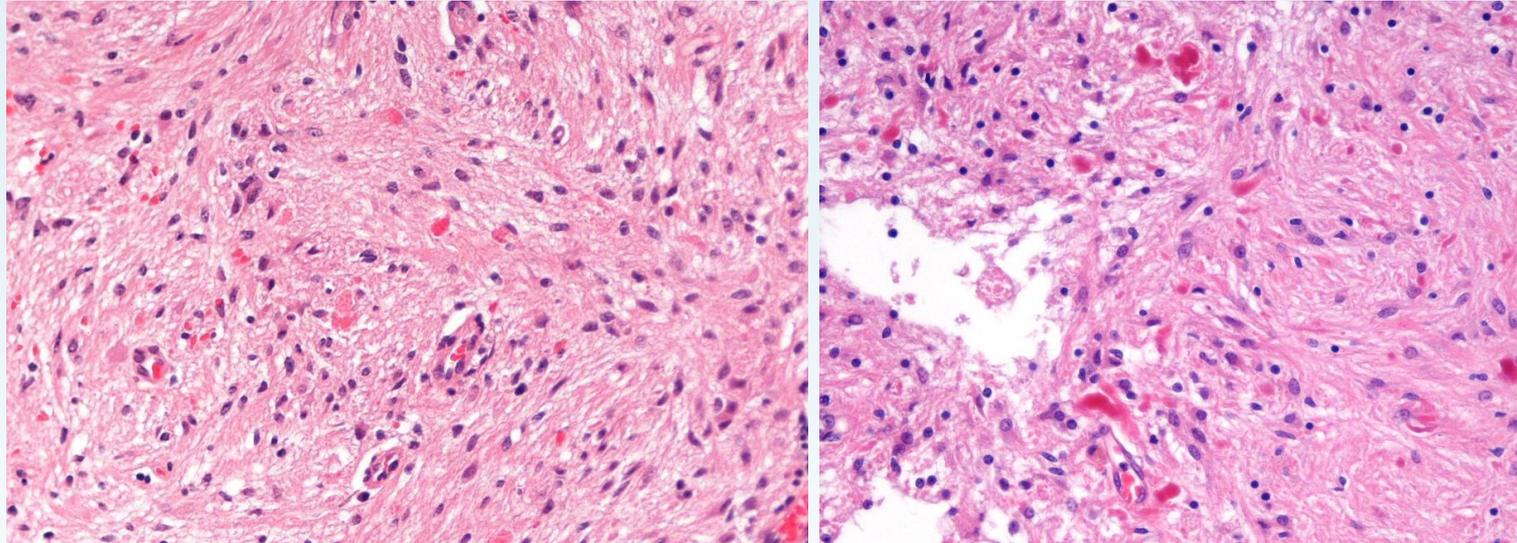
Pilocytic astrocytoma-Architecture



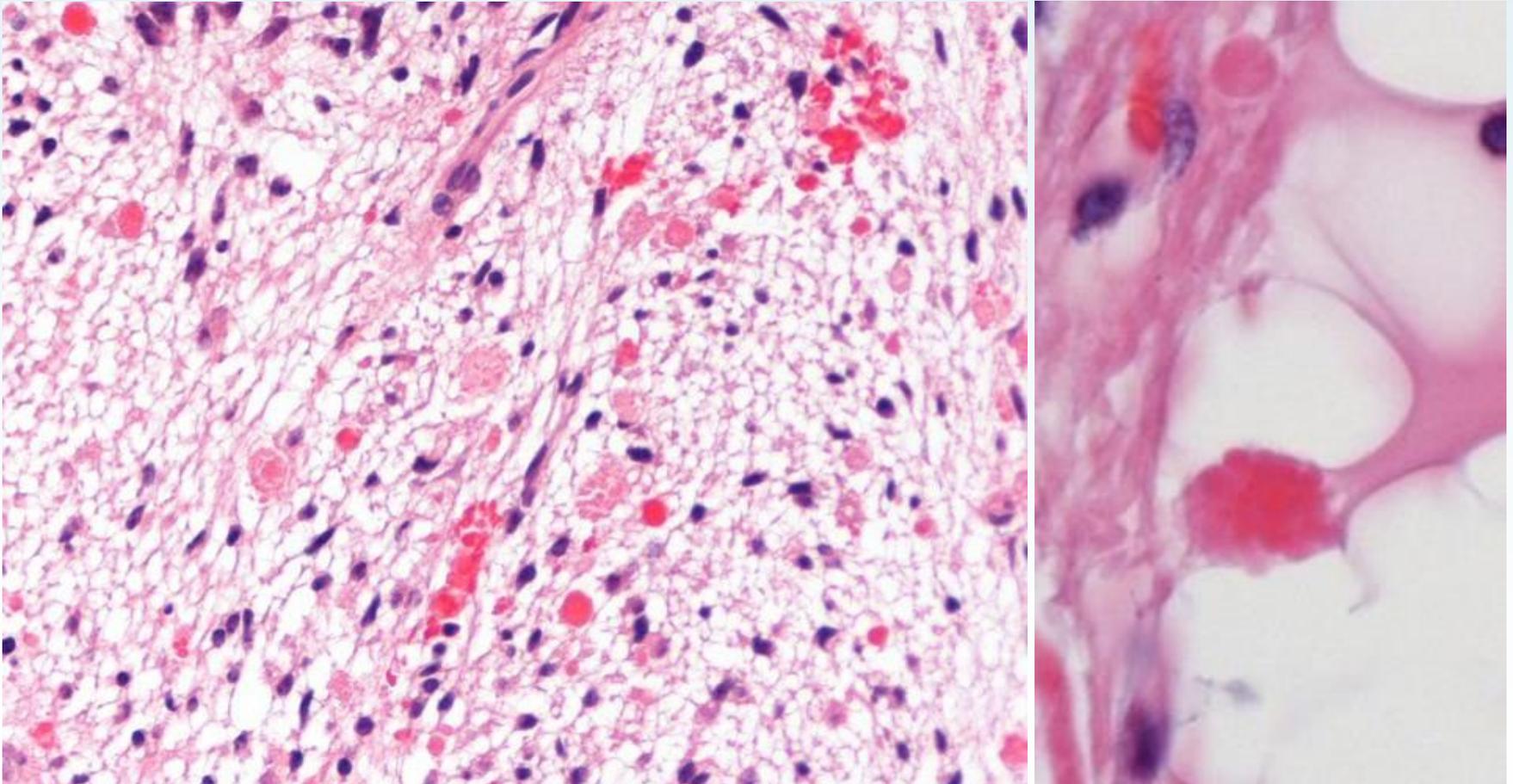
Pilocytic astrocytoma-Cytology



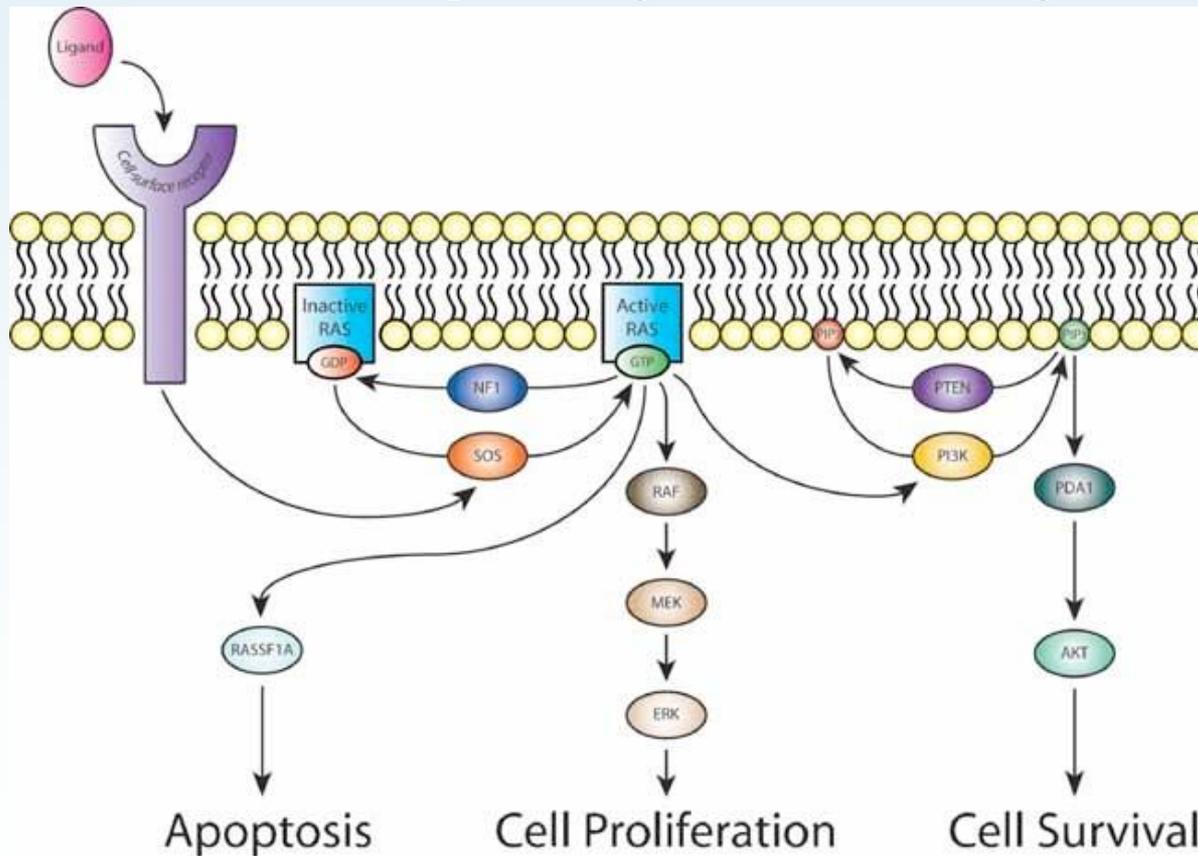
Pilocytic astrocytoma: Rosenthal fibres



