

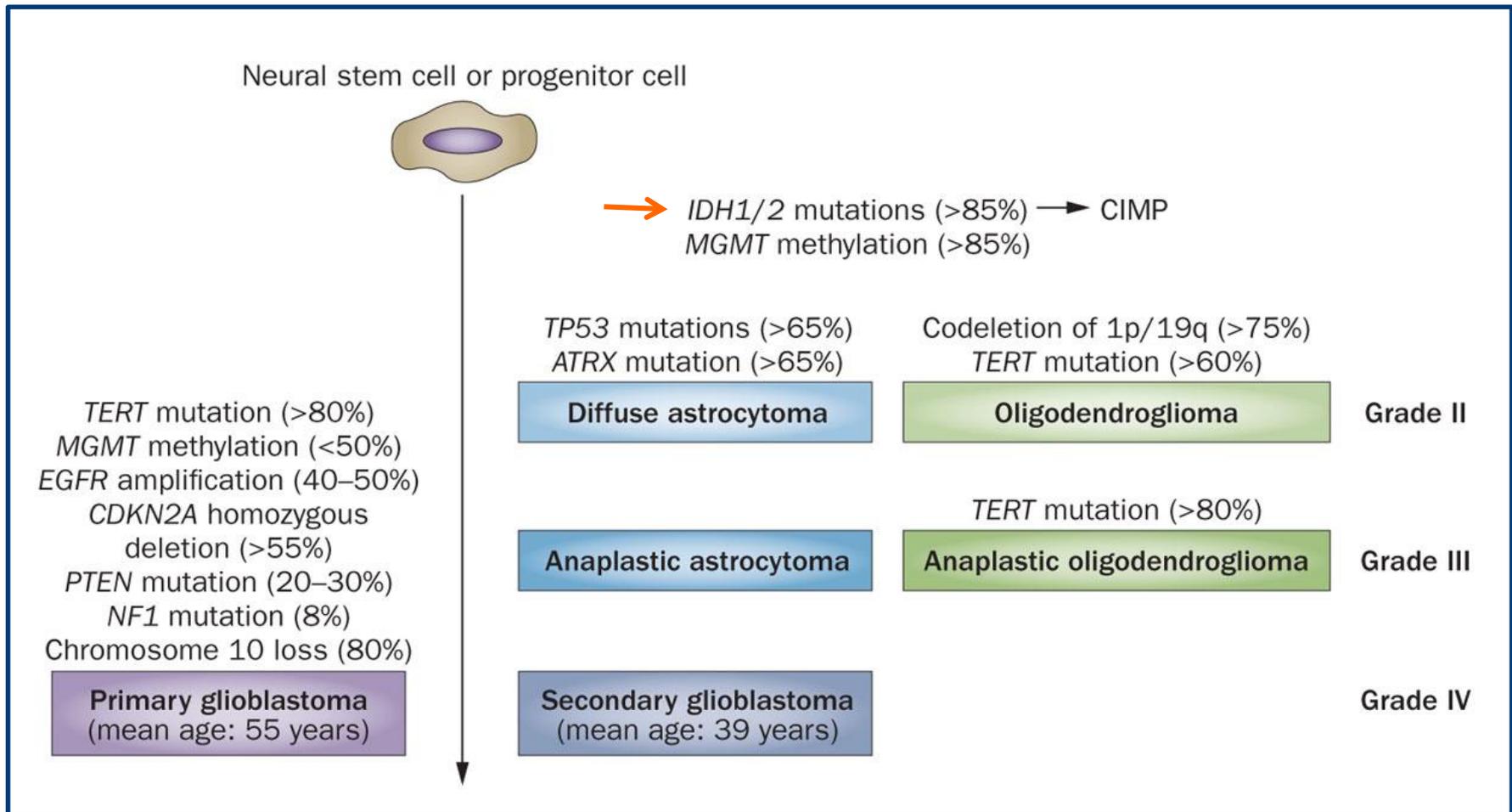
Lessons from translational research in EORTC glioma trials

Monika Hegi

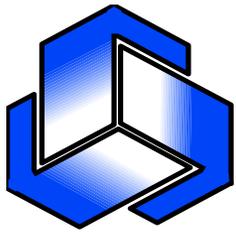
**Laboratory of Brain Tumor Biology and Genetics
Neurosurgery and Neuroscience Research Center
Department of Clinical Neurosciences
Lausanne University Hospital (CHUV), Lausanne, Switzerland**

ICTR-PHE , CIGG Geneva, February, 2016

Pathogenetic and Epigenetic Evolution of Glioma in Adults



Hegi, M. E. & Stupp, R. (2013) *Nat. Rev. Neurol.* doi:10.1038/nrneurol.2013.127; PMID 23817351



Predictive Factors for Benefit from the Addition of Temozolomide to RT EORTC-NCIC Trial

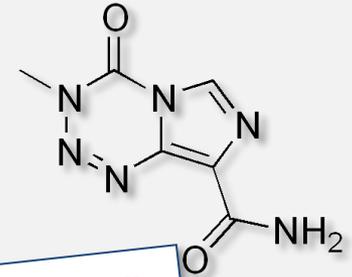


EORTC26981/22981 NCIC CE.3

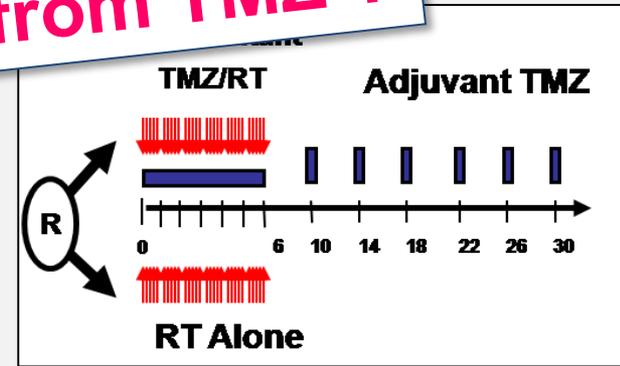
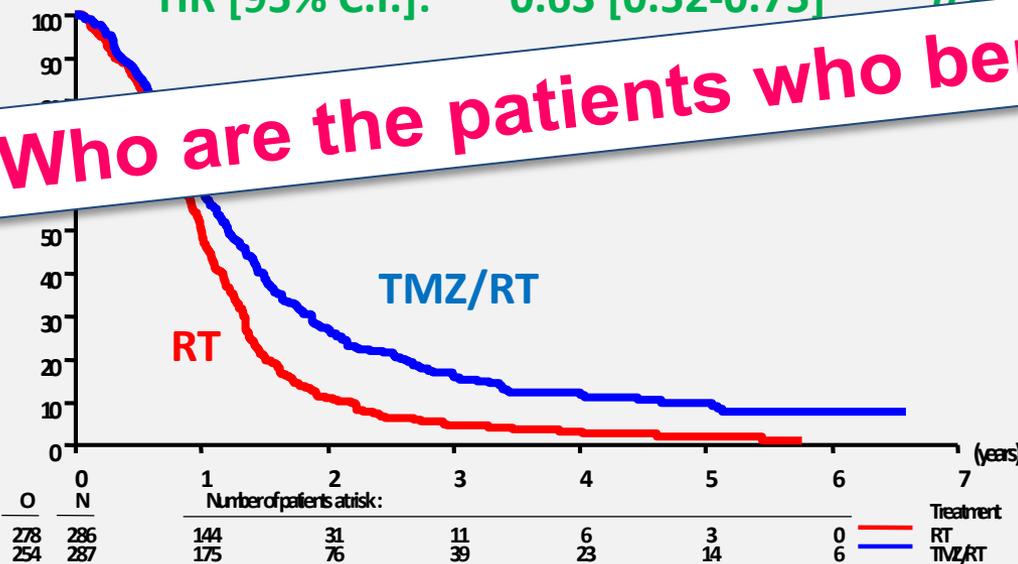
	<u>RT</u> (95% ci)	<u>TMZ/RT</u> (95% ci)
Median OS, mo:	12.1 (11.2, 13.0)	14.6 (13.2, 16.8)
2-yr survival:	10.9% (7.6-14.8)	27.2% (22.2-32.5)
5-yr survival:	1.9% (0.6-4.4)	9.8% (6.4-14.0)

HR [95% C.I.]: 0.63 [0.52-0.75]

$p < 0.0001$



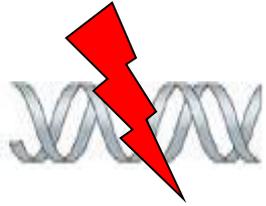
Who are the patients who benefit from TMZ ?



Stupp *et al.* N Engl J Med, 352:987-996, 2005
Stupp *et al.* Lancet Oncol, 10:559-66, 2009

Silencing of *MGMT* by promoter methylation renders tumor cells sensitive to alkylating agents

TREATMENT

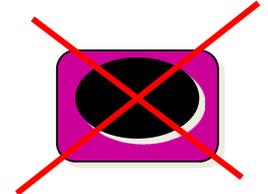
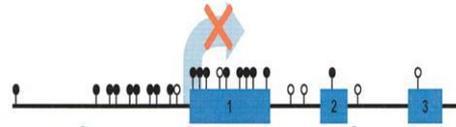


- Alkylating Agent
Temozolomide



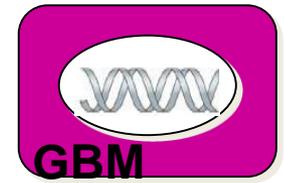
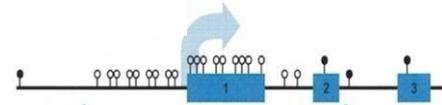
TUMOR TISSUE

Methylated *MGMT*



DNA damage
cell death via MMR

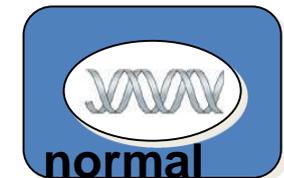
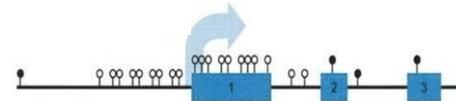
Unmethylated *MGMT*



