



*Neuro-news: innovazioni diagnostiche e terapeutiche
in neurologia*

TUMORI CEREBRALI

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SIN Triregionale, Ivrea 6 Dicembre 2019

DISCLOSURES

I received consultancy and advisory board fees from :

UCB Pharma

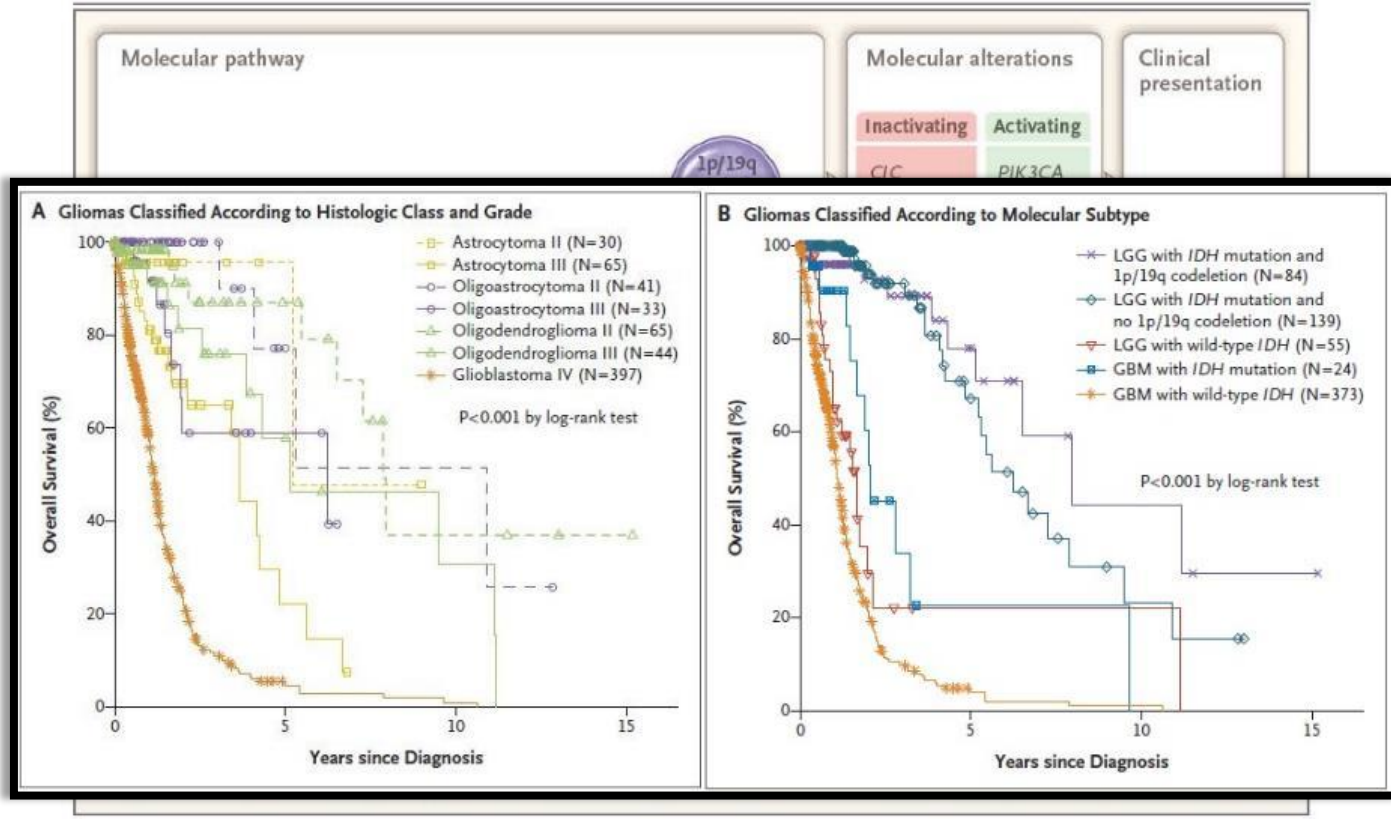
EISAI

Mundipharma

Novocure

OUTLINE

- Innovazioni diagnostiche: gli avanzamenti in biologia molecolare
- Innovazioni terapeutiche: i risultati dei trials clinici di fase III nei gliomi di alto grado e le terapie target
- Liquid biopsy nei tumori cerebrali: nuovo strumento per il clinico?



Comprehensive, Integrative Genomic Analysis of Diffuse LGG
 The Cancer Genome Atlas Research Network; NEJM, June 2015

Today we know that **different molecular arrangements can have stronger prognostic and predictive value** than the histotype itself.

WHO 4 Ed. 2007

Diffuse astrocytoma, (II)
 Fibrillary astrocytoma
 Gemistocytic astrocytoma
 Protoplasmatic astrocytoma

Anaplastic astrocytoma, (III)

Glioblastoma, (IV)
 Giant cell glioblastoma
 Gliosarcoma
 Gliomatosis cerebri

Oligodendroglioma, (II)
 Anaplastic oligodendroglioma, (III)

Oligoastrocytoma, (II)
 Anaplastic oligoastrocytoma, (III)

WHO 4 Ed.+ 2016

Diffuse astrocytoma, IDH-1 mutant (II)
 Gemistocytic astrocytoma, IDH-mutant (II)
 Diffuse astrocytoma, IDH-wildtype (II)
 Diffuse astrocytoma, NOS (II)

Anaplastic astrocytoma, IDH-mutant (III)
 Anaplastic astrocytoma, IDH-wildtype (III)
 Anaplastic astrocytoma, NOS (III)

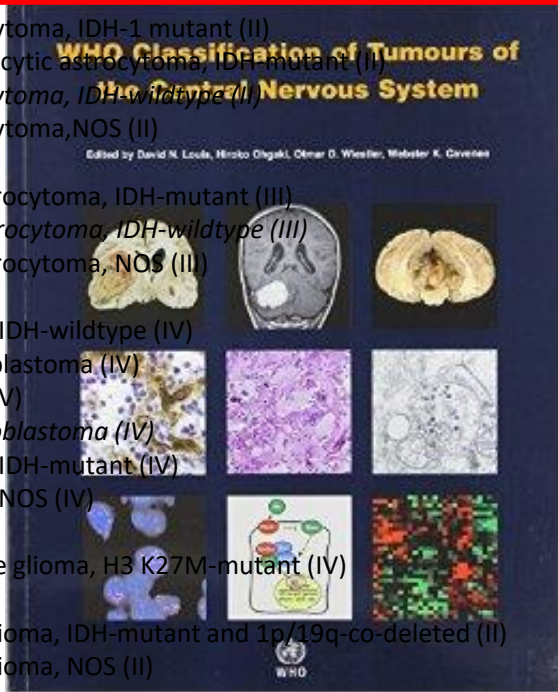
Glioblastoma, IDH-wildtype (IV)
 Giant cell glioblastoma (IV)
 Gliosarcoma (IV)
 Epithelioid glioblastoma (IV)
 Glioblastoma, IDH-mutant (IV)
 Glioblastoma, NOS (IV)