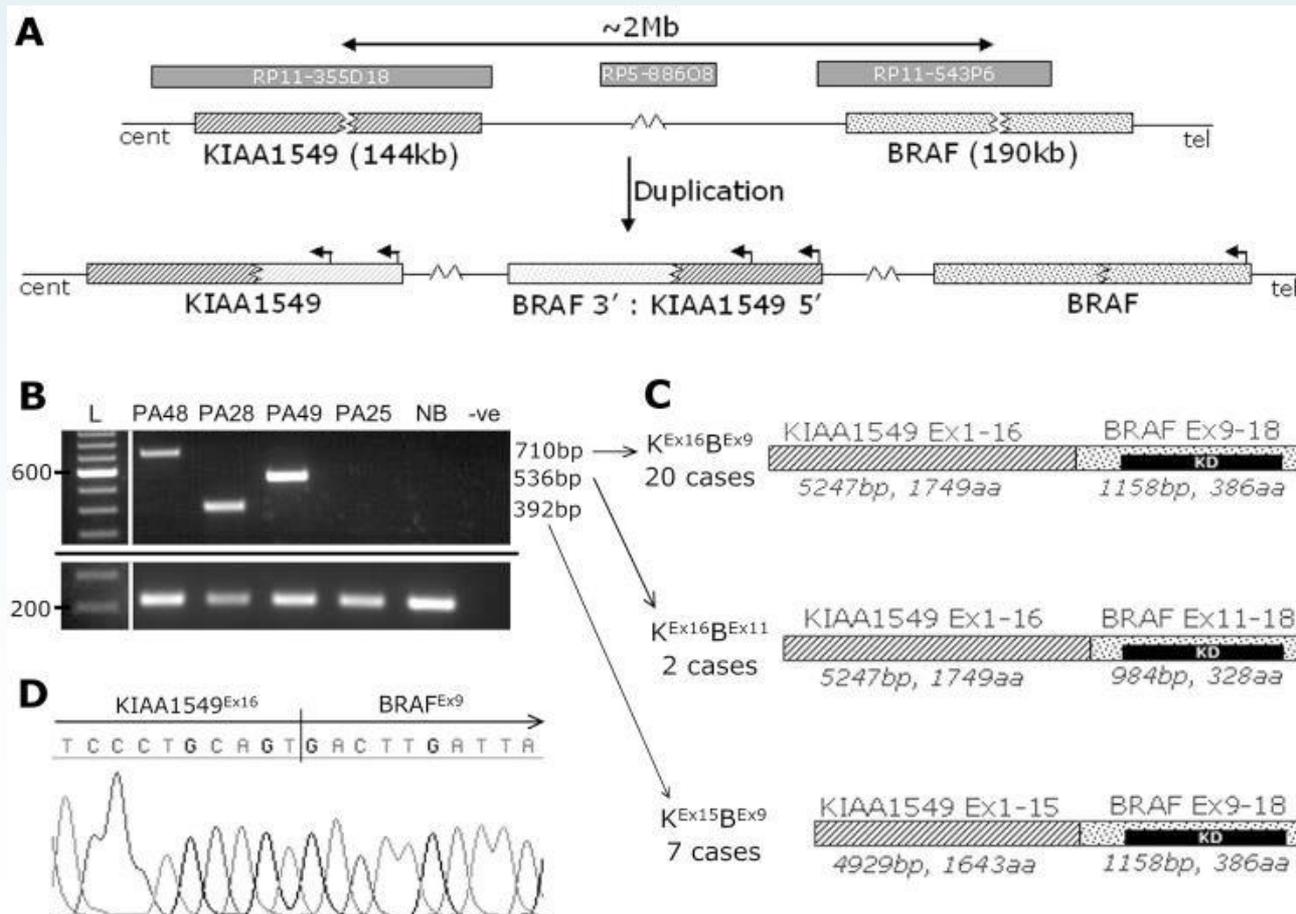
Pilocytic astrocytoma: Eosinophilic Granular Bodies (EGBs)



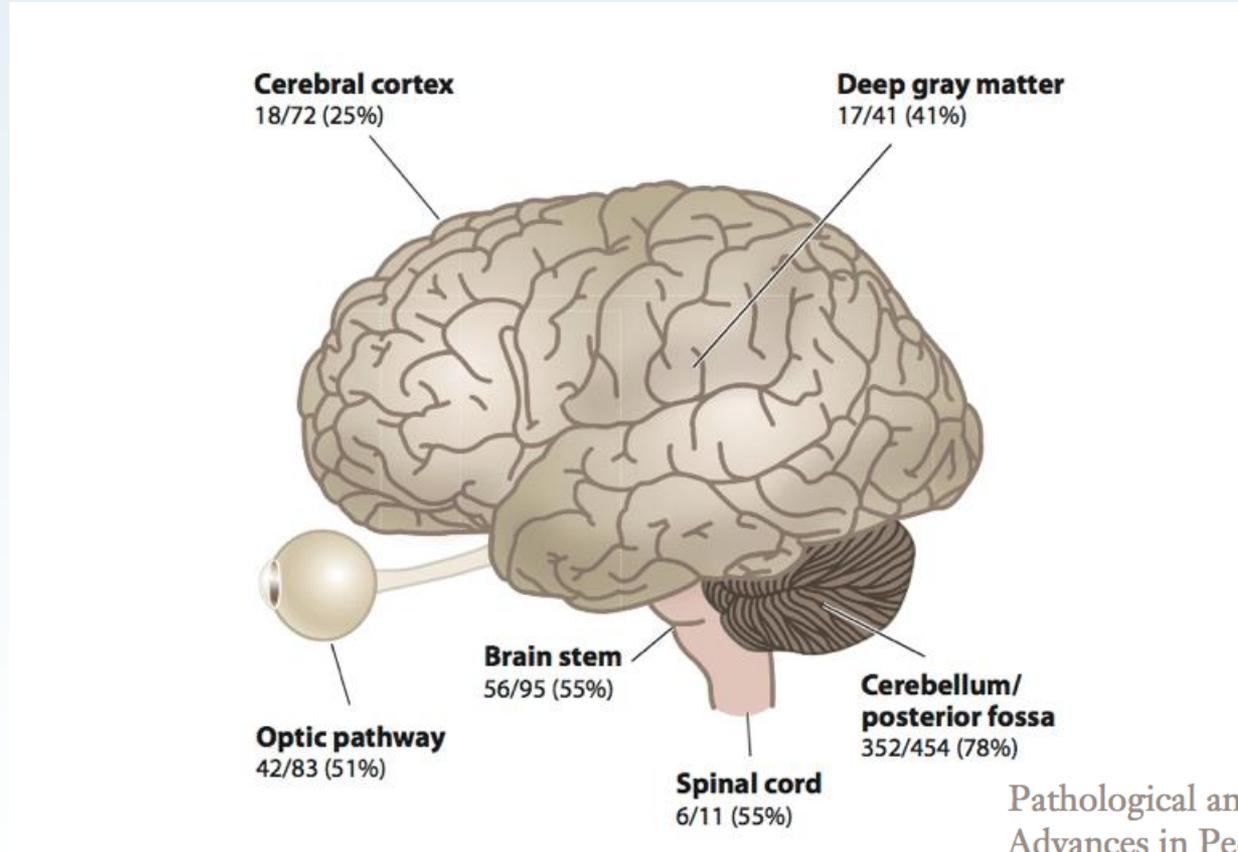
Abnormalities of the MAPK pathway characterise pilocytic astrocytoma