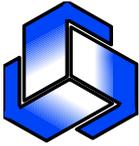
GBM
resistance

NORMAL TISSUE

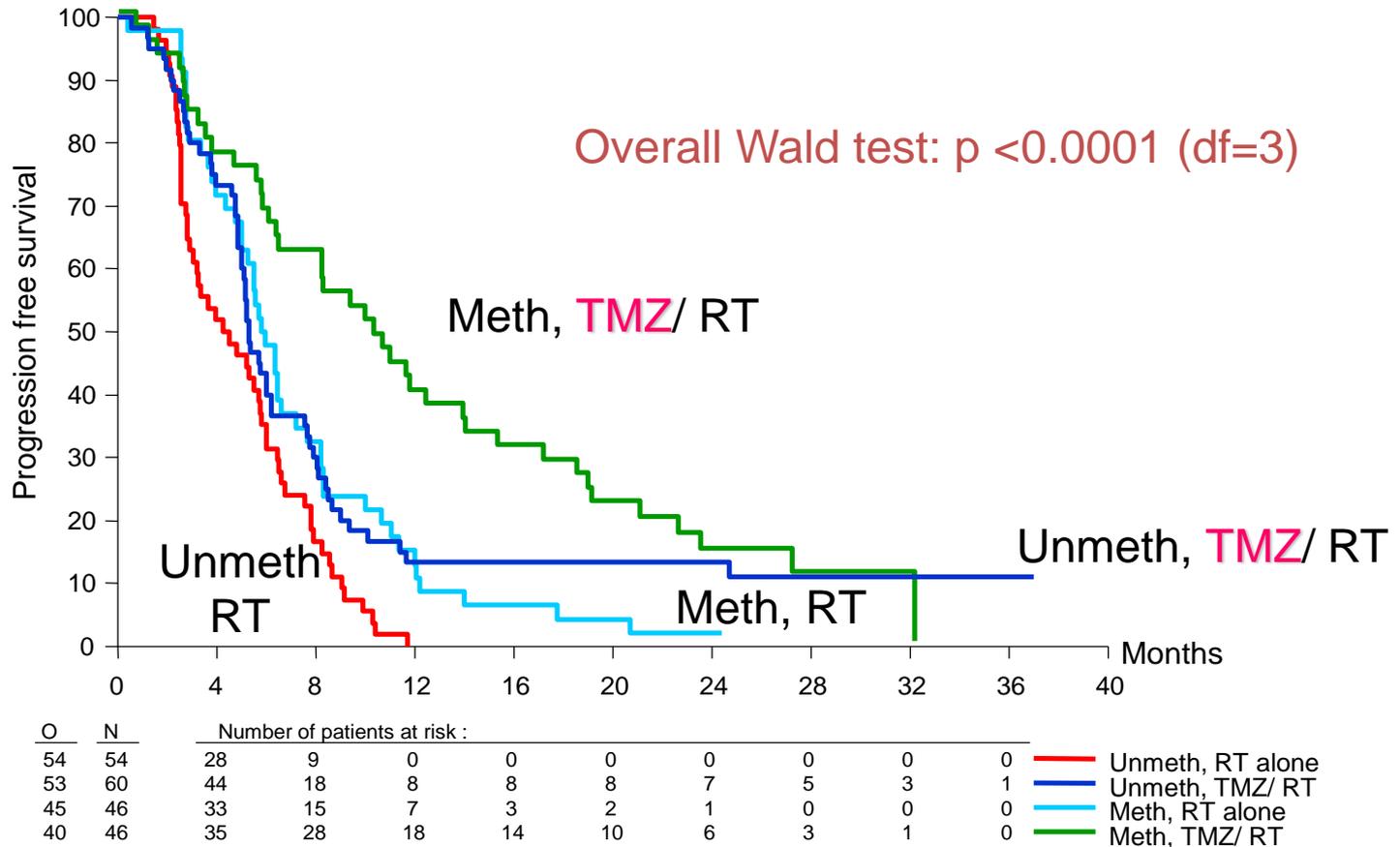
Unmethylated



normal
protected

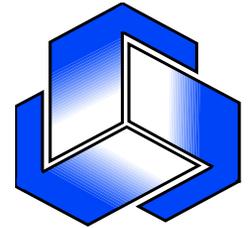


Progression-free survival supports MGMT methylation status as predictive factor for benefit from TMZ

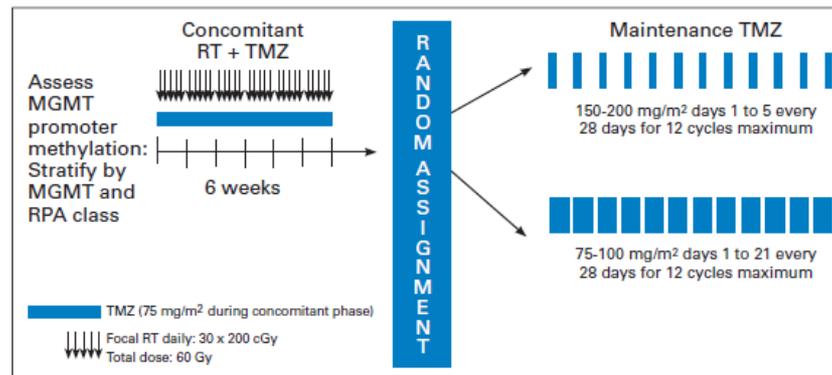
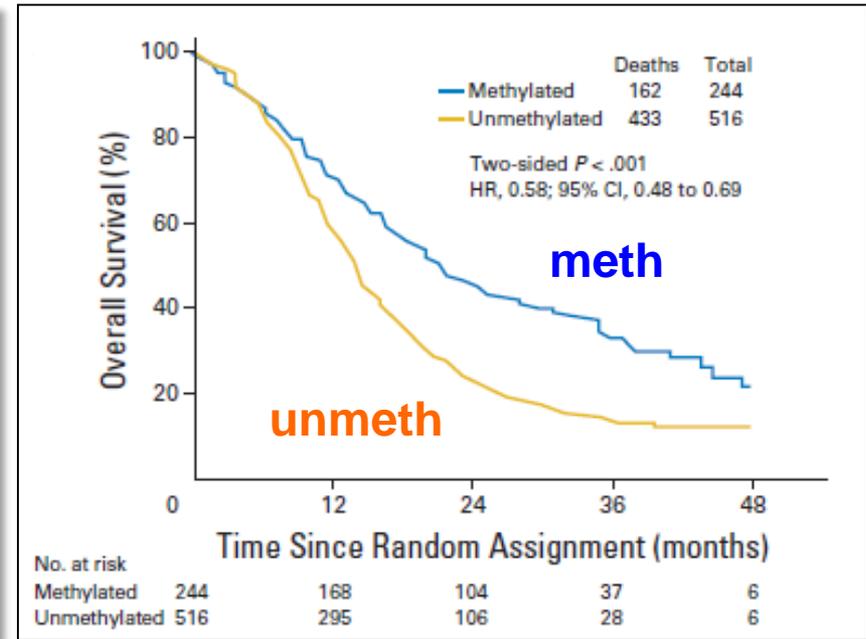
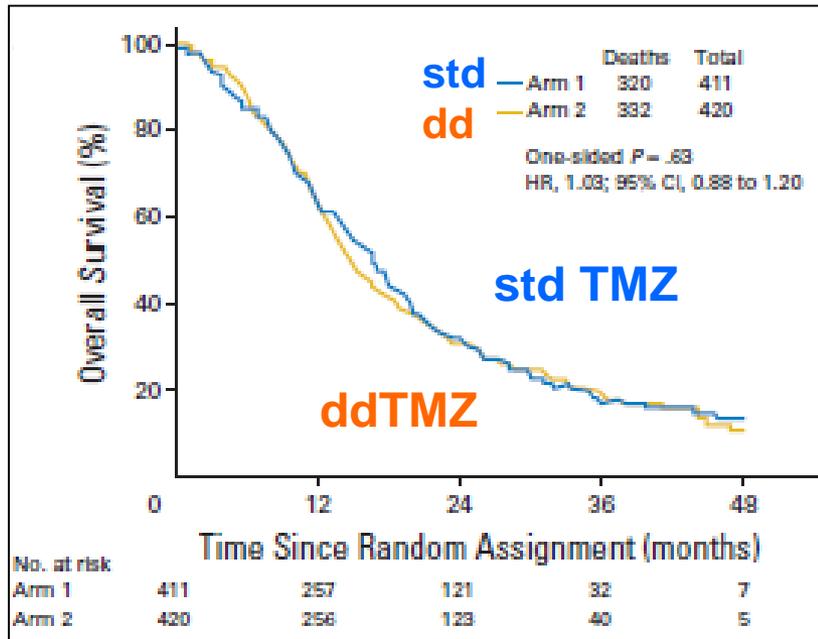




Testing dose dense TMZ in GBM



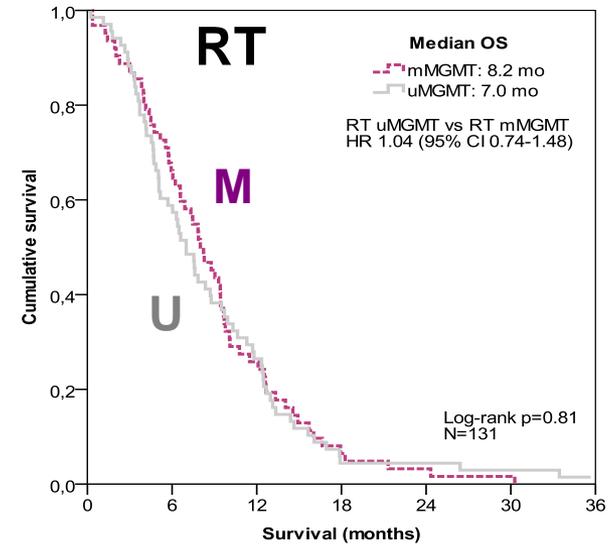
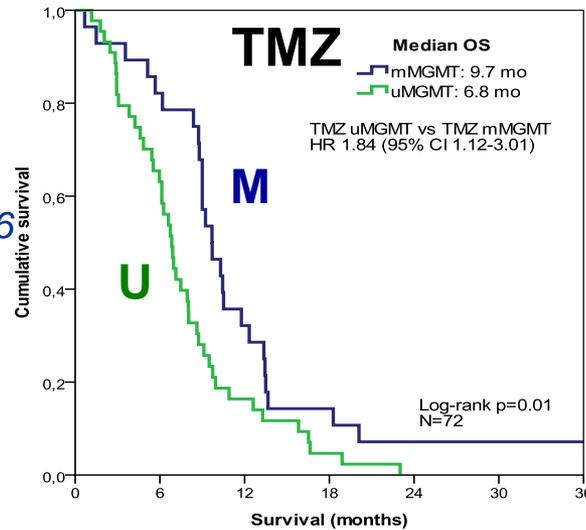
Prospective Validation of *MGMT* Methylation Status as Biomarker



