

Biomarkers of Brain Tumors to Temozolomide Treatment

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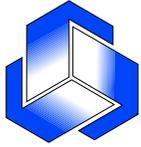
&

« NCCR Molecular Oncology », ISREC

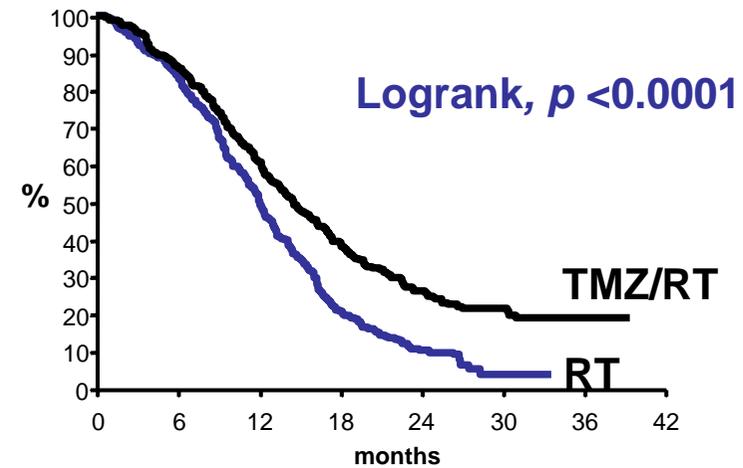
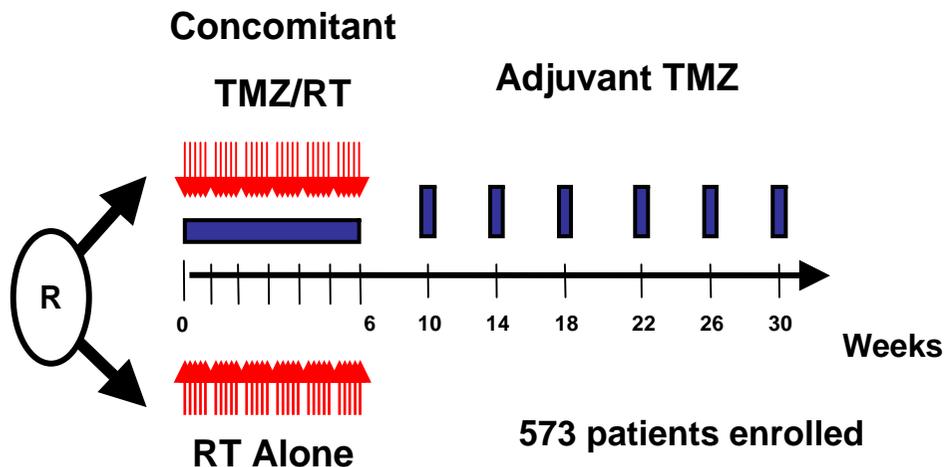


ENASCO Meeting, November 15-17, 2007





Predictive Factor(s) for Temozolomide (TMZ) Derived Benefit for Glioblastoma Patients



Who are the patients who benefit from TMZ ?

| | RT | TMZ/RT |
|----------------|------------------|--------|
| Median OS, mo: | 12.1 | 14.6 |
| 2-yr survival: | 10% | 26% |
| HR [95% CI]: | 0.63 [0.52-0.75] | |

O⁶-Methylguanine-DNA Methyltransferase (MGMT)

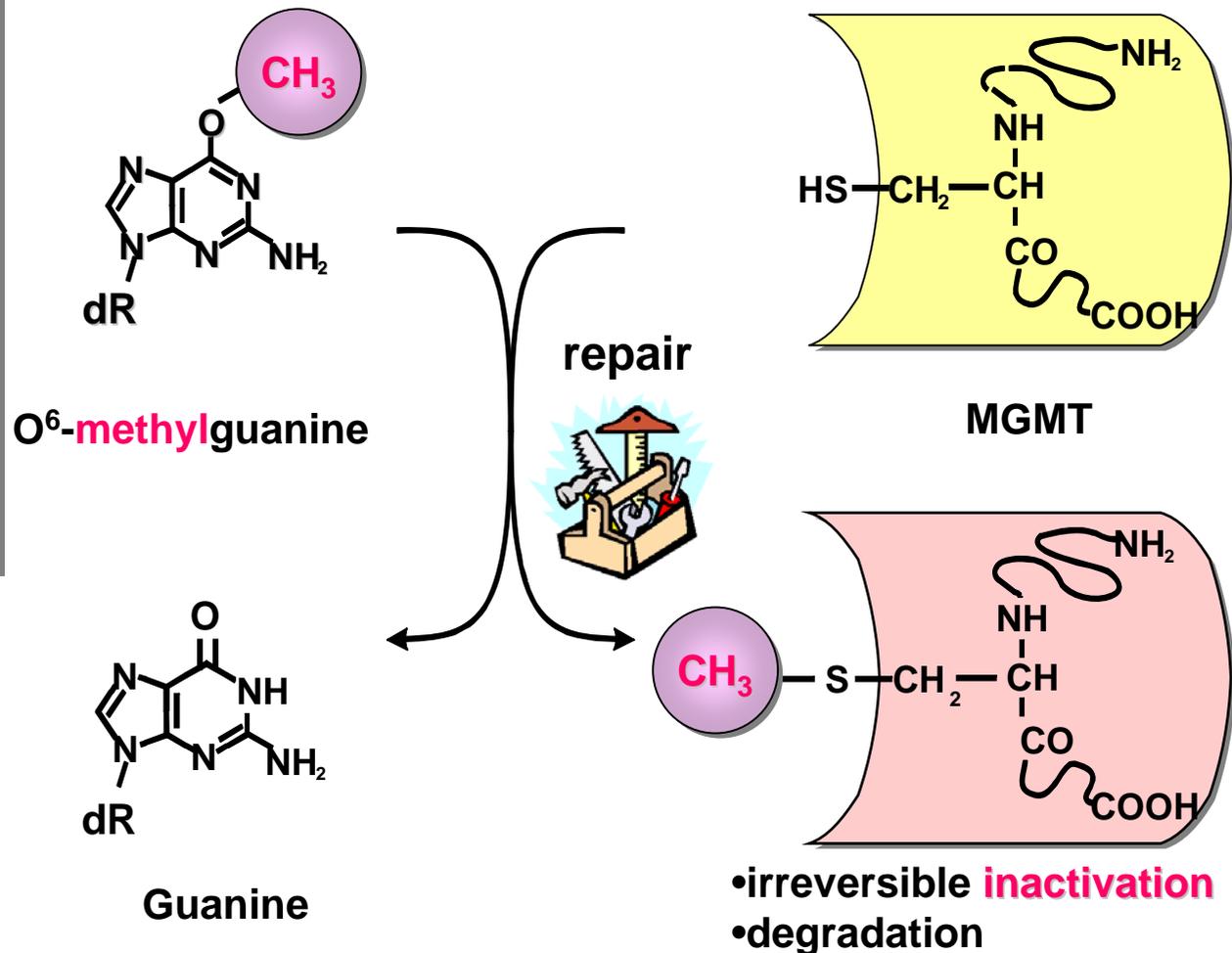
•TMZ, alkylating agent

MGMT

- CHR 10q26
- linked with resistance to alkylating agent therapy
- inducible :
 - RT
 - genotoxic agents
 - glucocorticosteroids