BRAF fusions are characteristic of pilocytic astrocytoma



The frequency of *BRAF* fusions vary with anatomical site



Pathological and Molecular
Advances in Pediatric
Low-Grade Astrocytoma

Fausto J. Rodriguez,¹ Kah Suan Lim,¹
Daniel Bowers,⁴ and Charles G. Eberhart^{1,2,3}

Departments of ¹Pathology, ²Ophthalmology, and ³Oncology, Johns Hopkins University
School of Medicine, Baltimore, Maryland 21205; email: ceberha@jhmi.edu

⁴Department of Pediatrics, University of Texas Southwestern Medical School, Dallas,
Texas 75390

***BRAF* genetics are diagnostically useful**

Acta Neuropathol (2010) 120:271–273

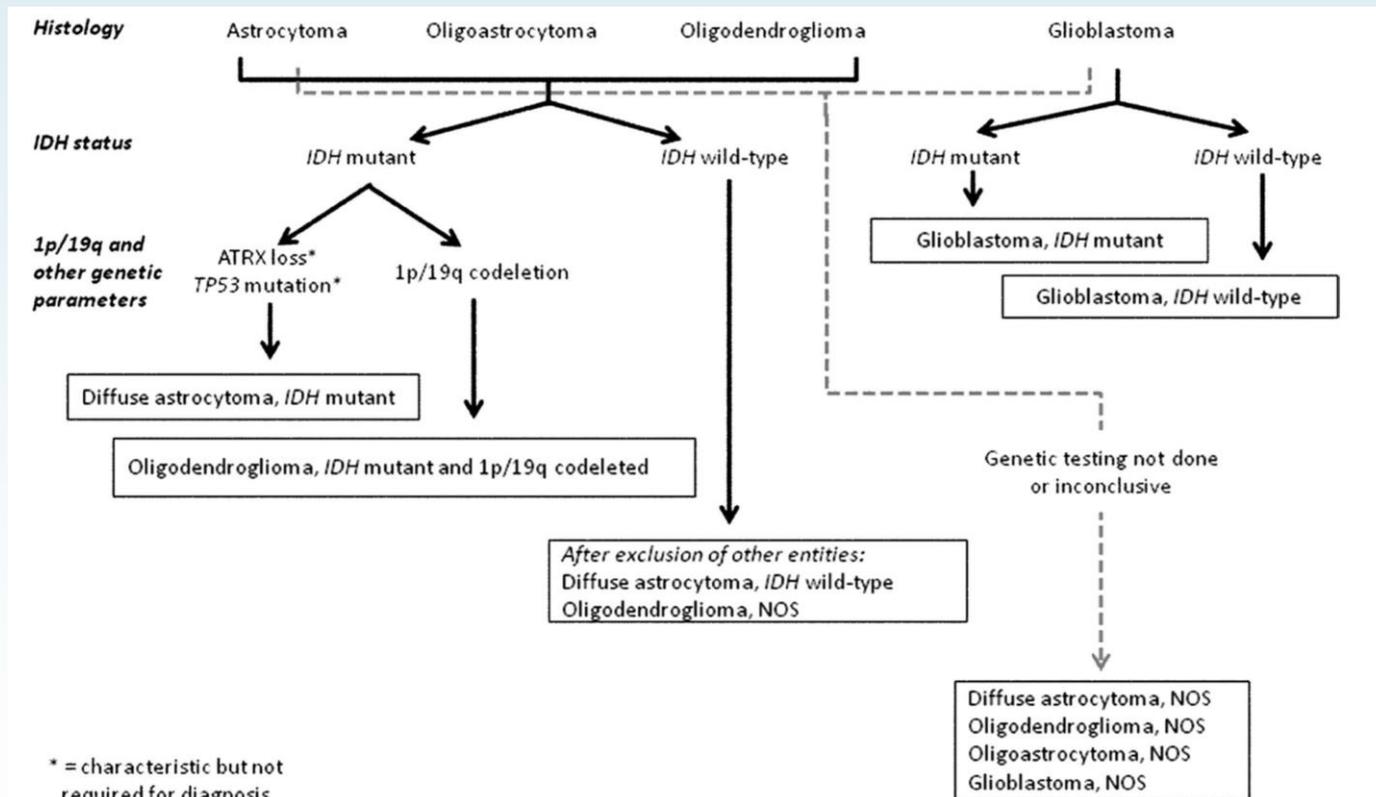
DOI 10.1007/s00401-010-0693-y

CORRESPONDENCE

***RAF* gene fusions are specific to pilocytic astrocytoma in a broad paediatric brain tumour cohort**

Andrew R. J. Lawson · Ruth G. Tatevossian ·
Kim P. Phipps · Simon R. Picker · Antony Michalski ·
Denise Sheer · Thomas S. Jacques · Tim Forshew

Diffuse gliomas in the WHO are defined by IDH mutations and 1p19q co-deletion



..but they are different tumours

Acta Neuropathol (2011) 121:753–761
DOI 10.1007/s00401-011-0810-6

ORIGINAL PAPER

Adult grade II diffuse astrocytomas are genetically distinct from and more aggressive than their paediatric counterparts

David T. W. Jones · Shani A. Mulholland · Danita M. Pearson ·
Deborah S. Malley · Samuel W. S. Openshaw · Sally R. Lambert ·
Lu Liu · L. Magnus Bäcklund · Koichi Ichimura · V. Peter Collins

Acta Neuropathol (2005) 109: 387–392
DOI 10.1007/s00401-004-0976-2

REGULAR PAPER

Portia A. Kreiger · Yoshifumi Okada · Scott Simon
Lucy B. Rorke · David N. Louis · Jeffrey A. Golden

Losses of chromosomes 1p and 19q are rare in pediatric oligodendrogliomas

Paediatric diffuse tumours in the WHO

Paediatric diffuse astrocytoma

Although the histopathology of paediatric diffuse astrocytoma resembles that of adult diffuse astrocytoma, there are many important distinctions between the disease in children and in adults.

