

**First line therapy of glioblastoma.
Chemo-radiotherapy integration: for some
but not for all?**

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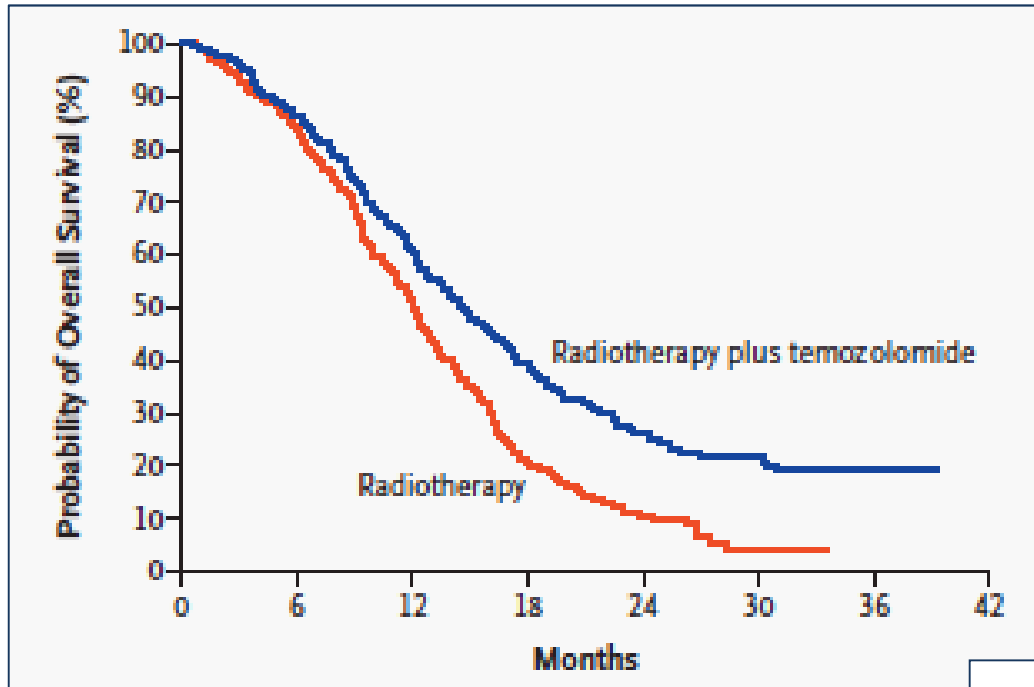
Outline

- the standard
- why should we choose the patients?
- how could we choose the patients?
- which are the choices?
- the choice of choosing: is it the right choice?



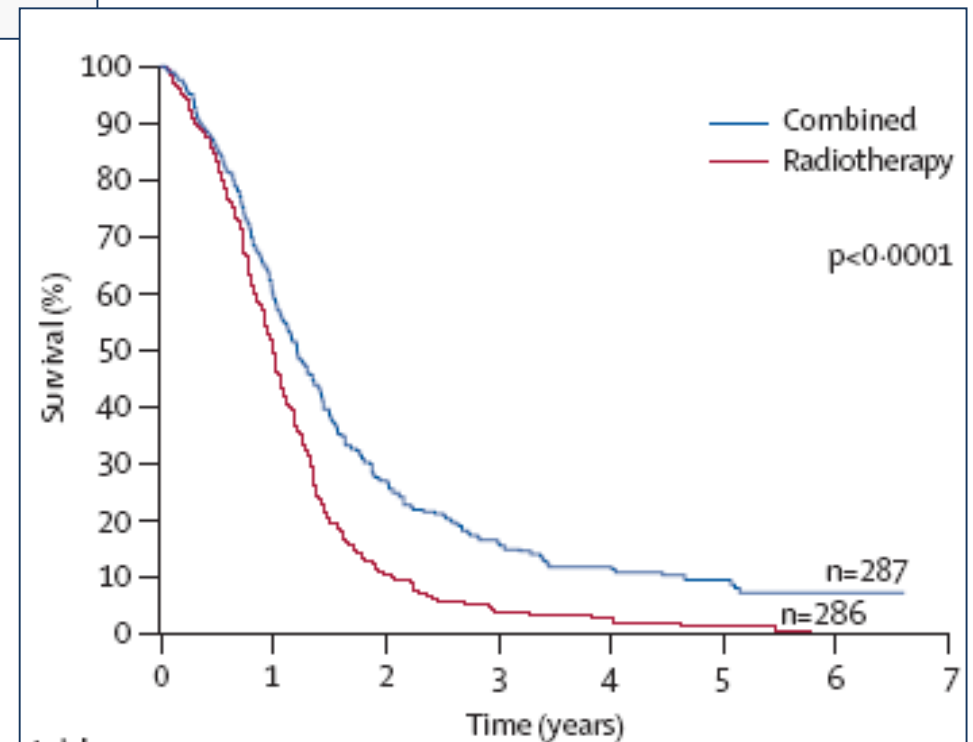
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Stupp 2005 NEJM

Stupp 2009 Lancet Oncol





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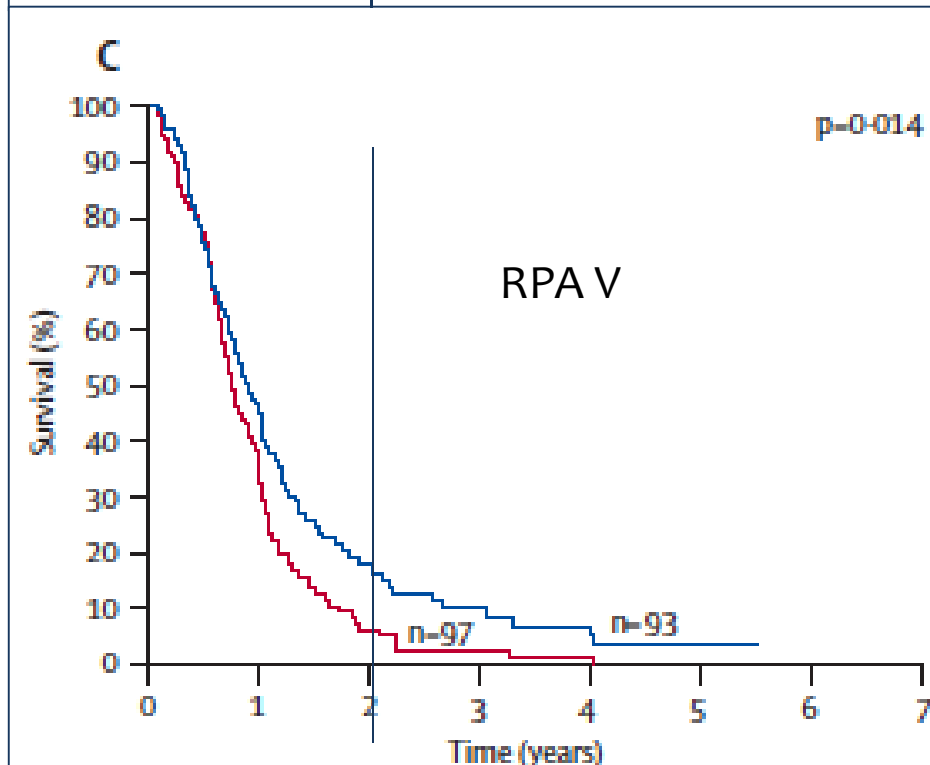
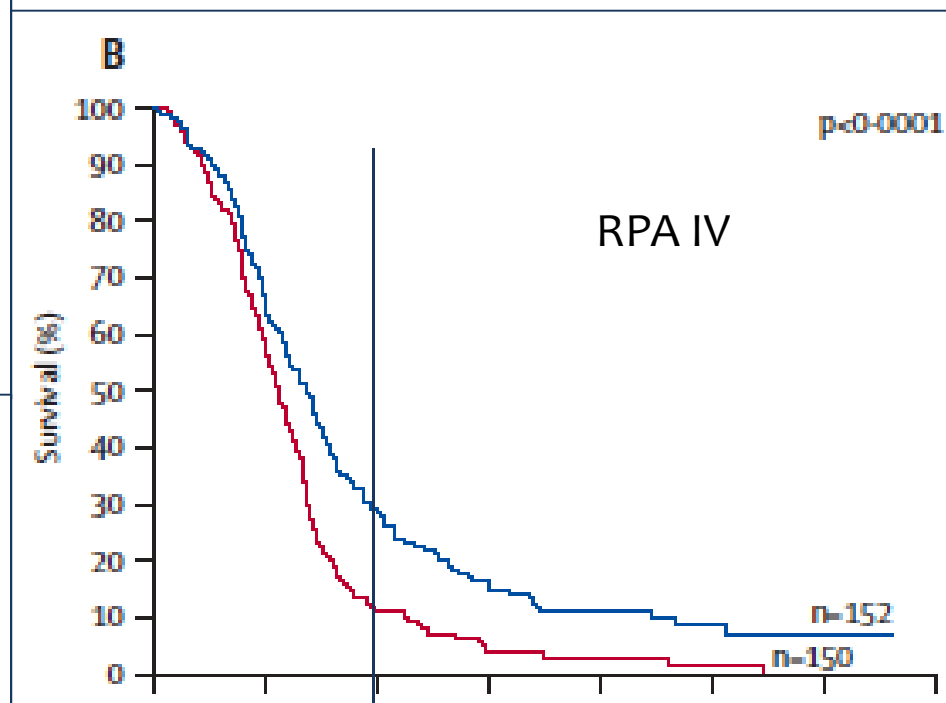
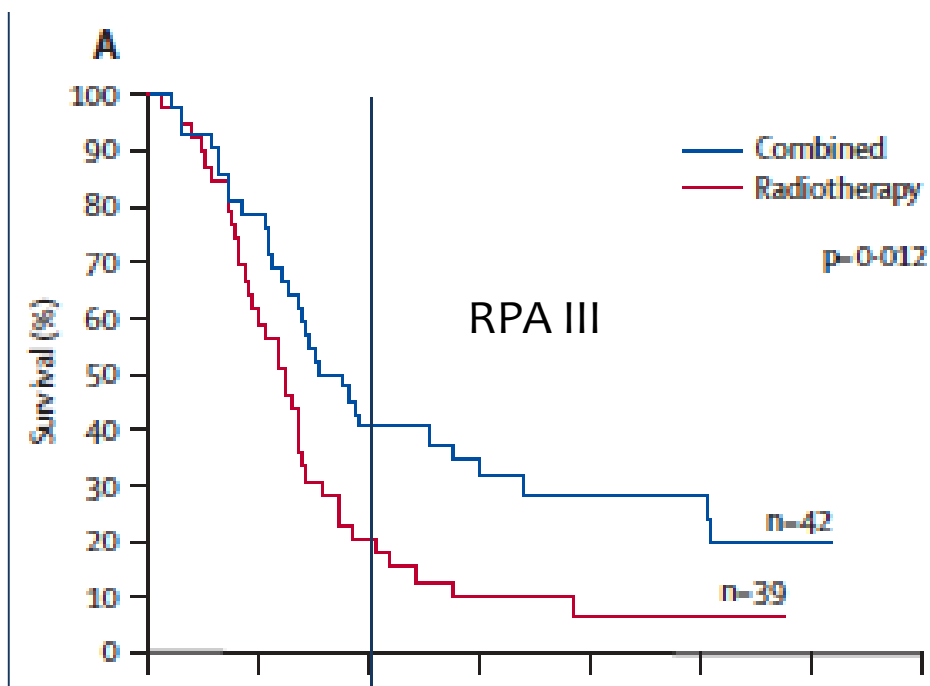


Why?