Predictive Effect of *MGMT* in Elderly GBM Patients

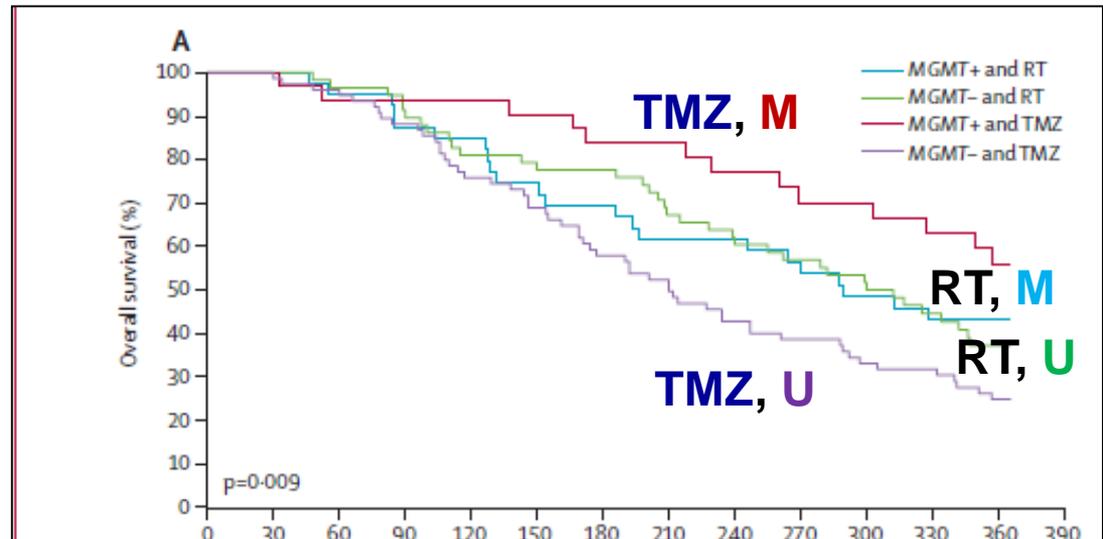
Nordic Trial RT vs TMZ

Malmström et al.
Lancet Oncol **2012**; 13: 916-926



NOA-08 RT vs TMZ

Wick et al.
Lancet Oncol **2012**; 13(7):707-15



***MGMT* methylation is a predictive factor in GBM**

- The presence of a methylated *MGMT* promoter is associated with benefit from alkylating agent therapy
- Patients with an unmethylated *MGMT* promoter profit very little from TMZ, if at all.
- Elderly patients should be treated according to their *MGMT* status
- Patients with unmethylated *MGMT* should be recruited into trials with new promising drugs
- **Terms:** predictive versus prognostic
 - **predictive**, treatment dependent ; impact on choice of therapy
 - **prognostic**, independent of treatment; no direct impact on treatment

Patient Selection by *MGMT* Status randomized Phase I/II and III Trials for GBM

NO TMZ

- GLARIUS Irinotecan & Bev
NCT00967330, rand. Phase II
- TORISEL Temsirolimus
NCT01019434, rand. Phase I/II
- PPX and Concurrent Radiation
NCT01402063, rand. phase II

With TMZ

- CENTRIC Cilengitide Phase III
EORTC26071-22072 EMD 121974-011

Diagnosis

Step 1: central *MGMT* methylation status assay

MGMT unmethyl.

Step 2: Randomization

versus RT/TMZ-TMZ

+ New Agent

Radiotherapy

Concomitant Phase Adjuvant (maintenance) Phase

MGMT methyl.

Step 2: Randomization

versus RT/TMZ-TMZ

+ New Agent

TMZ

Radiotherapy

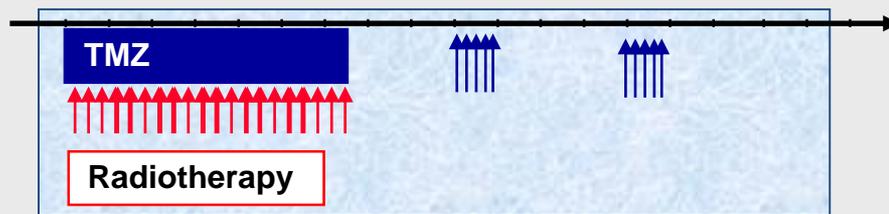
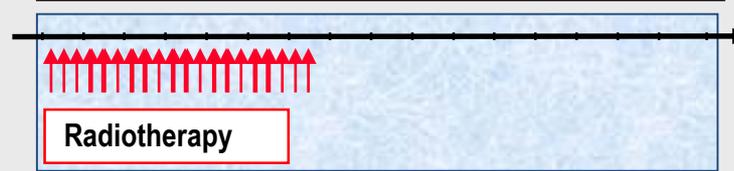
Concomitant Phase Adjuvant (maintenance) Phase

Registration

Randomiz.

R

R



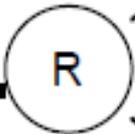
EORTC 26082/22081: Study Design

for newly diagnosed GBM with unmethylated *MGMT*

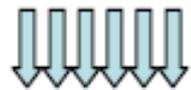
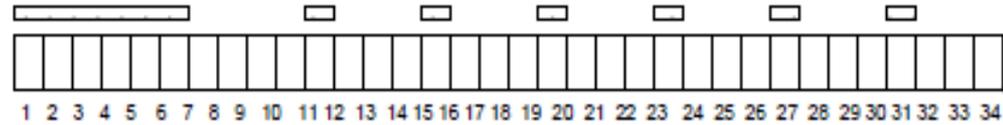
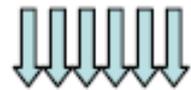
Randomized phase II – 2 arms

MGMT testing
(ratio < 0.6)
Pathology review

Surgery/open
biopsy
7 weeks



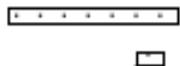
RT start: within 8 days from randomization



CCI-779 once every week at 25 mg.



RT regimen 5 X per week (30x 2 Gy, total dose 60 Gy)



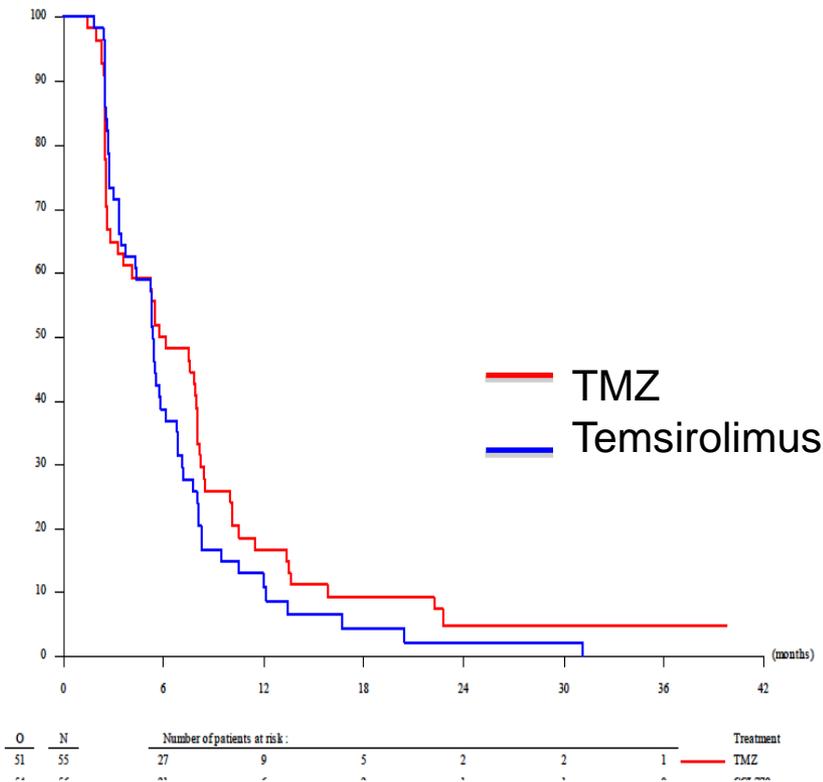
TMZ 75 mg/m² daily for the whole period of RT including
TMZ 150-200mg/m² D1-D5 every 4 wks – max 12 cycles

- 54 pts per arm
- Rule to Phase III: at least 39 eligible pts alive at 12 months in the Temsirolimus arm (>72%)