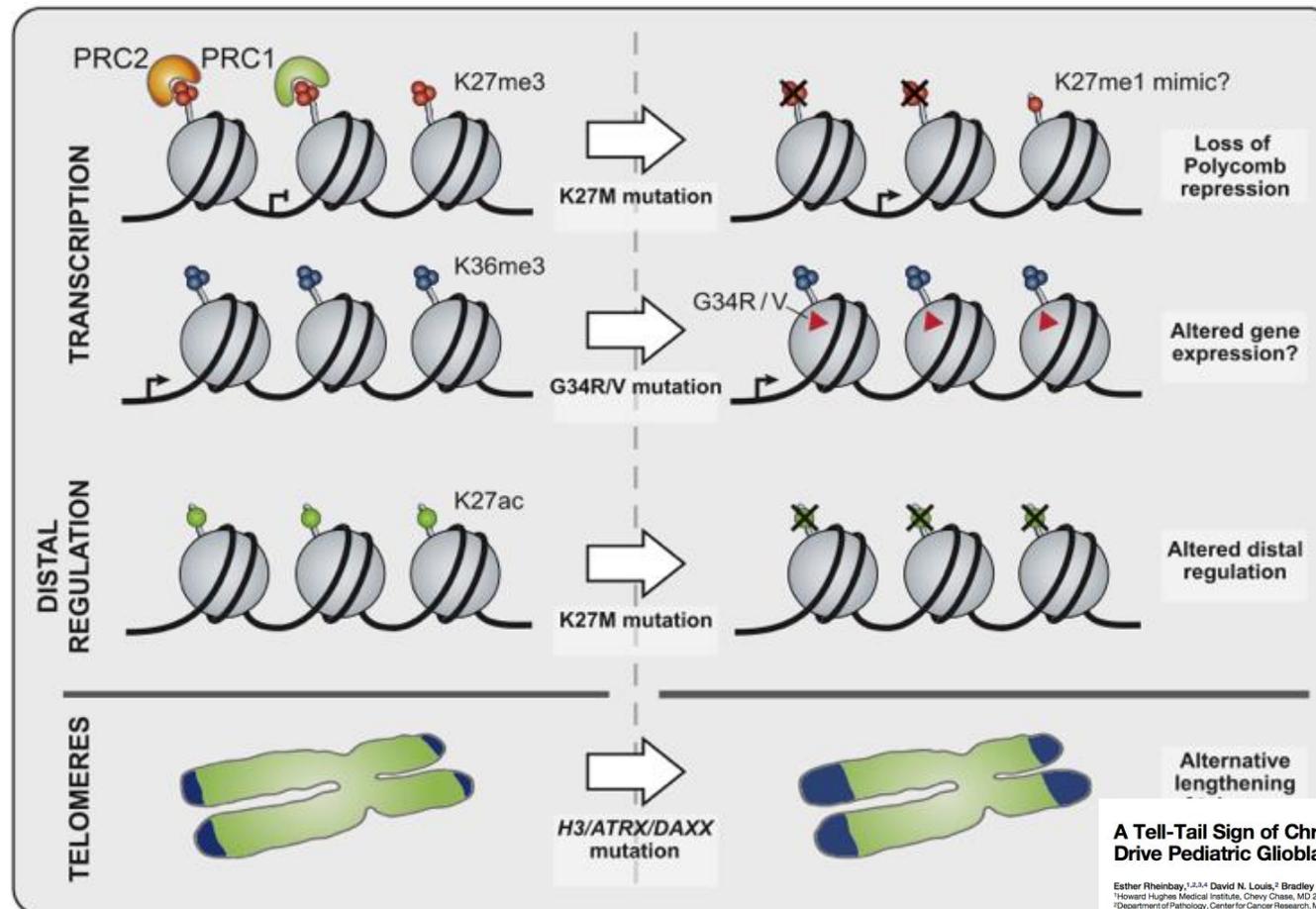
Clinicopathological aspects

The annual incidence of paediatric diffuse astrocytoma (defined by patient age < 20 years at diagnosis) is 0.27 cases per 100 000 population; lower than that of adult diffuse astrocytoma, which is 0.58 per 100 000 {1863}. Most paediatric diffuse astrocytomas are located in the cerebral hemispheres, but a significant proportion present in the thalamus, which is an unusual site for adult diffuse astrocytoma. Anaplastic progression occurs in approximately 75% of adult lesions, but is rare in paediatric tumours {284}.

Genetic aspects

Diffuse astrocytomas in children and adults have distinct genetic profiles. However, diffuse astrocytomas with genetically defined so-called adult-type disease can present in adolescents, and so-called paediatric-type disease can present in young adults. Paediatric diffuse astrocytomas are characterized mainly by alterations in *MYB* and *BRAF*. Amplification or rearrangements of *MYB* are detected in approximately 25% of paediatric diffuse astrocytomas {2518, 2855}. Rearrangements of *MYBL1* have also been described {2068}. Other paediatric diffuse astrocytomas harbour *BRAF* V600E mutations, *FGFR1* alterations, or *KRAS* mutations {2855}. Rare paediatric diffuse astrocytomas contain the H3 K27M mutation usually found in paediatric high-grade gliomas {2855}. The mutations in *IDH1*, *IDH2*, *TP53*, and *ATRX* that are frequently found in adult diffuse astrocytomas are not present in the paediatric tumours {2443}.

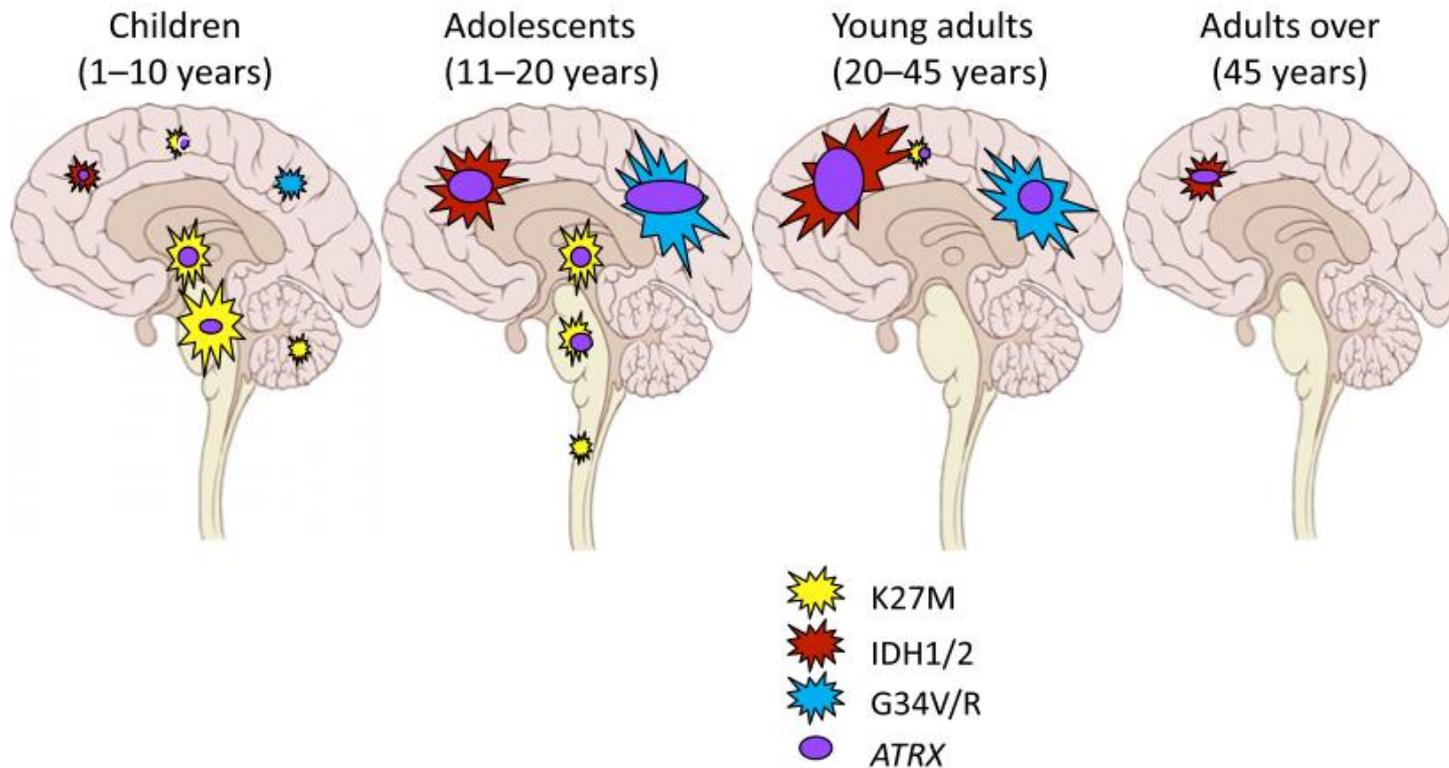
Paediatric high grade gliomas have mutations in histone genes