	Deaths/ patients	Hazard ratio (95% CI)	Median (months; 95% CI)	2 years (%)	3 years (%)	4 years (%)	5 years (%)
Age <50 years							
Radiotherapy	83/88	1.0	13.6 (11.6-15.6)	14.8 (8.3-23.0)	6.5 (2.5-13.1)	4.9 (1.5-11.3)	4.9 (1.5-11.3)
Combined	79/95	0.6 (0.4-0.8)	17.4 (15.3-21.5)	34.7 (25.3-44.3)	25.4 (17.0-34.7)	20.1 (12.4-29.1)	17.0 (9.8-25.9)
Age ≥50 years							
Radiotherapy	195/198	1.0	11.9 (10.6-12.6)	9.1 (5.6-13.7)	3.4 (1.4-6.7)	2.3 (0.8-5.2)	0.7 (0.1-3.5)
Combined	175/192	0.7 (0.5-0.8)	13.6 (11.8-15.1)	23.5 (17.7-29.7)	11.4 (7.3-16.5)	8.2 (4.7-12.9)	6.4 (3.2-11.0)
Age 50-60 years							
Radiotherapy	109/111	1.0	12.0 (10.0-14.2)	11.8 (6.6-18.6)	4.2 (1.5-9.4)	2.1 (0.4-6.6)	1.1 (0.1-5.1)
Combined	101/109	0.7 (0.5-0.9)	14.6 (13.6-17.9)	24.8 (17.1-33.2)	11.0 (6.0-17.7)	8.0 (3.8-14.2)	6.4 (2.6-12.6)
Age >60 years							
Radiotherapy	86/87	1.0	11.8 (10.4-12.7)	5.7 (2.1-12.0)	2.3 (0.4-7.2)	2.3 (0.4-7.3)	0
Combined	74/83	0.7 (0.5-0.97)	10.9 (8.9-14.9)	21.8 (13.5-31.2)	12.3 (6.1-20.8)	8.8 (3.6-16.9)	6.6 (2.1-14.7)

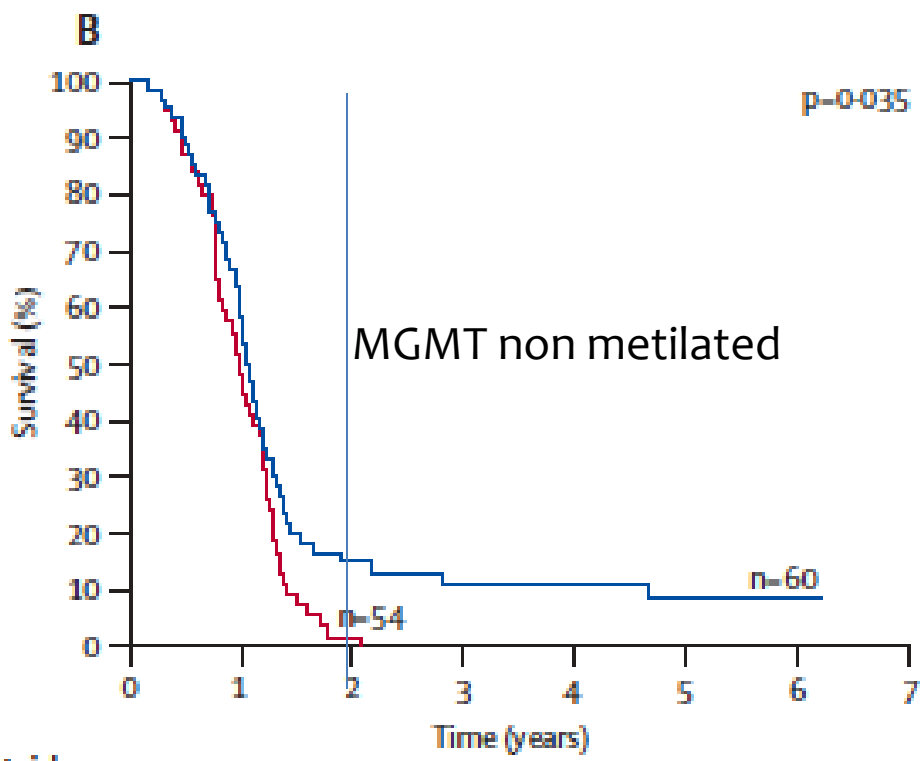
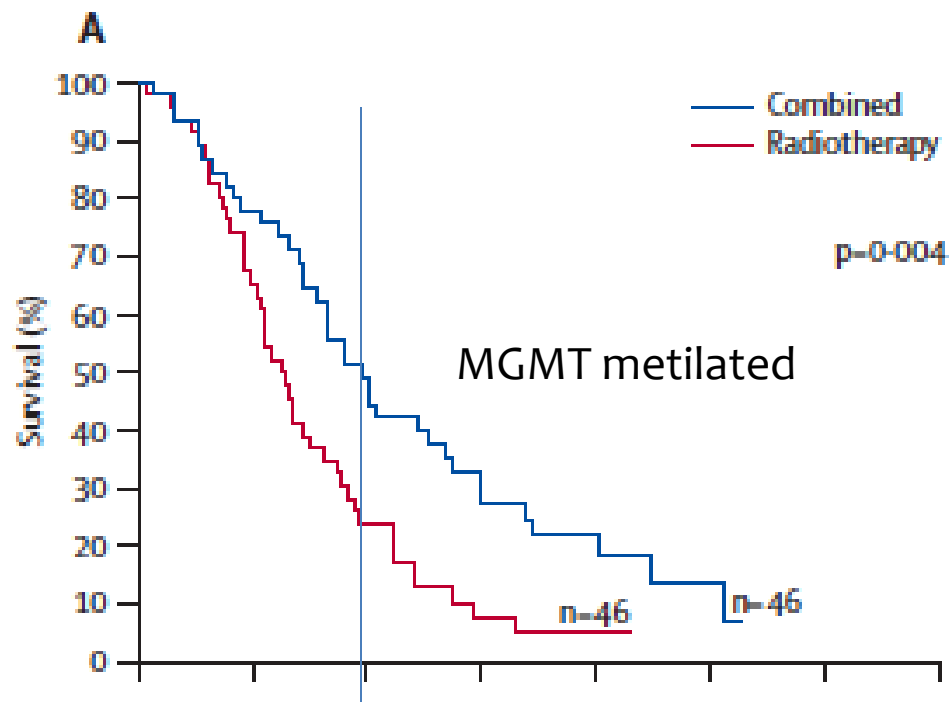


Why?





Why?





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How? Age



The life expectancy of elderly pts with GBL is significantly shorter than in younger pts

How? Age



Age itself?

Why the life expectancy of elderly pts with GBL is significantly shorter than younger pts ?



How? Age

Age itself?

Biological factors?
-Less IDH1 mutation
-Less m-MGMT
-TP53 mutation
-EGFR amplification

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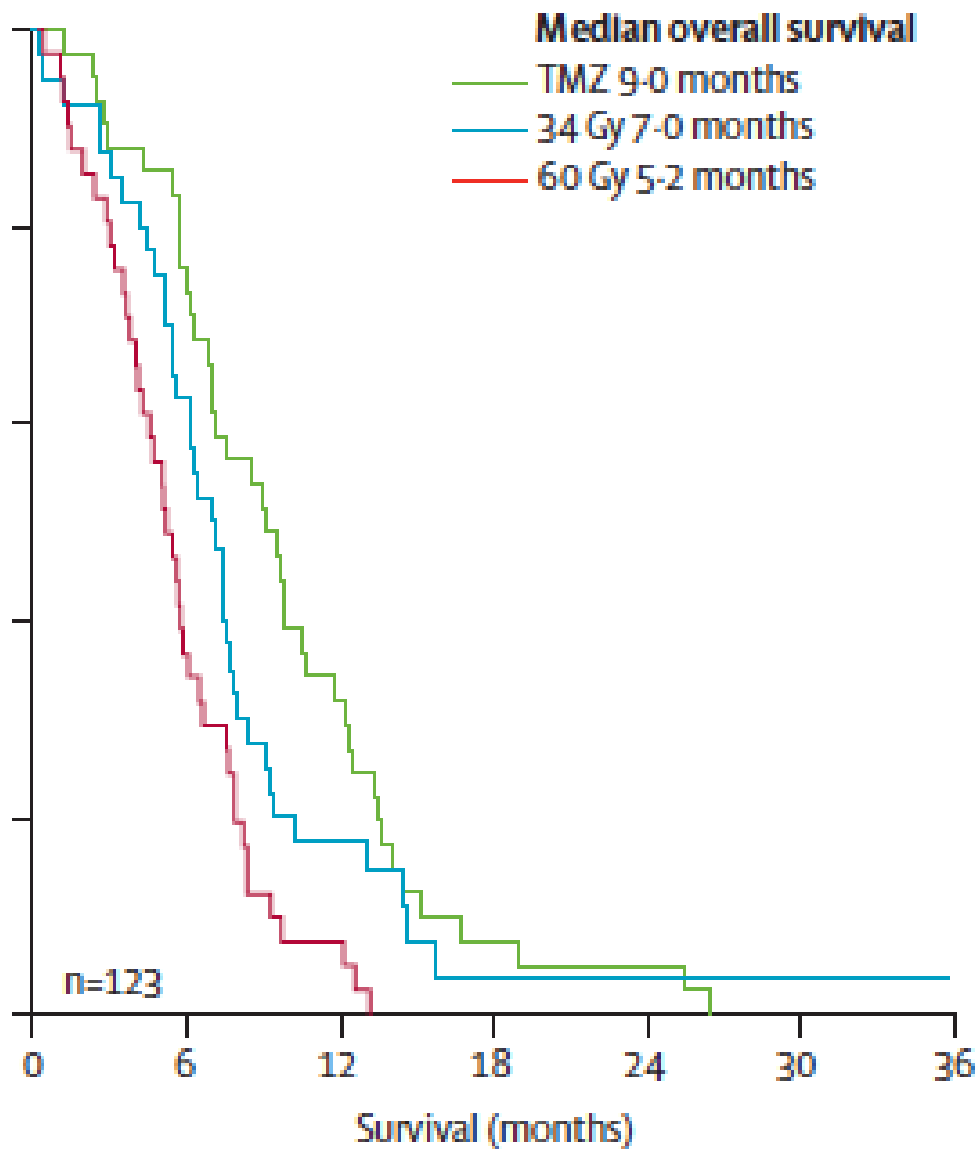
Less therapy?
-Less resection
-Less radiotherapy
-Less cht
Iwamoto FM 2008 Ann Neurol
-More need of
supportive care
-More frequent toxicity

Why the life expectancy of elderly pts with GBL is significantly shorter than younger pts ?