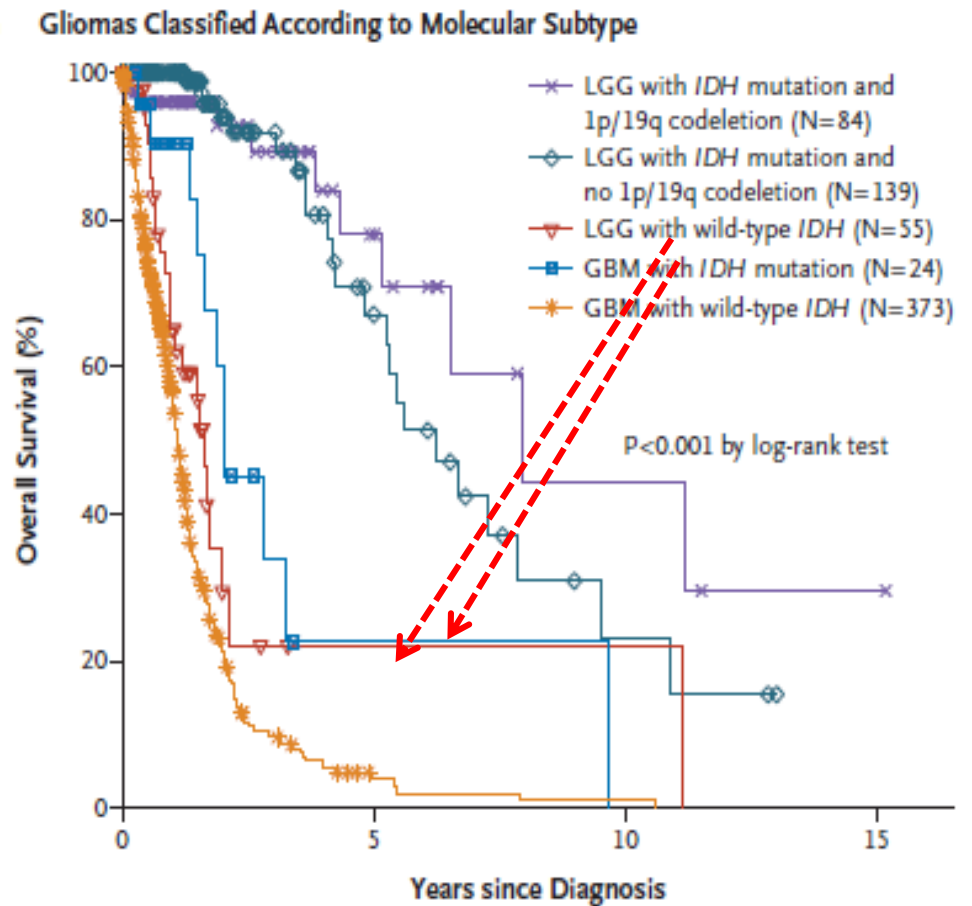
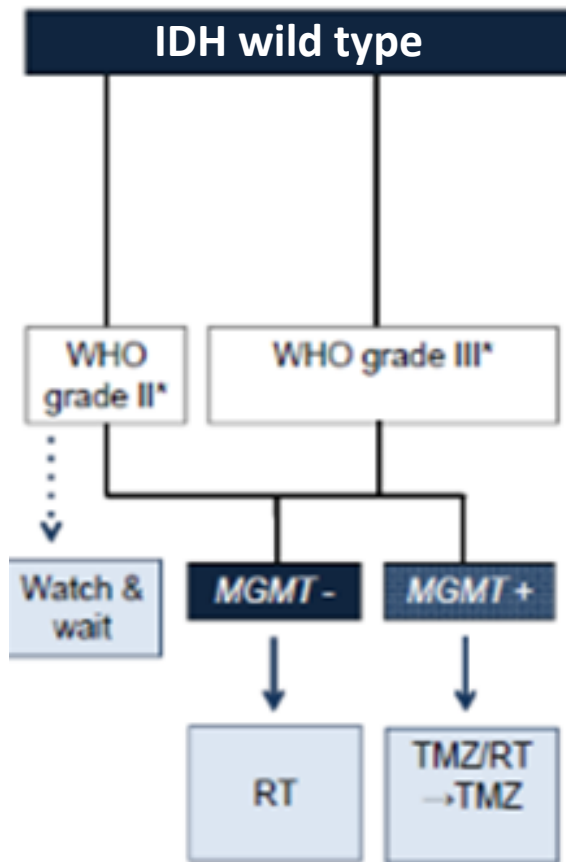
Diffuse midline glioma, H3 K27M-mutant (IV)

Oligodendroglioma, IDH-mutant and 1p/19q-co-deleted (II)
 Oligodendroglioma, NOS (II)

Anaplastic oligodendroglioma, IDH-mutant and 1p/19q-codeleted (III)
 Anaplastic oligodendroglioma, NOS (III)

Oligoastrocytoma (II)
 Anaplastic oligoastrocytoma (III)

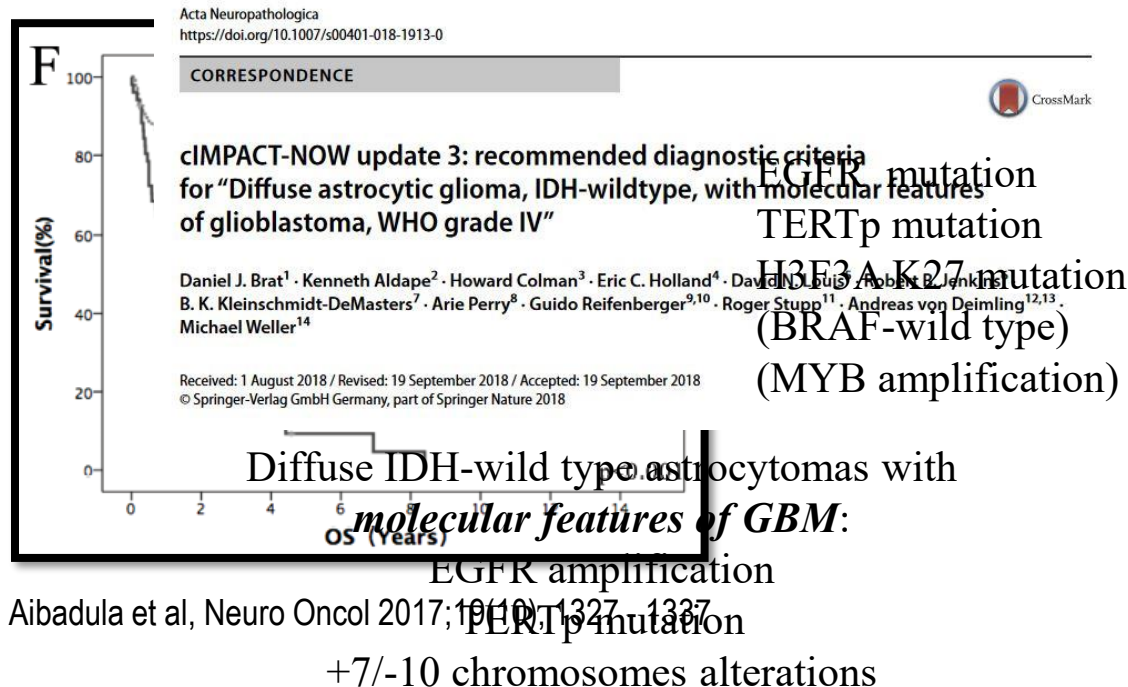




Grade II IDH-wild type

With the current 2016 WHO Classification **all IDH-wild type tumours are classified as astrocytomas**, regardless of the tumour grade. Thus, cases originally diagnosed as either oligodendrogliomas or oligoastrocytomas according to WHO 2007 need to be reclassified as astrocytomas.

There is now increasing evidence of the **molecular heterogeneity** of the IDH-wild type grade II astrocytomas.



CORRESPONDENCE



cIMPACT-NOW update 1: Not Otherwise Specified (NOS) and Not Elsewhere Classified (NEC)

David N. Louis¹ · Pieter Wesseling^{2,3,4} · Werner Paulus⁵ · Caterina Giannini⁶ · Tracy T. Batchelor⁷ · J. Gregory Cairncross⁸ · David Capper^{9,10} · Dominique Figarella-Branger¹¹ · M. Beatriz Lopes¹² · Wolfgang Wick¹³ · Martin van den Bent¹⁴

Acta Neuropathologica (2018) 135:639–642
<https://doi.org/10.1007/s00401-018-1826-y>

CORRESPONDENCE



cIMPACT-NOW update 2: diagnostic clarifications for *diffuse midline glioma, H3 K27M-mutant* and *diffuse astrocytoma/anaplastic astrocytoma, IDH-mutant*

David N. Louis¹ · Caterina Giannini² · David Capper³ · Werner Paulus⁴ · Dominique Figarella-Branger⁵ · M. Beatriz Lopes⁶ · Tracy T. Batchelor⁷ · J. Gregory Cairncross⁸ · Martin van den Bent⁹ · Wolfgang Wick^{10,11,12} · Pieter Wesseling^{13,14}

Acta Neuropathologica (2018) 136:805–810
<https://doi.org/10.1007/s00401-018-1913-0>

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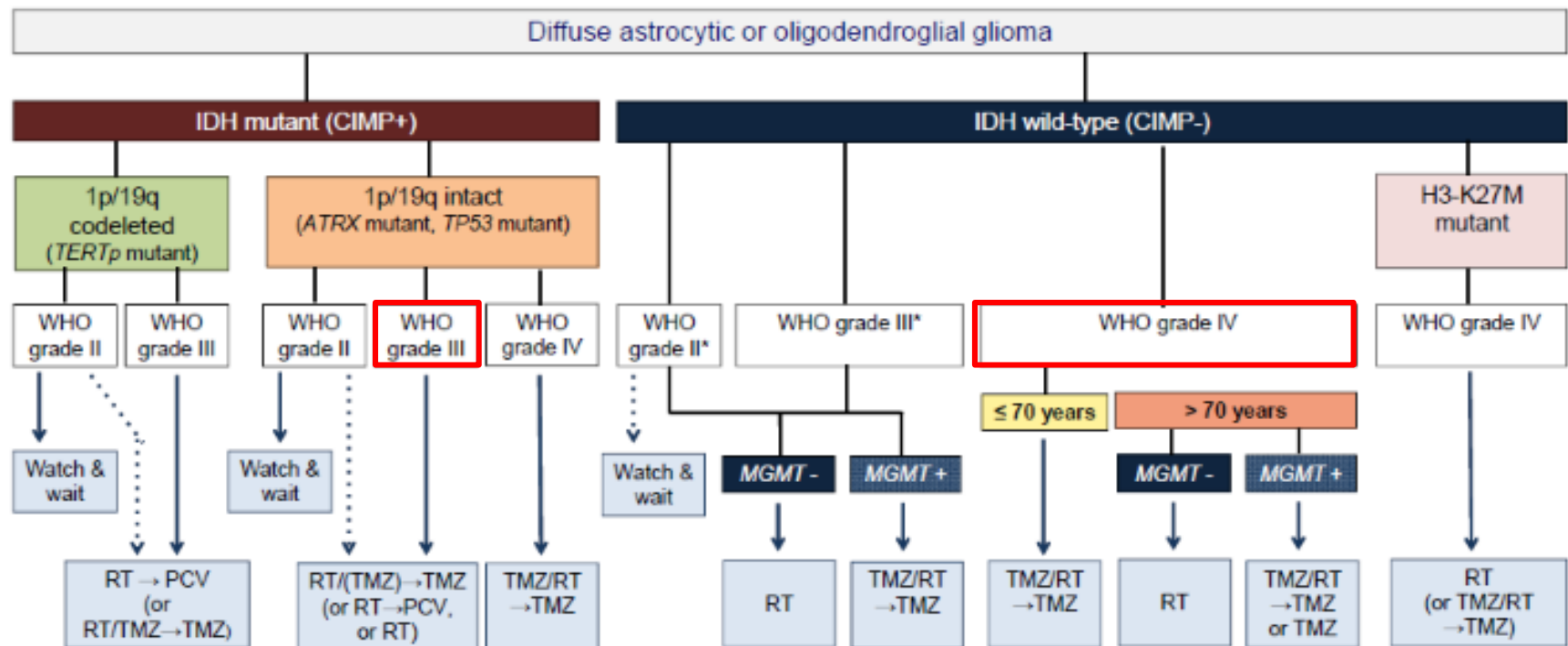
cIMPACT-NOW update 3: recommended diagnostic criteria for “Diffuse astrocytic glioma, IDH-wildtype, with molecular features of glioblastoma, WHO grade IV”