A Tell-Tail Sign of Chromatin: Histone Mutations Drive Pediatric Glioblastoma

Esther Rheinbay,^{1,2,3,4} David N. Louis,² Bradley E. Bernstein,^{1,2,3,4} and Mario L. Suva^{1,2,3}
¹Howard Hughes Medical Institute, Chevy Chase, MD 20815, USA
²Department of Pathology, Center for Cancer Research, Massachusetts General Hospital and Harvard Medical School, Boston, MA 02114, USA
³Broad Institute of Harvard and MIT, Cambridge, MA 02142, USA
⁴Bioinformatics Program, Boston University, Boston, MA 02215, USA
 *Correspondence: berstein.brady@bmg.harvard.edu
 DOI: 10.1016/j.ccr.2012.03.001

Mutations in paediatric glioma relate to location and age



Therapeutic Intervention in Brain Tumors?

Chromatin Remodeling Defects in Pediatric and Young Adult Glioblastoma: A Tale of a Variant Histone 3 Tail

Adam M. Fontebasso¹; Xiao-Yang Liu²; Dominik Sturm³; Nada Jabado^{1,2,4}

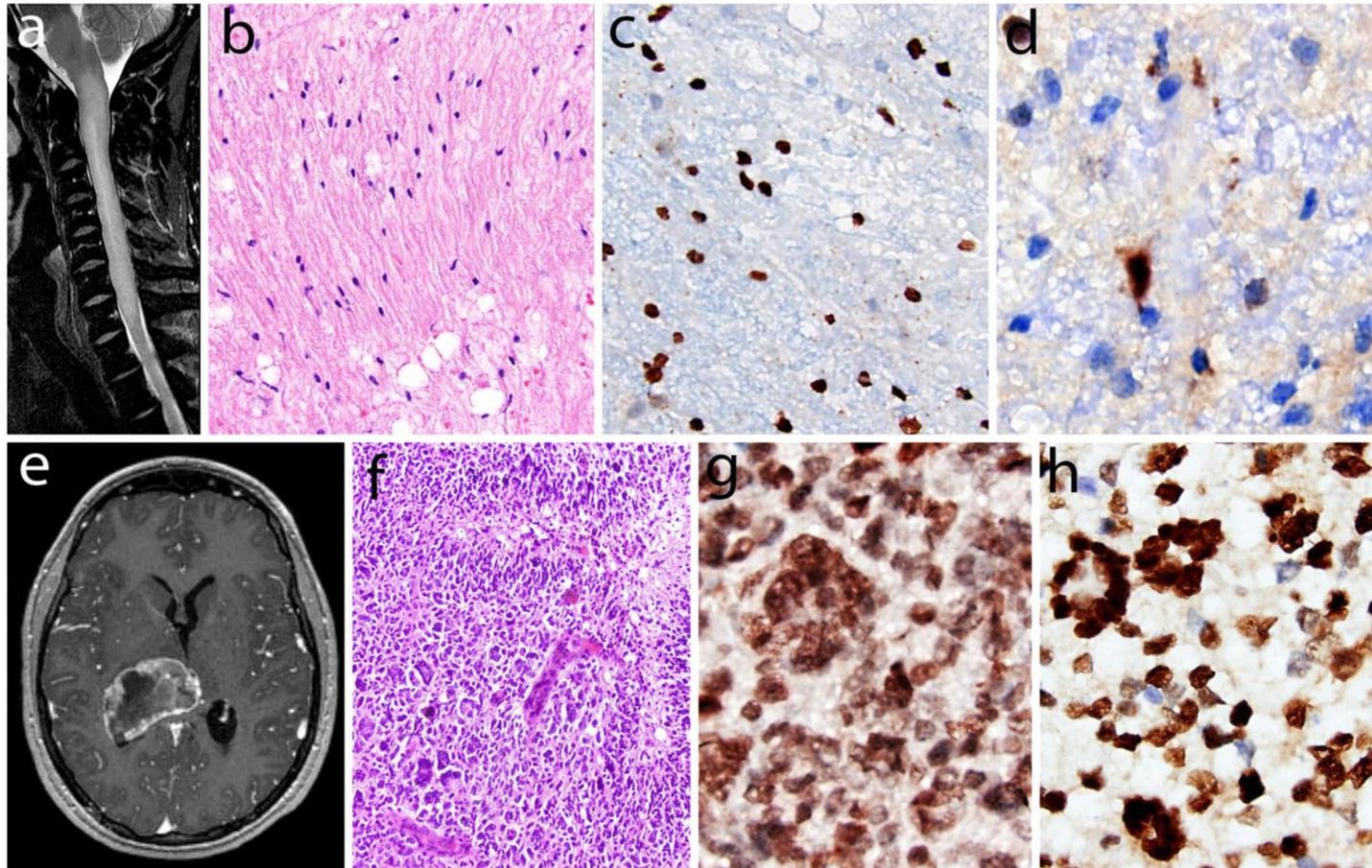
¹ Division of Experimental Medicine, McGill University and McGill University Health Centre, Montreal, QC, Canada.

² Department of Human Genetics, McGill University and McGill University Health Centre, Montreal, QC, Canada.

³ Division of Pediatric Neuro-oncology, German Cancer Research Centre (DKFZ), Heidelberg, Germany.

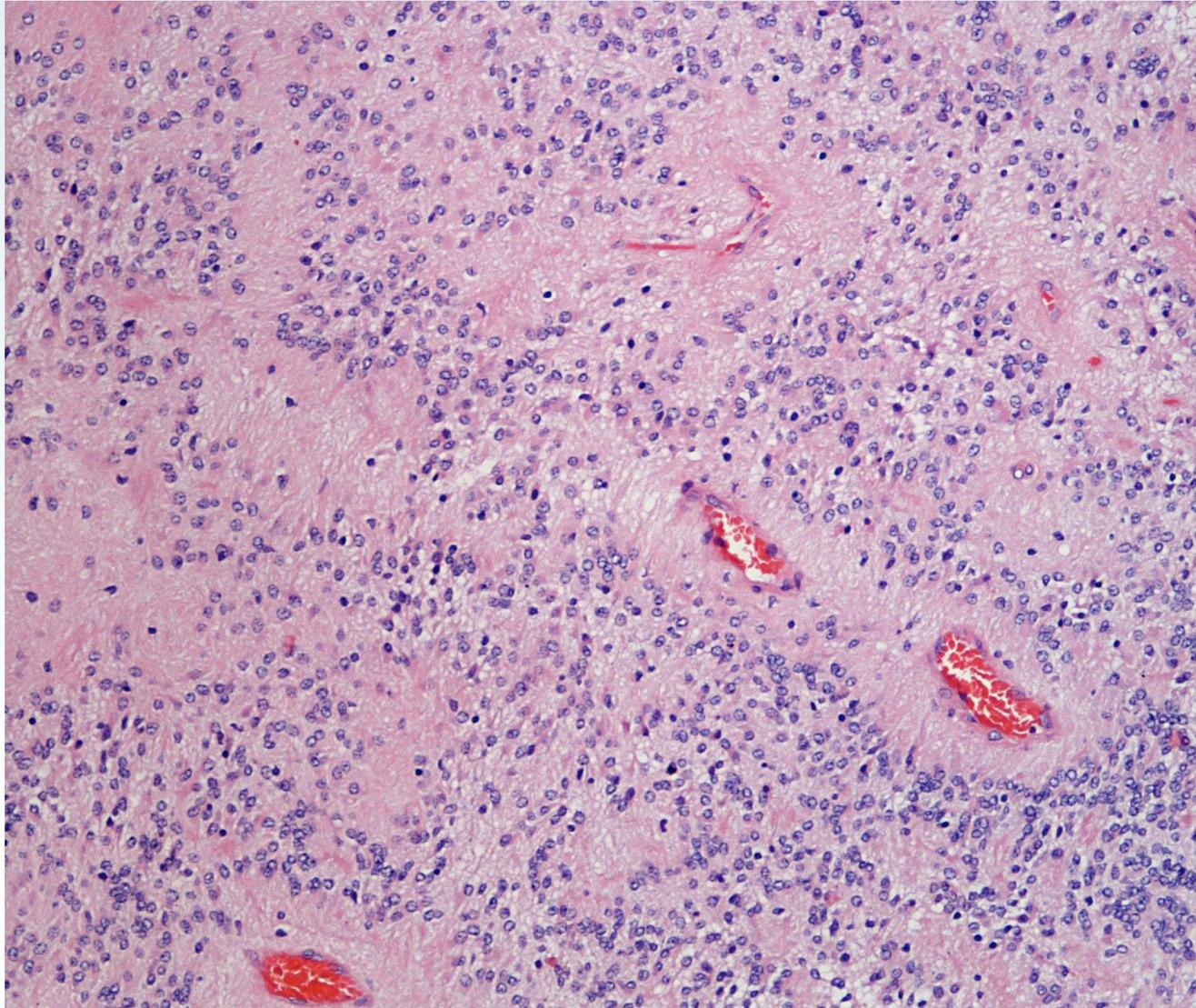
⁴ Department of Pediatrics, McGill University and the McGill University Health Centre Research Institute, Montreal, QC, Canada.

