

Completeness of required site-specific factors for brain and CNS tumors in the Surveillance, Epidemiology and End Results (SEER) 18 database

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The Central Brain Tumor Registry of the United States



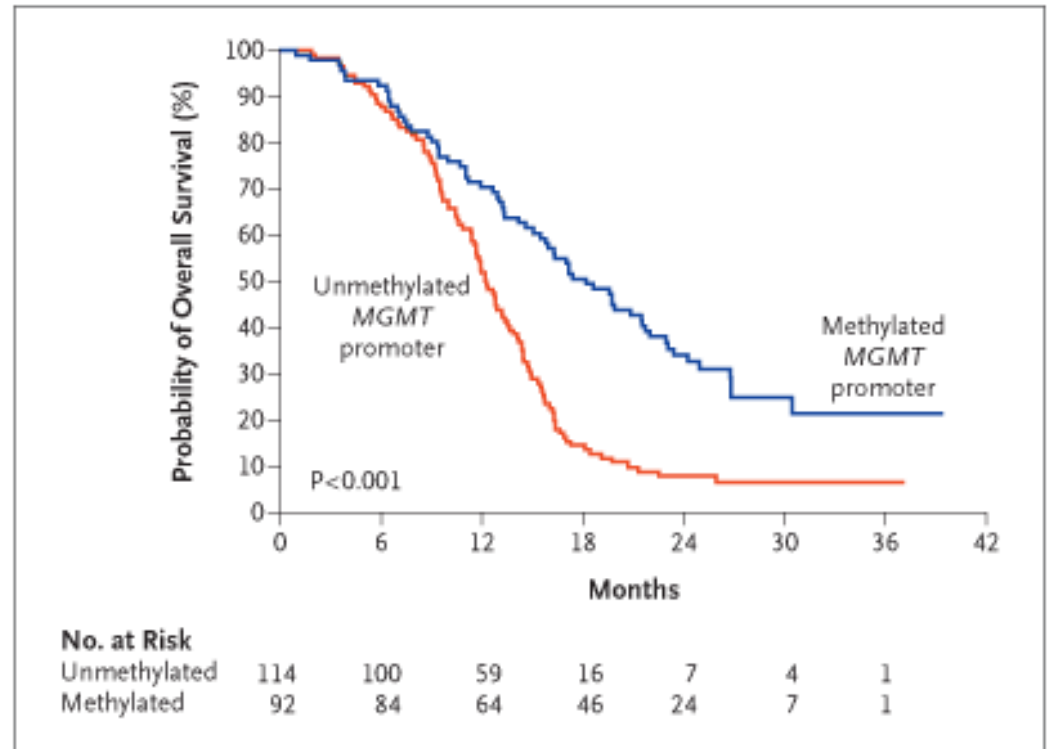
Background

- Cancer classification and diagnosis are becoming increasingly based on molecular features, or ‘biomarkers’
- With increasing focus on ‘precision medicine,’ it is critical to measure tumor markers on a population level
- Brain tumors are highly heterogeneous, and relevant biomarkers vary significantly by histology



BT Prognostic Biomarkers: Methylation of MGMT

- Hypermethylation at MGMT – responsive to alkylating agents (i.e. Temozolomide)
- More responsive to both chemo and radiation if methylated

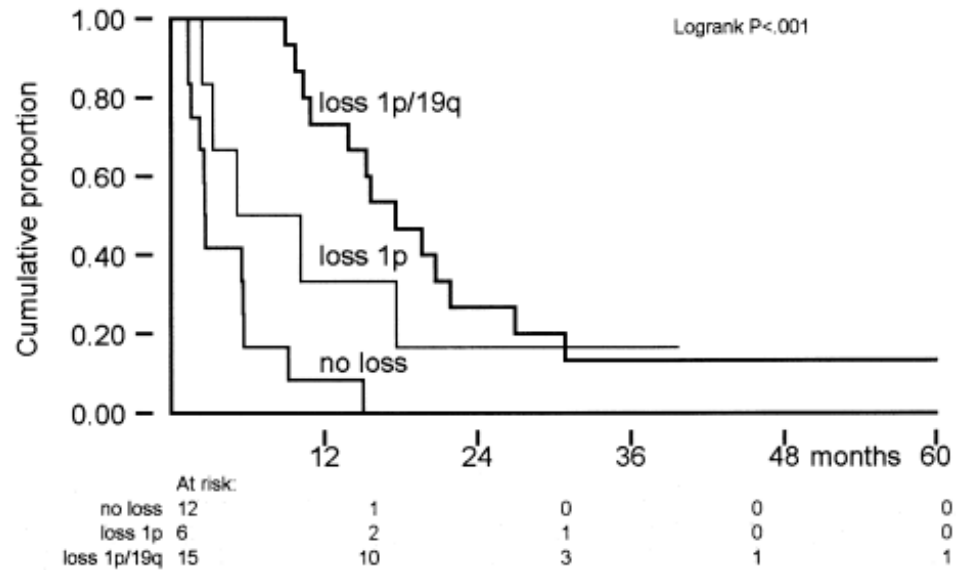


Hegi et al., NEJM 2005



BT Prognostic Biomarkers: Codeletion of 1p and 19q

- Concurrent loss of 1p and 19q is common in oligodendrogliomas
 - Also known as loss of heterozygosity (LOH)
- Collection is part of standard clinical care in LGG
- WHO has revised histologic criteria to include these markers as necessary for diagnosis of oligodendroglioma



Van den Bent et al, Cancer 2003



Are these markers complete at the population level?