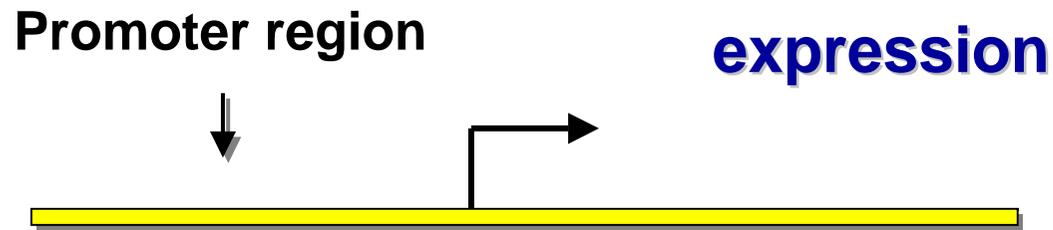
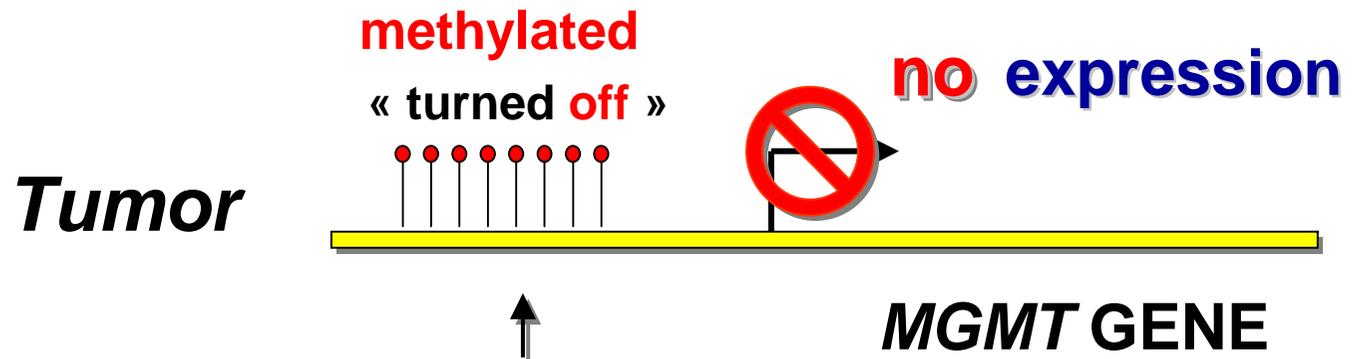
MOUSE MODELS & Alkylating Agents

- *MGMT*^{-/-} hypersensitive
- *MGMT*^{Tg} resistant to tumor formation



Silencing of the *MGMT* Repair Gene by Methylation of the Gene Promoter

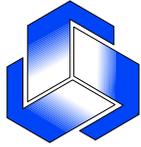
0⁶-Methylguanine-DNA Methyltransferase (MGMT)



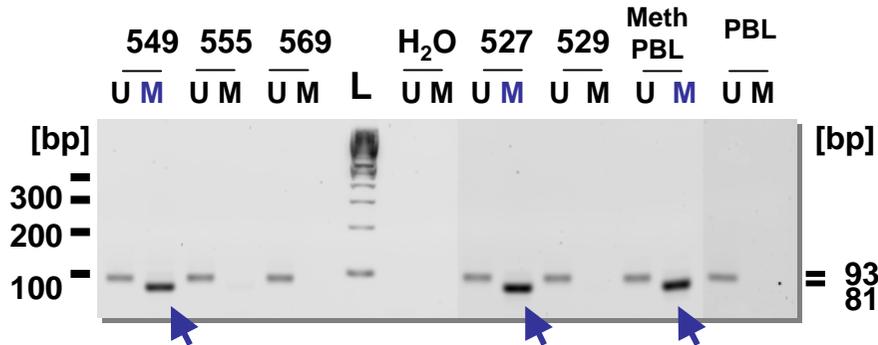
Tumor

- **No** repair protein
- **No repair** of TMZ treatment induced DNA damage

- **Response to tumor treatment**
- **Improved survival of glioblastoma patient**

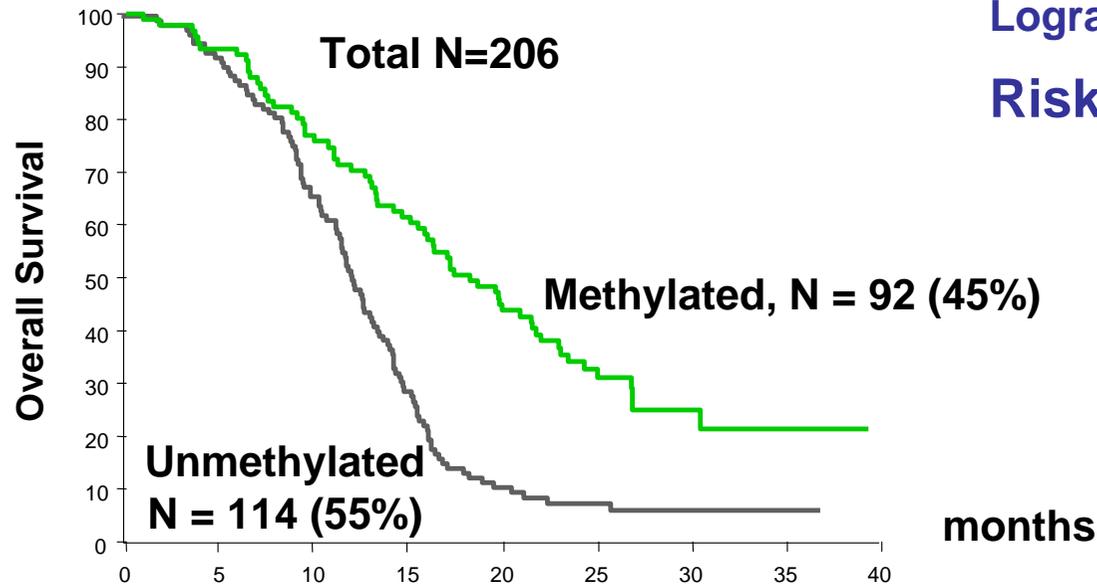


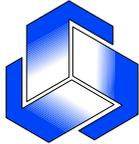
MGMT Promoter Methylation Predicts Better Outcome in Glioblastoma Patients of this Trial



| <i>MGMT</i> | Unmeth | Meth |
|----------------|------------------|-------|
| Median OS, mo: | 12.2 | 18.2 |
| 2-yr survival: | 7.8% | 34.1% |
| HR [95% CI]: | 0.45 [0.32-0.61] | |
| Logrank test: | $p < 0.0001$ | |