How? Age

Nordic
randomized
phase III
trial



> 70 yy

How? Extent of surgery



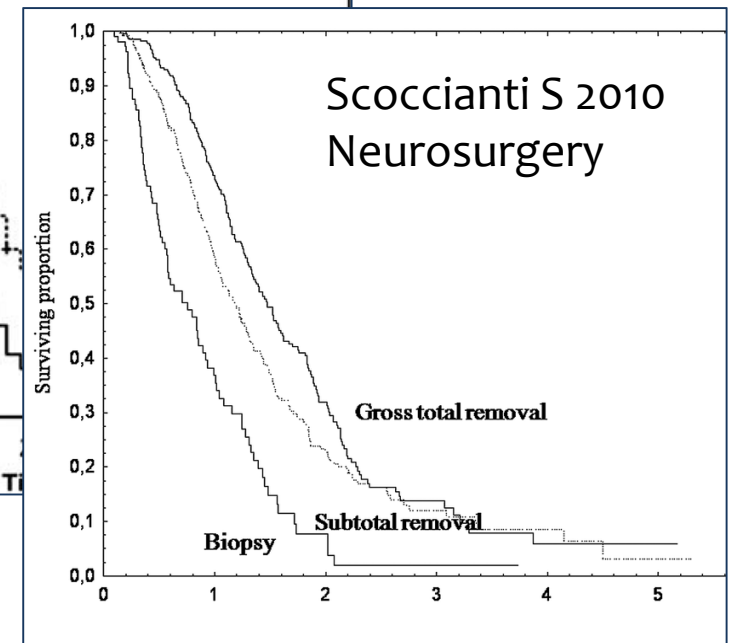
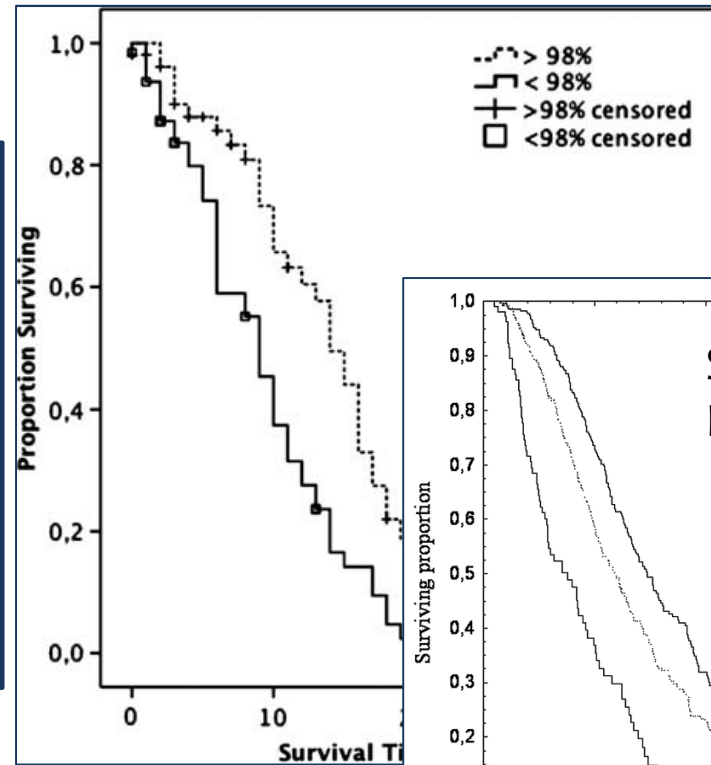
Radical surgery is better than partial surgery or biopsy



How? Extent of surgery

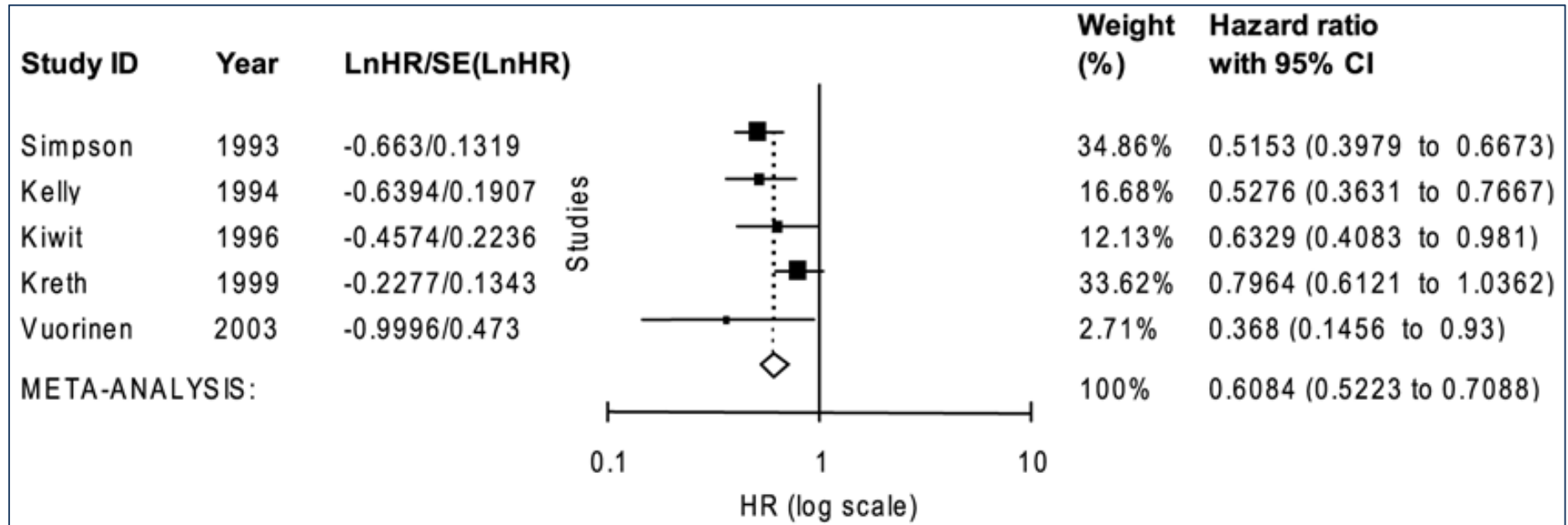
A complete surgical excision of HGG is impossible

Is the extent of resection really an independent factor in predicting survival?



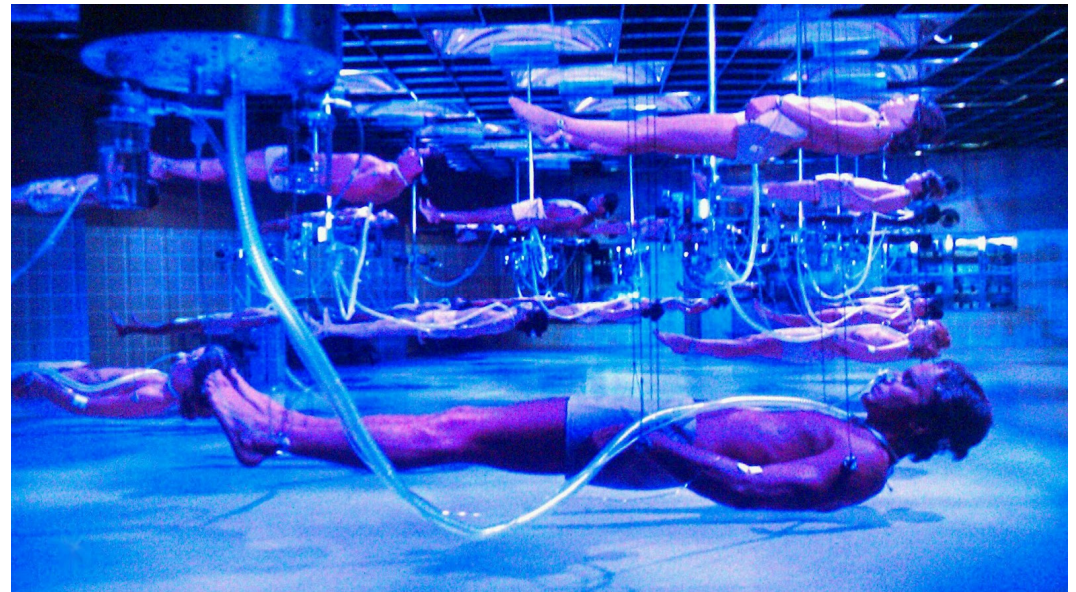
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How? Extent of surgery



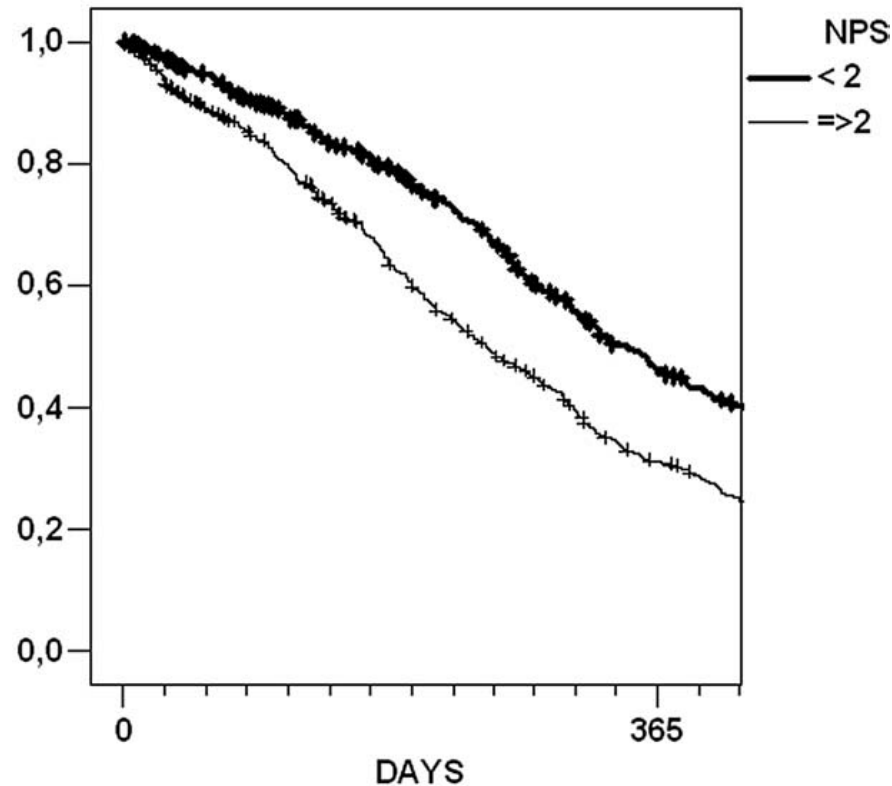
Radical surgery is better than partial surgery or biopsy

How? General and neurological PS



Patients with poor general and neurological performance have a worse prognosis

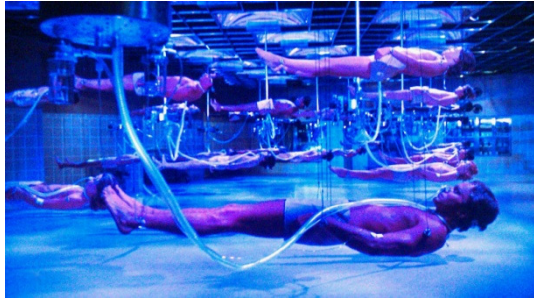
How? General and neurological PS



Patients with poor general and neurological performance have worse prognosis than the others



