

Neurorradiología en la Patología Tumoral Cerebral

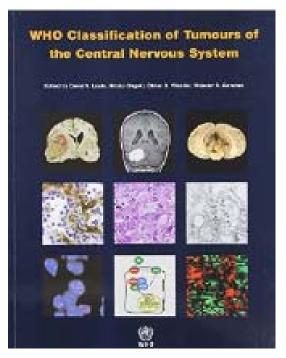
20-21 febrero 2020 | Madrid



RADIOGENÓMICA: APROXIMACIÓN Y AVANCES "FENOTIPOS DE IMAGEN DE LOS GENOTIPOS DE GLIOMAS"

Cristina Auger Acosta
Unidad de RM. Servicio de Radiología.
Hospital General Universitario Vall D'Hebron





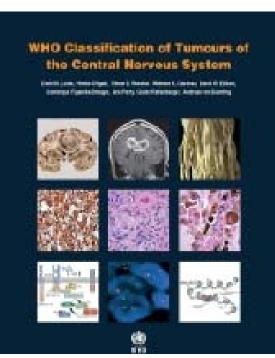
4ª edición

Acta Neuropathol (2007) 114:97–109 DOI 10.1007/s00401-007-0243-4

REVIEW

The 2007 WHO Classification of Tumours of the Central Nervous System

David N. Louis · Hiroko Ohgaki · Otmar D. Wiestler · Webster K. Cavenee · Peter C. Burger · Anne Jouvet · Bernd W. Scheithauer · Paul Kleihues



4ª edición revisada

Acta Neuropathol (2016) 131:803–820 DOI 10.1007/s00401-016-1545-1

REVIEW

The 2016 World Health Organization Classification of Tumors of the Central Nervous System: a summary

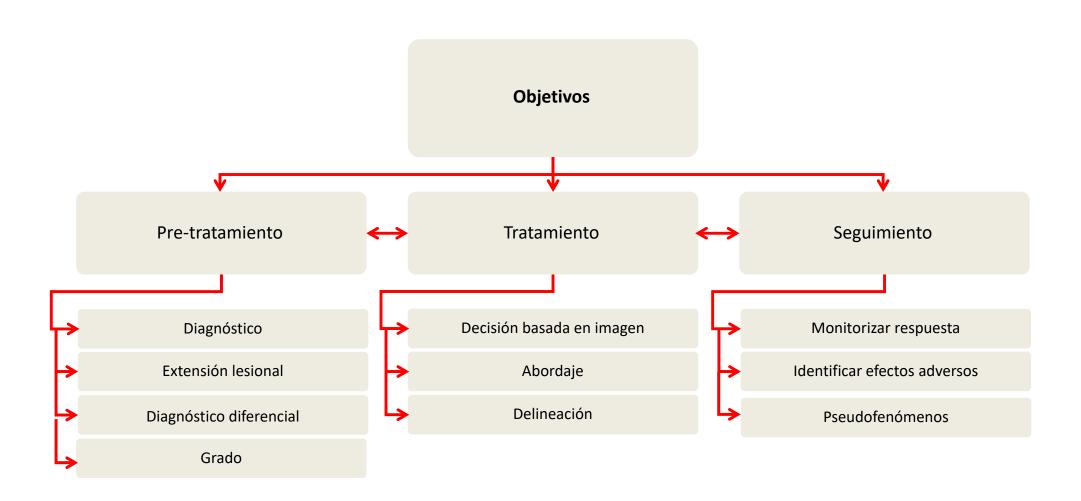
David N. Louis¹ · Arie Perry² · Guido Reifenberger^{3,4} · Andreas von Deimling^{4,5} · Dominique Figarella-Branger⁶ · Webster K. Cavenee⁷ · Hiroko Ohgaki⁸ · Otmar D. Wiestler⁹ · Paul Kleihues¹⁰ · David W. Ellison¹¹

Diffuse astrocytic and oligodendroglial tumours	
	9400/3
	9411/3 <i>9400/3</i>
	9400/3
	9401/3
	9401/3
Anapiastic astrocytoma, NOS	9401/3
Glioblastoma (DH-wildtype	9440/3
Giant cell gliobiasioma	9441/3
Gliosarcoma	9442/3
	9440/3
	9445/3* 9440/3
Gilobiastoria, NOS	9440/3
Diffuse midline glioma H3 K27M-mutant	9385/3*
	0.450/0
	9450/3 9450/3
Oligoderia oglioria, 1903	9430/3
Anaplastic oligodendroglioma, IDH-mutant	
and 1p/19q-codeleted	9451/3
Anaplastic oligodendroglioma, NOS	9451/3
Oligoastroaytoma NOS	9382/3
	9382/3
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Other astrocytic tumours	
	9421/1
	9425/3 9384/1
	9424/3
Anaplastic pleomorphic xanthoastrocytoma	9424/3
14/1/0 2016	
	Diffuse astrocytoma (DH-mutant Gemistocytic astrocytoma, EH-mutant Diffuse astrocytoma, IDH-wildtype Diffuse astrocytoma, IDH-wildtype Diffuse astrocytoma, NOS Anaplastic astrocytoma (DH-mutant Anaplastic astrocytoma, NOS Glioblastoma (DH-wildtype Giant cell gliobiastoma Gliosarcoma Epithelioid gliobiastoma Glioblastoma, NOS Diffuse midline glioma H3 K27M-mutant Glioblastoma, NOS Diffuse midline glioma IDH-mutant and 1p/19q-codeleted Oligodendroglioma, NOS Anaplastic oligodendroglioma, NOS Anaplastic oligodendroglioma, NOS Oligoastrocytoma, NOS Anaplastic oligodendroglioma, NOS Other astrocytic tumours Pilocytic astrocytoma Pilomyxoid astrocytoma Subependymal giant cell astrocytoma Pleomorphic xanthoastrocytoma