Risk of death reduced by 55%

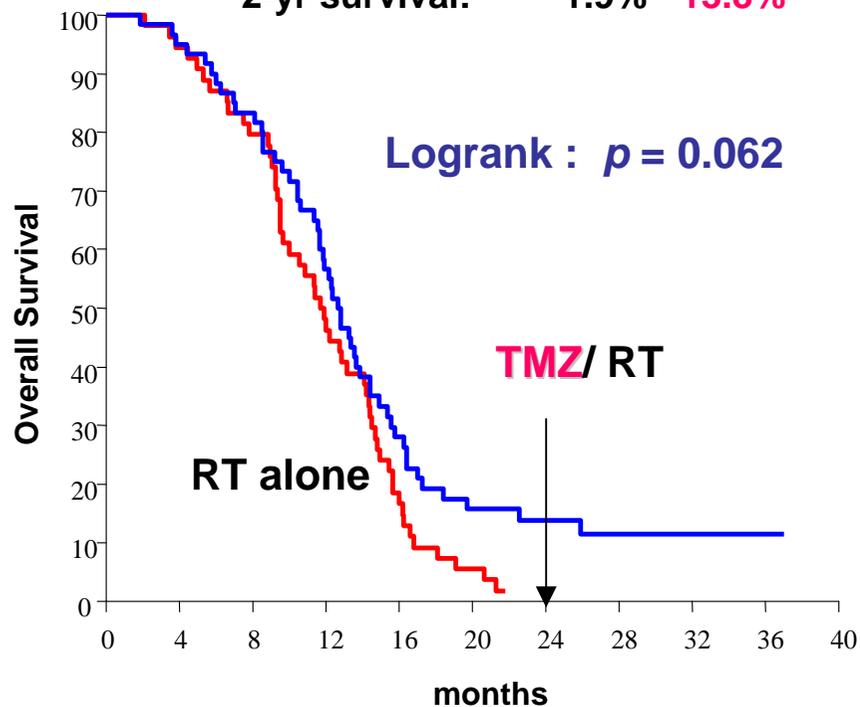




MGMT Promoter Methylation Predicts Benefit from TMZ Treatment

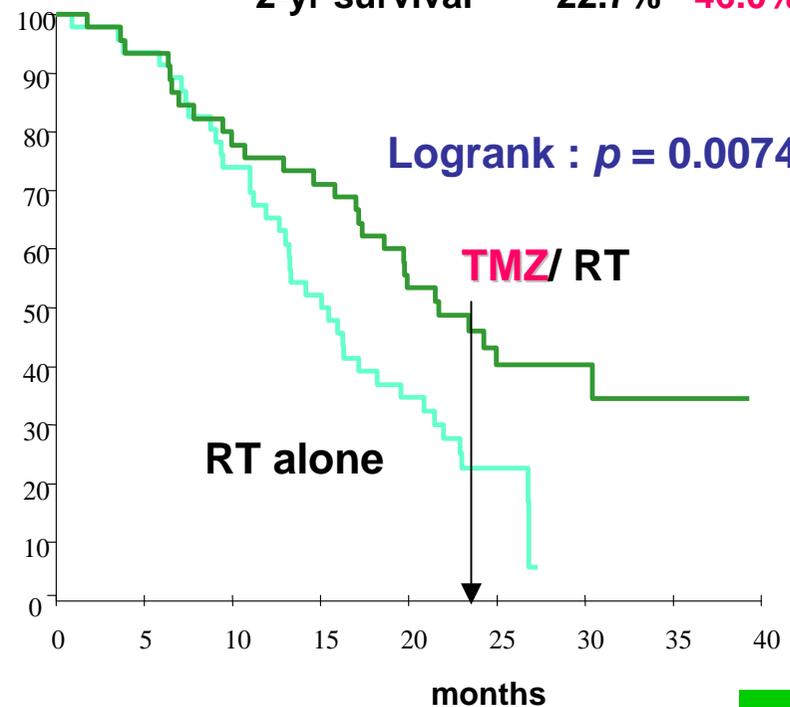
Unmethylated MGMT

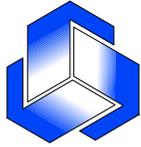
Randomization: RT **TMZ/RT**
Median OS mo: 11.8 **12.7**
2-yr survival: 1.9% **13.8%**



Methylated MGMT

Randomization: RT **TMZ/RT**
Median OS mo: 15.3 **21.7**
2-yr survival 22.7% **46.0%**