How? RPA “new version”

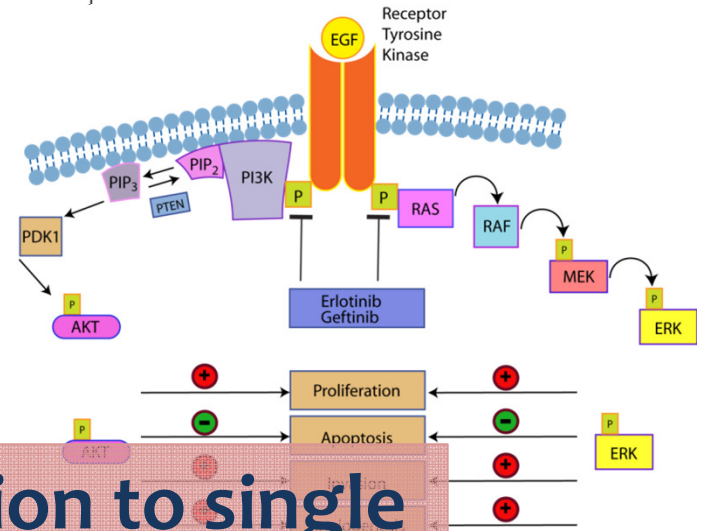
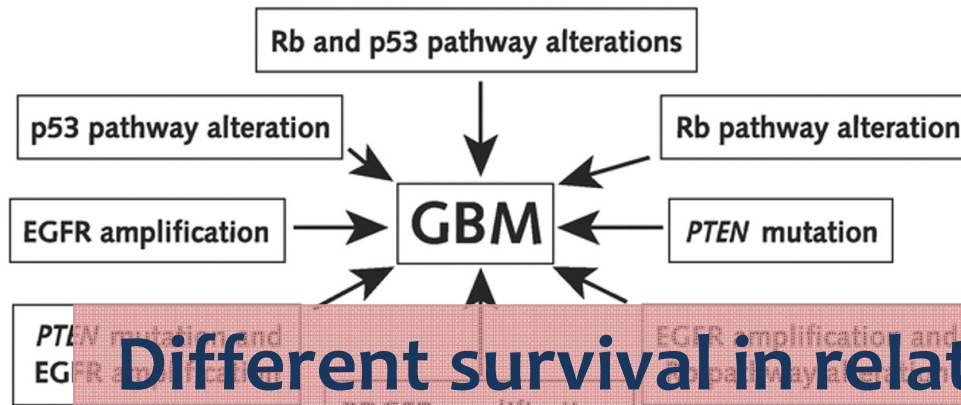
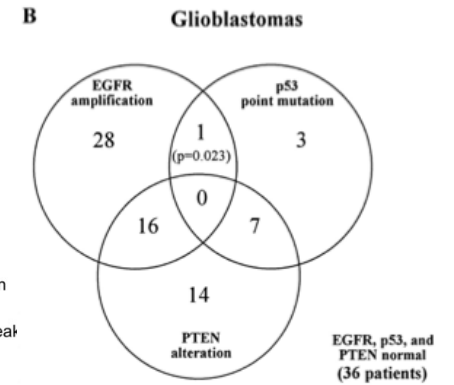
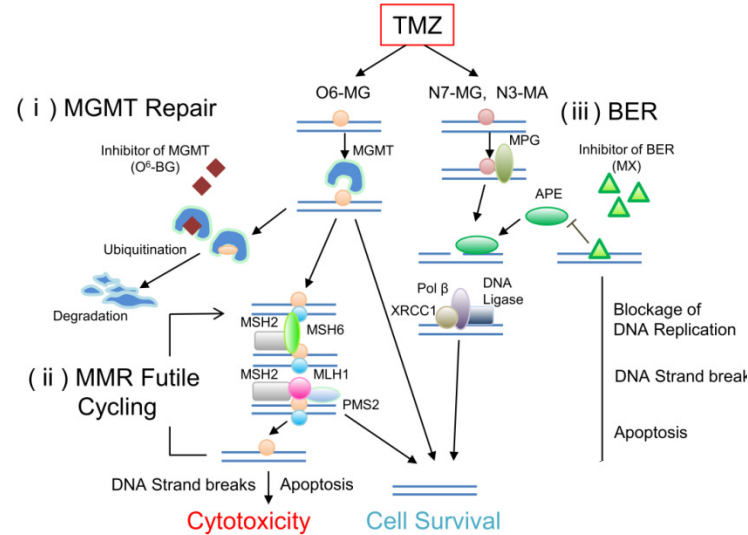
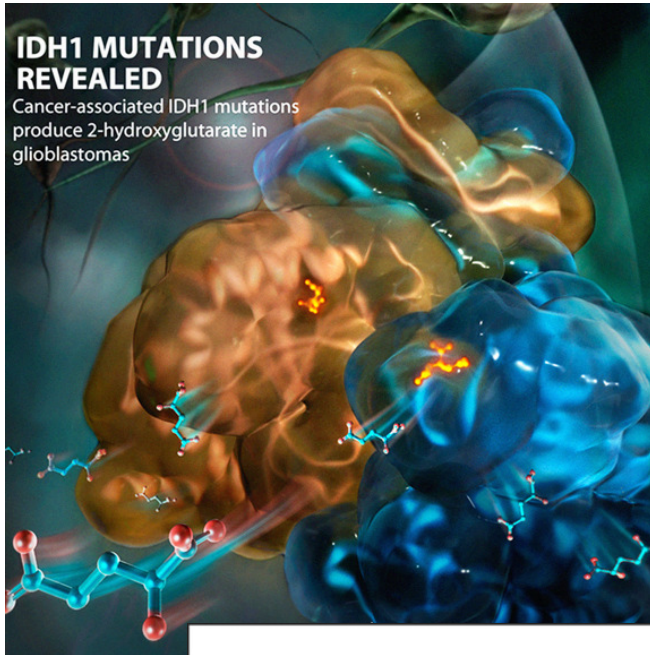


RPA class	Definition variables	Survival (mo)
III	<50 y and KPS \geq 90	17.1
IV	<ul style="list-style-type: none"> <50 y and KPS <90 \geq50 y, KPS \geq70, resection, and working 	11.2
V	<ul style="list-style-type: none"> \geq50 y, KPS \geq70, resection, and not working \geq50 y, KPS \geq70, biopsy only \geq50 y, KPS < 70 	7.5