Daniel J. Brat¹ · Kenneth Aldape² · Howard Colman³ · Eric C. Holland⁴ · David N. Louis⁵ · Robert B. Jenkins⁶ · B. K. Kleinschmidt-DeMasters⁷ · Arie Perry⁸ · Guido Reifenberger^{9,10} · Roger Stupp¹¹ · Andreas von Deimling^{12,13} · Michael Weller¹⁴



The 2017 EANO guideline

Lancet Oncology 2017;18:e315-e329



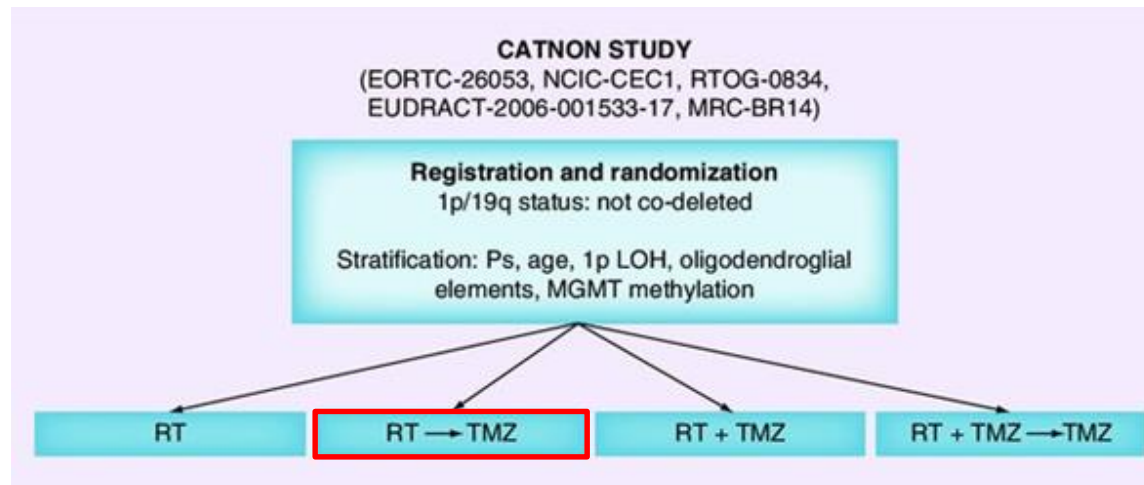
* Refers to the provisional entities of diffuse astrocytoma, IDH wildtype, and anaplastic astrocytoma, IDH wildtype.

CATI
con
delet

Interim results from the CATNON trial (EORTC study 26053-22054) of treatment with concurrent and adjuvant temozolomide for 1p/19q non-co-deleted anaplastic glioma: a phase 3, randomised, open-label intergroup study

with
-co-
label

Martin J van den Bent, Brigitta Baumert, Sara C Erridge, Michael A Vogelbaum, Anna K Nowak, Marc Sanson, Alba Ariela Brandes, Paul M Clement, Jean Francois Baurain, Warren P Mason, Helen Wheeler, Olivier L Chinot, Sanjeev Gill, Matthew Griffin, David G Brachman, Walter Taal, Roberta Rudà, Michael Weller, Catherine McBain, Jaap Reijneveld, Roelien H Enting, Damien C Weber, Thierry Lesimple, Susan Clenton, Anja Gijtenbeek, Sarah Pascoe, Ulrich Herrlinger, Peter Hau, Frederic Dhermain, Irene van Heuvel, Roger Stupp, Ken Aldape, Robert B Jenkins, Hendrikus Jan Dubbink, Winand N M Dinjens, Pieter Wesseling, Sarah Nuyens, Vassilis Gofinopoulos, Thierry Gorlia, Wolfgang Wick, Johan M Kros



- **OS** HR 0.645 (95% CI 0.450, 0.926; p= 0.0014) adj TMZ
- **PFS** HR 0.586 (95% CI 0.472, 0.727; p < 0.0001) adj TMZ
- Benefit from adjuvant (adj) temozolomide (TMZ) on overall survival (OS) but remained **inconclusive about concurrent (conc) TMZ**.

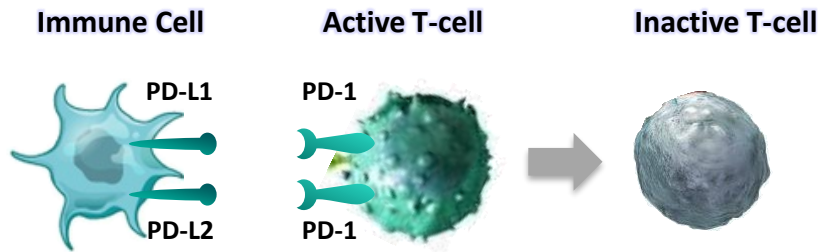
Notable phase III novel agents targeting glioblastoma multiforme

Project	Mechanism	Company	Trial ID	Notes
Opdivo	Anti-PD-1 MAb	Bristol-Myers Squibb	NCT02617589 NCT02667587	Checkmate-498, vs Temodar: failed May 2019. Checkmate-548, on top of Temodar: data due this year.
Depatuxizumab mafodotin	Anti-EGFR ADC	Abbvie	NCT02573324	Intelligence-1 study, on top of Temodar in patients with EGFR amplification: failed May 2019.
Toca 511 & Toca FC	Gene therapy & pyrimidine analogue	Tocagen	NCT02414165	Toca 5 study, on top of Temodar or Avastin. In May 2019 cleared to continue to final readout, due by end 2019.
DCVax-L	Cancer vaccine	Northwest Biotherapeutics	NCT00045968	On top of Temodar. Recruitment halted 2015; still no final readout.
Trans sodium crocetinate	Vitamin A analogue	Diffusion Pharmaceuticals	NCT03393000	Intact study, on top of Temodar, biopsy-only patients: primary completion 2021.
Marizomib	Proteasome inhibitor	Celgene	NCT03345095	EORTC-1709-BTG study, on top of Temodar: primary completion 2022.

Source: EvaluatePharma and Clinicaltrials.gov.

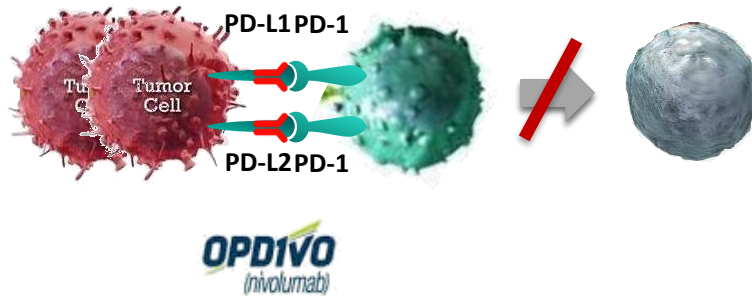
Programmed Death Pathway and Nivolumab

Normal Homeostatic Mechanism



- Normal function of PD-1 pathway is to attenuate immune response to avoid immune system attack of “self”
- A “brake” to prevent overreaction & overproliferation

Tumor Microenvironment



- Tumor cells “co-opt” the PD-1 pathway to evade T-cell immune responses

- Nivolumab occupies the PD-1 receptor of T-cells → prevents inhibitory ligand binding & T-cell inactivation

CheckMate 498

CA209-498 (NCT02617589): Phase 3 Randomized, Open-Label Study of RT in Combination With Nivolumab or TMZ in Newly Diagnosed *MGMT*-Unmethylated GBM¹

Estimated enrollment N = 550

Key inclusion criteria

- Newly diagnosed brain cancer or tumor called GBM or GBM
- Males and females ≥ 18 years old
- Tumor test result shows *MGMT* unmethylated type
- KPS ≥ 70%

R

Experimental:

Nivolumab IV infusion + RT Q2W (dose as specified); then nivolumab Q4W

Active Comparator:

Standard therapy with TMZ + RT (dose as specified)