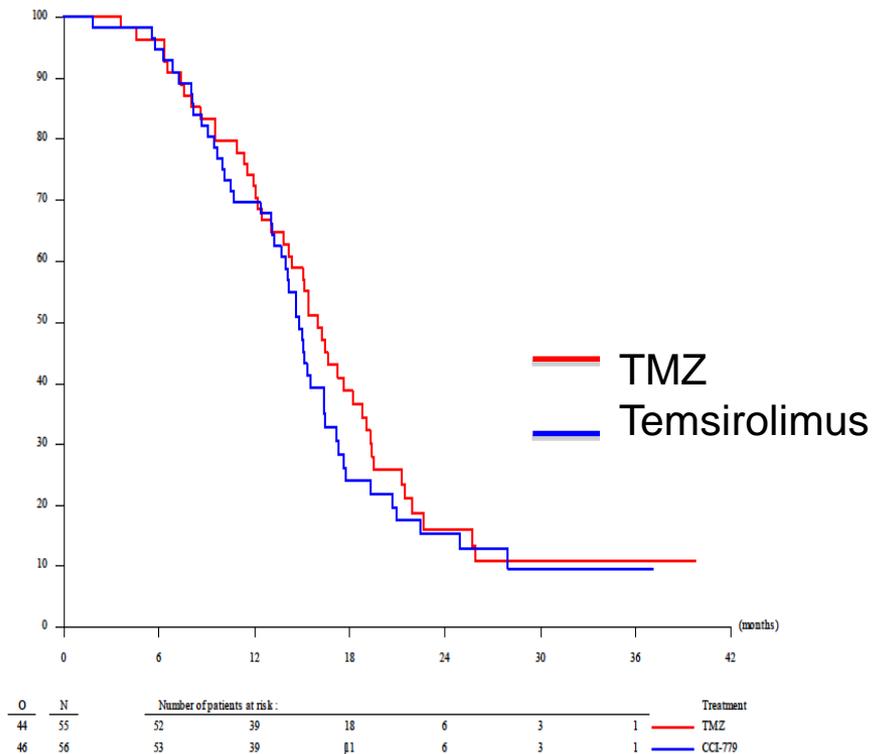
Temsirolimus/RT → Temsirolimus vs TMZ/RT→TMZ (EORTC26082)

Non-comparative

Progression-free survival

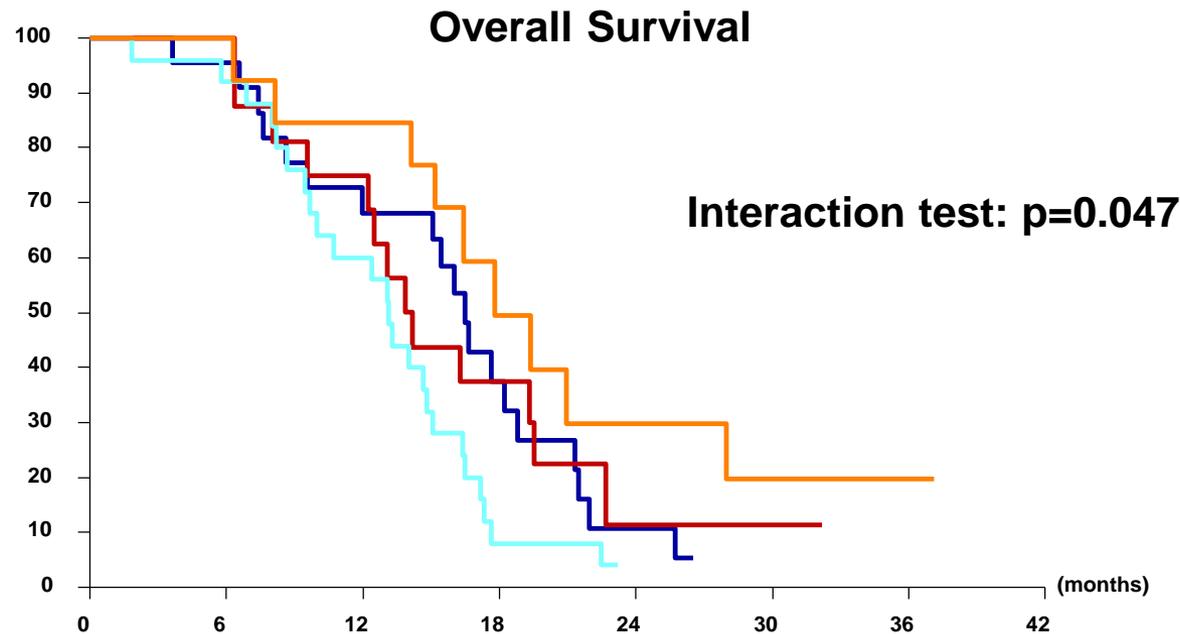


Survival

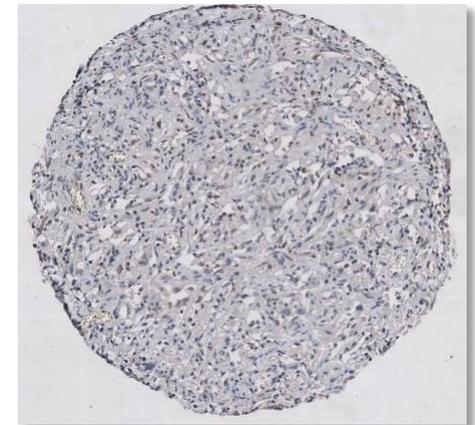
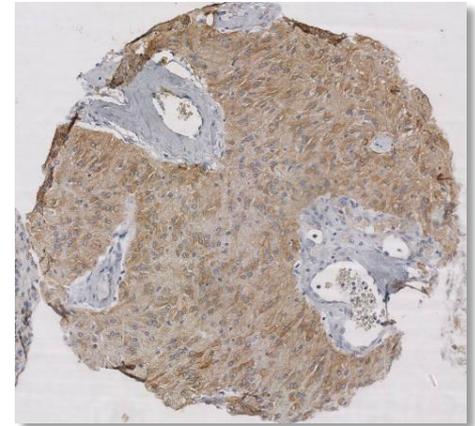


OS by p-mTOR expression and treatment

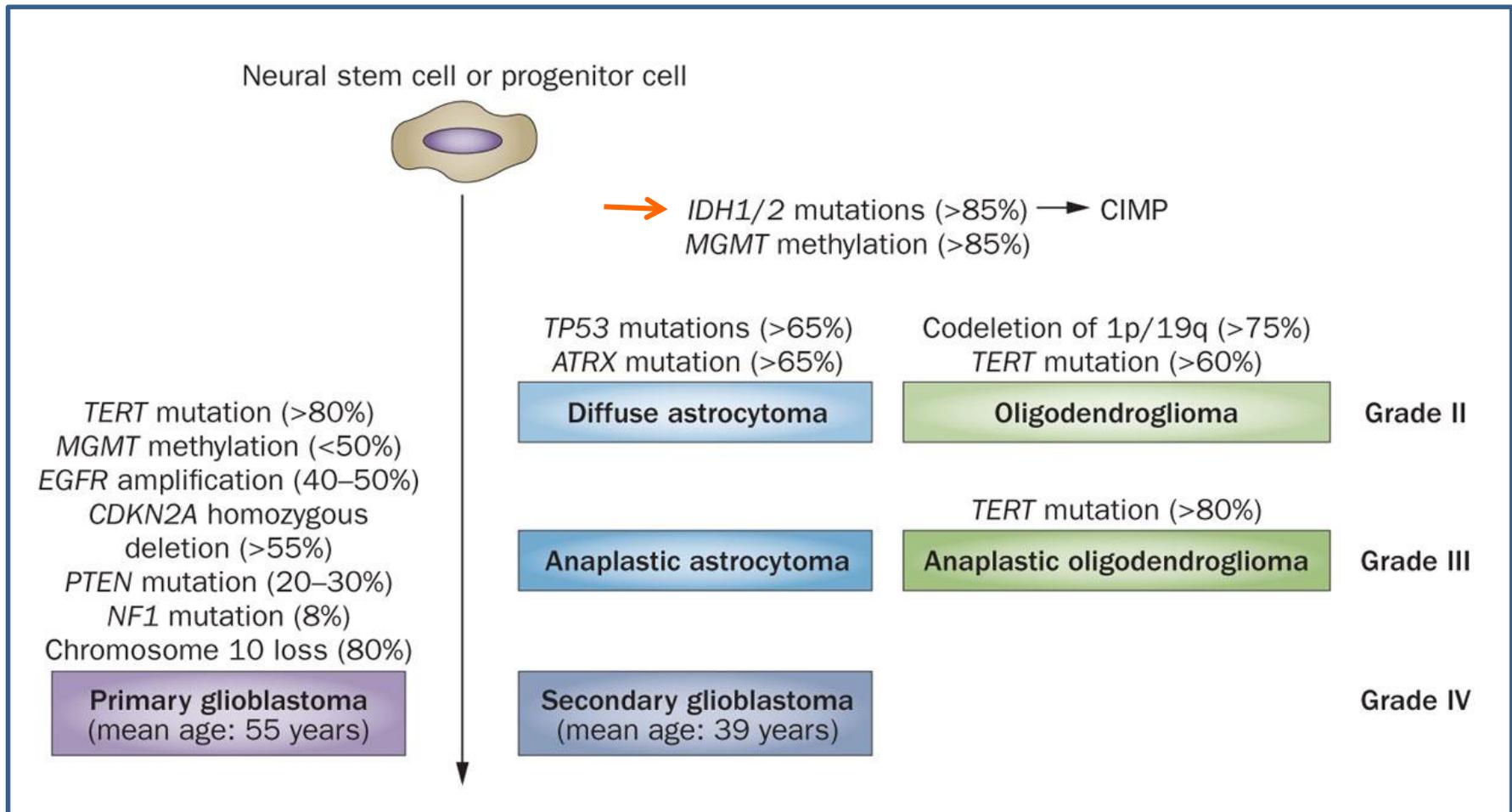
p-mTOR is predictive for benefit from temsirolimus



O	N	Number of patients at risk :						Trt/p-mTOR
19	23	21	15	7	2	0	0	— TMZ/Neg
13	16	16	12	5	1	1	0	— TMZ/Pos
24	25	23	15	2	0	0	0	— CCI-779/Neg
9	13	13	11	5	3	2	1	— CCI-779/Pos