New entity: Diffuse midline glioma, H3 K27M-mutant

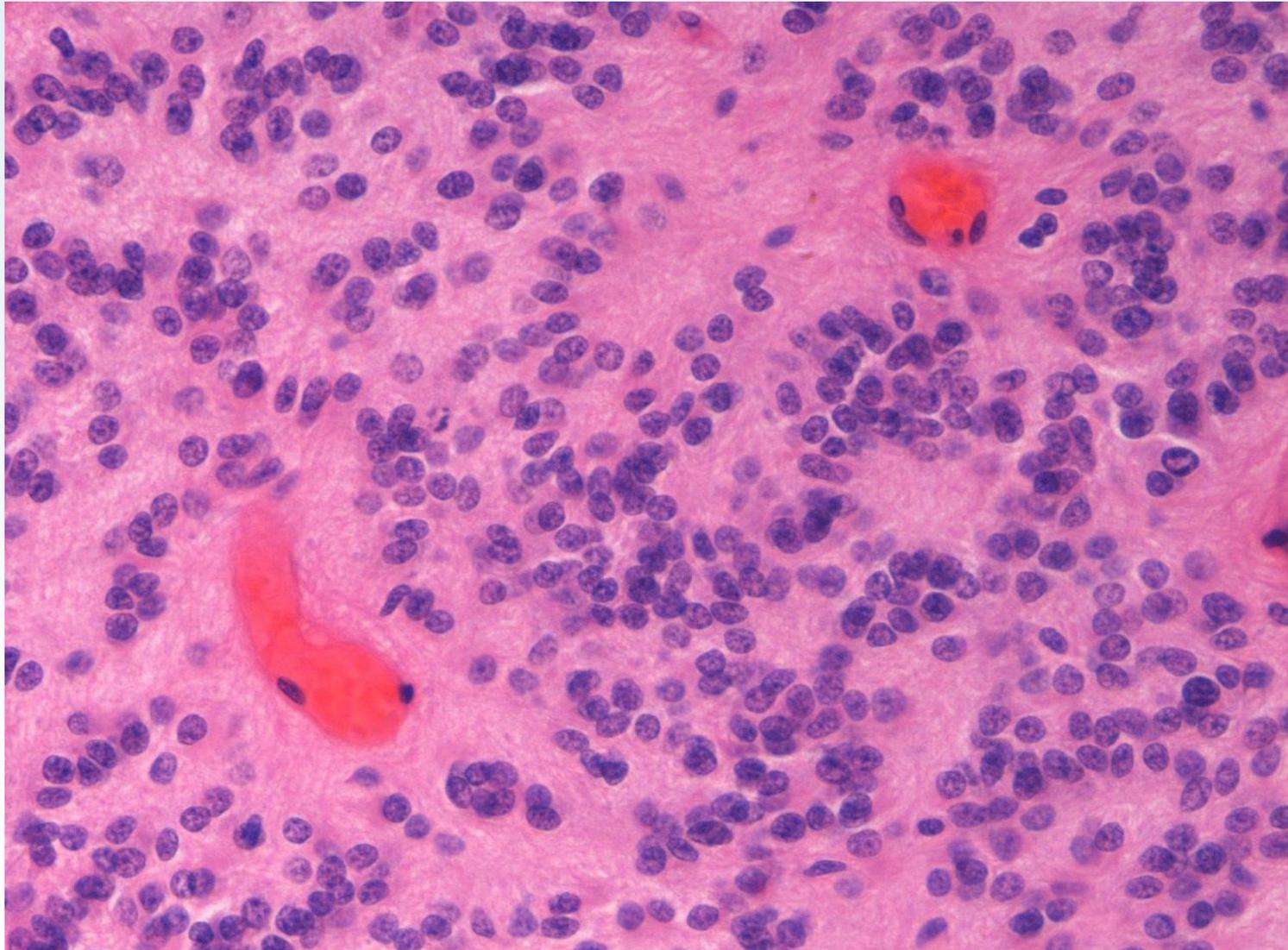


Ependymoma

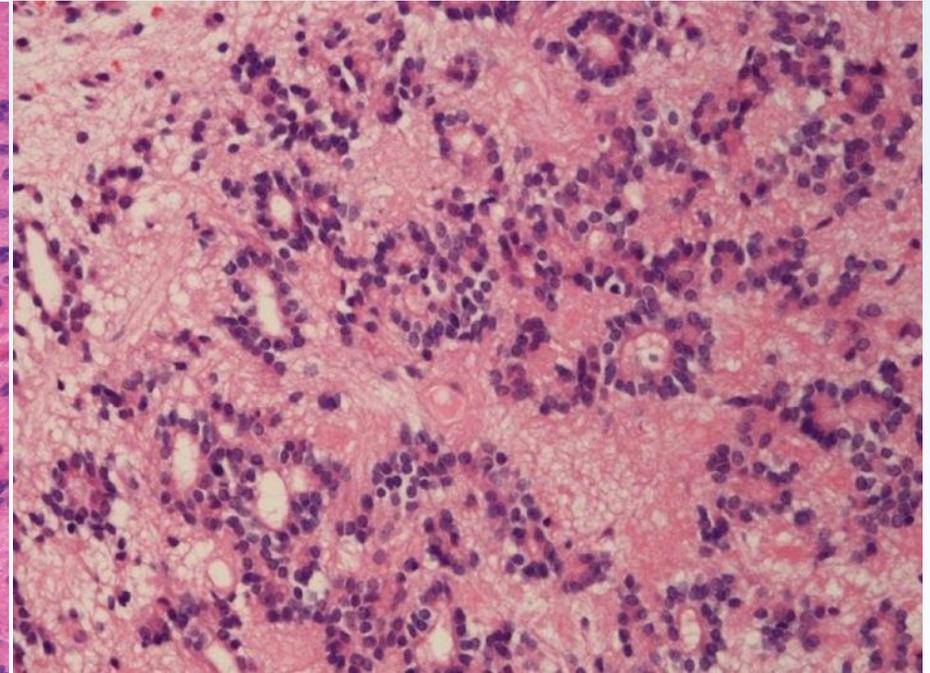
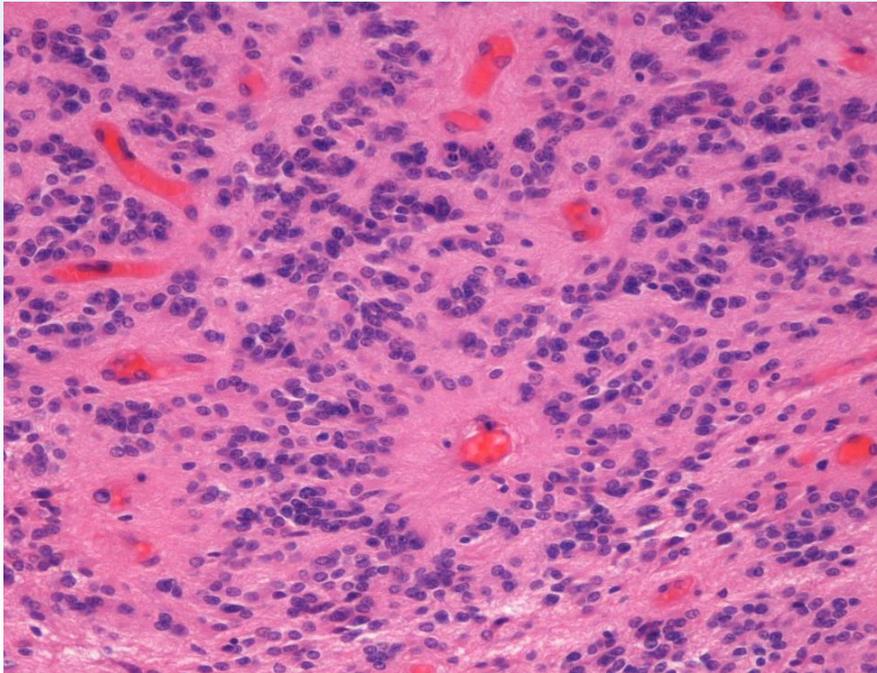
Ependymoma-Architecture



Ependymoma-Cytology



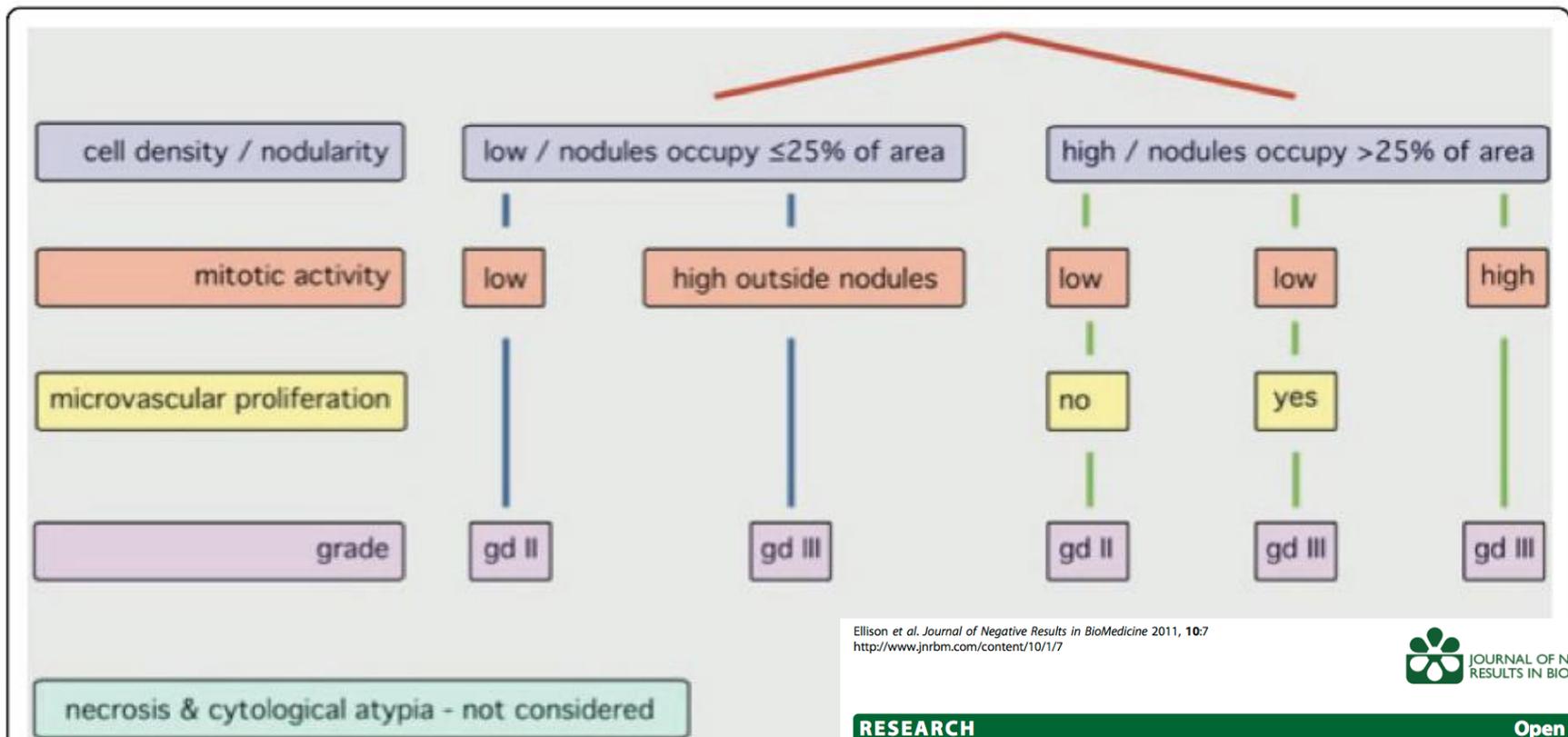
Ependymoma-Secondary structures



Challenges for ependymoma

- Prognosis in children is poor
- Criteria for anaplasia are poorly defined and subjective
- Large effect of surgical clearance on prognosis
- Variability in histology
- Conflicting studies