Start Date: February 2016

Estimated Study Completion Date: October 2019

Estimated Primary Completion Date: March 2019

Status: currently recruiting participants

Study Sponsor: Bristol-Myers Squibb

Collaborator: Ono Pharmaceutical Co. Ltd

- Primary Outcome Measure: 3-year OS
- Secondary Outcome Measure: PFS; 2-year OS

CheckMate 498

CA209-498 (NCT02617589): Phase 3 Randomized, Open-Label Study of RT in Combination With Nivolumab or TMZ in Newly Diagnosed *MGMT*-Unmethylated GBM¹

Estimated enrollment N = 550

Key inclusion criteria

- Newly diagnosed brain cancer or tumor called GBM or GBM
- Males and females ≥ 18 years old
- Tumor test result shows *MGMT* unmethylated
- KPS ≥ 2

Experimental:

Nivolumab IV + RT
Stereotactic RT



Bristol-Myers Squibb Announces Phase 3 CheckMate -498 Study Did Not Meet Primary Endpoint of Overall Survival with Opdivo (nivolumab) Plus Radiation in Patients with Newly Diagnosed *MGMT*-Unmethylated Glioblastoma Multiforme

Category:

[Corporate/Financial News](#)

Thursday, May 9, 2019 6:59 am EDT

Primary Outcome Measure: 3-year OS

Secondary Outcome Measure: PFS; 2-year OS

CheckMate 548

CA209-548 (NCT02667587): Phase 3, Randomized, Single-Blind Study of TMZ + RT With Nivolumab in Newly Diagnosed *MGMT*-Methylated GBM¹

Estimated enrollment N = 693

Key inclusion criteria

- Males and females ≥ 18 years old
- Newly diagnosed brain cancer or tumor called GBM or GBM
- Tumor test result shows *MGMT* methylated / indeterminate tumor
- KPS $\geq 70\%$
- Substantial recovery from surgical resection

R

Experimental:

- Nivolumab IV infusion specified dose on specified days
- RT: 2 gray units 5 times/week for 6 weeks
- TMZ 75 mg/m² daily during RT, followed by a 4-week treatment break; then TMZ 150 mg/m² days 1-5 of cycle 1 and increased to 200 mg/m² days 1-5 of cycles 2-6 as tolerated (additional cycles may be permitted with approval of sponsor)

Placebo Comparator:

- Placebo IV infusion specified dose on specified days
- RT: 2 gray units 5 times/week for 6 weeks
- TMZ 75 mg/m² daily during RT, followed by a 4-week treatment break; then TMZ 150 mg/m² days 1-5 of cycle 1 and increased to 200 mg/m² days 1-5 of cycles 2-6 as tolerated (additional cycles may be permitted with approval of sponsor)

Start Date: May 2016

Estimated Study Completion Date: August 2023

Estimated Primary Completion Date: February 2021

Status: currently recruiting participants

Study Sponsor: Bristol-Myers Squibb

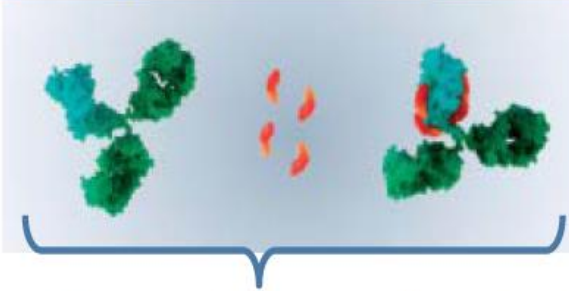
Collaborator: Ono Pharmaceutical Co. Ltd

- Primary Outcome Measures: OS, PFS^a
- Secondary Outcome Measures: OS, PFS^b

^aDetermined by BICR; ^bDetermined by investigator.

1. [Clinicaltrials.gov. NCT02667587](https://clinicaltrials.gov/ct2/show/study/NCT02667587). Accessed April 17, 2018.

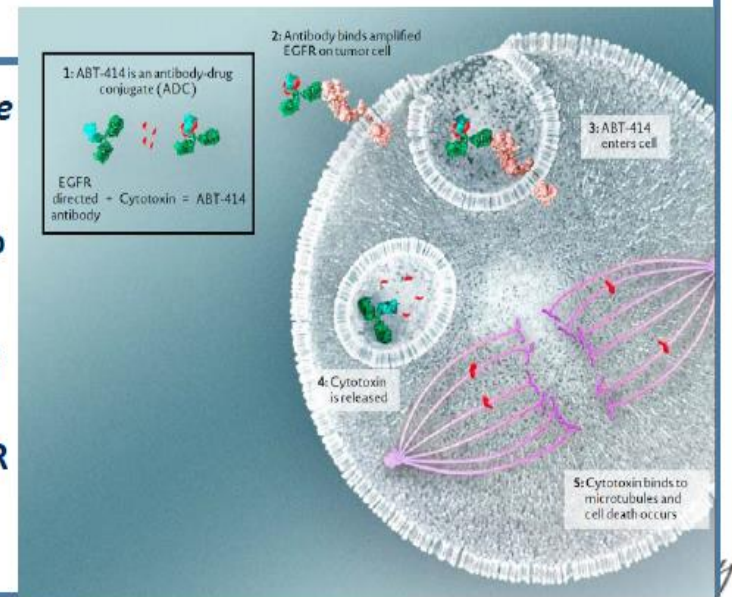
Depatux-M (ABT-414) is a monoclonal Antibody Drug Conjugate (ADC) directed against EGFR



Antibody + Toxin = Antibody Drug Conjugate
(ABT-806) (MMAF) (Depatux-M)

Depatux-M is an antibody-drug conjugate (ADC), comprised of an antibody that *selectively targets activated EGFR* and a cytotoxin that is only released inside the *tumor cell*

- EGFR amplification (~50% of GBM) leads to **preferential exposure of a unique epitope** of the EGFR protein that binds Depatux-M
- Unlike other EGFR directed therapies, there is **limited binding to EGFR in normal tissue** such as skin and other epithelial tissue.
- Depatux-M uses **activated EGFR** only as a target for **intracellular toxin delivery** and does not inhibit EGFR signaling; therefore, it can work in glioblastoma cells that are **resistant to classical EGFR inhibition**

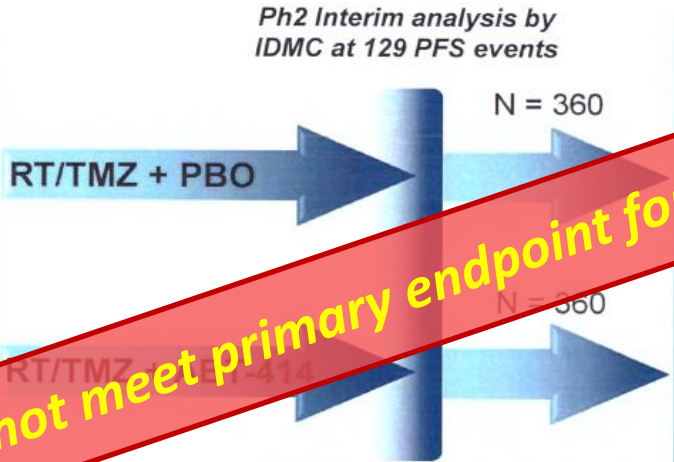


Abbvie

Patient Population

- Histologically confirmed *de novo* glioblastoma (primary) or gliosarcoma
- EGFR amplification
- Chemoradiation therapy start **within 7 weeks** of surgery/biopsy
- Baseline MRI
- Karnofsky Performance Score ≥ 70

1:1 Randomization Placebo-controlled



Endpoints

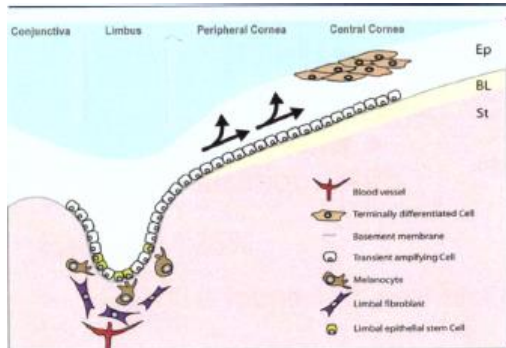
- ### Primary Objective
- Overall Survival (OS)
- ### Secondary Objectives
- Progression-Free Survival (PFS)
 - OS and PFS in EGFRvIII subgroup
 - Time to deterioration in:
 - Verbal memory and executive function (HVL-R and COWA)
 - Symptom severity (MDASI-BT)
 - Symptom interference (MDASI-BT)

The study did not meet primary endpoint for OS (May 2019)

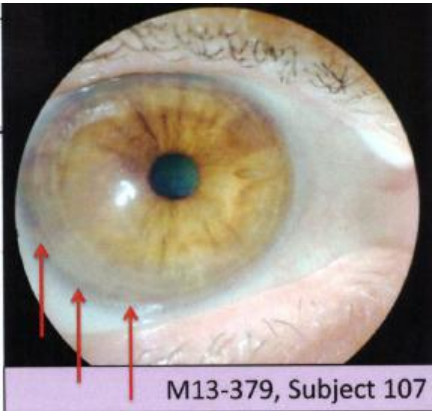
Stratification Factors

- EGFRvIII status
- MGMT methylation status
- RPA (recursive partitioning analysis) score
- Region of World