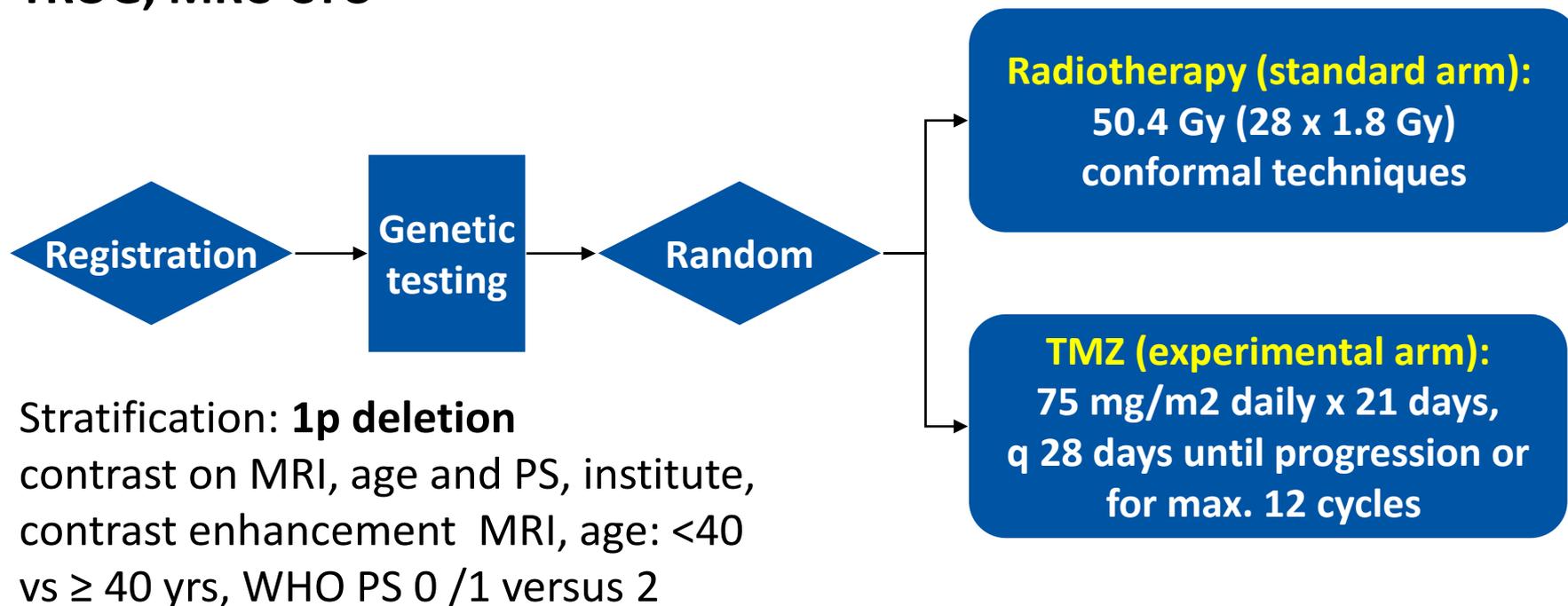
Pathogenetic and Epigenetic Evolution of Glioma in Adults



Hegi, M. E. & Stupp, R. (2013) *Nat. Rev. Neurol.* doi:10.1038/nrneurol.2013.127; PMID 23817351

EORTC/NCIC-CTG/TROG/MRC-CTU (EORTC 22033-26033) in patients with a high risk low-grade glioma

Participating groups: EORTC ROG and BTG, NCI-CTG,
TROG, MRC-CTU



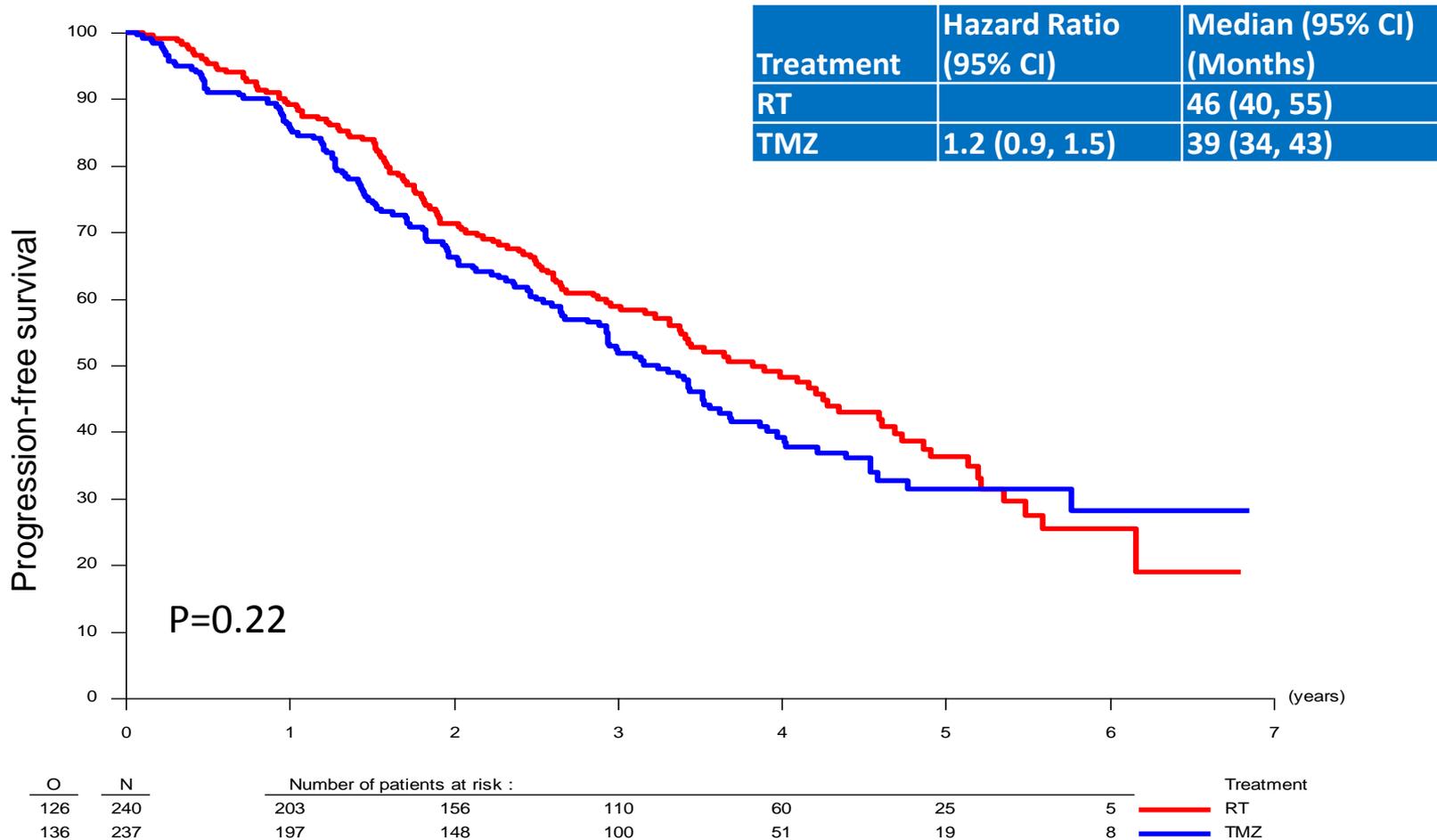
Accrual: **2005 – 2012**

707 patients registered

477 randomized

Baumert, Hegi,...Stupp. Submitted

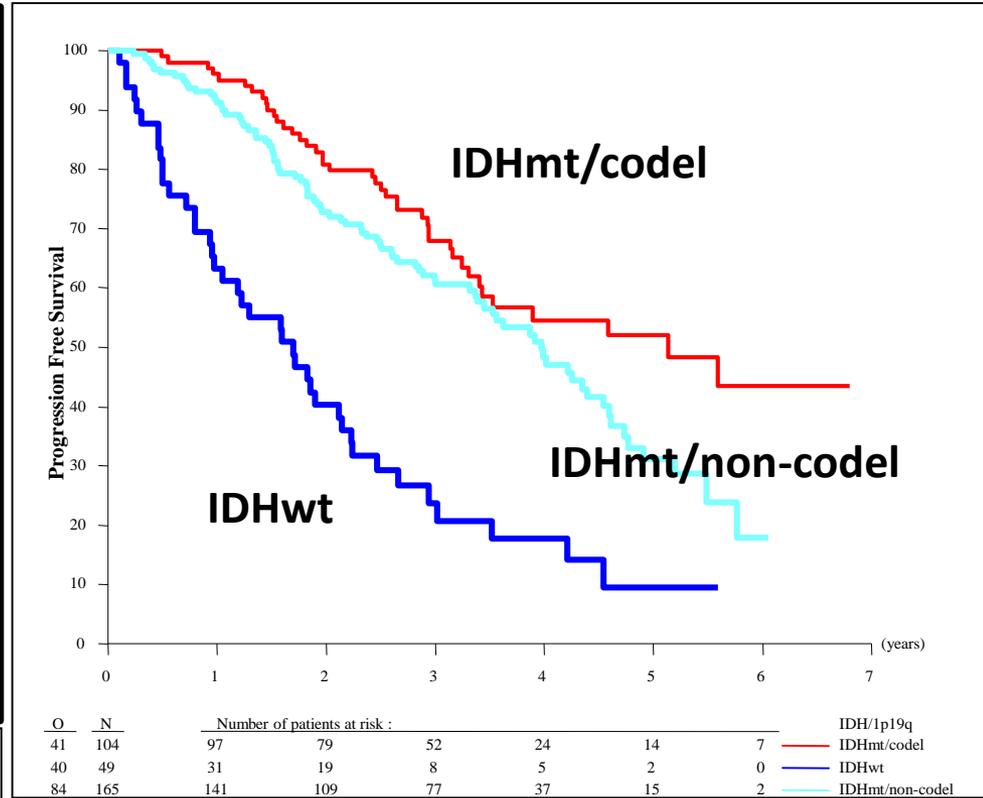
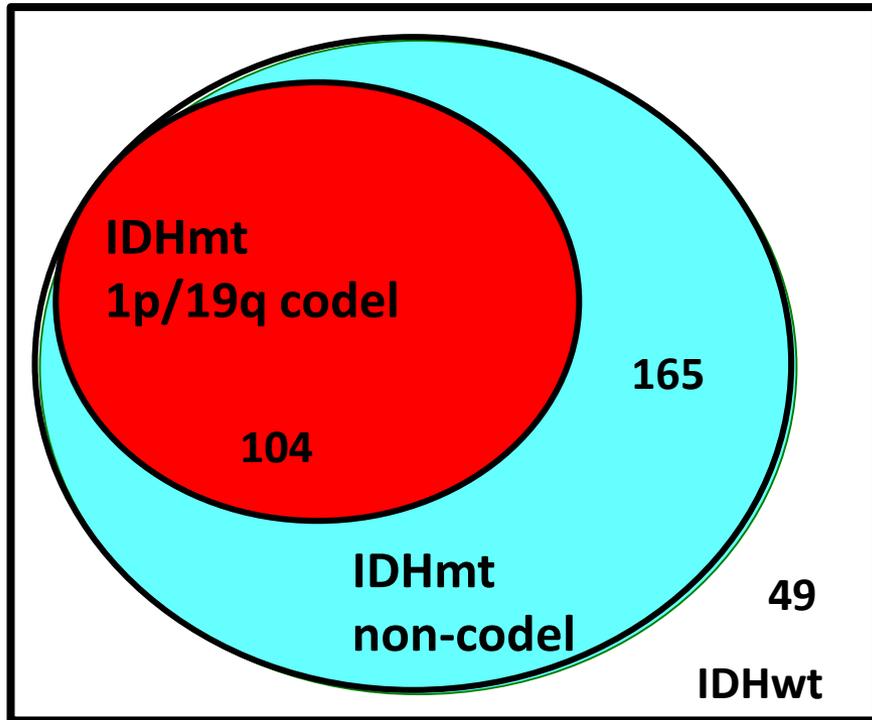
Progression-Free Survival