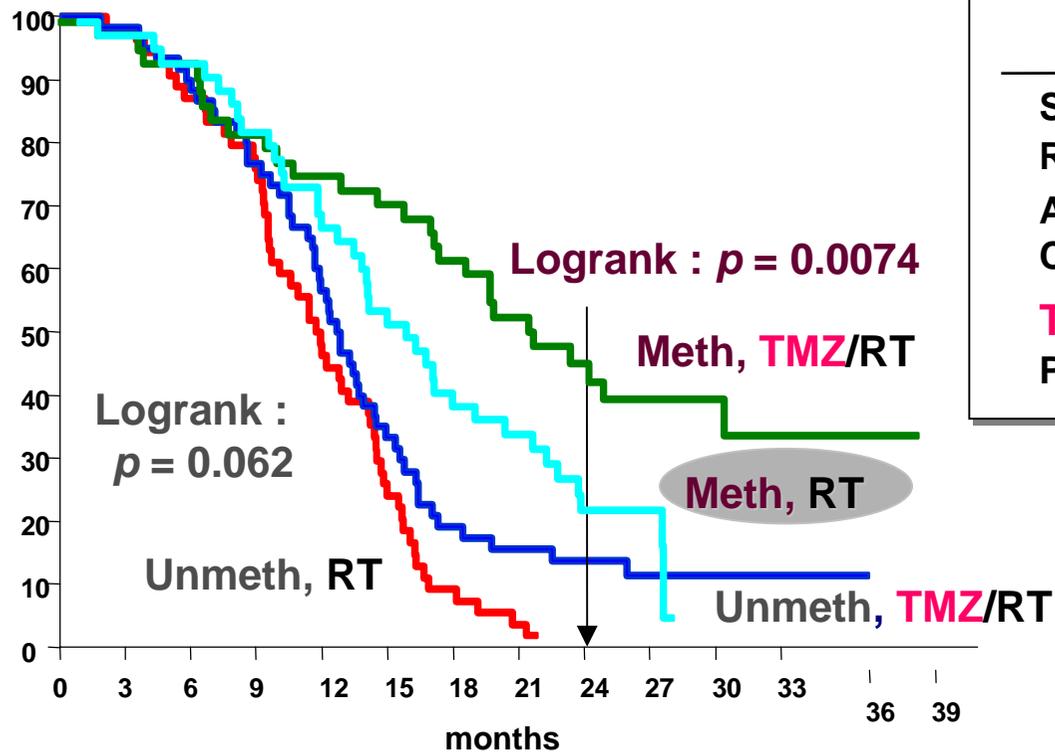




Predictive Value of *MGMT* Methylation for Overall Survival



Overall Wald test : $p < 0.0001$ (df=3)



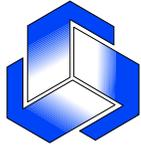
Treatment at progression
at discretion of treating physician

| Randomization | RT [%] n=286 | TMZ/RT [%] n=287 |
|--------------------------------|-----------------|---------------------|
| Surgery | 23 | 23 |
| Repeat RT | 4 | 5 |
| Any Additional Chemotherapy | 72 | 58 |
| Temozolomide | 60 | 25 |
| Palliative Care only | 17 | 22 |