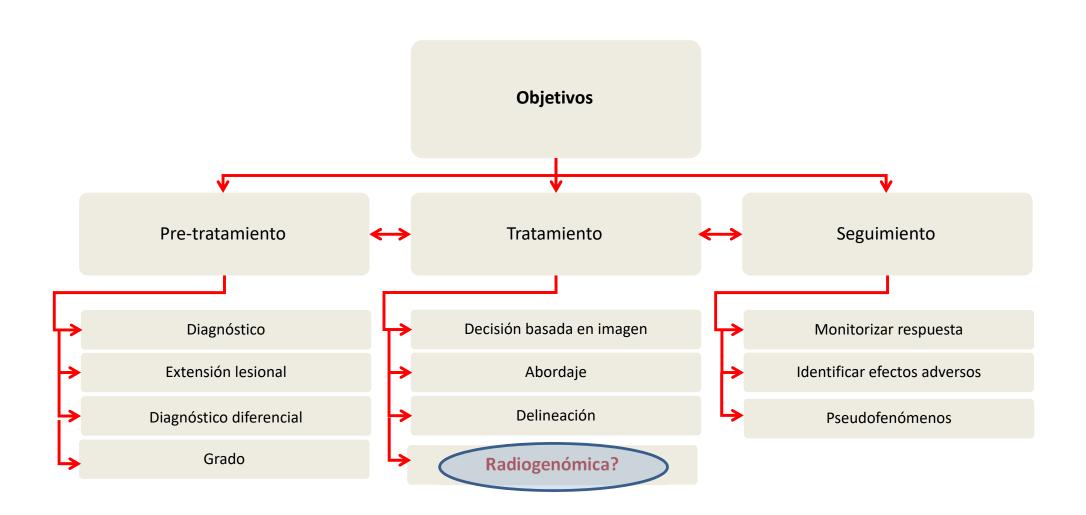
WHO 2007

WHO 2016

Objetivos de la NR en los tumores cerebrales

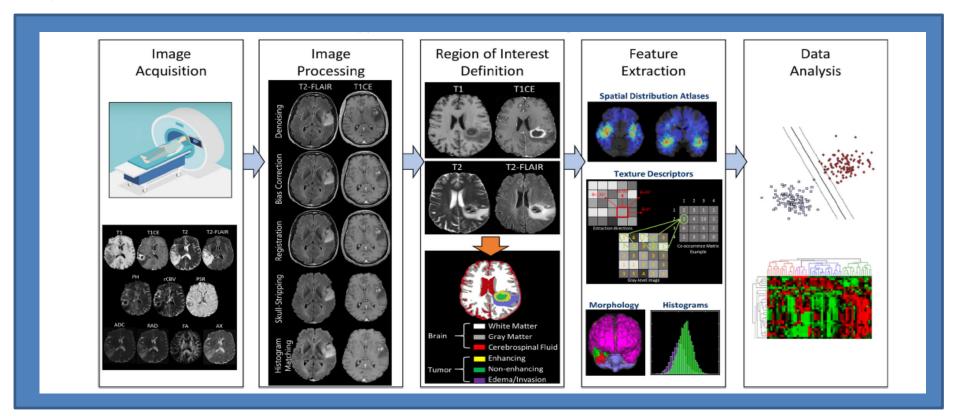


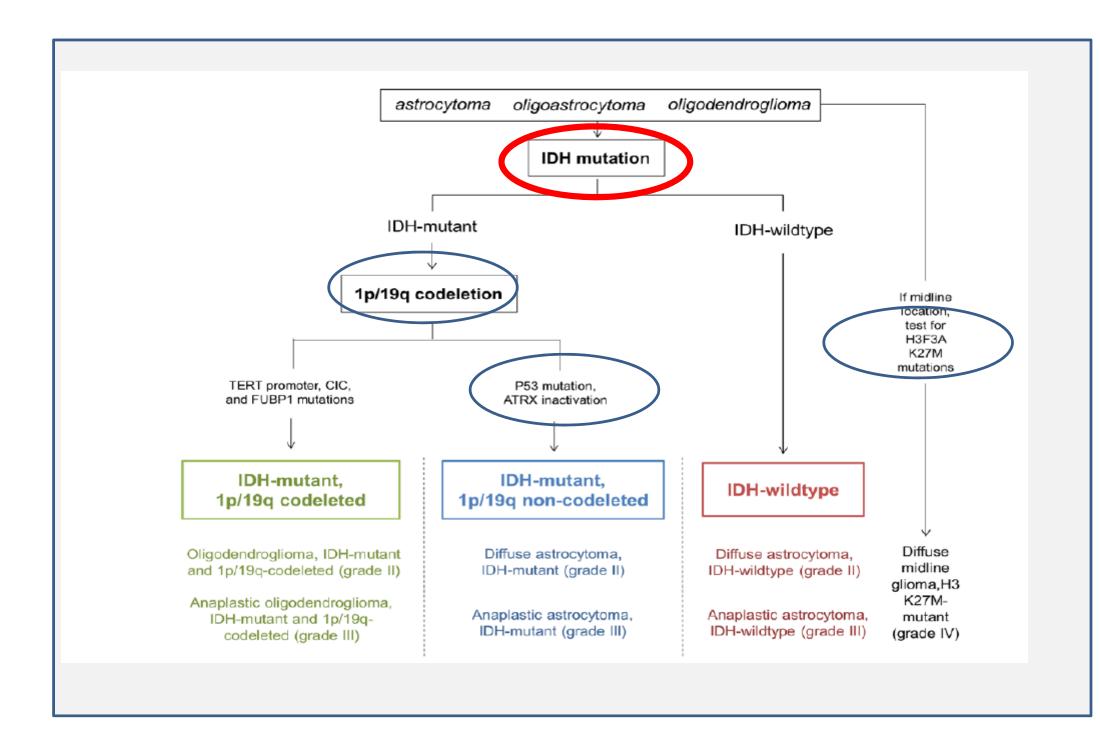
Objetivos de la NR en los tumores cerebrales



Relación entre la Imagen y Genómica en Tumores cerebrales (Radiogenómica) Objetivo

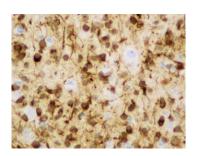
- Comprender bases moleculares del cáncer a través de la aplicación de análisis genómico, histopatología e imagen
- Identificar características radiológicas que predigan genómica, (mutaciones, expresión)

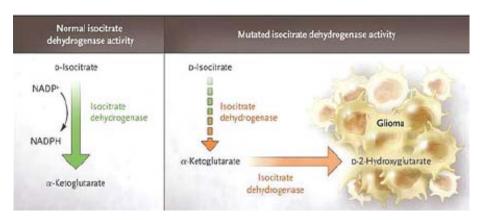




Radiogenómica en gliomas

Isocitrato DesHidrogenasa 1&2





•Responsable de la conversión del isocitrato a alfa-ketoglutarato. Su mutación produce una reducción del alfa-ketoglutarato a 2-hidroxiglutarato

Presente en:

Pacientes jóvenes

Gliomas grado II/III (>75%)

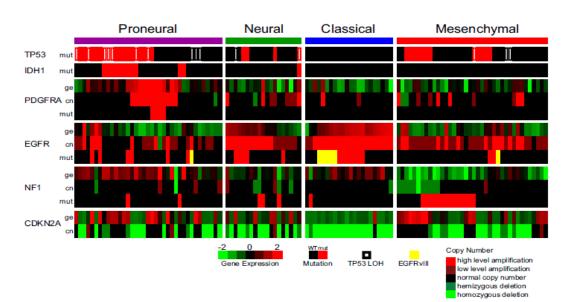
GBM secundarios (aprox 83%)

- Asociado con un perfil de expresión génica (proneural)
- Mayor supervivencia (3,8 vs 1,1 años)

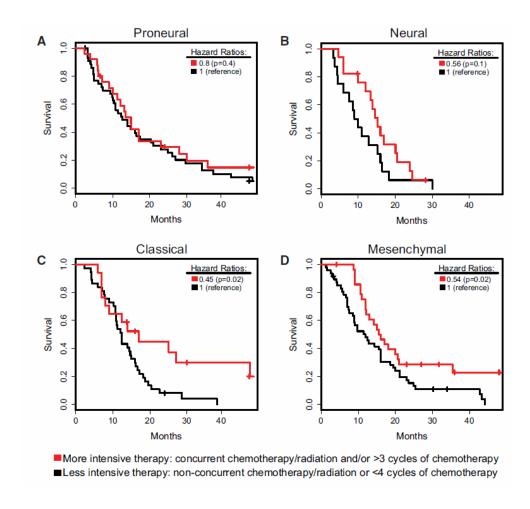
Gliomas

Clasificación basada en perfiles de expresión génica *Vias de señalización: RTK/P13K/PTEN; P53; RB1*