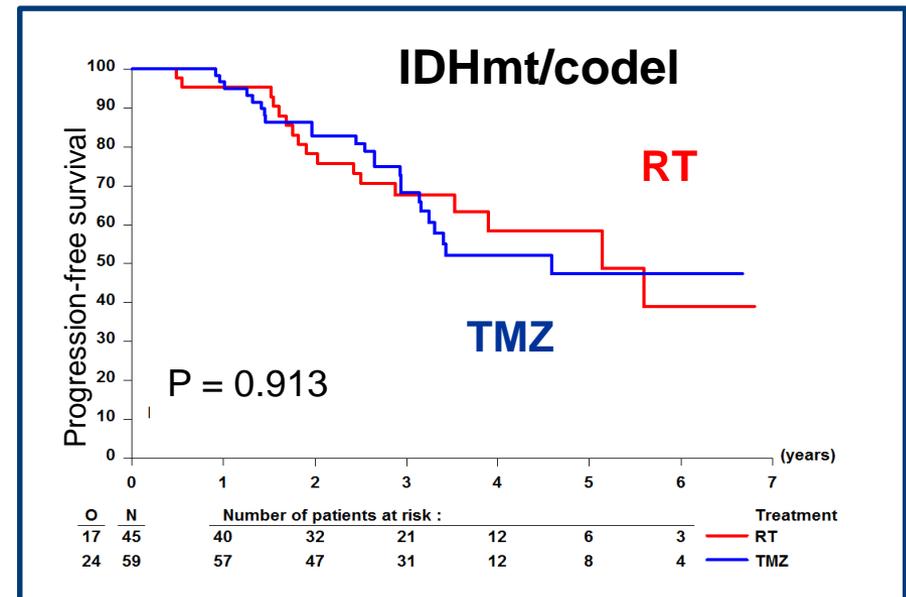
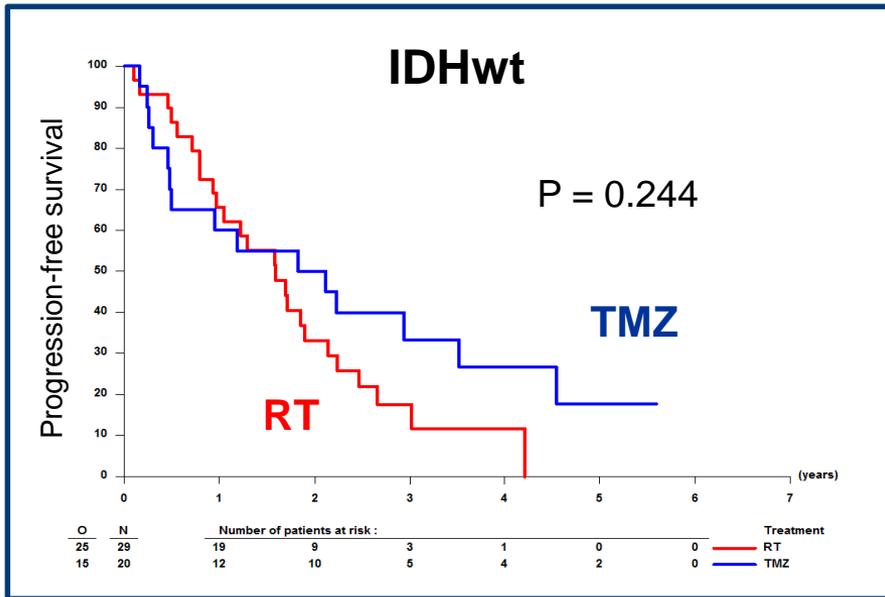
Baumert et al., Proc ASCO 2013 (abstr #2007)

PFS by Molecular Marker Subgroups: Prognostic value

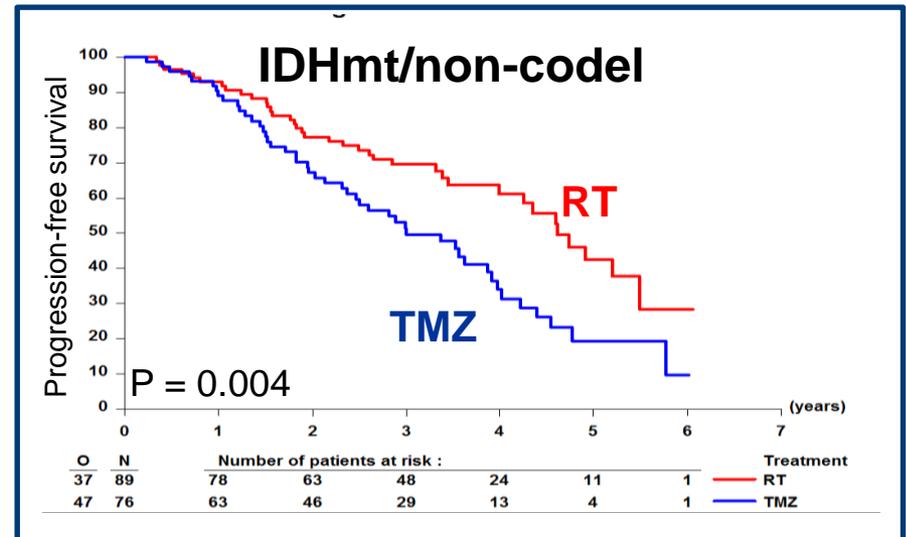
EORTC 22033



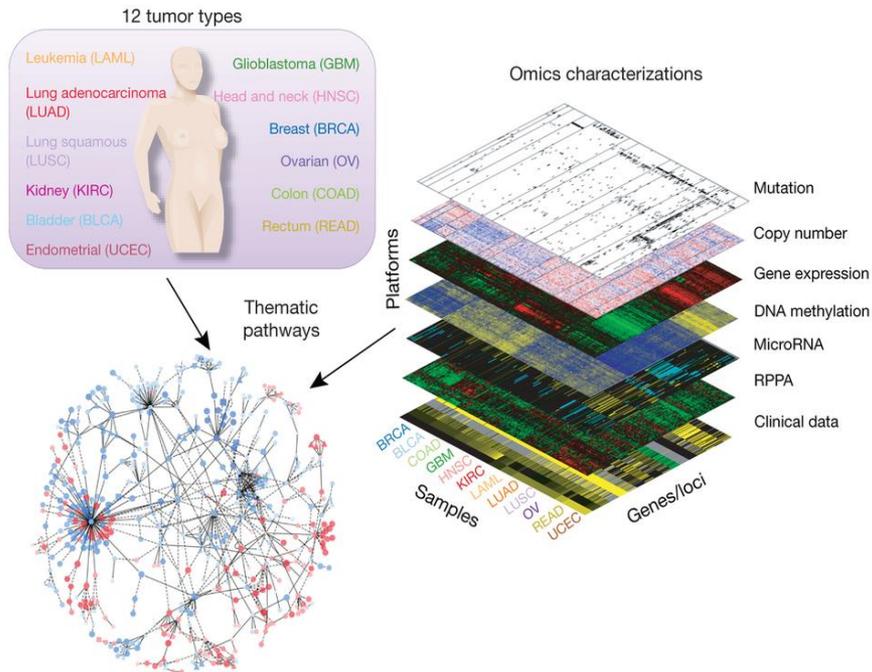
Molecular subgroups and treatment



Subgroup	Patients (N)	Events (O)	Median (95% CI) (Months)
RT-IDHmt/codel	45	17	61.63 (42.32, N)
RT-IDHwt	29	25	19.09 (11.27, 25.69)
RT- IDHmt/non-codel	89	37	55.36 (47.87, 65.87)
TMZ-IDHmt/codel	59	24	55.03 (37.95, N)
TMZ-IDHwt	20	15	23.69 (5.55, 42.25)
TMZ-IDHmt/non-codel	76	47	36.01 (28.42, 46.95)