Microcystic keratopathy and corneal epithelial microcysts



Secker *et al.*, 2010

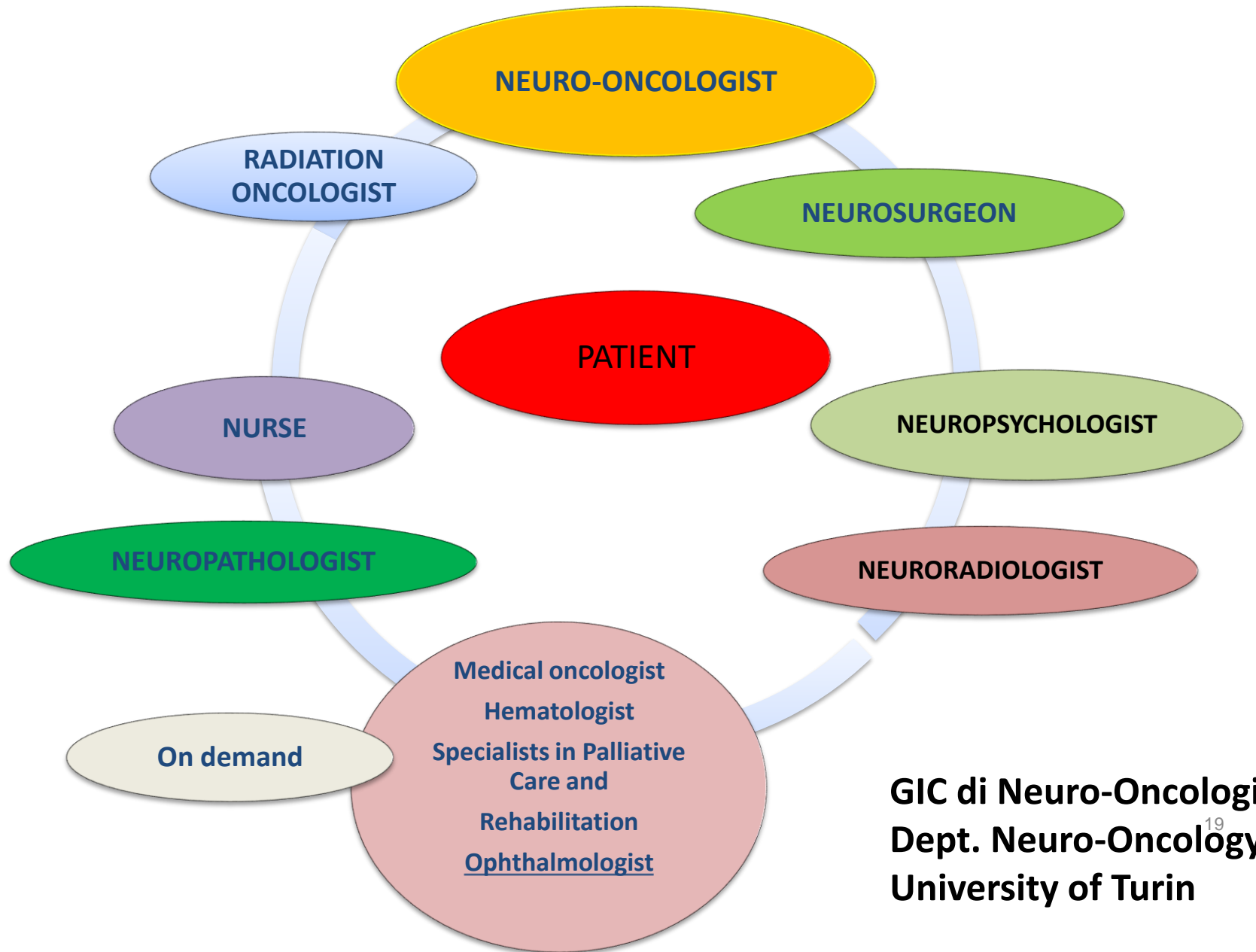


M13-379, Subject 107

Microcystic Keratopathy – Clinical Presentation

- Both eyes typically affected
- Symptoms develop 7 – 28 days from 1st ABT-414 infusion
 - Blurred vision
 - Photophobia
 - Foreign body sensation
 - Dry Eye
 - Eye Pain
- Symptoms are reversible, resolving by 4 – 6 weeks
- Prophylactic steroid eye drops mitigate but do not prevent

Only 3% discontinued ABT-414 due to eye toxicity



GIC di Neuro-Oncologia
Dept. Neuro-Oncology¹⁹,
University of Turin



Regorafenib compared with lomustine in patients with relapsed glioblastoma (REGOMA): a multicentre, open-label, randomised, controlled, phase 2 trial

Giuseppe Lombardi, Gian Luca De Salvo, Alba Ariela Brandes, Marica Eoli, Roberta Rudà, Marina Faedi, Ivan Lolli, Andrea Pace, Bruno Daniele, Francesco Pasqualetti, Simona Rizzato, Luisa Bellu, Ardi Pambuku, Miriam Farina, Giovanna Magni, Stefano Indraccolo, Marina Paola Gardiman, Riccardo Soffietti, Vittorina Zagonel

Lancet Oncol 2019; 20: 110–119

Published **Online**

December 3, 2018

<http://dx.doi.org/10.1016/>

S1470-2045(18)30675-2

REGOMA: study design

A randomized, multicenter, controlled open-label phase II clinical trial

rGBM after RT/TMZ (Stupp protocol)

- *PD by RANO criteria at least 12 weeks after completion of radiotherapy, unless the recurrence is outside the radiation field or has been histologically documented*
- *At least 1 bi-dimensionally measurable target lesion with 1 diameter of at least 10mm*
- *Histologically confirmed GBM*
- *ECOG PS 0-1 (KPS \geq 70)*

R
1:1

Regorafenib
160mg/day (3 weeks on, 1 week off)

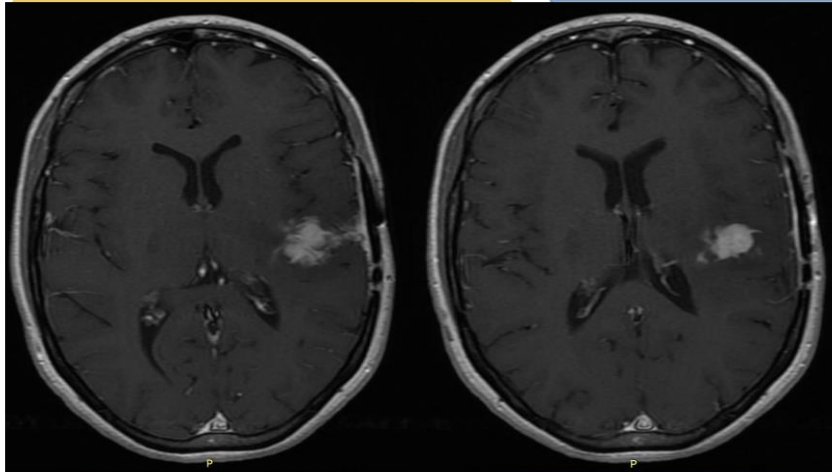
Lomustine
110mg/m² day1 (every 6 weeks)

Treat
until PD
(RANO criteria)

- Stratification factors: center and surgery at recurrence
- Study location: 10 centers in Italy

Lancet Oncology, 2019

From personalised to precision therapy

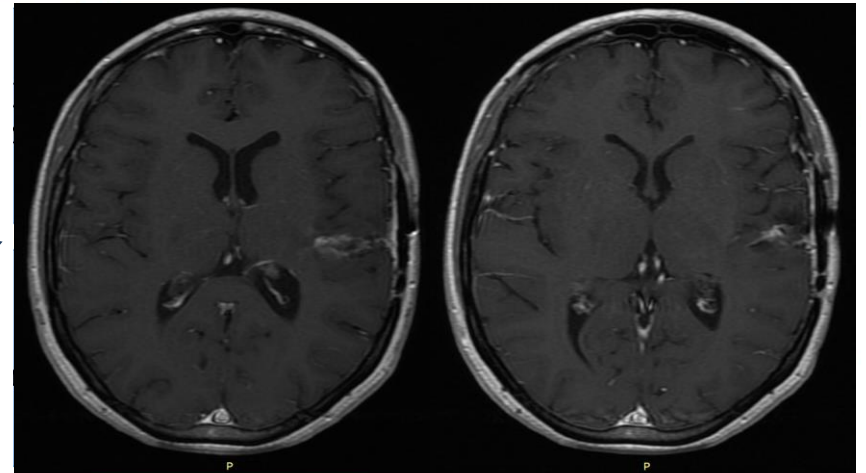


Anaplastic ganglioglioma

Relapse after chemoradiation
before dabrafenib



Major partial response following 1 year of
dabrafenib





National Comprehensive
Cancer Network®

NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines®)

Central Nervous System Cancers

Version 2.2019 — September 16, 2019

- There are no identified targeted agents with demonstrated efficacy in glioblastoma. However, the panel encourages molecular testing of tumor because if a driver mutation is detected, it may be reasonable to treat with a targeted therapy on a compassionate use basis and/or the patient may have more treatment options in the context of a clinical trial. Molecular testing also has a valuable role in improving diagnostic accuracy and prognostic stratification that may inform treatment selection.