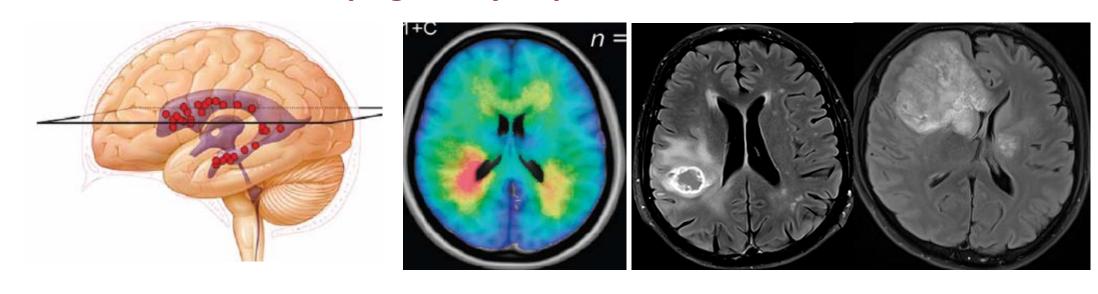
GBM

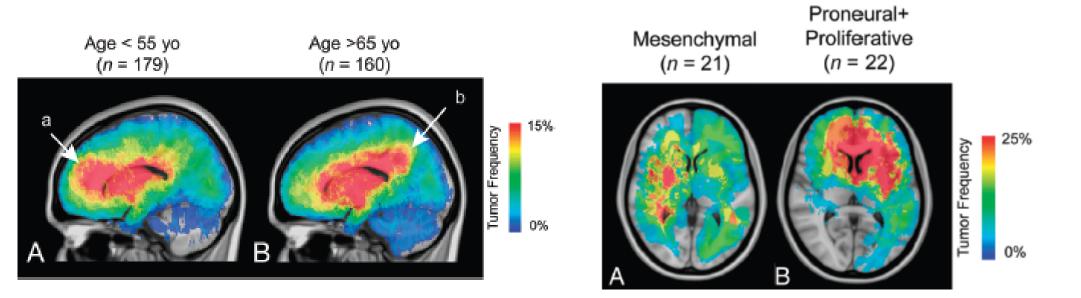


- Proneural (mejor pronóstico)
 - ✓ Menor volumen de realce tumoral
- Mesenquimal
 - ✓ Más necrosis



Radiogenómica en gliomas Topografía y supervivencia

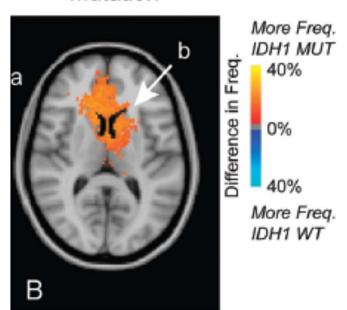




LOCALIZACIÓN

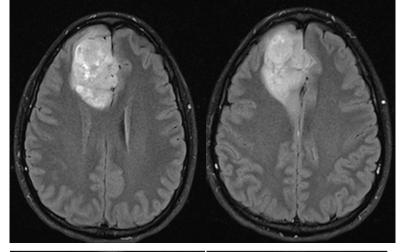
Lóbulo frontal Unilobares

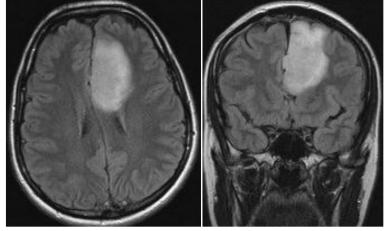
IDH1 Mutation

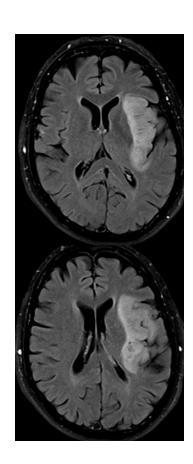


Radiogenómica en gliomas Mutaciones IDH

• Hallazgos radiológicos predicen la mutación IDH1 (precisión 97,5%)



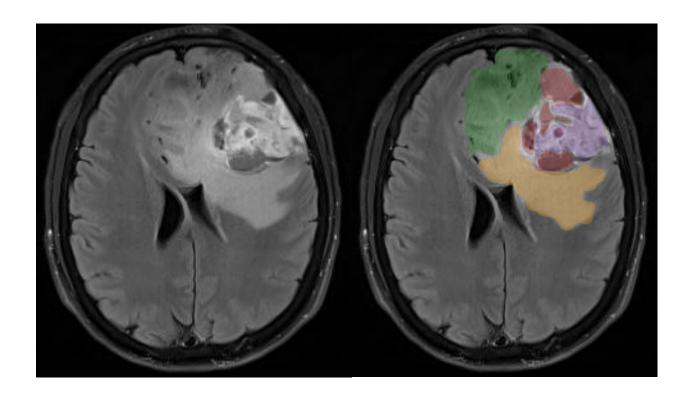




• Hallazgos radiológicos predicen la mutación IDH1 (precisión 97,5%)

APARIENCIA

- •Alto porcentaje sin realce
- •Gran tamaño
- •Presencia de quistes
- Presencia satélites



Qi et al.Oncol Lett 2014 Carrillo et al. AJNR 2012 Wang et al.Eur J Neurol 2015

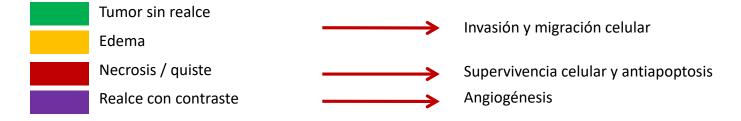


Table II. Analyzing the frequency of IDH1/2 mutations and tumor location according to anatomical structures.

Histology	F, n	T, n	P or O, n	Multilobes, n	I or BG, n	D or BS, n	CB, n
DA							
n	45	9	9	11	14	12	11
IDH mutation	38	7	8	3	6	1	9
IDH wild-type	7	2	1	8	8	11	2
AA							
n	21	16	8	17	4	14	2
IDH mutation	15	13	4	5	3	4	1
IDH wild-type	6	3	4	12	1	10	1
Overall							
n	66	25	17	28	18	26	13
IDH mutation	53	20	12	8	9	5	10
IDH wild-type	13	5	5	20	9	21	3

LOCALIZACIÓN

Lóbulo frontal Unilobares

Table IV. Analyzing the frequency of IDH1/2 mutations and different MRI features of gliomas.

MRI features	Alla, n (%)	P-value	DAa, n (%)	P-value	AAa, n (%)	P-value
Pattern of growth		<0.001		0.007		0.001
Unilateral	116/178 (65.2)		71/104 (68.3)		45/74 (60.8)	
Bilateral	1/15 (6.7)		1/7 (14.3)		0/8 (0.0)	
Tumor margins		< 0.001		0.001		0.012
Sharp	66/85 (77.6)		44/55 (80.0)		22/30 (73.3)	
Indistinct	51/108 (47.2)		28/56 (50.0)		23/52 (44.2)	
Tumor signal intensity		< 0.001		0.003		< 0.001
Homogeneous	70/89 (78.7)		45/58 (77.6)		25/31 (80.6)	
Heterogeneous	47/104 (45.2)		27/53 (50.9)		20/51 (39.2)	
Contrast enhancement		< 0.001		0.001		0.003
Absent or slight	74/97 (76.3)		47/60 (78.3)		27/37 (73.0)	
Significant	43/96 (44.8)		25/51 (49.0)		18/45 (40.0)	
Mass effect		0.654		0.320		0.216
Absent or moderate	47/75 (62.7)		38/54 (70.4)		9/21 (42.9)	
Severe	70/118 (59.3)		34/57 (59.6)		36/61 (59.0)	
Edema		0.181		0.533		0.375
Absent or moderate	71/109 (65.1)		49/73 (67.1)		22/36 (61.1)	
Severe	46/84 (54.8)		23/38 (60.5)		23/46 (50.0)	