Three new simplified RPA classes



How? Biological aspects

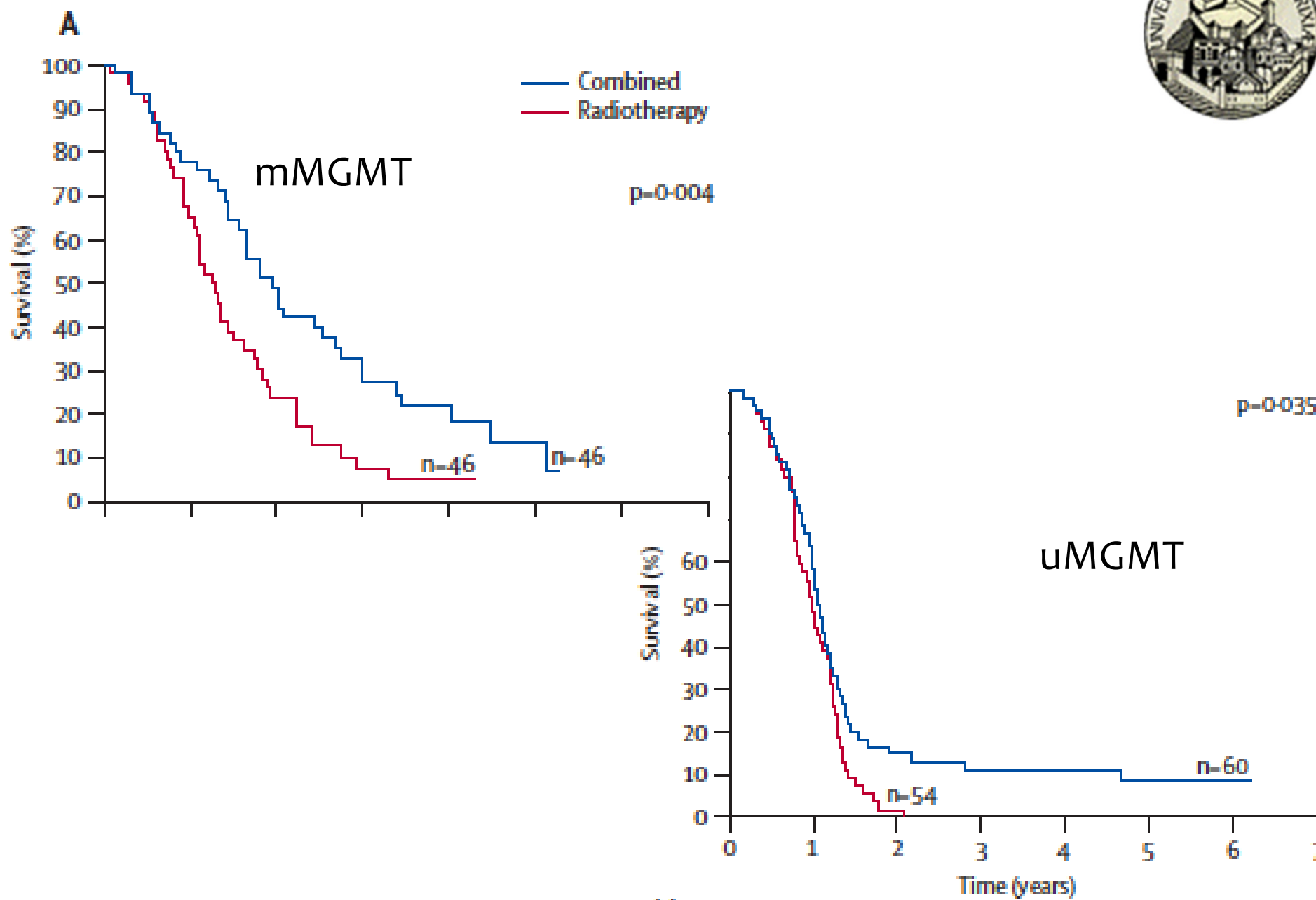


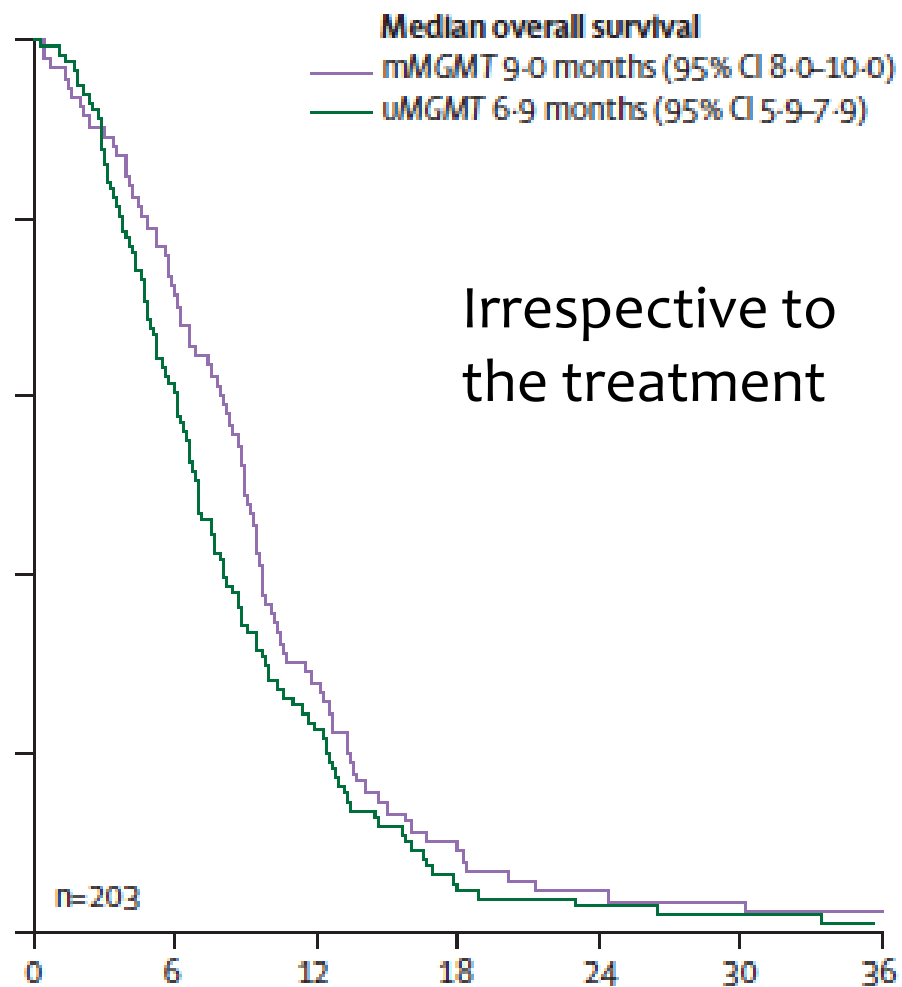
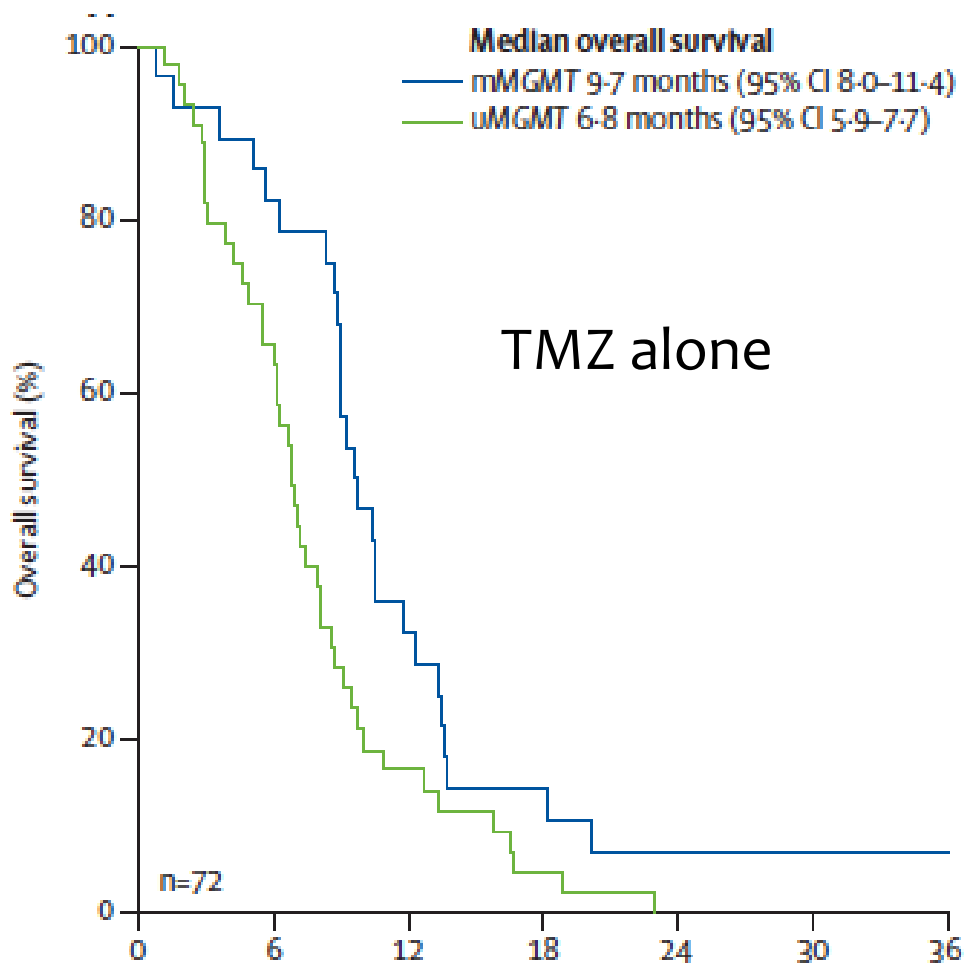
Different survival in relation to single biological factors and their combination

How? Biological aspects “MGMT methylation”



- Biological effect → reduced DNA repair, association with G-CIMP phenotype in IDH1/2 mutant tumours
- Better response to chemotherapy → better OS and PFS
- Problems related to the identification methods have been debated
- None of the present or ongoing trials answering this question: patients with MGMT methylation should be treated with TMZ (alone or concomitant and adjuvant to RT) or not?





How? Biological aspects “IDH mutation”

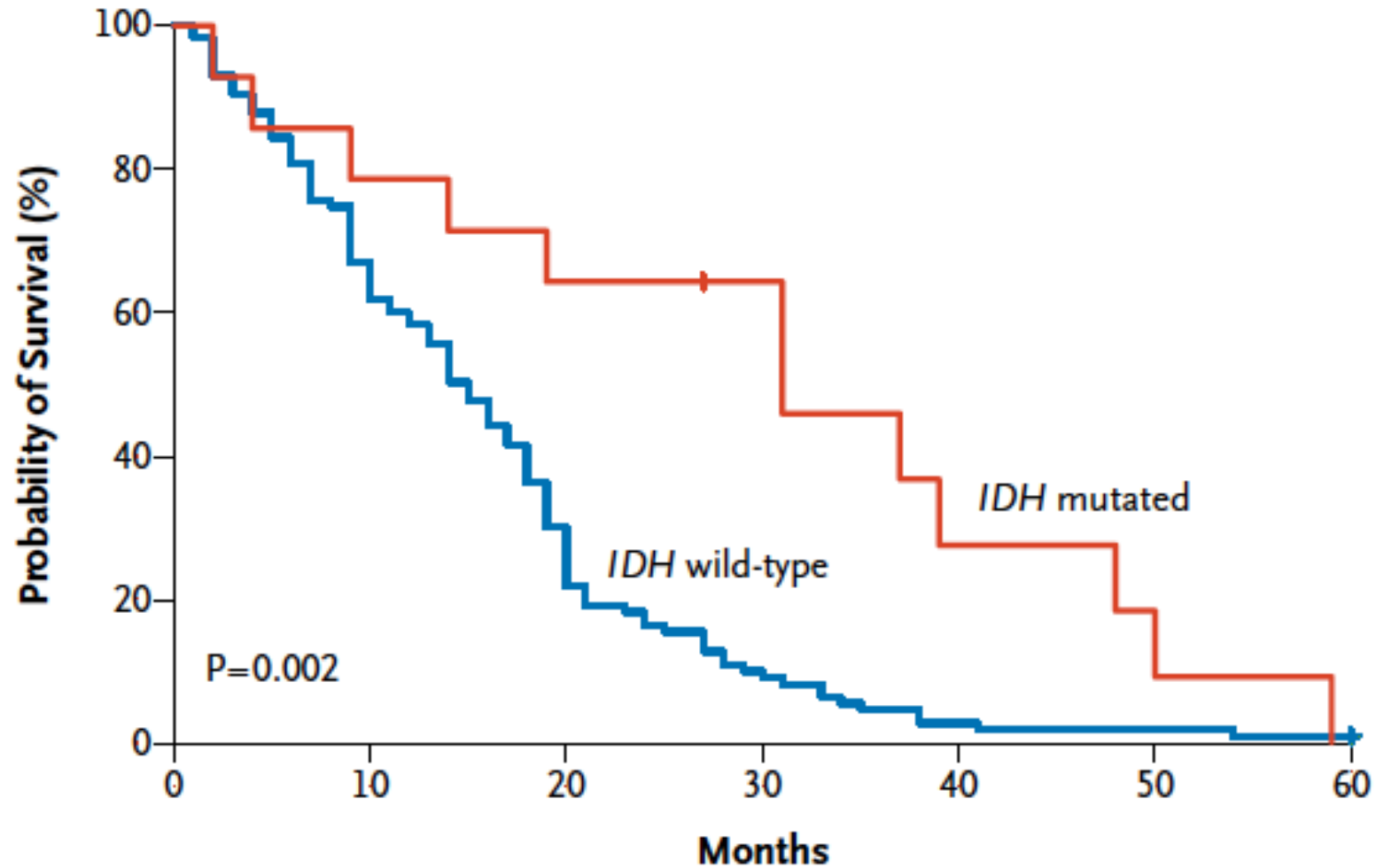


- Biological effect → increased concentrations of 2-hydroxyglutarate, association with G-CIMP phenotype
- Differentiate IDH-wild type vs IDH mutant glioma
worse vs better prognosis
- IDH status could be included in future classification
- IDH-mutant tumours are driven by specific epigenetic alterations, phenotypically characterized by a status (G-CIMP-positive) suitable for specific therapeutic interventions
- **IT HASN'T A DEFINED ROLE IN CLINICAL DECISION MAKING**

How? Biological aspects



A Glioblastoma



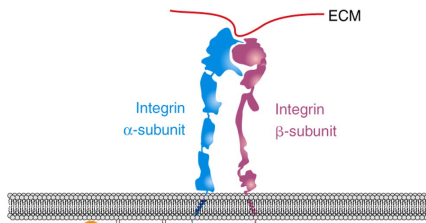
How? Biological aspects “EGFR v III rearrangement”



- Biological effect → deletion of the gene EGFR: results in a constitutive and ligand independent oncogenic mutation
- The mutation can probably be considered a negative prognostic factor (reduces long term survival)
- Target treatments against EGFR are not effective
- EGFR v III mutation is an immunogenic factor that could possibly be used as target for “*vaccination*”
- EGFRvIII mRNA has been detected in the serum of patients with EGFR vIII positive-glioblastoma → it could be useful to monitor response to therapy and to detect relapse