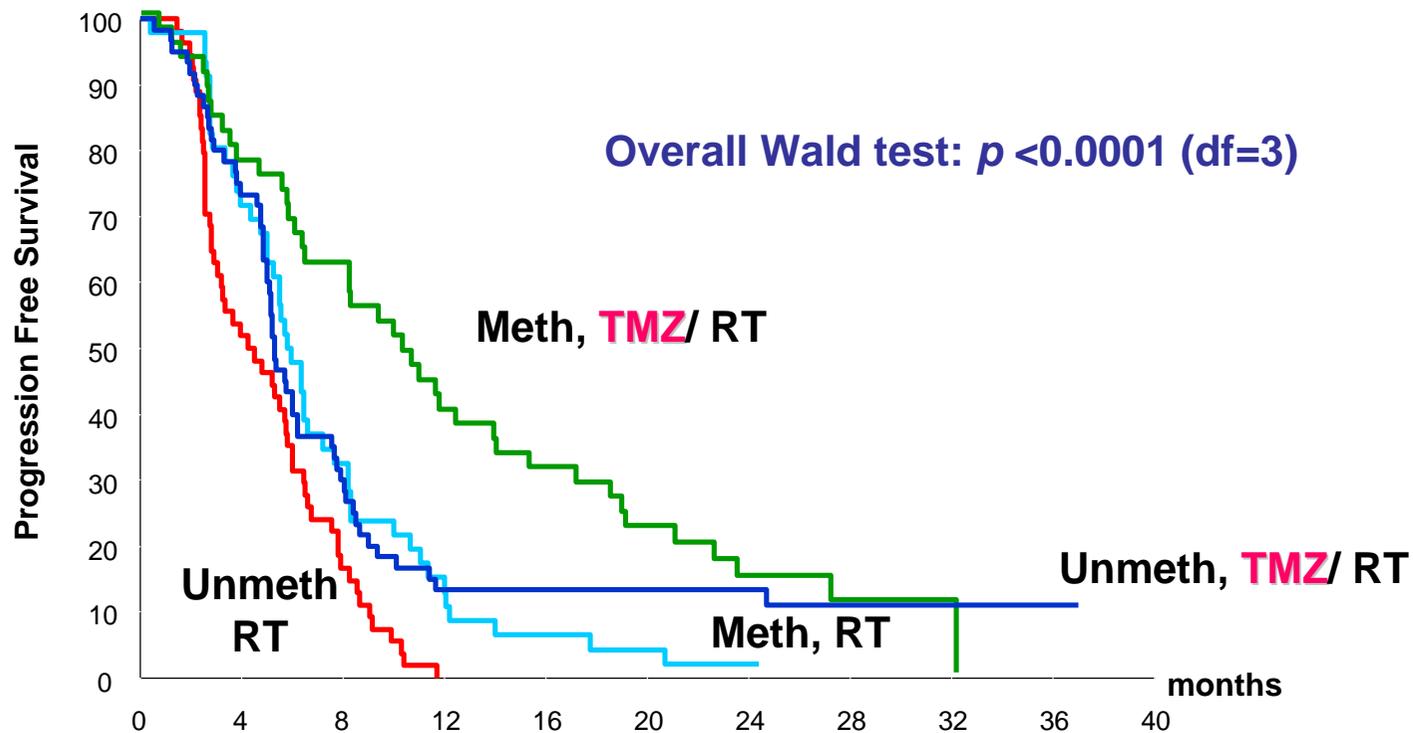
Stupp *et al*/ NEJM 2005

Hegi *et al.* N Engl J Med, 352: 997-1003, 2005





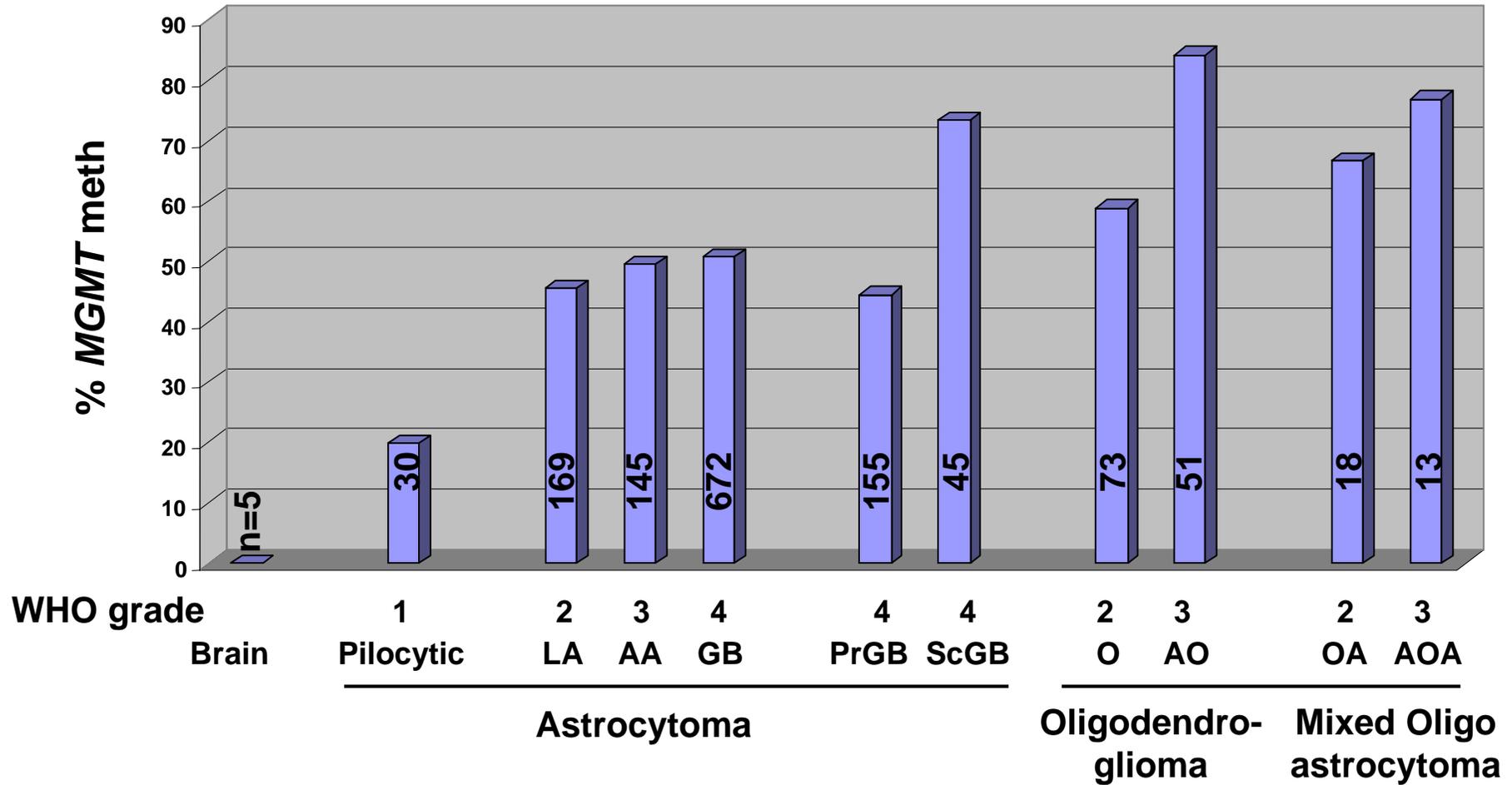
Progression Free Survival Supports *MGMT* Methylation Status as Predictive Factor for Benefit from TMZ



| O | N | Number of patients at risk : | | | | | | | | | |
|----|----|------------------------------|----|----|----|----|---|---|---|---|---|
| 54 | 54 | 28 | 9 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| 53 | 60 | 44 | 18 | 8 | 8 | 8 | 7 | 5 | 3 | 1 | |
| 45 | 46 | 33 | 15 | 7 | 3 | 2 | 1 | 0 | 0 | 0 | |
| 40 | 46 | 35 | 28 | 18 | 14 | 10 | 6 | 3 | 1 | 0 | |

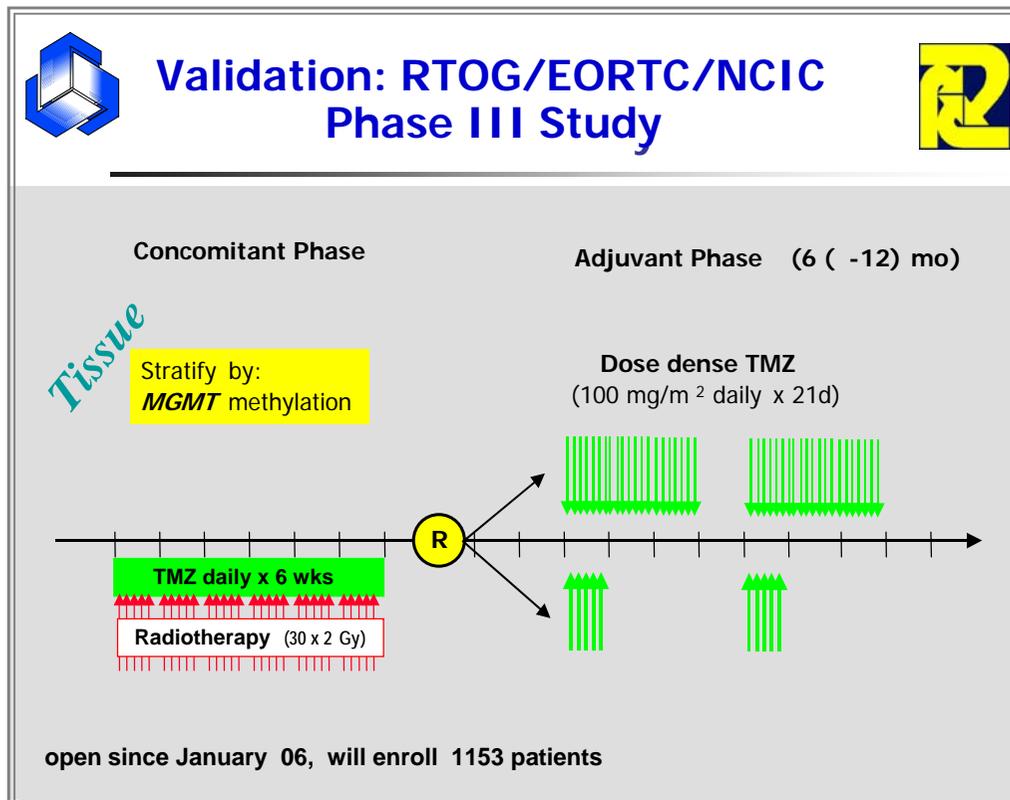
- Unmeth, RT alone
- Unmeth, TMZ/ RT
- Meth, RT alone
- Meth, TMZ/ RT

***MGMT* Promoter Methylation Ranges from 20 to >80% Depending on Glioma Subtype**



Review of the literature, August 2007
 data from 1376 patients from 15 publications

Validation of *MGMT* as Predictive Factor Depletion of *MGMT* in Tumor Cells by a Dose Dense Schedule



Integrated Translational Research Program:

**Identification of other resistance
factors and new targets**

Study Chairs:

Mark R. **Gilbert**, M.D. (Medical Oncology)

Minesh **Mehta**, M.D. (Radiation Oncology)

Ken **Aldape**, M.D. (Neuropathology and Correlative Biology)

Arnab **Chakravarti**, M.D. (Neuropathology and Correlative
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EORTC

Roger **Stupp**, M.D. (Medical Oncology)

Monika **Hegi**, Ph.D. (Neuropathology and Correlative
Biology)

Trials in GBM based on the RT/TMZ→TMZ scheme

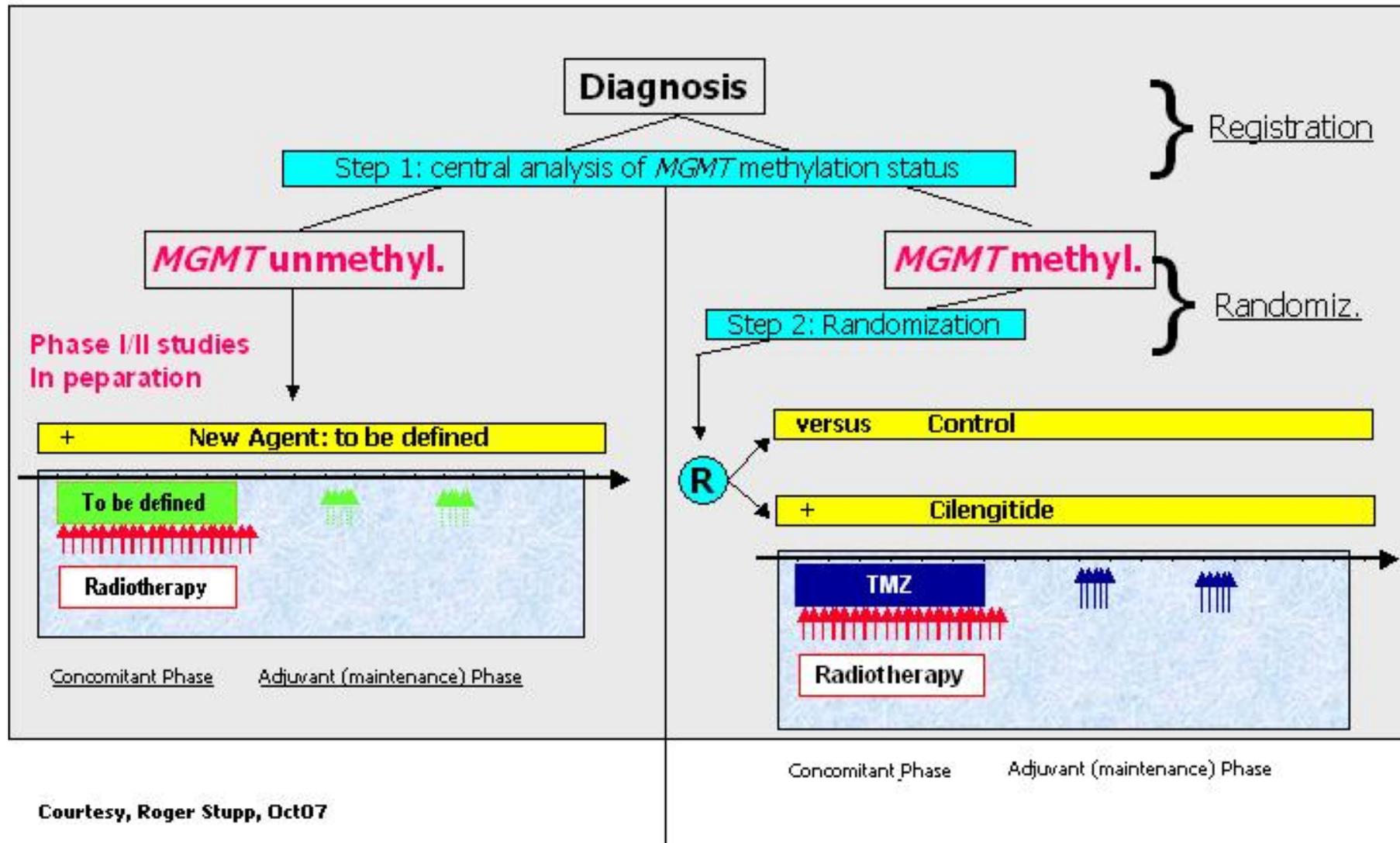
Table 1. Ongoing Trials of Novel Agents With Concomitant to Radiotherapy

| Investigational Agent | Standard Treatment | Phase | No. of Patients | End Point(s) | Sponsor | Remarks |
|-----------------------|--------------------|------------------|-----------------|--------------------------|-----------------------------------|---|
| Cilengitide | TMZ/XRT | II | 60 | PFS | Marck Serono | Assess completed, multicenter (Europe), ¹⁰⁰ phase III trial in preparation |
| Cilengitide | TMZ/XRT | I/II | 112 | OS | NABTT | Ongoing |
| Tamoxifen | TMZ/XRT | I | 46 | Safety and toxicity | NCCTG | Started May 2008 |
| Cetuximab | TMZ/XRT | I/II | 46 | Feasibility, PFS, OS | University of Heidelberg, Germany | No maintenance TMZ |
| CDK-119 | TMZ/XRT | II/III | 90-376 | PFS | Callex | Tumor-specific vaccine for EGFRvIII expressing tumors |
| Valproic acid | TMZ/XRT | III (randomized) | 180 | Survival | EORTC | Phase I completed; further drug development still continued |
| Imatinib/hydroxyurea | TMZ/XRT | I/II | 2 x 30T | Safety and toxicity, PFS | Novartis Germany | Phase II separate strata with/without EIAED |
| Bevacizumab | TMZ/XRT | II | 70 | Survival | UCLA | Requires fresh-frozen tumor tissue |
| Bevacizumab | TMZ/XRT | III | 800 | Survival | Berenson | Start planned summer 2007 |
| Ercosulfurin | TMZ/XRT | I/II | 72 | Survival (ph1) | Lilly | Started September 2006 |
| Ercosulfurin | XRT | II | 64 | PFSS | Lilly Germany | Only for patients with an unmethylated MGMT |
| Tipifarnib | TMZ/XRT | I | 30 | Safety and toxicity | NABTT | Completed |
| Tipifarnib | XRT | II | 27 | Survival | Instituto Claude Regaud, Toulouse | France: Toulouse, Clermont-Ferrand |
| Larotidine | XRT | I/II | 80 | Survival | DFCI | Phase I in phase II in combination with TMZ/radiation therapy planned |
| Venlafaxine | TMZ/XRT | III (randomized) | 160 | Survival | DFCI | Start planned summer 2007. Harvard affiliate, MSKCC, University of Virginia, University of Pittsburgh |
| Valproic acid | TMZ/XRT | II | 41 | PFS, survival | National Cancer Institute | Valproic acid as a histone deacetylase inhibitor |
| Boreavertin | TMZ/XRT | II/III | 80 | Safety, PFSS | Novocis | Biosynthetic drug targeting hypoxic cells |
| Carmustine water | TMZ/XRT | II | 72 | Survival | Johns Hopkins | |