APARIENCIA

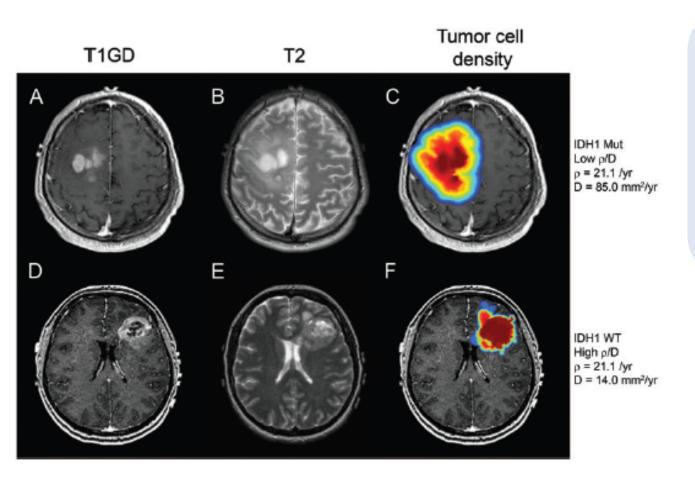
- •Alto porcentaje sin realce
- •Gran tamaño
- •Presencia de quistes
- Presencia satélites

Qi et al.Oncol Lett 2014

Radiogenómica en gliomas

IDHmut vs IDHwt

Ratio proliferación/Dispersión: p/D

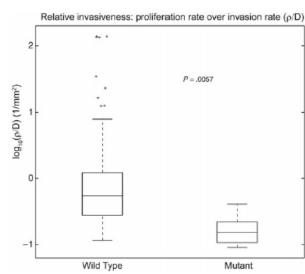


76% precisión

IDHmut<IDHwt



IDHmut: + difusos - agresivos

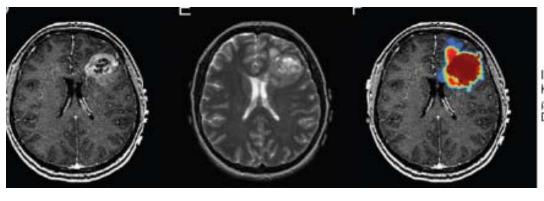


Radiogenómica en gliomas

Baldock et al Neuro-Oncology 2014

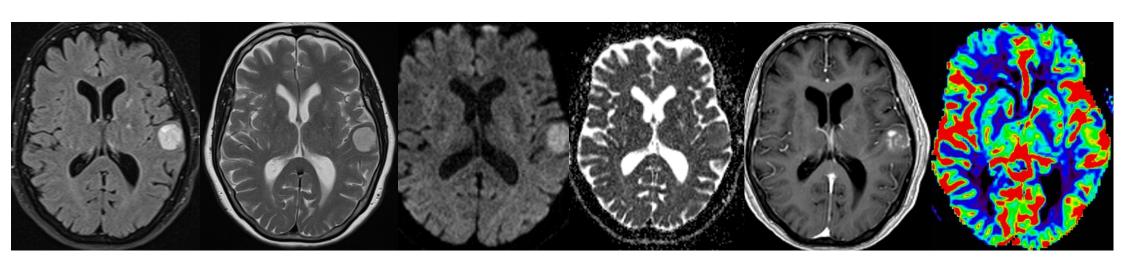






IDH1 WT ρ = 21.1 /yr D = 14.0 mm²/yr

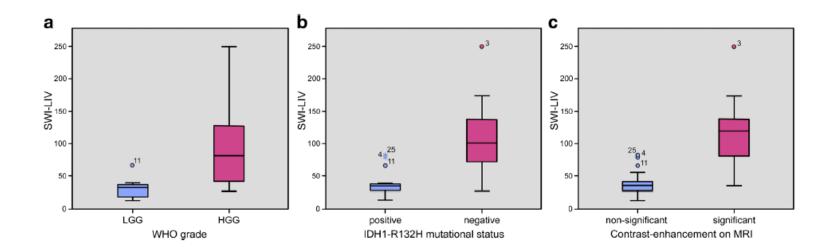
IDHwt

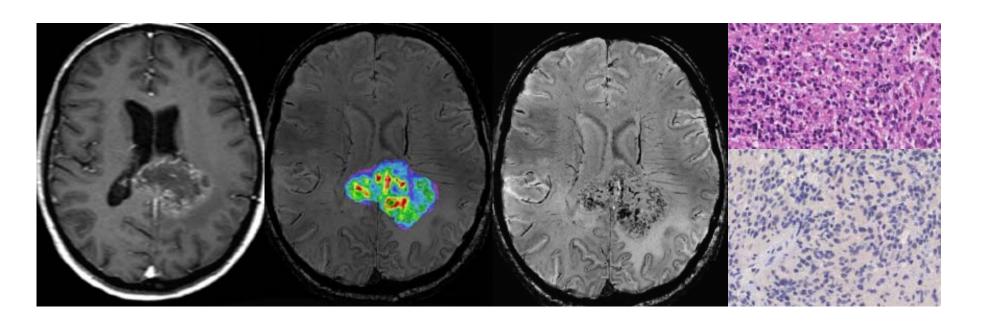


Vall d'Hebron

Mutaciones IDH1/Susceptibilidad magnética

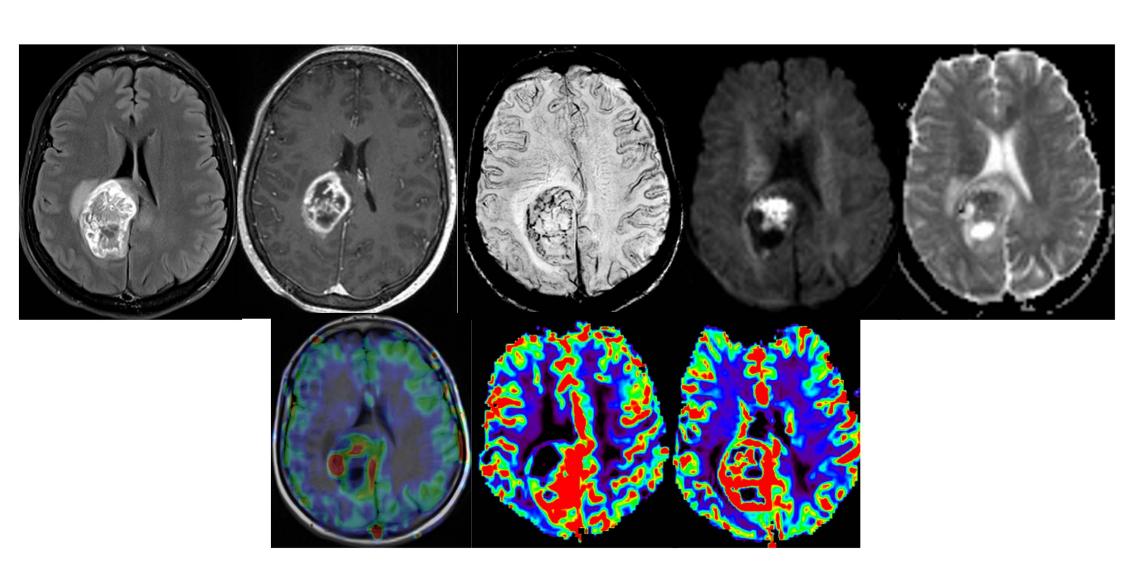
30 pacientes SWI 7T

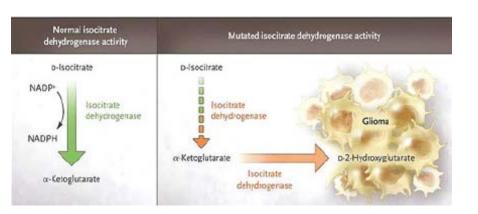




Radiogenómica en gliomas

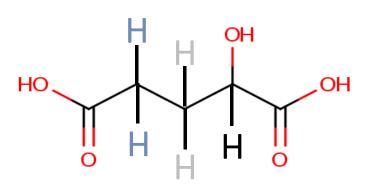
IDHwt

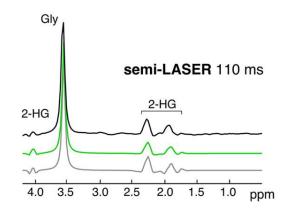




Espectroscopia y 2-hidroxiglutarato (2-HG)