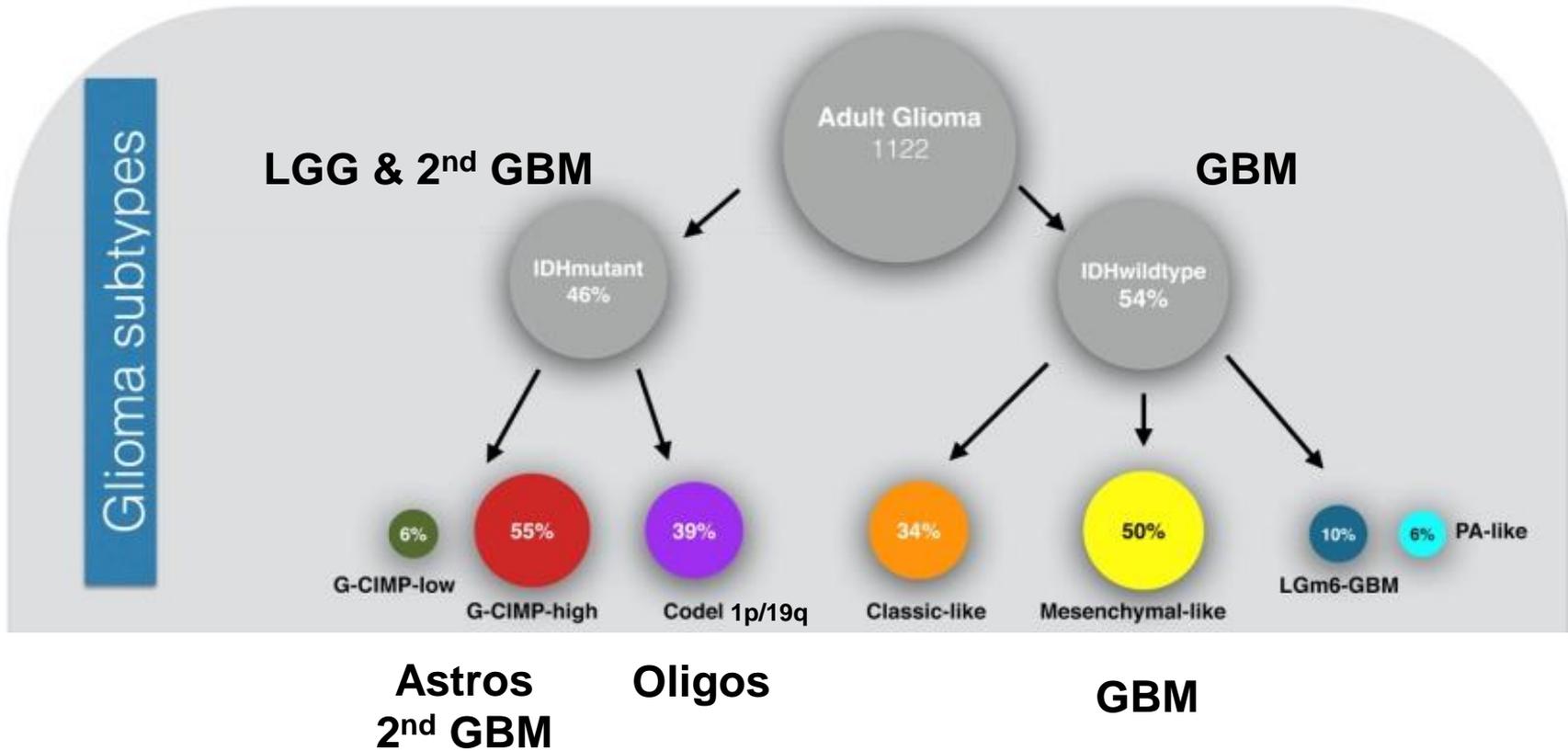
Translation of molecular knowledge into patient care



- Classification
- Outcome prediction
- Targets /treatment options
- Biomarkers for treatment choices
- Tumor heterogeneity

Adapted from : TCGA network, Nature Genetics, 1113–1120 (2013)

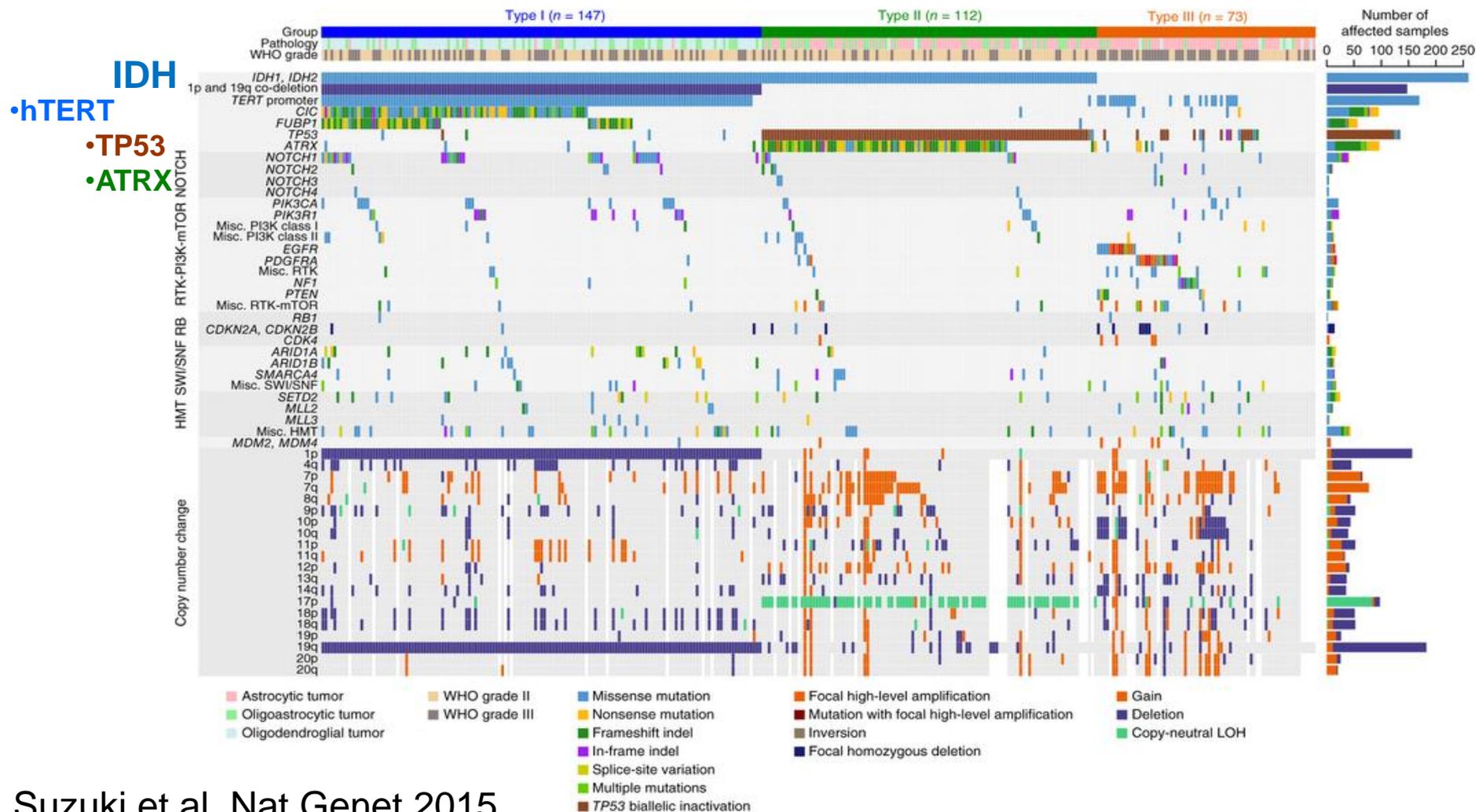
Molecular Glioma Subtypes



Adapted from Ceccarelli et al., Cell 2016 164, 550-563 DOI: (10.1016/j.cell.2015.12.028)

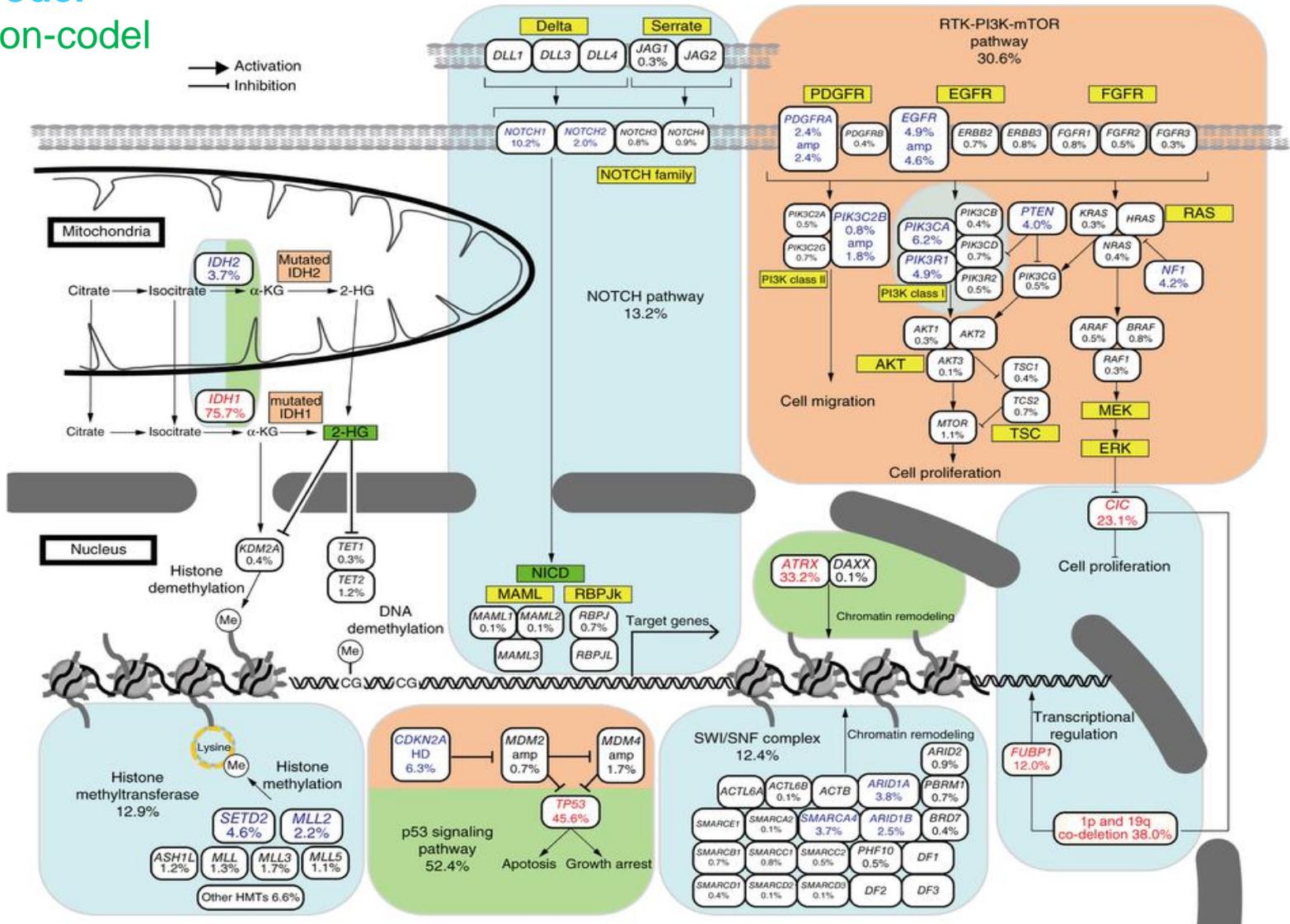
Landscape of genetic lesions in grade II and III gliomas

IDH mutant
IDH wild-type
1p/19 codel
1p/19 non-codel



Driver genes and functional pathways involved in grade II and III gliomas.