Neuro-Oncology

XX(XX), 1–11, 2019 | doi:10.1093/neuonc/noz119 | Advance Access date 4 July 2019

**The medical necessity of advanced molecular testing in
the diagnosis and treatment of brain tumor patients**

Craig Horbinski, Keith L. Ligon, Priscilla Brastianos, Jason T. Huse, Monica Venere, Susan Chang, Jan Buckner, Timothy Cloughesy, Robert B. Jenkins, Caterina Giannini, L. Burt Nabors, Patrick Y. Wen, Kenneth J. Aldape, Rimas V. Lukas, Evanthia Galanis, Charles G. Eberhart, Daniel J. Brat, and Jann N. Sarkaria

IDH mutations and brain tumour related epilepsy

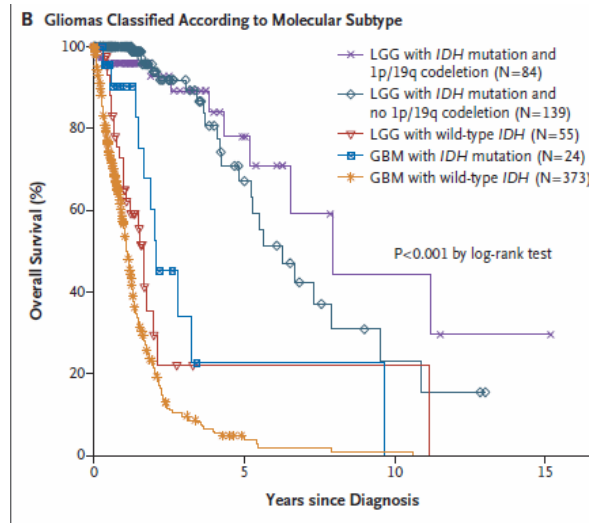
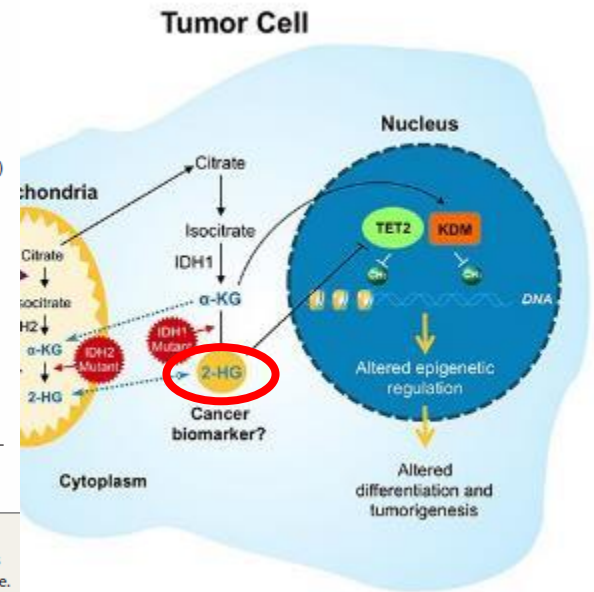


Figure 6. Clinical Outcomes.

Panel A shows Kaplan–Meier estimates of overall survival among patients with LGGs that are classified according to traditional histologic type and grade. GBM samples (from previously published Cancer Genome Atlas data²²) are also included for comparison. Panel B shows overall survival among patients with LGGs that are classified according to *IDH* mutation and 1p/19q codeletion status. GBM samples classified according to *IDH* mutation status are also included. The results of an age-adjusted analysis are provided in Table S2 in Supplementary Appendix 1. Further details on molecular subtype, grade, and molecular subtype is shown in Fig. S22 in Supplementary Appendix 1.



Luibinas et al. Epilepsia 2014

2-HG may increase BTR epilepsy by mimicking glutamate on NMDA receptors

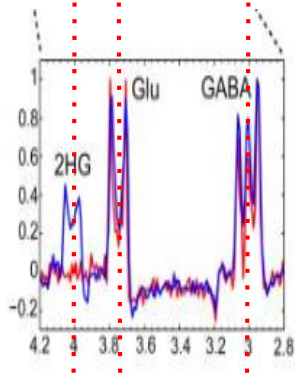
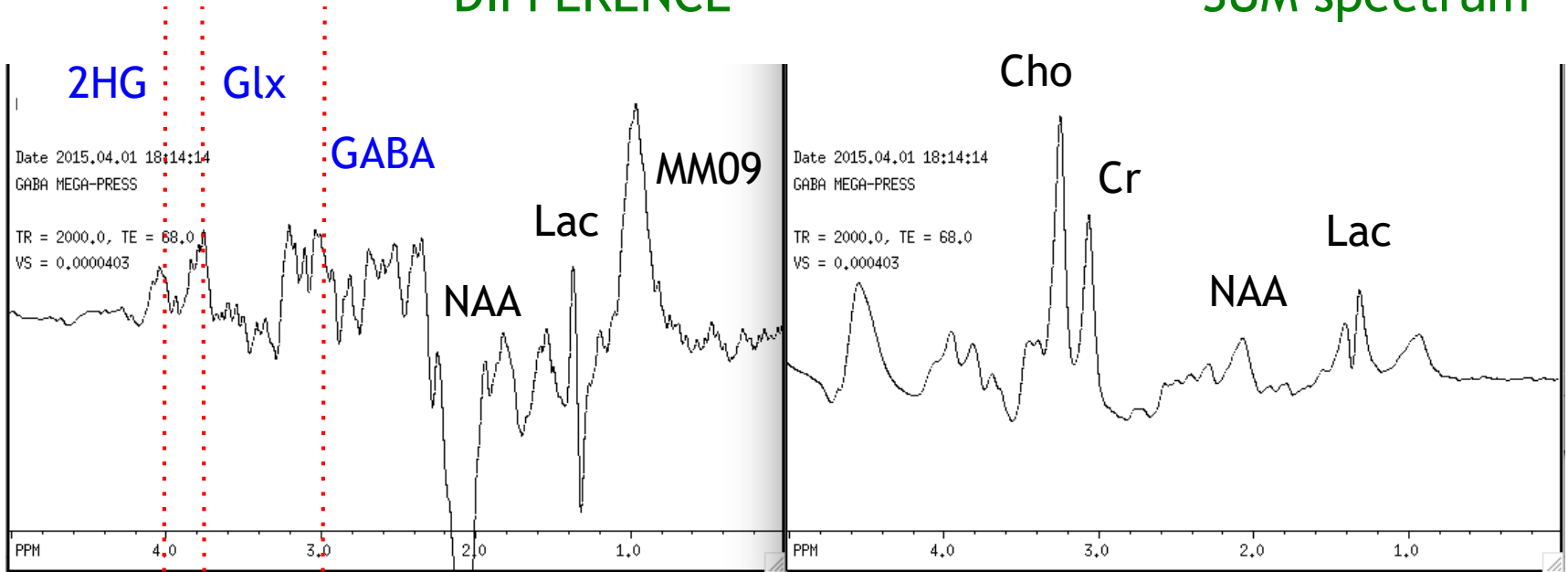
Brat et al N Engl J Med. 2015 ;372(26):2481-98.

H. Chen et al, Neurology 2017

2HG and GABA in astrocytoma WHO-II (IDH mutant)

DIFFERENCE

SUM spectrum



MEGA-PRESS sequence

TR/TE= 2000/68 msec

VOI = 30 ml

NEX = 240

TA = 8 min



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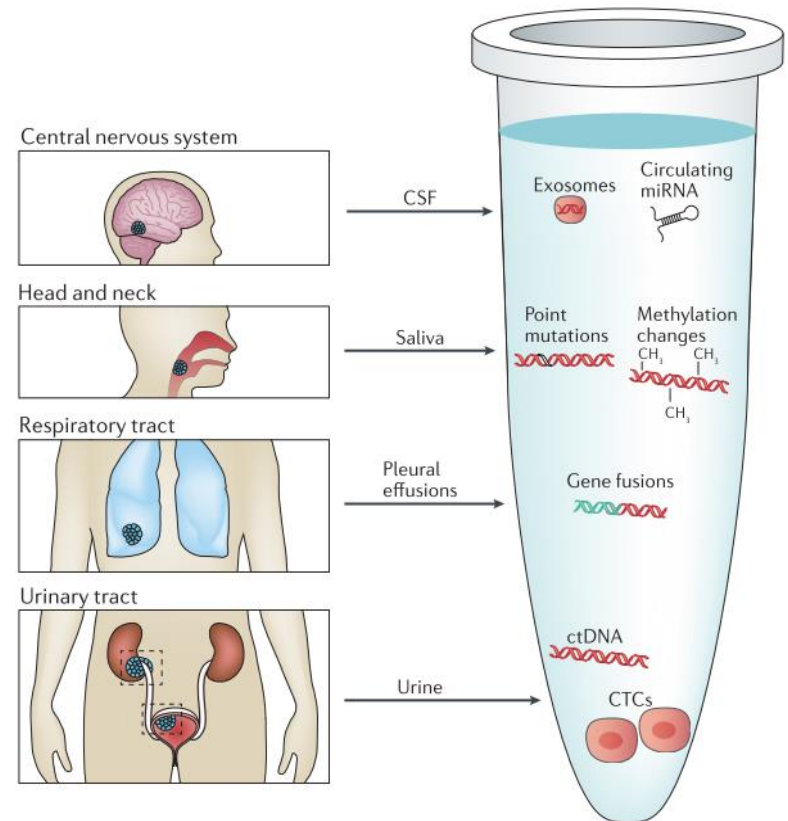
Riccardo Pascuzzo

LIQUID BIOPSY IN PRIMARY BRAIN TUMOURS

Which source for liquid biopsy?