2-HG es una molécula estructuralmente compleja que da diferentes señales en espectro centradas sobre 1,9, 2,25 y 4,02 ppm





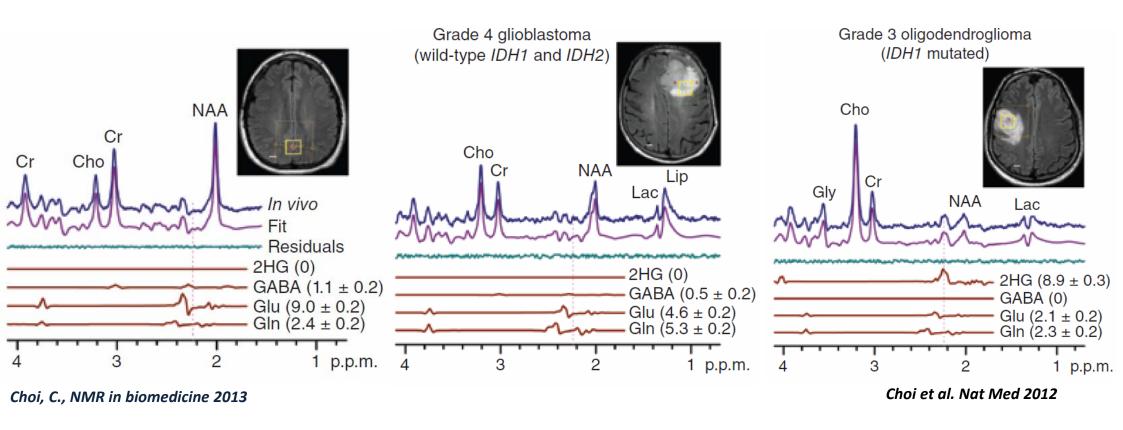
Estas señales se solapan con las debidas a otros metabolitos como el NAcc GABA, Glu, Gln, GSH, mIns.

ERM 2-hidroxiglutarato (2-HG)

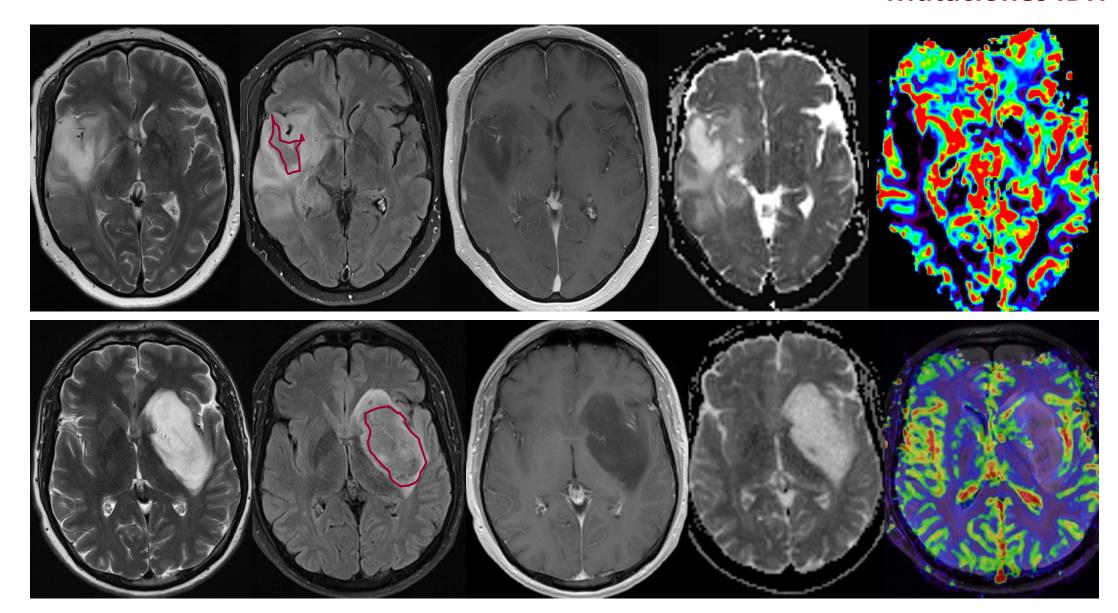
Técnicas para estudiar la presencia de 2-HG "in vivo"

técnica	Detección directa	ventajas	limitaciones
SVS TE 30ms	NO "fitting"	comercial	"fitting"
SVS TE 97ms	NO "fitting"	modificación secuencia comercial	"fitting"
"editing" SVS o CSI	SI eliminación señales otros compuestos	detección 2-HG No falsos positivos	NO comercial Volumen y TA 个
2D-COSY	SI presencia pico conexión	detección 2-HG No falsos positivos	NO comercial procesado cuantificación volumen y TA 个

- Marcador diagnóstico y pronóstico en gliomas (inmunohistoquimia)
- 2HG (2-hydroxyglutarate) se produce por todos los enzimas de la mutación *IDH*
- 2HG puede detectarse con espectroscopía (3Tesla) (2.25 ppm)



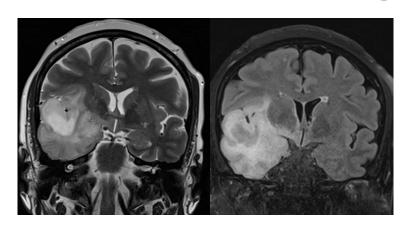
Mutaciones IDH



T2-FLAIR—mismatch sign

T2-FLAIR—mismatch sign

Mutaciones IDH



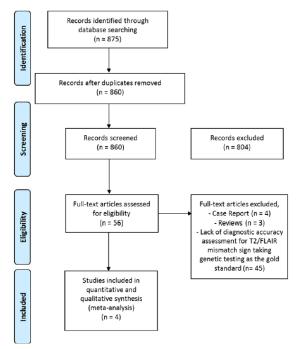
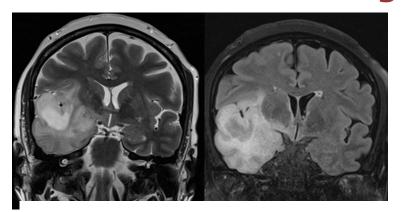


TABLE 2. Diagnostic performance of the T2-FLAIR-mismatch sign in predicting IDH mutation

		ADH Mutation Diagnostic Performance Parameter								
Authors & Year	Sensitivity, 9	6 Specificity, %	PPV, %	NPV, %	LR+	LR-	Posttest Probability (+), %*	Posttest Probability (-), %*		
Patel et al., 2017 (cases)	14.7	100	100	20.9	14.7	0.9	98.3	77.7		
Patel et al., 2017 (validation)	18.9	100	100	14.0	18.9	8.0	98.7	76.8		
Lasocki et al., 2018	58.1	100	100	14.8	58.1	0.4	99.6	63.2		
Broen et al., 2018	26.8	100	100	10.3	26.8	0.7	99.1	75.0		
Overall	32.1	100	100	15.1	32.1	0.7	99.2	73.5		

^{*} Positive and negative posttest probabilities were determined using Bayes theorem with a pretest probability of 80.2%.