Abbreviations: TMZ, temozolomide; XRT, irradiation; NABTT, New Approaches to Brain Tumor Therapy CNS Consortium; NCCTG, North Central Cancer Treatment Group; EORTC, European Organization for Research and Treatment of Cancer; EIAED, enzyme inducing antiepileptic drugs; UCLA, University of California, Los Angeles; NABTT, North American Brain Tumor Consortium; DFCI, Dana-Farber Cancer Institute; MSKCC, Memorial Sloan-Kettering Cancer Center, New York; PFS, progression-free survival; OS, overall survival; PFSS, 6-month PFS rate.



Cilengitide Phase III for GBM



Courtesy, Roger Stupp, Oct07

Frequency of *MGMT* Methylation in Glioblastoma

Range published for GBM 34 to 68%
(gel based, mostly on frozen tissue)

| Tumor type | # samples | MGMT-meth | % | Reference |
|--------------|------------|------------|-----------|------------------------|
| GBM | 29 | 10 | 34 | {Watanabe, 2005 #3670} |
| GBM | 21 | 8 | 38 | {Balana, 2003 #3672} |
| GBM | 29 | 12 | 41 | {Esteller, 2000 #1455} |
| GBM | 12 | 5 | 42 | {Yu, 2004 #2165} |
| GBM | 74 | 33 | 45 | {Kamiryo, 2004 #3677} |
| GBM | 206 | 92 | 45 | {Hegi, 2005 #2000} |
| GBM | 44 | 30 | 68 | {Blan, 2004 #3674} |
| GBM | 38 | 26 | 68 | {Hegi, 2004 #1721} |
| GBM | 219 | 126 | 58 | {Criniere, 2007 #6708} |
| TOTAL | 672 | 342 | 51 | |

Comparison of qMSP and Classic Gel Based Nested MSP for determination of the *MGMT* status

Quantitative MSP

OncoMethylome Sciences

Ilse Vlassenbroeck
Stéphane Califice
Josef Straub
Ivano Di Stefano
Fabrice Moreau
Isabelle Renard,
Bruno Flamion
James DiGuseppi
Katja Bierau

Gel Based MSP

Lab of Tumor Biology and Genetics, Neurosurgery, CHUV

Annie-Claire Diserens
Marie-France Hamou
Monika E. Hegi

Statistics

NCCR Molecular Oncology & Swiss Institute of Bioinformatics

Eugenia Migliavacca
Mauro Delorenzi

**Tissues from Trials : Lausanne, R. Stupp; Rotterdam, M. van den Bent
Regensburg, P. Hau**

Experimental Workflow of the Assays

- **Evaluation of tissue** (H&E slide, tumor content, amount)
- **4 sections / sample for each center**

GEL BASED ASSAY **CHUV**

- **Bisulfite treated DNA**
- **Nested MSP** : 1st PCR (298bp)
- 2nd PCR (discriminating)
 - PCR for **meth** *MGMT* (81bp)
 - PCR s for **unmeth** *MGMT* (93bp)
- **Results** visualized on gel

qMSP **OncoMethylome Sciences**

- **Bisulfite treated DNA**
- **quantitative MSP** :
 - **meth** *MGMT* (136bp)
 - beta Actin (125bp)
- **Ratio of mMGMT/Actin *1000**

Comparison of results

Non-disclosed Unpublished Data

Vlassenbroeck *et al* submitted

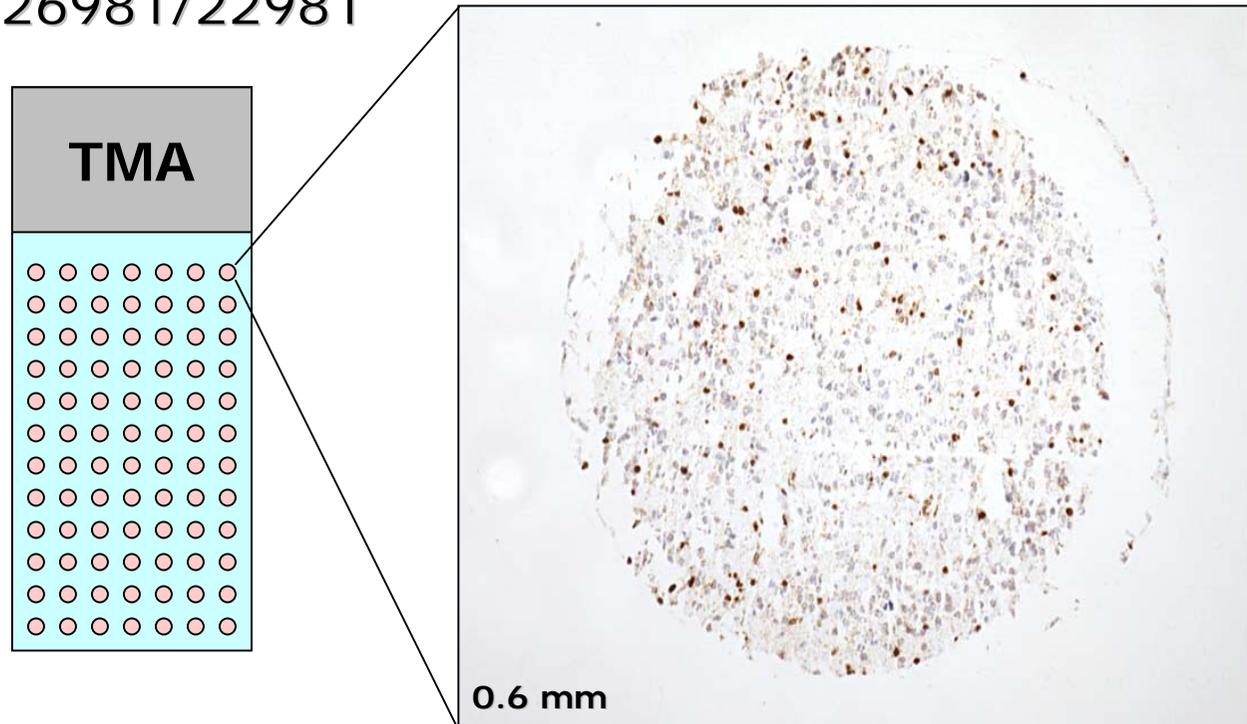
MGMT immunohistochemistry in GBM: Interobserver agreement in EORTC/NCIC trial26981/22981