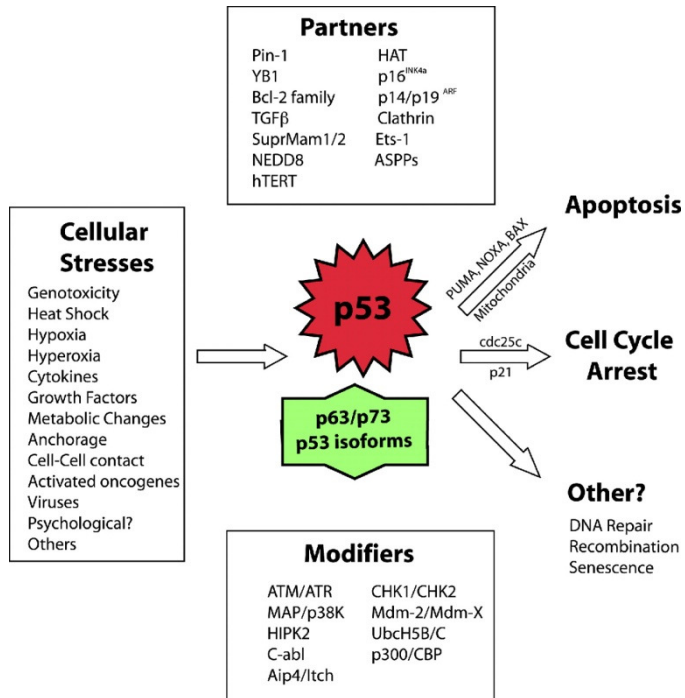


How? Biological aspects “integrins”



- Integrins → cell adhesion molecules involved in glioma cell migration/invasion and angiogenesis
- Any integrin, if overexpressed, is involved in multi-drug resistant glioma cells and is responsible for their increased adhesive and invasive capacities.
- Others are up-regulated on the endothelium cells during tumour angiogenesis and are rapidly accessible in tumour blood vessels; they stimulate endothelium cells proliferation, migration and lumen formation
- Elevated levels of integrins were found in glioma stem cells

How? Biological aspects “p53 in glioma”



- P53 expression is related with p53 mutation;
- Problems are evident regard the prognostic value of p53 in GBL ; it is not validated as independent prognostic factor
- p53 is involved in regulation of neural stem cells → its alteration can increase loss of cell differentiation and increase in neurosphere renewal
- conflicting results are evident about the relationship between p53 and response to TMZ



How? Biological aspects “gene profile”

- Determination of gene expression profile derived from classic tumour samples \longleftrightarrow clinical outcome
- The HOX signature and EGFR expression \longrightarrow independent negative prognostic factors
- The functional association of HOX gene signature with glioblastoma stem cells have been confirmed and the negative prognostic effect was confirmed

Murat A 2008 JCO

Gallo M 2013 Cancer Res

- A new classification of GBL based on supervised gene expression profiling, guided by patients outcome:
 - a) pro-neural
 - b) proliferative
 - c) mesenchymal

glioblastoma

Phillips HS 2006 Cancer Cell
Verhaak RGV 2010 Cancer Cell



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- a) pro-neural
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- c) classic
- d) mesenchymal



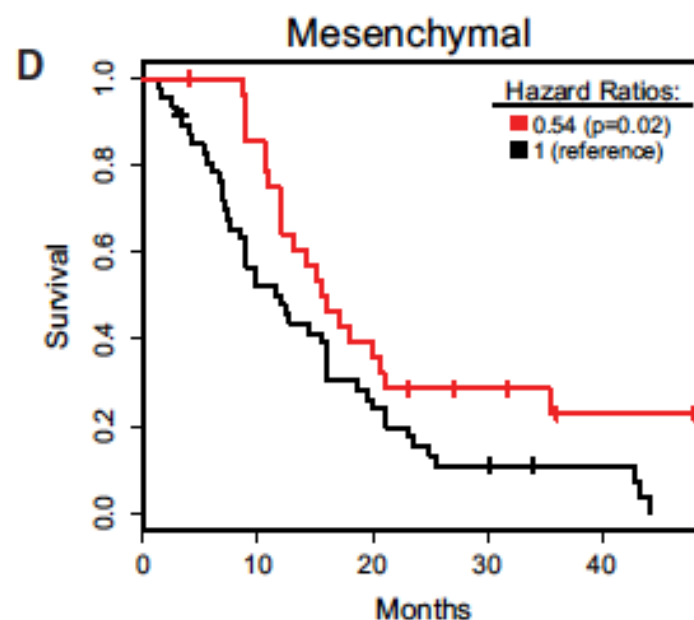
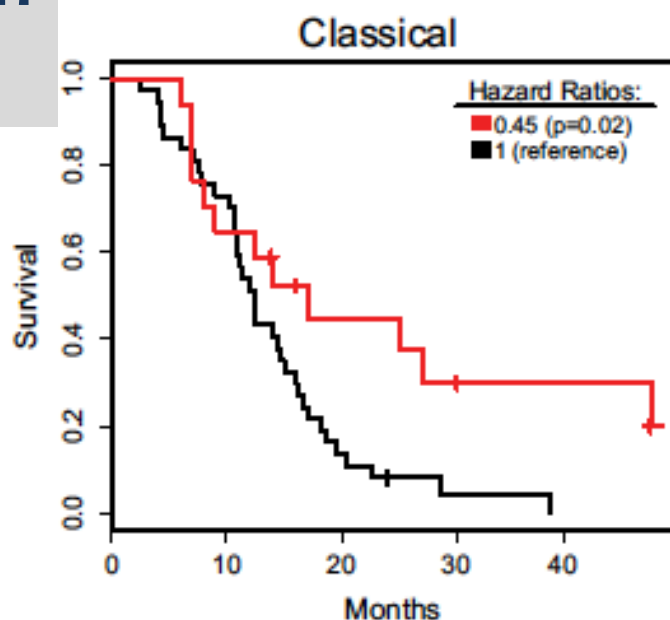
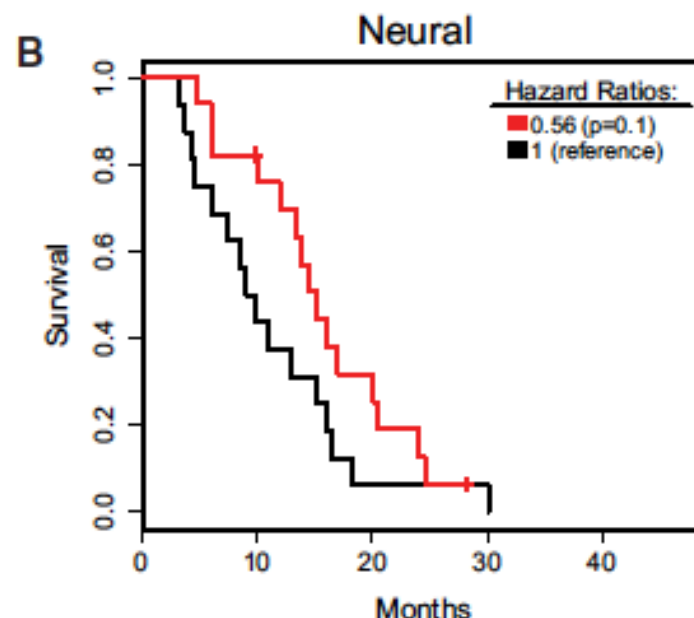
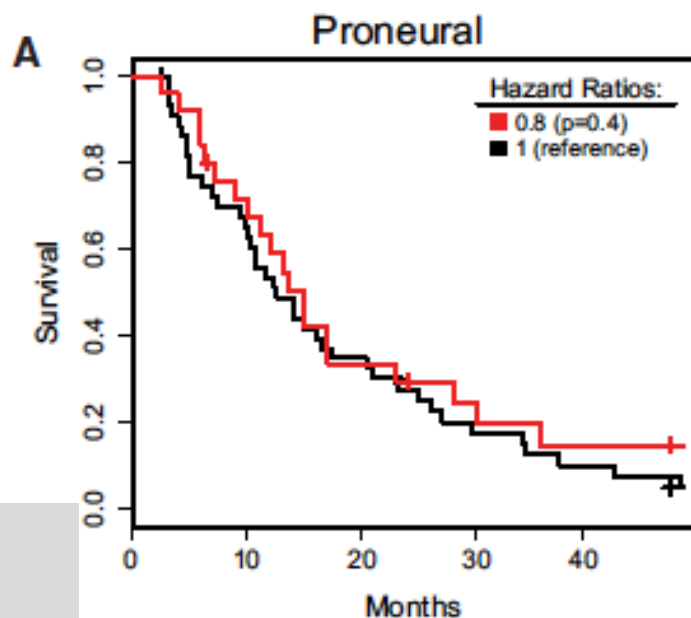
glioblastoma

Phillips HS 2006 Cancer Cell

Verhaak RGV 2010 Cancer Cell



Gene expression profile



■ More intensive therapy: concurrent chemotherapy/radiation and/or >3 cycles of chemotherapy
■ Less intensive therapy: non-concurrent chemotherapy/radiation or <4 cycles of chemotherapy



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Which choices?



- Combined treatment



- Single treatment
 - a. Radiotherapy alone
 - b. Chemotherapy alone
 - c. Others ?