T2-FLAIR—mismatch sign



Records identified through database searching (n = 875) Records sententified through database searching (n = 875) Records screened (n = 860) Records screened (n = 804) Records screened (n = 804) Full-text articles assessed for eligibility (n = 55) Full-text articles assessed for eligibility (n = 55) Full-text articles excluded, - Case Report (n = 4) - Reviews (n = 3) - Reviews (n = 4) - Reviews (n

Mutaciones IDH

TABLE 4. Diagnostic performance of the T2-FLAIR-mismatch sign in predicting IDHmut-Noncodel

	IDH Mutation Diagnostic Performance Parameter								
Authors & Year	Sensitivity, %	Specificity, %	PPV, %	NPV, %	LR+	LR-	Posttest Probability (+), %*	Posttest Probability (-), %*	
Patel et al., 2017 (cases)	22.1	100	100	51.8	22.1	8.0	95.7	43.8	
Patel et al., 2017 (validation)	45.5	100	100	76.0	45.5	0.5	97.8	35.3	
Lasocki et al., 2018	31.4	87.5	78.6	46.7	2.5	8.0	71.5	43.9	
Broen et al., 2018	50.7	100	100	68.1	50.7	0.5	98.1	33.0	
Overall	33.7	98.5	95.5	60.7	22.5	0.7	95.7	40.2	

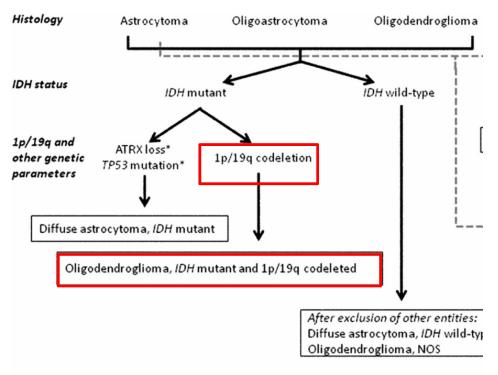
TABLE 3. Diagnostic performance of the T2-FLAIR-mismatch sign in predicting IDHmut-Codel

	IDH Mutation Diagnostic Performance Parameter							
Authors & Year	Sensitivity, %	Specificity, %	PPV, %	NPV, %	LR+	LR-	Posttest Probability (+), %*	Posttest Probability (-), %*
Patel et al., 2017 (cases)	0.0	84.3	0.0	70.4	0.01	1.2	0.4	33.9
Patel et al., 2017 (validation)	0.0	65.5	0.0	38.0	0.01	29.9	0.4	39.8
Juratli et al., 2019	73.2	76.0	83.3	63.3	3.1	0.4	56.9	13.2
Broen et al., 2018	0.0	62.0	0.0	53.4	0.01	1.6	0.4	41.1
Overall	29.9	72.7	44.4	58.7	1.1	1.0	32.2	29.4

^{*} Positive and negative posttest probabilities were determined using Bayes theorem with a pretest probability of 30.2%.



WHO2016





80% Gliomas de bajo grado son IDH mut: 37-50% codelección Son más sensibles al tratamiento

Mutaciones IDH/ Delección 1p/19q

102 LGG IDH mut

Lóbulo frontal Homogeneidad T2/flair mismatch sign T2* señal

Márgenes

Realce con contraste

Quistes

Necrosis

Diámetro máximo

Infiltración cortical

Edema

+ de 3 lóbulos

Desviación línea media

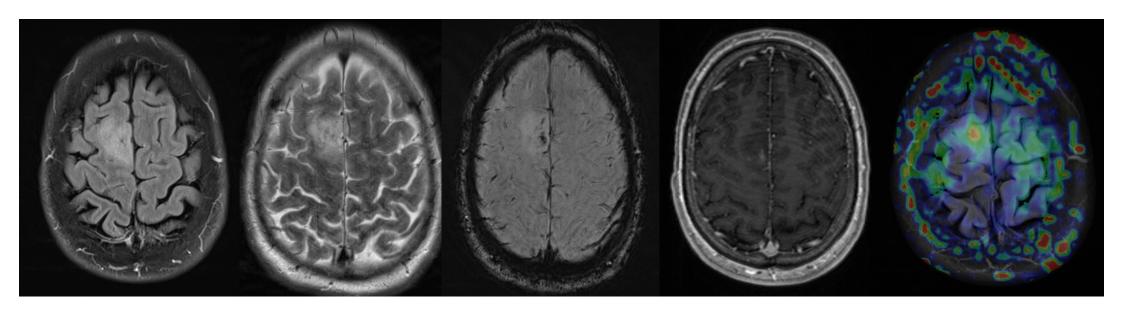
Hidrocefalia

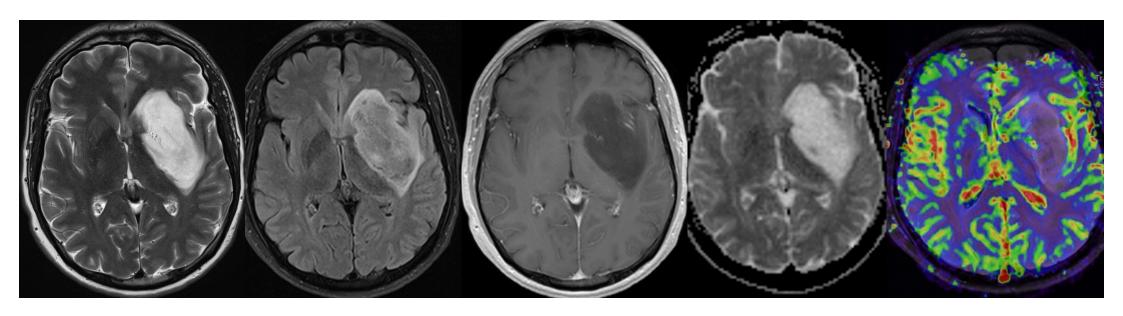


Table 1: Univariate logistic regression analyses for predicting 1p/19q codeletion among the training dataset

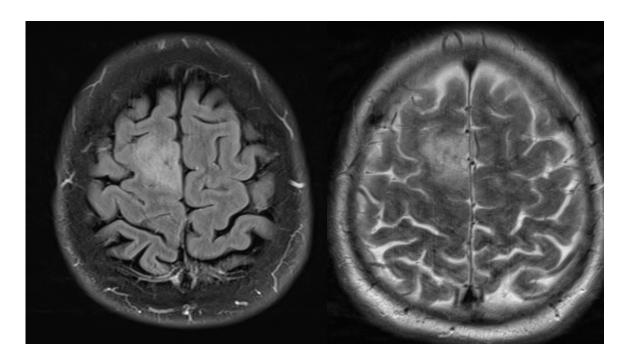
		Odds	
Predictor	Ratio	Ratio (95% CI)	P Value
Maximum diameter (cm)	1st Quartile: 3rd quartile	1.33 (0.70-2.49)	.381
Margins	Irregular: sharp	1.04 (0.47-2.33)	.917
Texture	Homogeneous (<75%):	12.33 (4.66-31.58)	$<.001^{a}$
	homogeneous (>75%)		
Peritumoral edema	Yes: no	1.42 (0.62-3.23)	.973
Hydrocephalus	No: yes	2.32 (0.88-6.11)	.089
Midline shift (cm) ^b	1st Quartile: 3rd quartile	4.27 (1.49-12.23)	.027ª
Enhancement	Yes: no	1.28 (0.57-2.86)	.555
Necrosis	Yes: no	2.18 (0.61–7.69)	.228
T2* blooming	Yes: no	6.97 (2.04-23.49)	.007ª
Cortical infiltration	Yes: no	2.02 (0.67-6.10)	.212
Cyst	No: yes	1.18 (0.48-2.91)	.715
T2 FLAIR mismatch sign ^c	No: yes	22.50 (6.26–∞)	<.001a
Gliomatosis	Yes: no	1.13 (0.18-7.08)	.896
Primary lobe	Frontal: nonfrontal	5.68 (2.08-15.44)	.001ª
Age	3rd Quartile: 1st quartile	3.38 (1.71–6.71)	$<.001^{a}$
Sex	Female: male	1.55 (0.69–3.50)	.283