- Blood
- CSF
- Other?

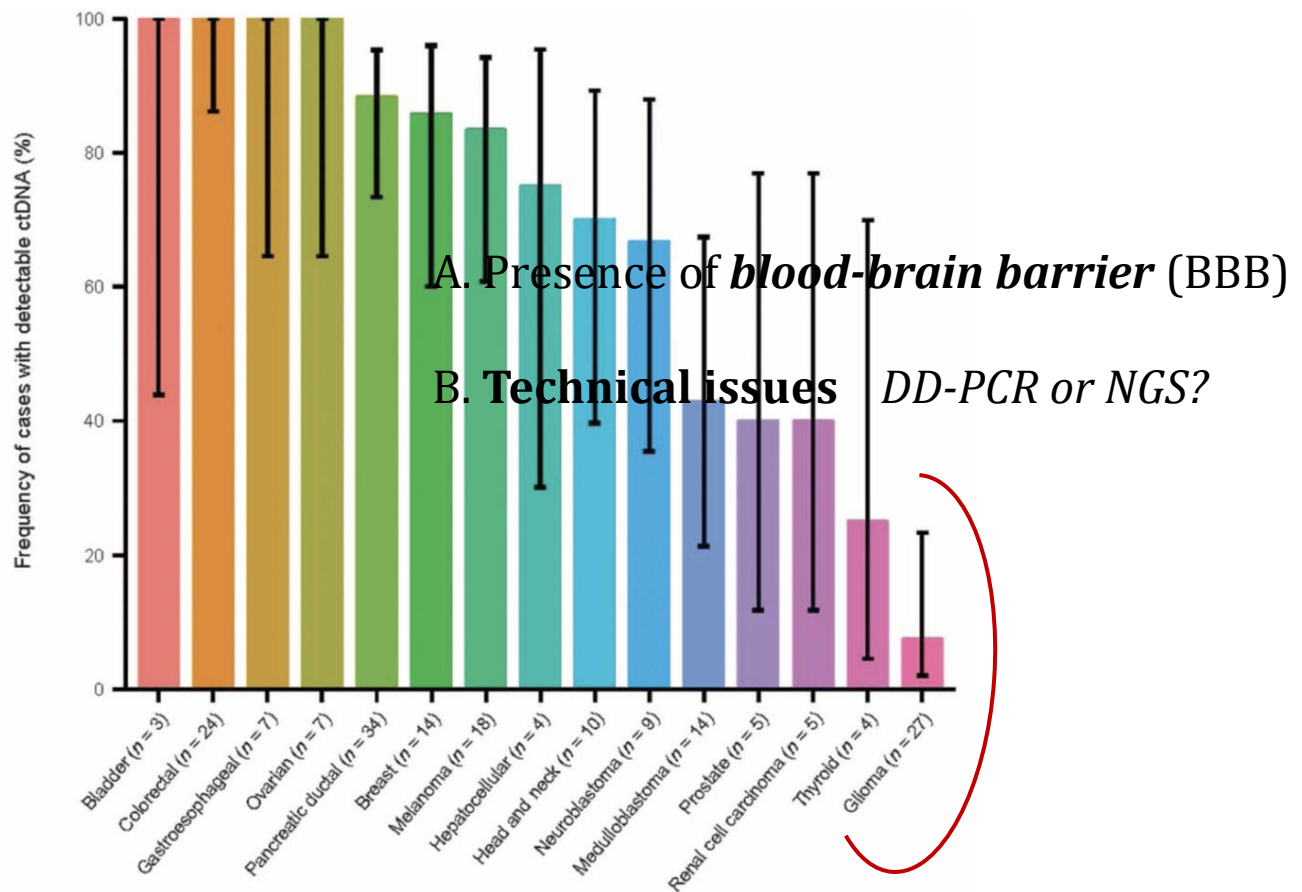
The source has to be *representative* of the specific disease of clinical interest



Siravegna G et al., *Nat Rev Clin Oncol* 2017

Circulating Tumour DNA (ctDNA)

Blood ctDNA of primary brain tumour patients is **low compared to other tumours** that are able to transfer ctDNA fragments into blood



ARTICLE

Received 25 Mar 2015 | Accepted 8 Oct 2015 | Published 10 Nov 2015

DOI: 10.1038/ncomms9839

OPEN

Cerebrospinal fluid ctDNA better than plasma ctDNA for the detection of brain tumour mutations

Leticia De Mattos-Arruda¹, Davis Torrejon¹, Mafalda Elena Martínez-Sáez⁴, Seana Vivancos¹, Vicente P. Josep Taberner^{1,3}, Enrique Jorge S. Reis-Filho² & Joa

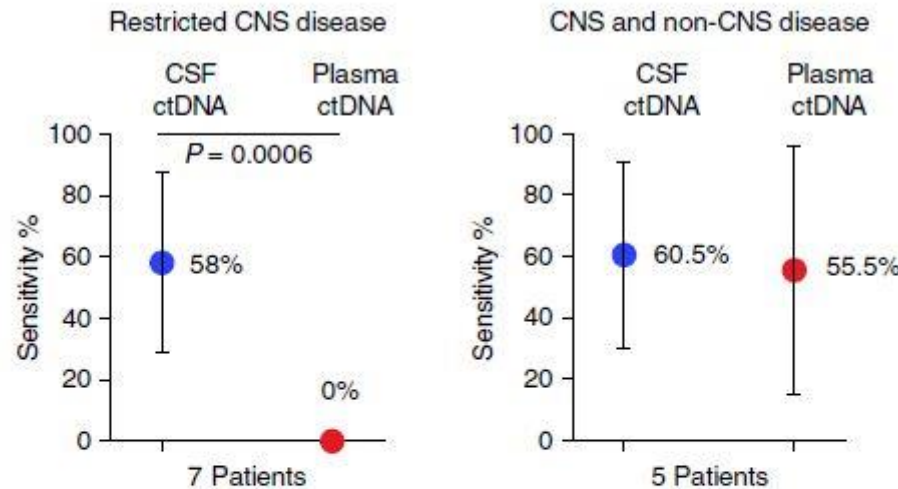


Figure 2 | Sensitivity analysis of CSF ctDNA and plasma ctDNA.

Sensitivity was inferred based on gene mutations detected in central nervous system (CNS) tumours, which were either identified in CSF or plasma ctDNA (Supplementary Table 5). Data were pooled and the mean with standard deviation error bars is shown. A Mann-Whitney test was used for the analysis and *P* value is shown.

co Martínez-Ricarte^{3,4},
ena Guerini-Rocco²,
oglio², Russel Towers⁷,
iría González-Cao⁸,
tes^{1,3},