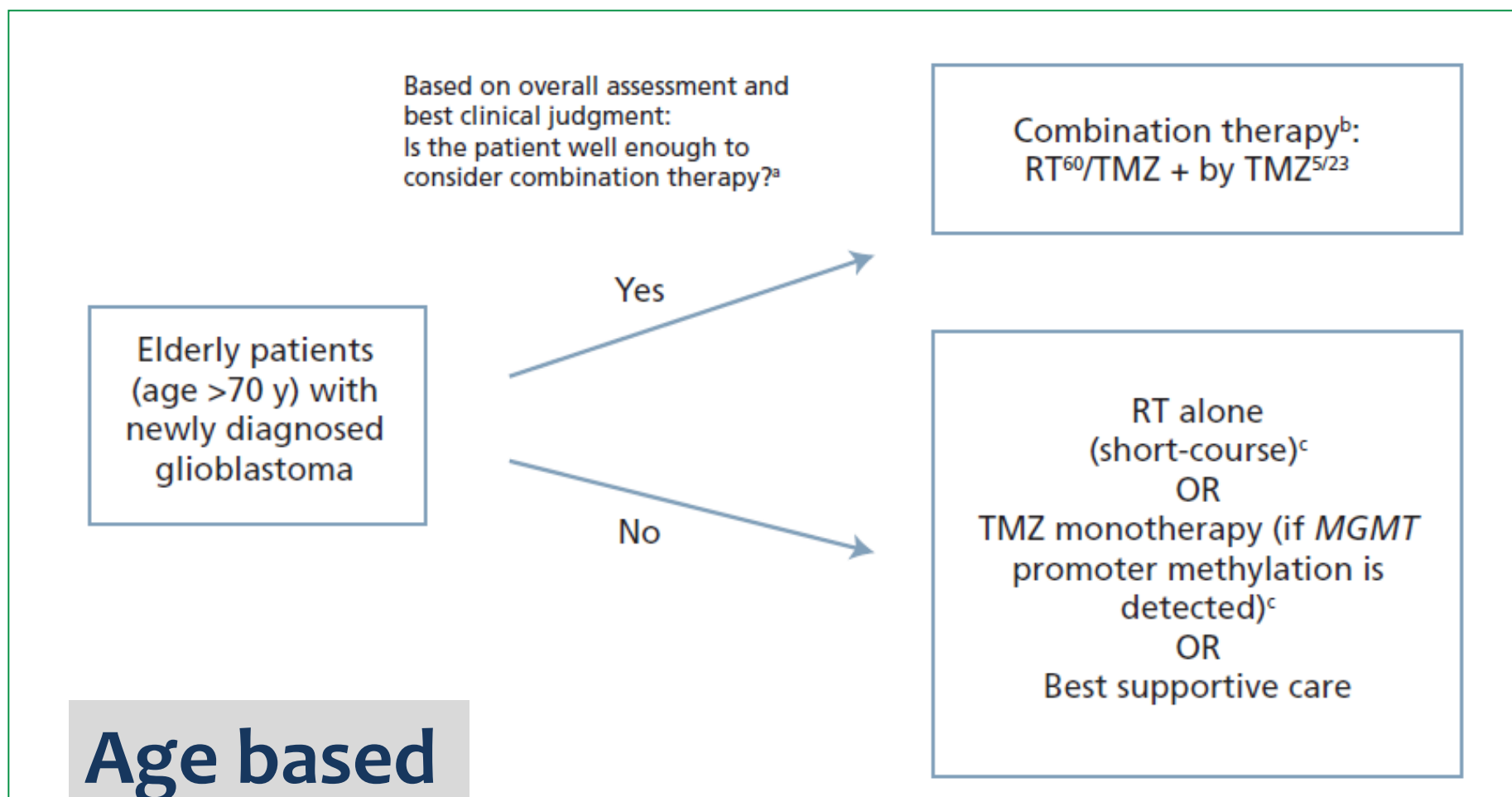


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The choice of choosing

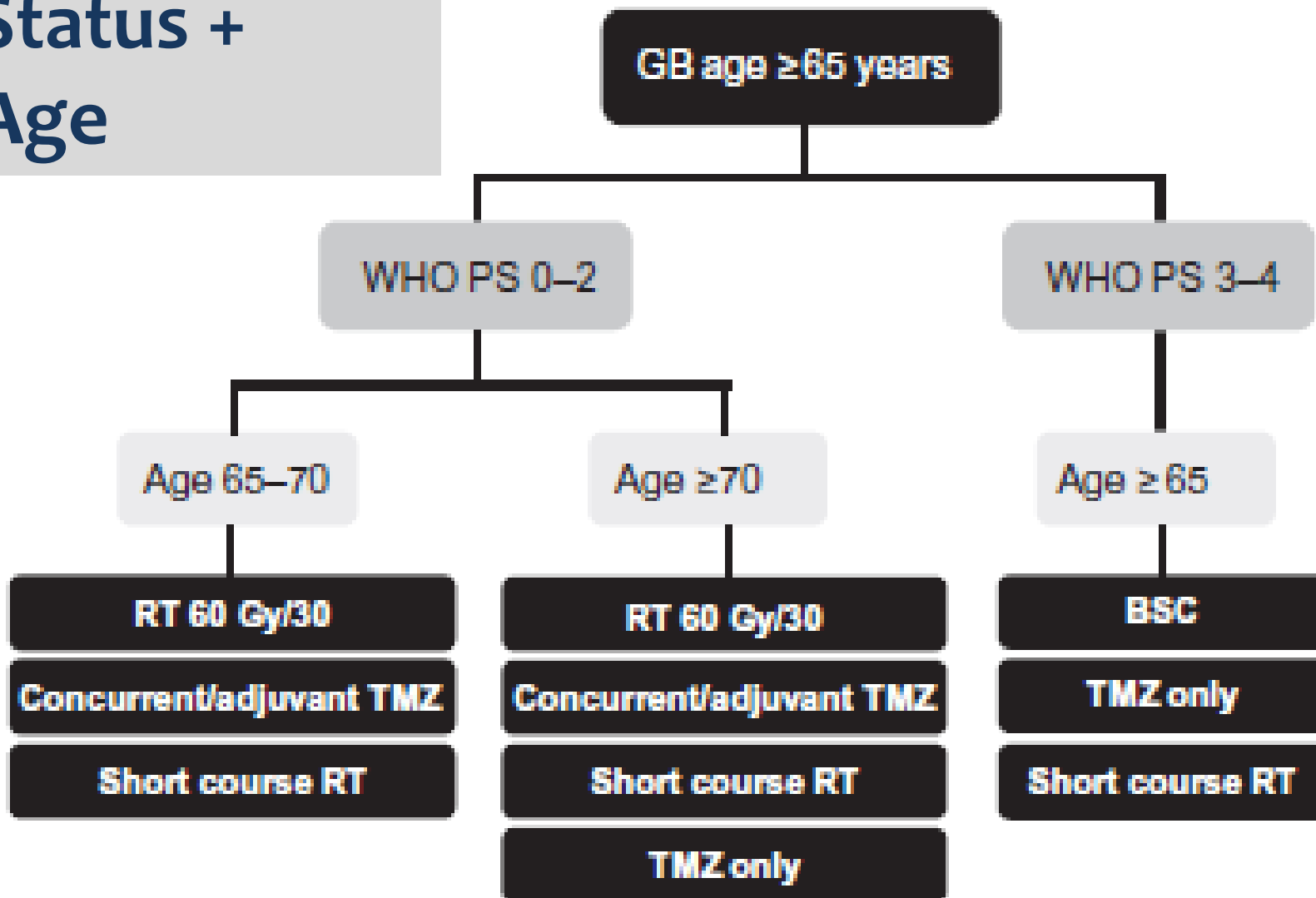


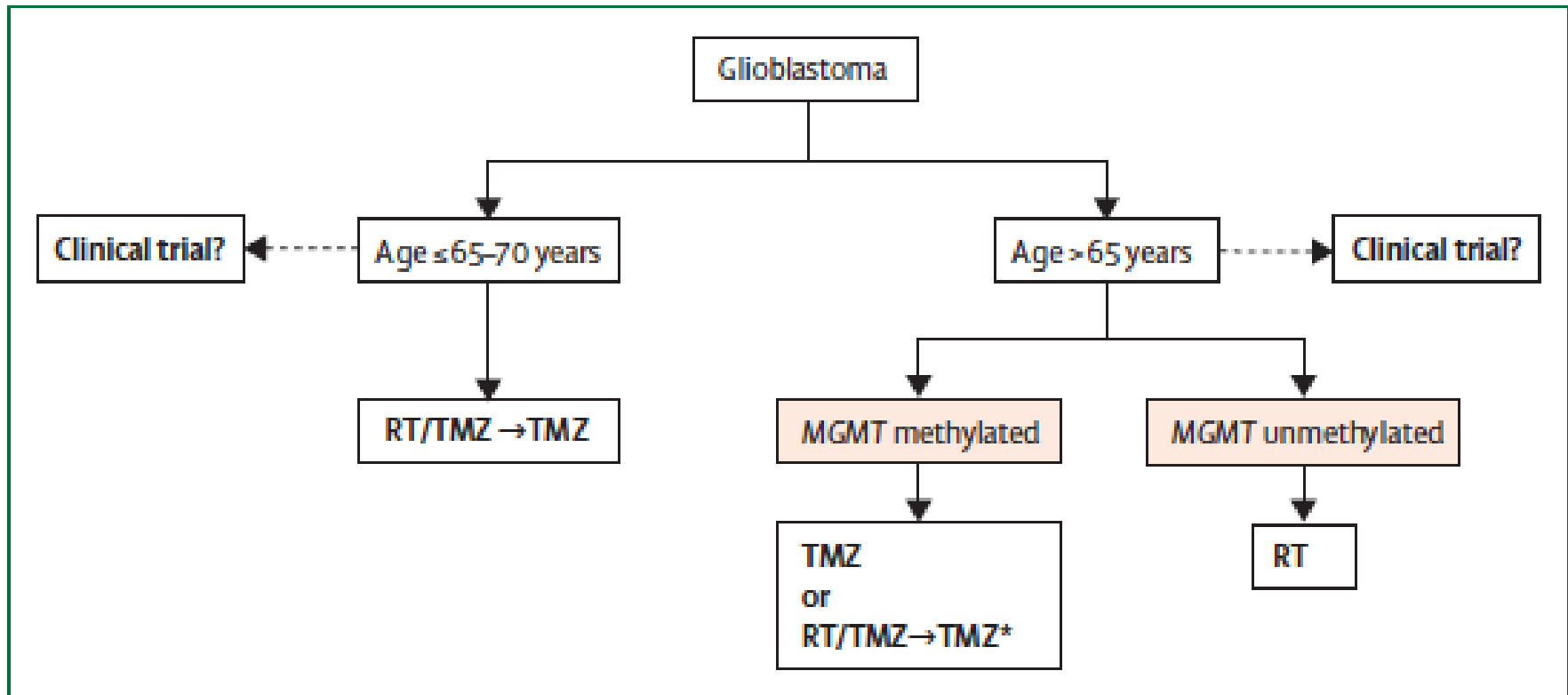


Age based approach

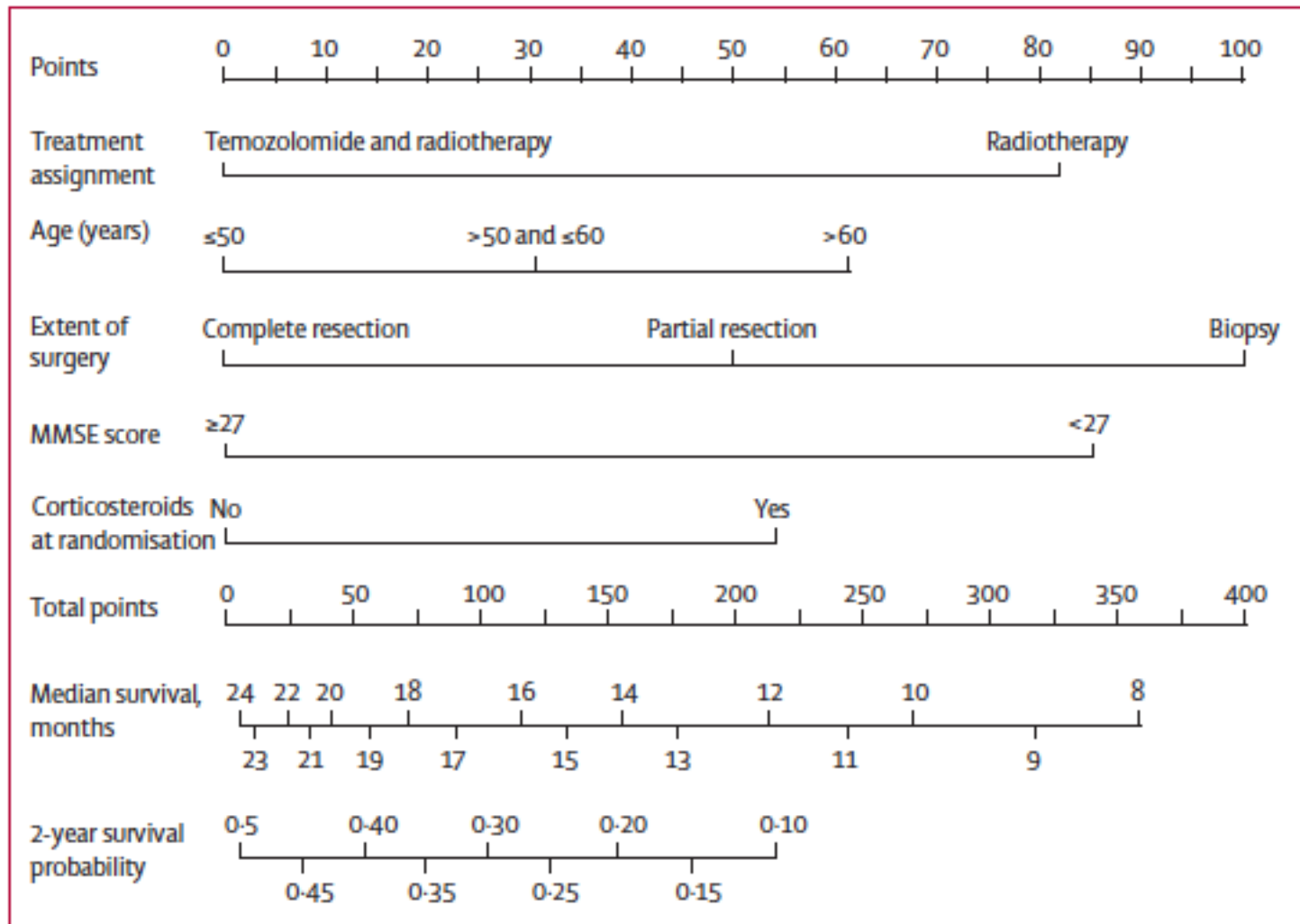


Performance Status + Age

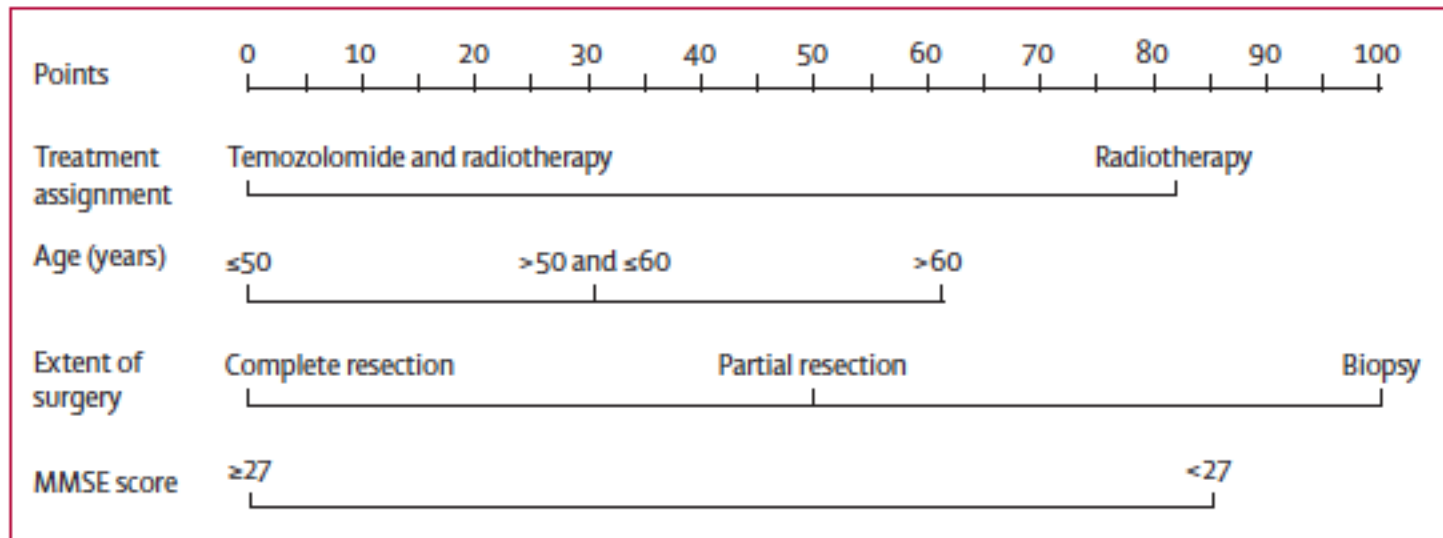




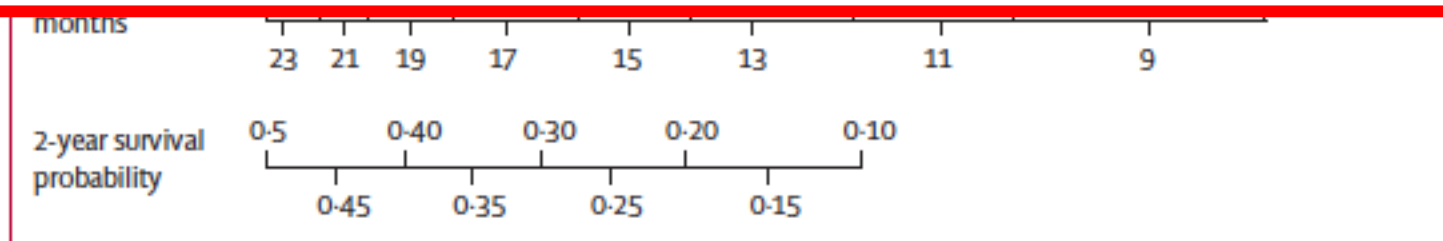
Biomarker based approach



Nomogram based approach



Conclusions. The authors would not recommend the use of this tool in patient counseling.