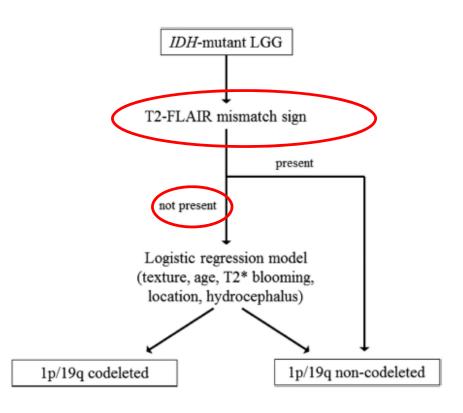
Mutaciones IDH/ Delección 1p/19q



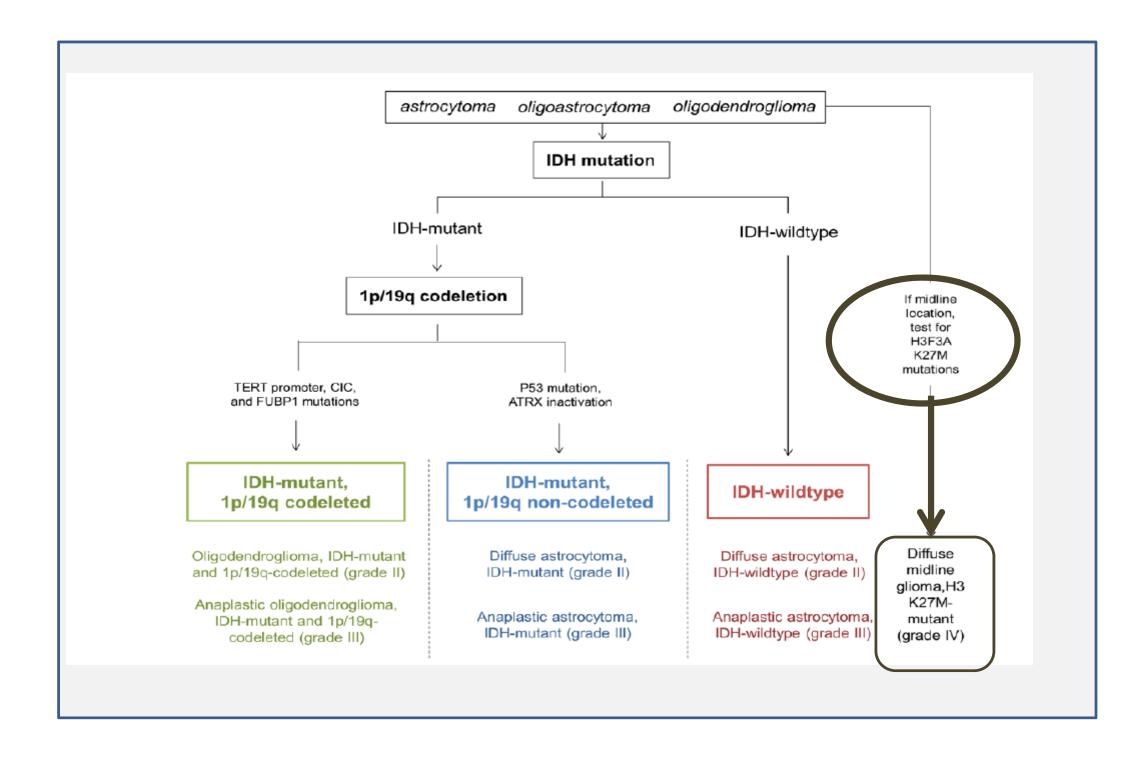


Mutaciones IDH/ Delección 1p/19q

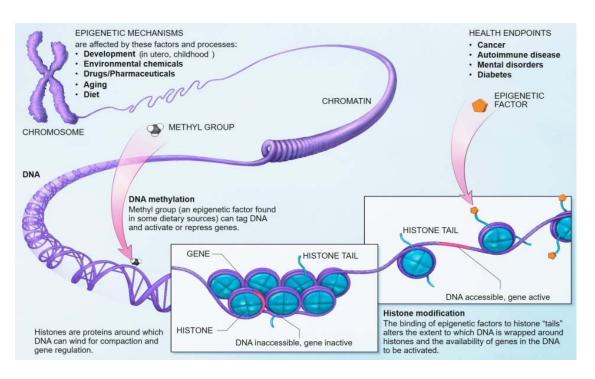


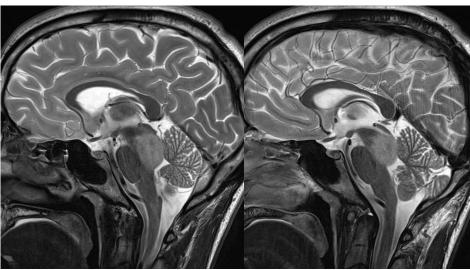


Batchala et al. AJNR 2019



Glioma difuso de línea media H3 K27 mutado





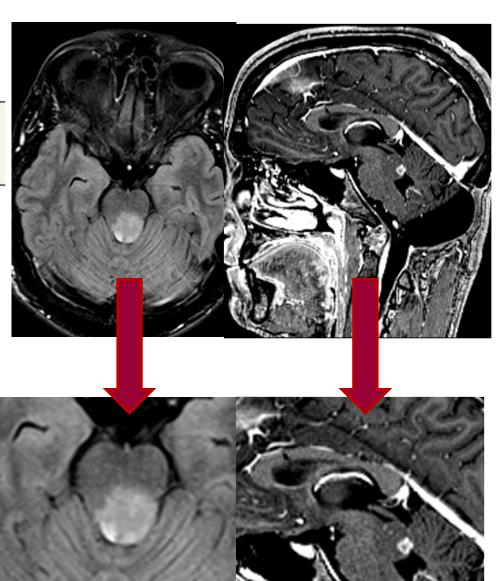
Tálamo
Hipotálamo
Pineal
III v
Cerebelo
Tronco
Médula espinal

Glioma difuso de línea media H3 K27 mutado

Table 2: MRI characteristics^a

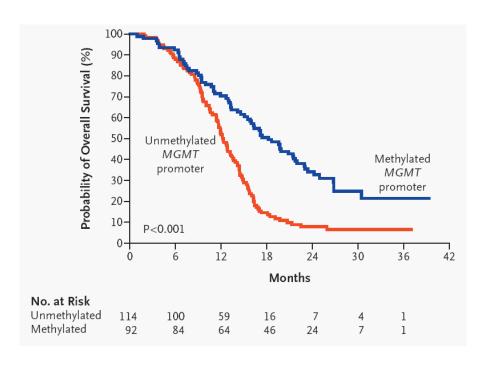
	All Patients (n = 33)	Histone H3 K27 Mutant (n = 24)	Histone H3 Wildtype (n = 9)
Multifocality	5	5	0
Contrast enhancement	22	(16)	6
Cystic component or necrosis	18	15	3
Edema	4	4	0
Infiltrative pattern	27	18	9
Mass effect	32	24	8
Irregular border	27	18	9
CSF-based metastases	7	6	1
Direct cortical invasion	12	9	3