- SEER registries have collected information on several tumor markers shown to be relevant in brain tumors since 2004
 - WHO grade (2004+)
 - MGMT Methylation (2008+)
 - 1p/19q deletion (2008+)
- Not **all** of these factors are relevant to **all** brain tumor histologies



AJCC Collaborative Staging Site-Specific Factors

- The AJCC Collaborative Staging system identified site-specific factors (SSF) for each cancer site that are relevant to determining stage of cancer
- Brain tumors are not staged, but AJCC includes several factors that are significantly associated with prognosis and survival in brain tumor



CoC and SEER Required Site-Specific Factors (SSF) for Collaborative Staging Version 02.04

Blue
Already coded in CSv01
Green
Additional clinically significant
White
Not required at this time

Schema Name	ICD-O-3 Primary Sites	SSF 1	SSF 2	SSF 3	SSF 4	SSF 5	SSF 6	SSF 7	SSF 8
Brain	C70.0, C71.0-C71.9	World Health Organization (WHO) Grade Classification	Ki-67/MIB-1 Labeling Index (LI): Brain	Functional Neurologic Status - Karnofsky Performance Scale (KPS)	Methylation of O6-Methylguanine-Methyltransferase (MGMT)	Chromosome 1p: Loss of Heterozygosity (LOH)	Chromosome 19q: Loss of Heterozygosity (LOH)	Surgical Resection	Unifocal vs Multifocal Tumor
Other CNS	C70.1, C70.9; C72.0-C72.5, C72.8-C72.9	World Health Organization (WHO) Grade Classification	Ki-67/MIB-1 Labeling Index (LI): Brain	Functional Neurologic Status - Karnofsky Performance Scale (KPS)	Methylation of O6-Methylguanine-Methyltransferase (MGMT)	Chromosome 1p: Loss of Heterozygosity (LOH)	Chromosome 19q: Loss of Heterozygosity (LOH)	Surgical Resection	Unifocal vs Multifocal Tumor
Intracranial Gland	C75.1-C75.3	World Health Organization (WHO) Grade Classification	Ki-67/MIB-1 Labeling Index (LI): Brain						



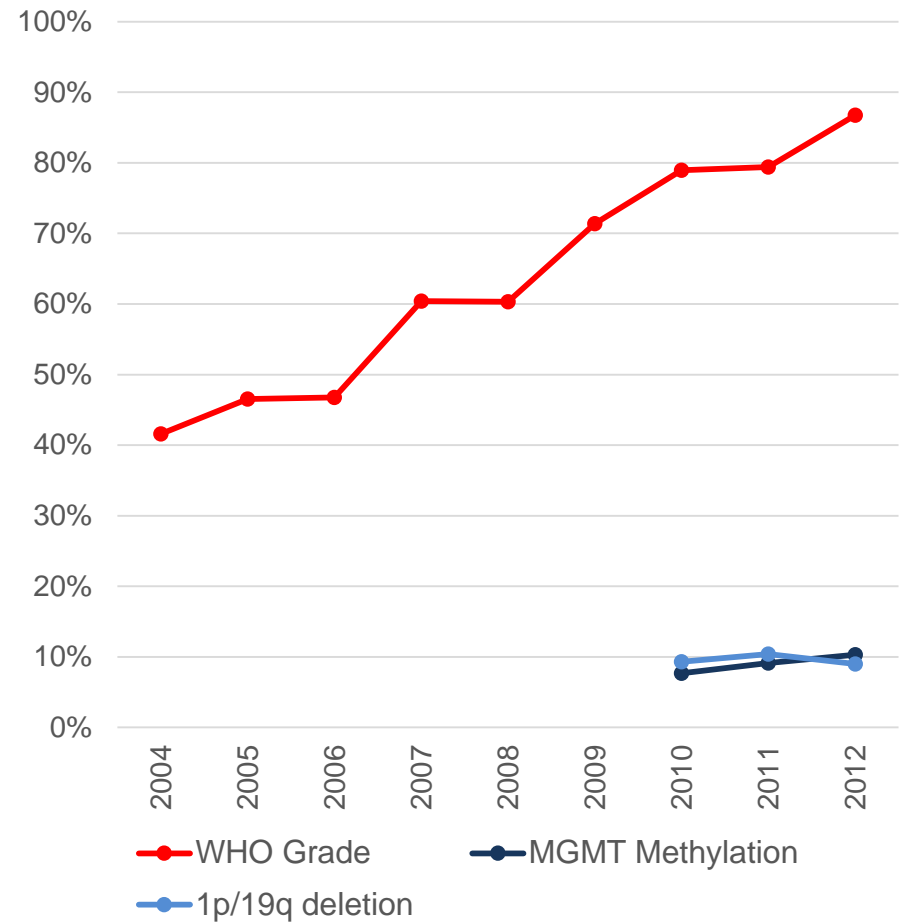
Methods

- Data were obtained via a SEER custom data request for four required factors
 - World Health Organization (WHO) grade
 - SSF
 - MGMT Methylation
 - 1p/19q loss of heterozygosity (LOH)