Approaches to grading in ependymoma



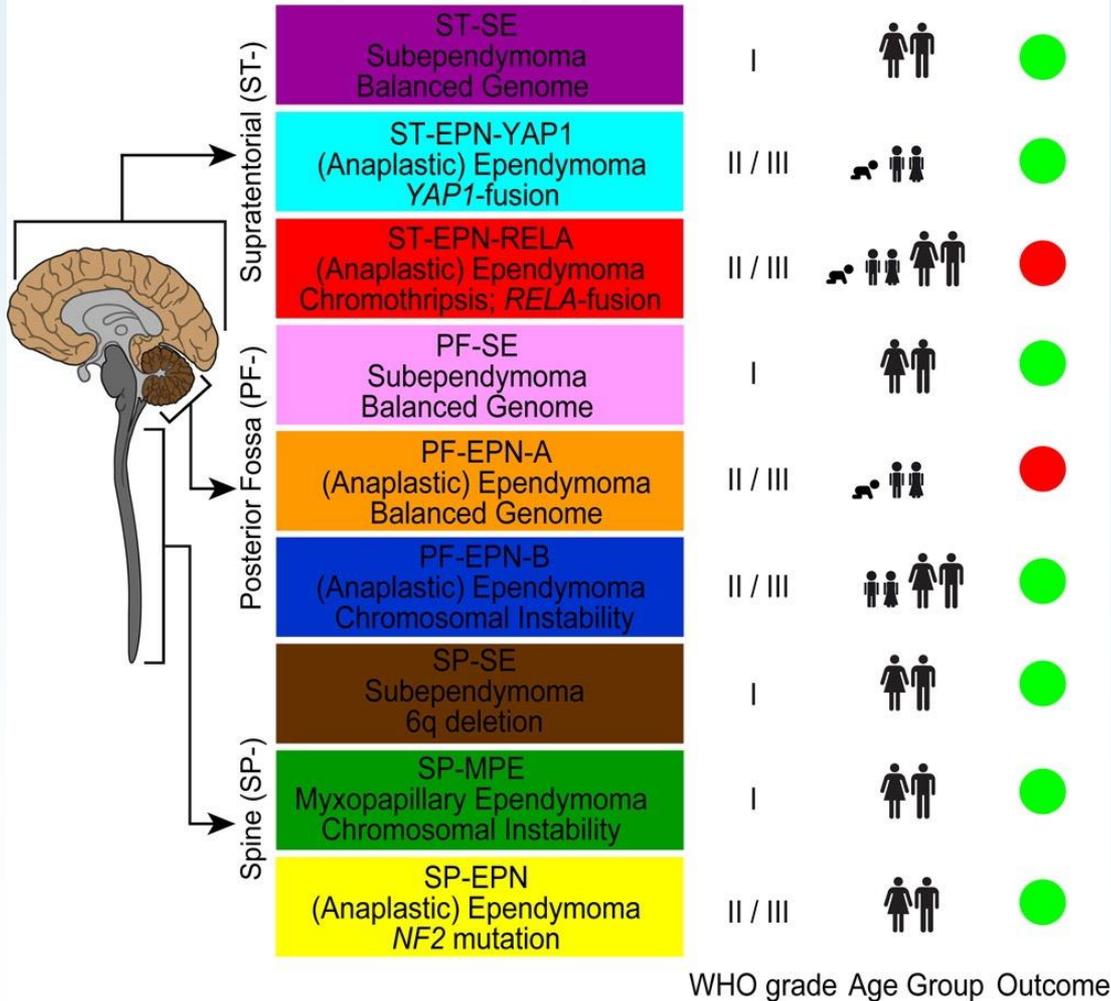
Ellison et al. *Journal of Negative Results in Biomedicine* 2011, 10:7
<http://www.jnrnm.com/content/10/1/7>

Figure 2 Novel grading scheme for pediatric intracranial classic (grade I are regions of high cell density).

Histopathological grading of pediatric ependymoma: reproducibility and clinical relevance in European trial cohorts

David W Ellison^{1*}, Mehmet Kocak², Dominique Figarella-Branger³, Giangaspero Felice⁴, Godfraind Catherine⁵, Torsten Pietsch⁶, Didier Frappaz⁷, Maura Massimino⁸, Jacques Grill⁹, James M Boyett² and Richard G Grundy¹⁰

Molecular Subgrouping of Ependymal Tumors is Superior to Histopathological Grading for Risk Stratification



Article

Cancer Cell

Molecular Classification of Ependymal Tumors across All CNS Compartments, Histopathological Grades, and Age Groups

Authors

Kristian W. Pajtler, Hendrik Witt, ..., Marcel Kool, Stefan M. Pfister

Anatomic Compartment	SPINE (SP-)			Posterior Fossa (PF-)			Supratentorial (ST-)		
Molecular Subgroup	SE	MPE	EPN	SE	EPN-A	EPN-B	SE	EPN-YAP1	EPN-RELA
Histopathology	sub-ependymoma (WHO I)	myxopapillary ependymoma (WHO I)	(anaplastic) ependymoma (WHO II/III)	sub-ependymoma (WHO I)	(anaplastic) ependymoma (WHO II/III)	(anaplastic) ependymoma (WHO II/III)	sub-ependymoma (WHO I)	(anaplastic) ependymoma (WHO II/III)	(anaplastic) ependymoma (WHO II/III)
Genetics	6q del.	CIN	CIN	balanced	balanced	CIN	balanced	aberr. 11q	aberr. 11q
Oncogenic Driver	?	?	<i>NF2</i>	?	?	?	?	<i>YAP1</i> -fusion	<i>RELA</i> -fusion
Tumor Location									
Age Distribution (years)									
Gender Distribution									
Patient Survival (OS; months)									

New entity: Ependymoma, *RELA* fusion-positive

Ependymoma, *RELA* fusion-positive

Ellison D.W.
Korshunov A.
Witt H.

Definition

A supratentorial ependymoma characterized by a *RELA* fusion gene.

The genetically defined *RELA* fusion-positive ependymoma accounts for approximately 70% of all childhood supratentorial tumours [1891] and a lower proportion of such ependymomas in adult patients [1880]. Ependymomas in the posterior fossa and spinal compartments do not harbour this fusion gene. *RELA* fusion-positive ependymomas exhibit a range of histopathological features, with or without anaplasia.

ICD-O code 9396/3

Grading

RELA fusion-positive ependymomas are classified according to their histopathological features into WHO grade II or grade III. No grade I ependymoma has been recorded as containing this genetic alteration.

Microscopy

RELA fusion-positive ependymomas do

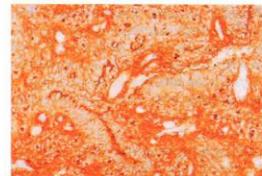


Fig. 3.15 *RELA* fusion-positive ependymoma. L1CAM protein expression correlates well with the presence of a *RELA* fusion gene.

not have a specified morphology [1891]. They exhibit the standard range of architectural and cytological features found in supratentorial ependymomas, but they often have a distinctive vascular pattern of branching capillaries or clear-cell change. Uncommon variants of ependymoma (e.g. tanyctytic ependymoma) do not tend to be *RELA* fusion-positive.

Immunophenotype

RELA fusion-positive ependymomas demonstrate the immunoreactivities for GFAP and EMA described in other ependymomas. Expression of L1CAM correlates well with the presence of a *RELA* fusion in supratentorial ependymomas [1891], but L1CAM can also be expressed by other types of brain tumours.

Genetic profile

The *C11orf95-RELA* fusion is the most common structural variant found in ependymomas [1880,1891,1974]. It forms in the context of chromothripsis, a shattering and reassembly of the genome that rearranges genes and produces oncogenic gene products [2852]. *RELA* fusion-positive ependymomas show constitutive activation of the NF-kappaB pathway, the *RELA*-encoded transcription factor p65 being a key effector in this pathway. Rarely, *C11orf95* or *RELA* can be fused with other genes as a result of chromothripsis [1891].

The presence of a *C11orf95-RELA* fusion gene can be detected by various methods, but a simple approach using

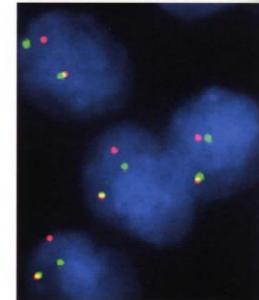


Fig. 3.16 *RELA* fusion-positive ependymoma. Interphase FISH with break-apart probes around the *RELA* gene. Overlapping probes (yellow) indicate an intact *RELA* gene, but probe separation (red/green) occurs with rearrangement of the *RELA* gene.

formalin-fixed, paraffin-embedded tissue is interphase FISH with break-apart probes around both genes. Rearrangement in the context of chromothripsis splits the dual-colour signals in probe sets for *C11orf95* and *RELA* [1891].

Prognosis and predictive factors

The data available to date (which come from only a single study) suggest that *RELA* fusion-positive ependymomas have the worst outcome of the three supratentorial molecular groups [1880].

Conclusions

- Childhood brain tumours can be cured but at the risk of long-term complications
- Real-time molecular diagnosis is required to stratify treatment



Jen Cotter



Jess Pickles



Tom Stone



Derek Li



Amy Fairchild



Alex Virasami



Fatma Scerif



Sherry Yasin



Simon Raphael Picker



Simon Paine



JP Martinez-Barbera
UCL-ICH



Sebastian Brandner
UCL-ION



Denise Sheer
QMUL



William Harkness
GOSH



Martin Tisdall
GOSH



Helen Cross
UCL-ICH



Jonathan Ham
UCL-ICH



Darren Hargrave
GOSH



Francois Guillemot
Francis Crick Institute



Glenn Anderson
GOSH



Mike Hubank
ICR



Steve Clifford
Newcastle