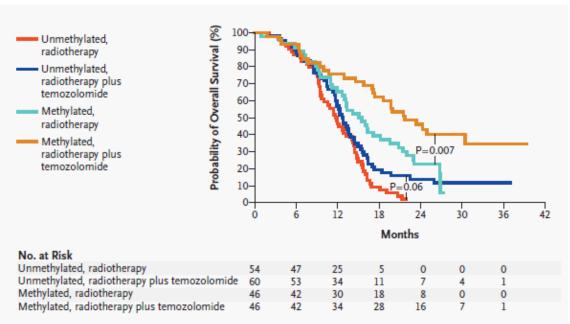
Aboian et al. AJNR2017



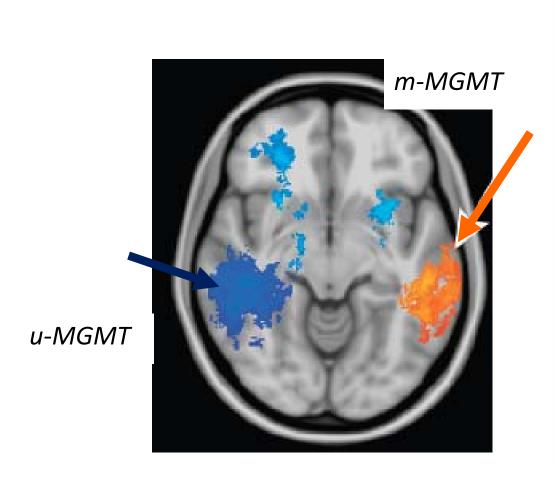
Enzima reparador MGMT

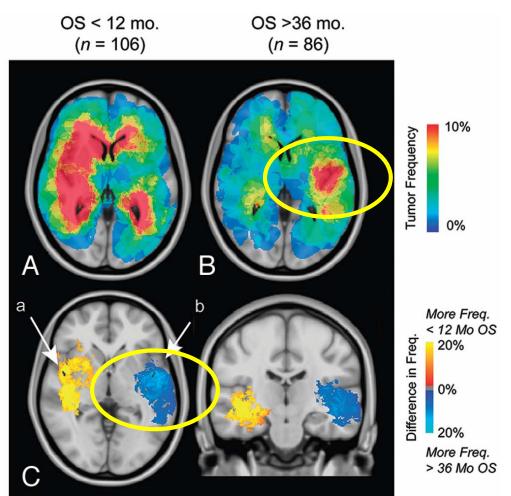
- Repara células tumorales (aumenta su supervivencia)
- Metilación MGMT inactiva su función reparadora (50% de GBMs)
- •M-MGMT mayor supervivencia pacientes, mejor respuesta a TMZ, mayor incidencia de PsP
- Pobre correlación entre hallazgos RMc y estado MGMT





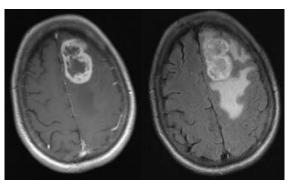
Radiogenómica en gliomas alto grado Topografía y supervivencia

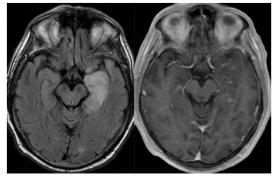


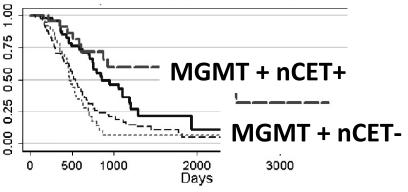


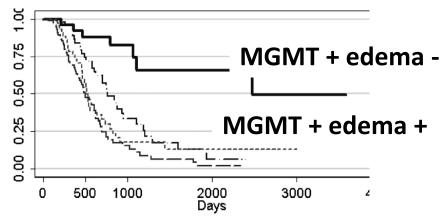
Radiogenómica en gliomas alto grado Enzima reparador MGMT

- Realce en anillo se asocia con MGMT no metilado
 - M-MGMT sin edema mayor supervivencia
 - M-MGMT con áreas sin realce mayor supervivencia



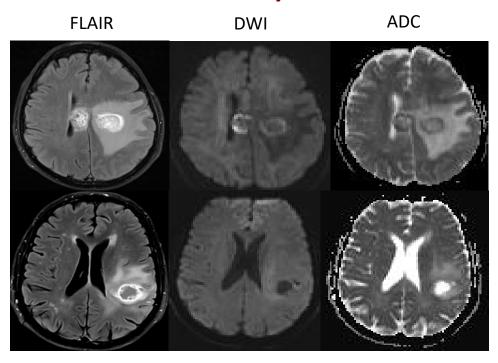




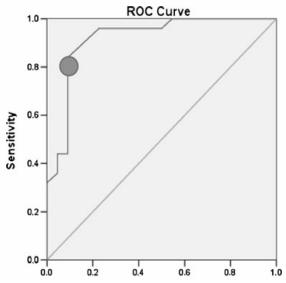


Radiogenómica en gliomas alto grado

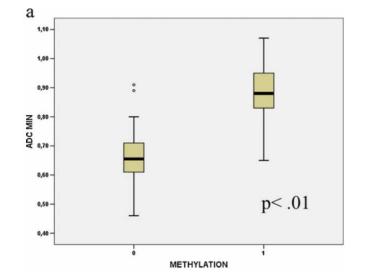
ADC y MGMT

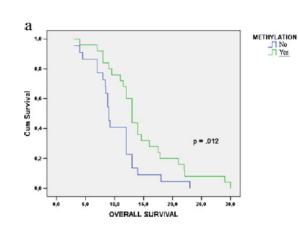


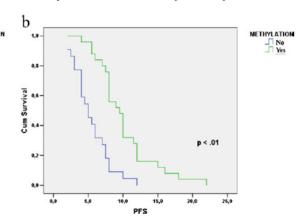
Diferenciación entre gliomas metilados v no-metilados



Valor ADC mínimo: 0,80×10-3 mm²/s, sensibilidad (84%) y especificidad (91%)



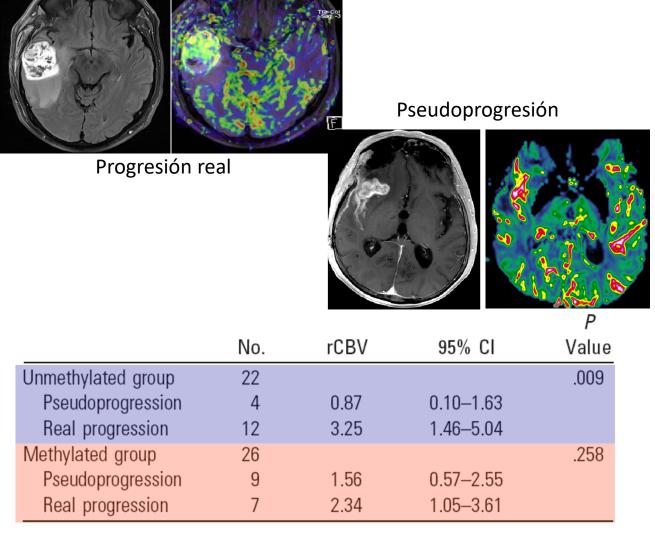


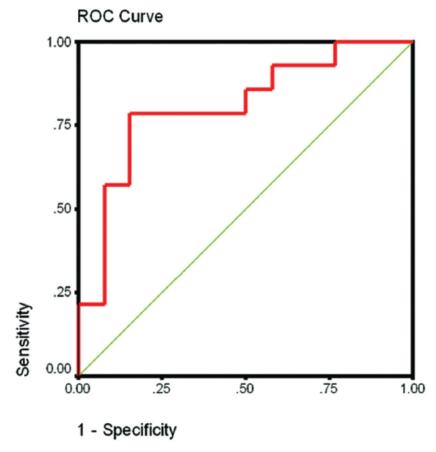


Romano et al. Eur Radiol 2012

Progresión precoz vs. Pseudoprogresión

Valor diagnóstico Perfusión (rCBV)





rCBV de 1,4 tiene 81,5% sensibilidad y 77,8% especificidad

Conclusiones

Técnicas de RM convencional y avanzada útiles para el diagnóstico y seguimiento de tumores cerebrales

- Diagnóstico
- Diagnóstico diferencial
- Gradación
- Planificación terapéutica
- Respuesta-progresión
- Pseudofenómeno: pseudoprogresión / radionecrosis / pseudorespuesta
- Radiogenómica



- Refinamiento clasificación
- Predicción pronóstica
- Tratamiento individualizado



Vall d'Hebron

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Servicio de Neurocirugía
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