**Nomogram
based approach**

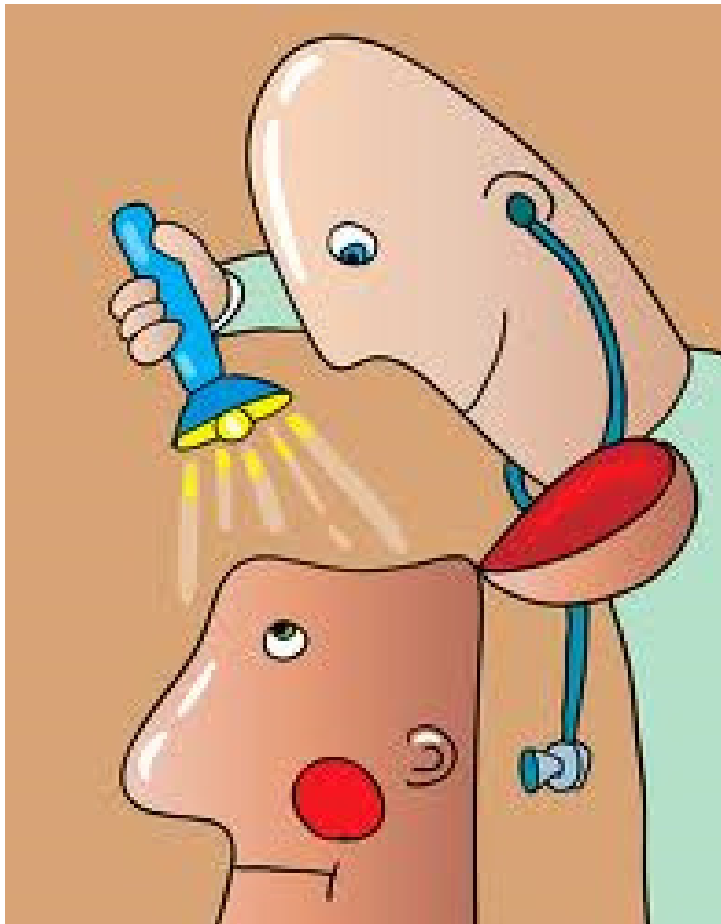


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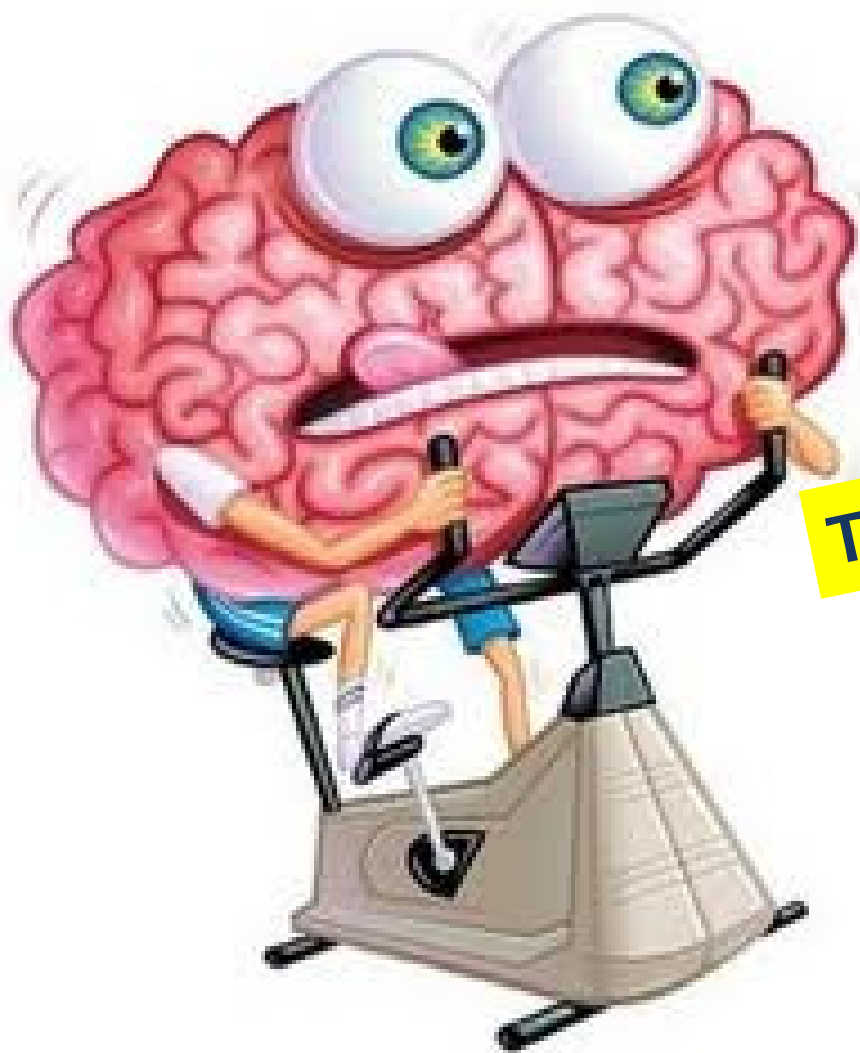


- for the **majority** of our patients
- mostly dependent on **performance status**
- for different reasons for the different patients
- of course, this is only a temporary choice...



Continuing to study the problem:

- **New biological target**
- **New target/non target therapies**
- **New integrations**



Thank you for your attention