**Preusser M, Janzer RC, Felsberg J, Reifenberger G, Hamou M-F,
Diserens A-C, Marosi C, Heinzl H, Stupp R, Hainfellner JA, Hegi ME**



Methods : 2 anti-MGMT antibodies, **Dako MT3.1, Zymed MT23.2**

- Tissue micro array (TMA): 163 tissue samples from GBM trial 26981/22981

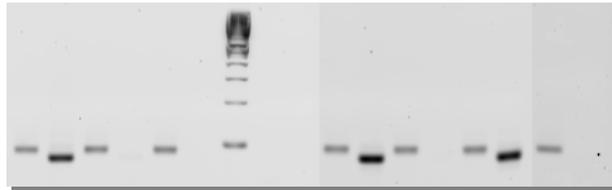
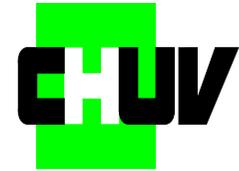


- 4 neuropathologists - 3 laboratories (RJ, GR, JAH, MP)
- Statistical analysis

Non-disclosed Unpublished Data

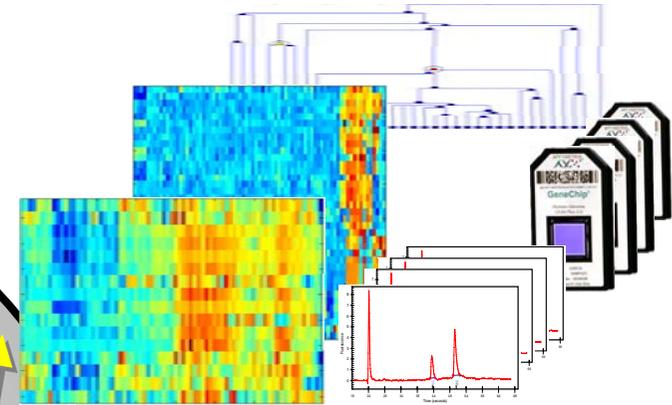
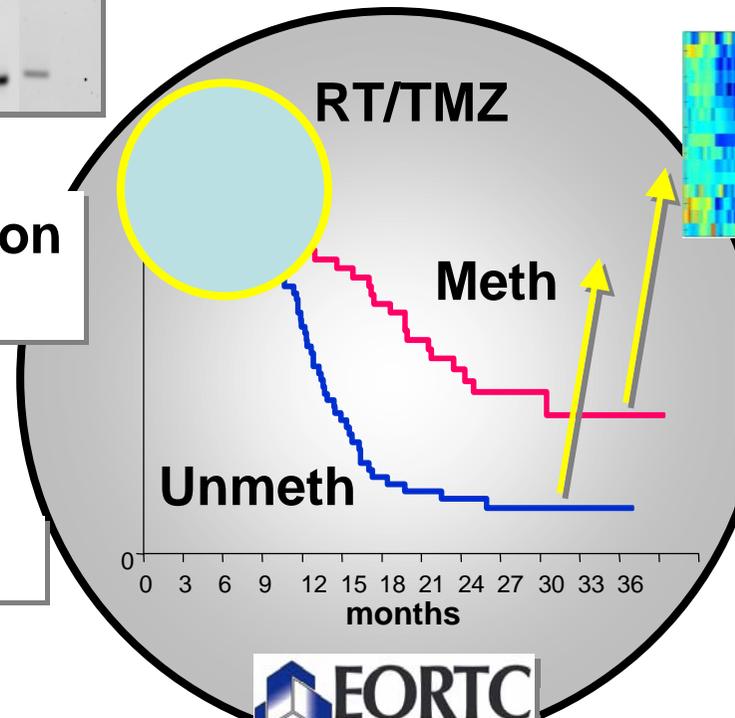
Preusser *et al* submitted

New Molecular Targets



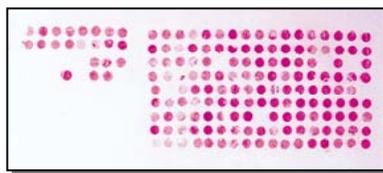
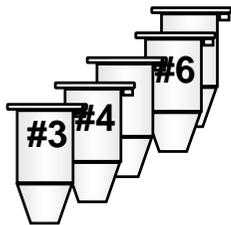
MGMT methylation Status

Frozen Tissue

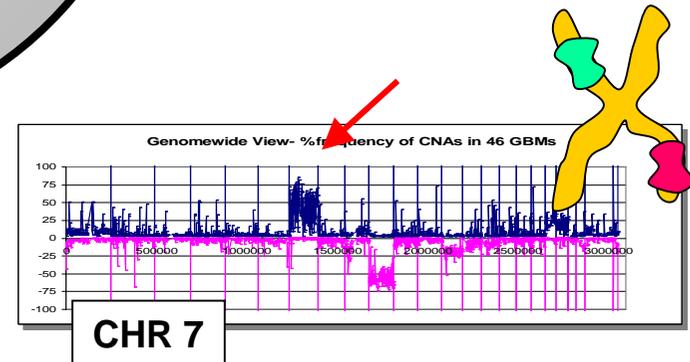


Gene Expression Profiles

Array-CGH



Tissue Array



Non-disclosed Unpublished Data

Murat *et al* submitted

Conclusions

- The *MGMT* methylation status predicts benefit from the alkylating agent TMZ
- Standardized *MGMT*-testing required
 - Quantitative MSP is reproducible, prospective testing ongoing
 - IHC is not useful for diagnostic *MGMT*-testing
- New trials will select patients based on *MGMT* status

The Team in the Lab



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Isabelle Desbaillets
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Marie-France Hamou
Yan Lachat
Sophie Shnaper

Monika Hegi
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Genève**

**Patrick Descombes
Didier Chollet**

**UCSF
Anjan Misra
Burt Feuerstein**



Oncology, CePO, CHUV

Roger Stupp

EORTC

Thierry Gorlia

**Patients
and their Families**

85 CENTERS



**Brain Tumor Group
Radiotherapy Group**



FONDATION NELIA ET AMADEO BARLETTA

Personalizing cancer treatment

85 CENTERS

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