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Luca Bertero



Carla Boccaccio

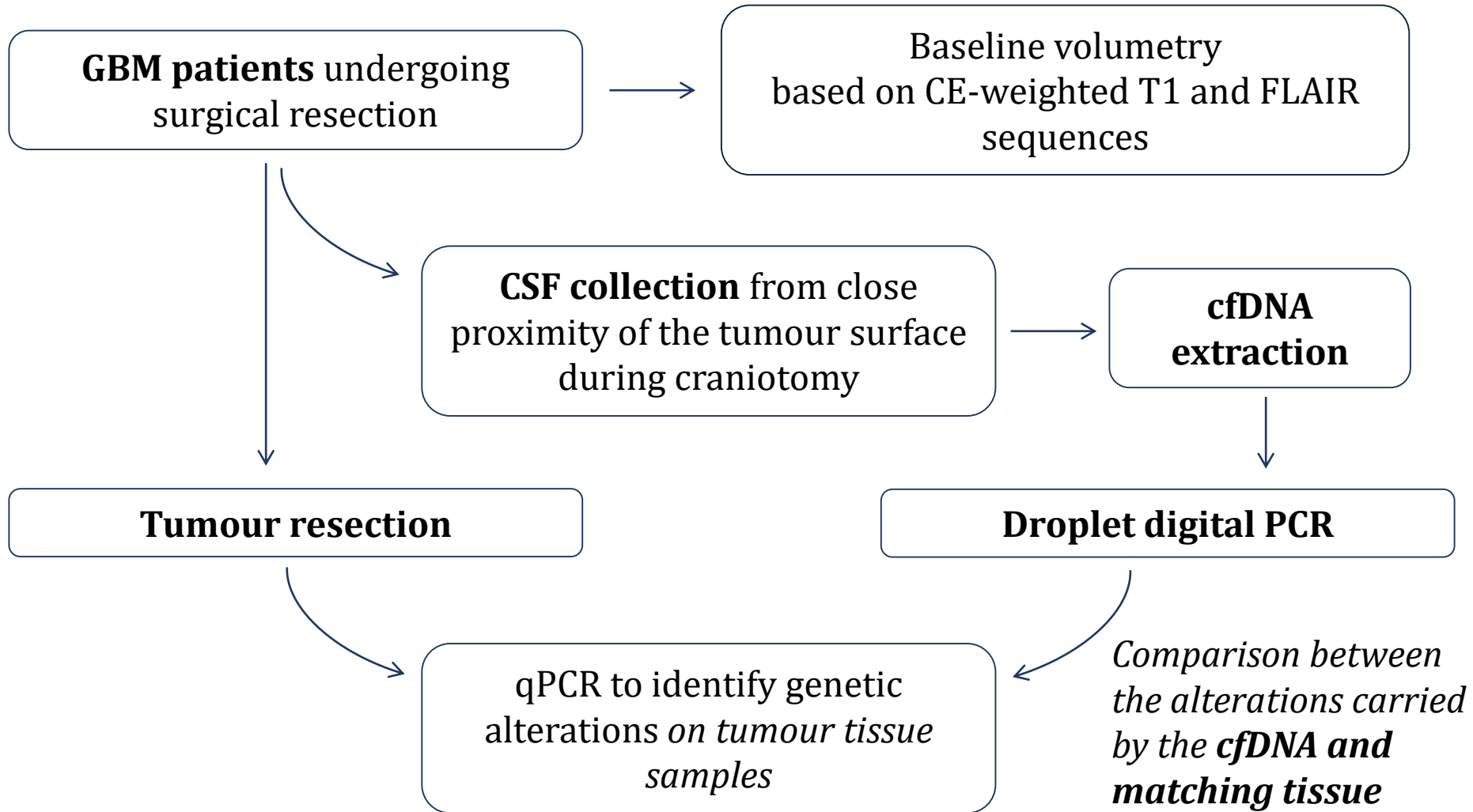


F. N. Orzan



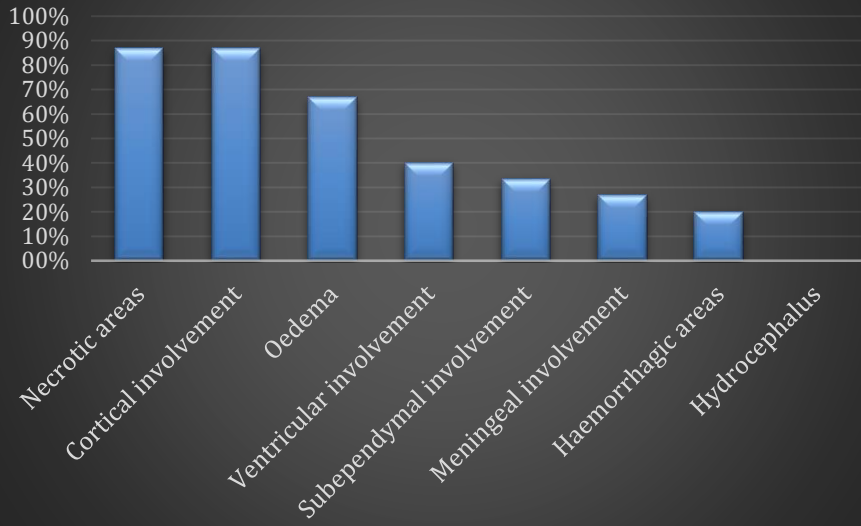
F. De Bacco

PATIENTS AND METHODS



RESULTS

Radiological features



(years)

59.1 (10-76)

Focal s

Ideomo

Seizure

Simple

Second

Comp

Prima

Intracr

Fronta

Tempo

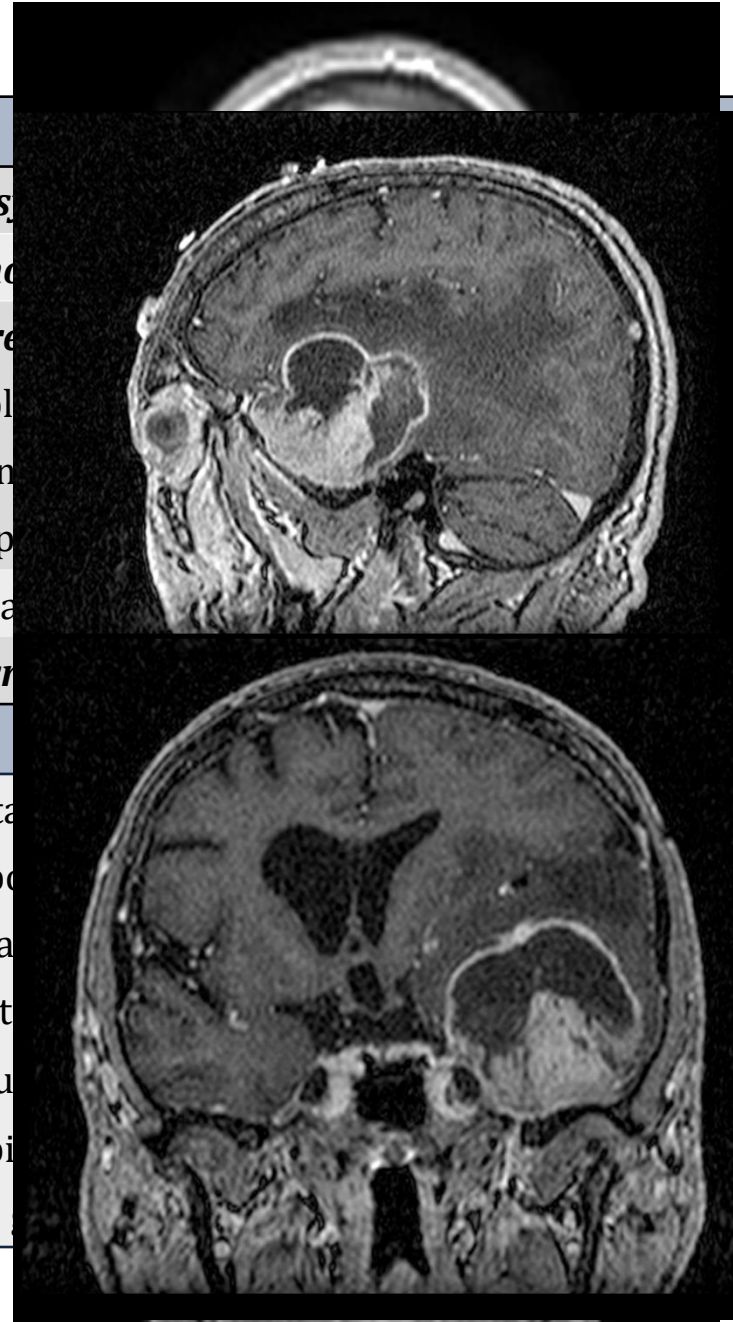
Insula

Pariet

Corpu

Occipi

Basal



RESULTS

Tissue

CSF

ID	Copy number variations					Point mutations						Alterations tested in CSF		Processable DNA
	EGFR	PDGFRA	CDK4	MDM2	CKDN2A	TP53	PTEN	IDH1	NRAS	PI3KR1	pTERT	Copy number	Mutation	
31002098	amp	0	0	0	del	NA	NA	NA	NA	NA	NA	EGFR		1
31002100	0	gain	0	0	del	0	0	0	0	1	NA		PI3KR1	1
31002108	0	0	amp	0	0	1	0	0	NA	NA	1	CDK4		1
31002112	amp	gain	amp	amp	del	0	0	0	NA	NA	1	EGFR / CDK4 / MDM2		1
31002107	0	amp	gain	0	del	1	1	0	NA	NA	NA	PDGFRA		1
31002093	0	0	0	0	del	0	0	0	0	0	NA	CKDN2A		1
31002088	gain	0	0	0	0	0	0	0	0	0	NA	EGFR		1
31002080	amp	0	amp	amp	0	0	1	0	NA	NA	NA	EGFR / CDK4 / MDM2		1
31002103	0	gain	amp	0	0	1	0	1	0	NA	NA	PDGFRA / CDK4	TP53 / IDH1	2
31002119	amp	0	0	0	del	NA	NA	NA	NA	NA	1	EGFR / CKDN2	pTERT	2
31002095	0	0	0	amp	del	0	0	0	0	0	1	MDM2 / CKDN2	pTERT	2
31002110	amp	0	0	0	del	NA	NA	NA	NA	NA	1	EGFR / CKDN2	pTERT	0
31002091	0	gain	0	0	del	1	1	0	NA	NA	NA	PDGFRA / CKDN2	TP53 / PTEN	0

RESULTS

CtDNA concentration in the CSF seems significantly related to **baseline contrast-enhancement volume** and **FLAIR/contrast enhancement ratio** ($p = 0.018$ and $p = 0.025$ respectively).

DNA concentration					
<i>Model</i>	Not standardised coefficients		Standardised coefficients		<i>Sig.</i>
	<i>B</i>	<i>Standard deviation Error</i>	<i>Beta</i>	<i>t</i>	
Constant	0.388	0.54		0.719	0.524
C.E. volume	0.029	0.006	0.98	4.766	0.018
FLAIR volume	-0.007	0.004	-0.292	-1.883	0.156
FLAIR / C.E.	0.124	0.029	0.716	4.208	0.025
CSF volume	-1.294	0.672	-0.298	-1.926	0.15