- IDHmt/codel
- IDHmt/non-codel
- IDHwt



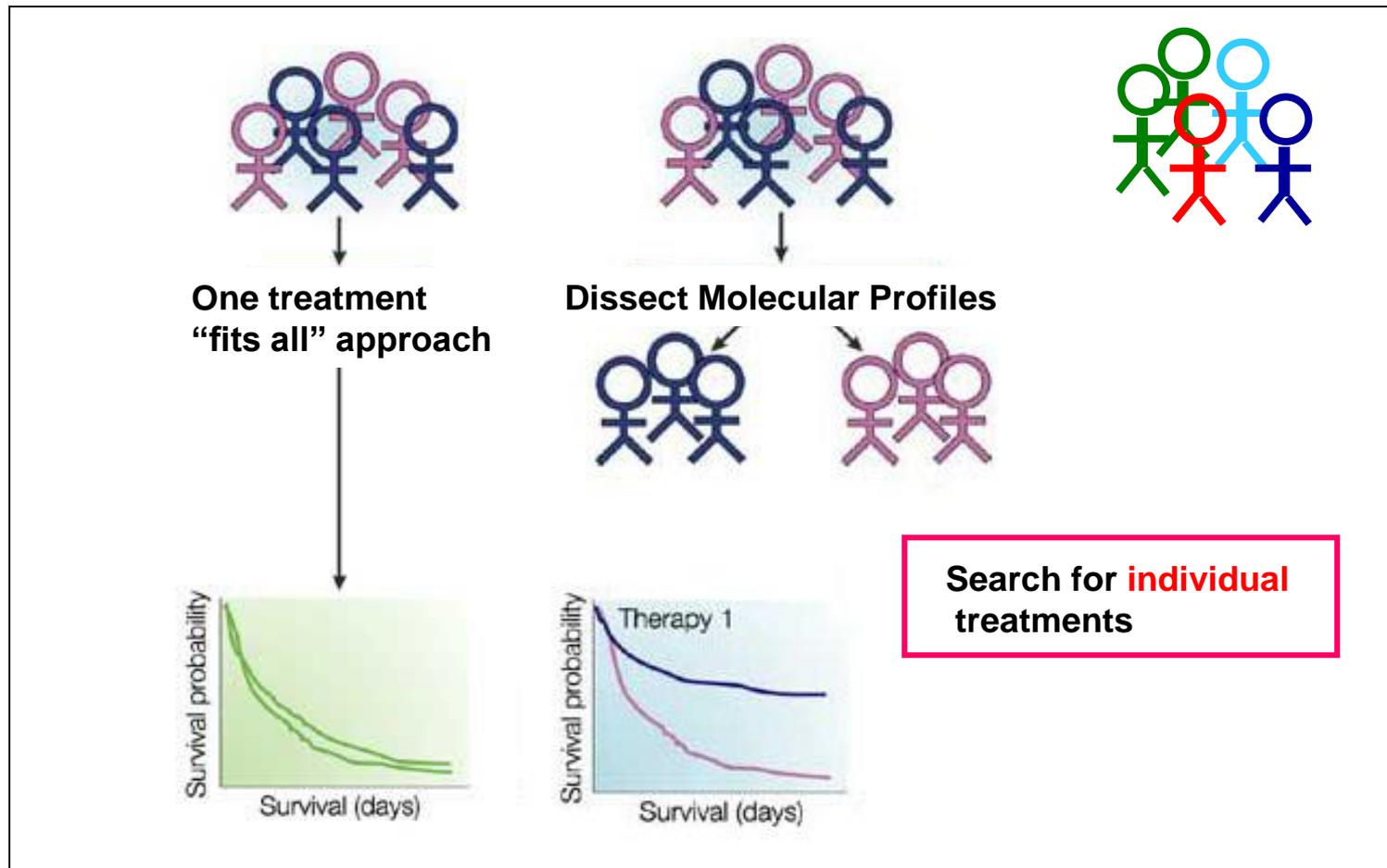
Treatment Recommendations

	Newly diagnosed	Progression
Anaplastic astrocytoma, WHO grade III	Resection or biopsy followed by radiotherapy or chemotherapy (or combined modality treatment)	Resection and chemotherapy, radiotherapy, or bevacizumab
Anaplastic oligodendroglioma or oligoastrocytoma, WHO grade III, with 1p/19q co-deletion	Resection or biopsy, followed by radiotherapy and chemotherapy, or chemotherapy	Resection and chemotherapy, radiotherapy, or bevacizumab
Anaplastic oligodendroglioma or oligoastrocytoma, WHO grade III, without 1p/19q co-deletion	Resection or biopsy, followed by radiotherapy or chemotherapy (or combined modality treatment)	Resection and chemotherapy, radiotherapy, or bevacizumab
Glioblastoma, WHO grade IV, (age <65–70 years)	Resection or biopsy, followed by radiotherapy plus concurrent temozolomide, followed by adjuvant temozolomide	Re-resection, reirradiation, rechallenge chemotherapy, or bevacizumab
Glioblastoma, WHO grade IV (age >65–70 years)	Resection or biopsy, followed by radiotherapy, or temozolomide with or without radiotherapy based on MGMT and performance status	Resection and chemotherapy or radiotherapy

Variations of RT with TMZ or other chemotherapy!!!

Table 1: Management options for newly diagnosed and progressive malignant gliomas

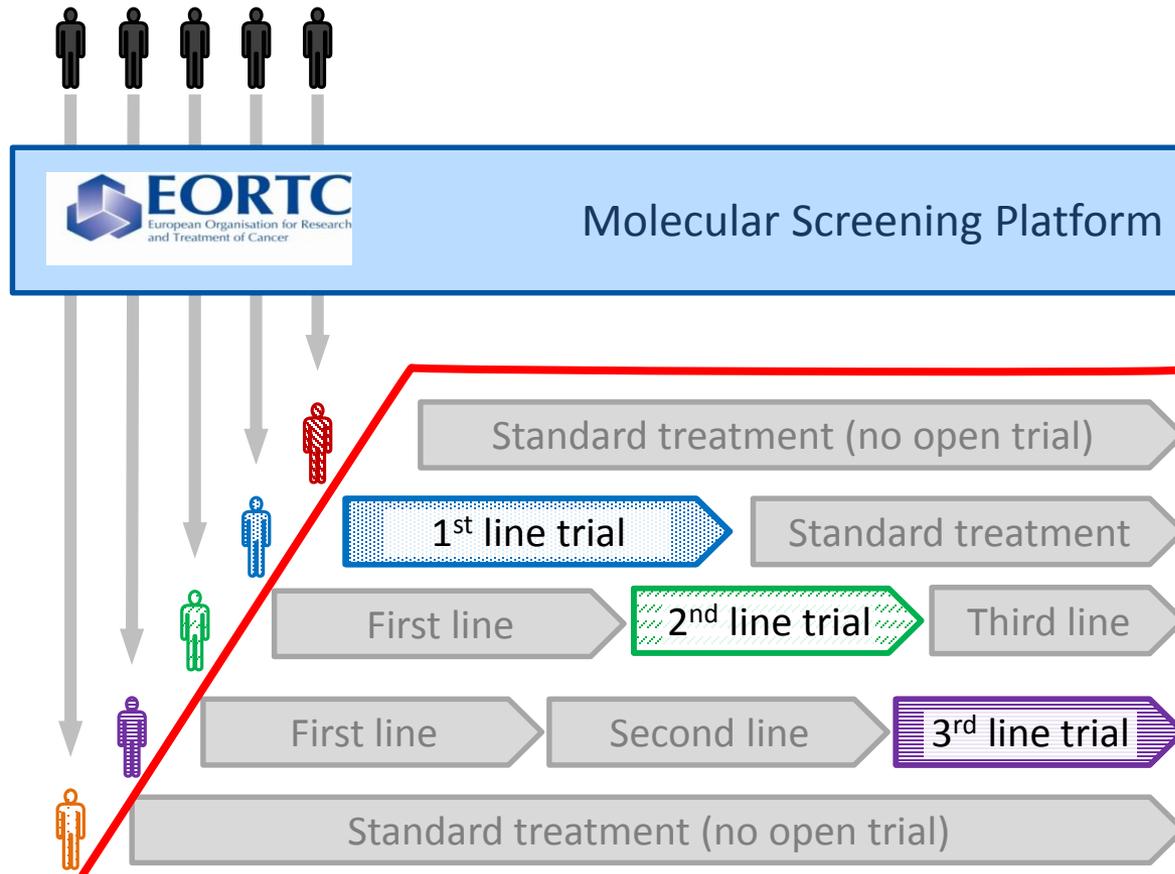
Molecular Profiles to Stratify Treatments *Towards Personalized Therapy*



Identification of Patients for personalized therapy

- **Activating mutations in proto-oncogenes, infrequent**
- **Driver vs Passenger mutations, function not always known**
- **Are they actionable (can they be targeted for therapy)**
- **Are respective targeted drugs available**
- **Testing for presence of target(s)**
- **Take advantage of other mechanisms (pathway)**
- **Target altered pathways at multiple hubs - combination therapies**

The SPECTA collaborative platform



Academia
investment

Industry
cooperation

SPECTA Platforms



Patient



Tumour tissue

Biobank



Biomarker analysis



Clinical data

Database



Patient enrolled in clinical trial



New treatment protocols



- Targeted Exome Seq (>400 genes)
- Gliomas, add HM450K (DNA meth)

Conclusions

- Molecular targeting agents require presence of the target → enriched population (dx become rare)
- International collaboration needed
- Partnership academia-industry indispensable

Tissue is the Issue