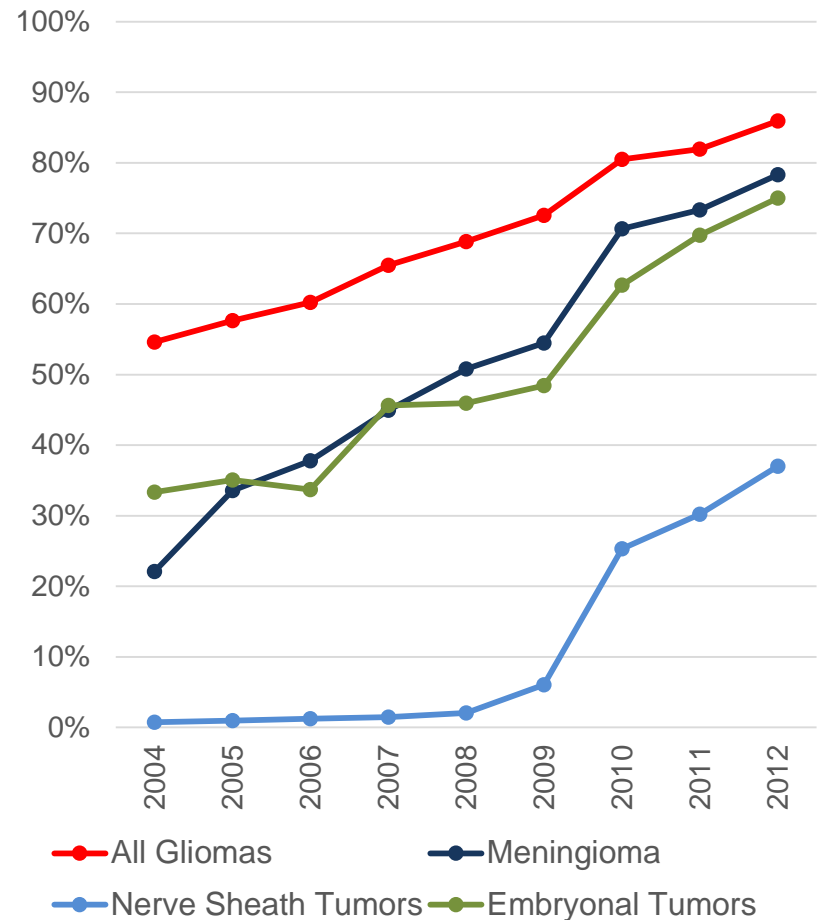
Overall Factor Completeness

- Overall percent complete:
 - WHO Grade: 46.5%
 - MGMT Methylation: 9.0%
 - 1p deletion: 9.5%
 - 19q deletion: 9.6%
- Completeness of all factors varied over time



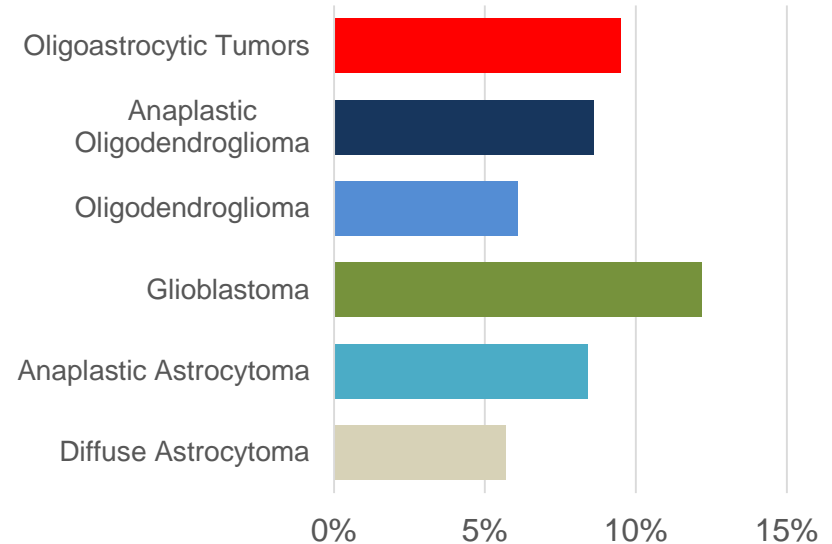
WHO Grade by Histology over Time

- WHO grade has been collected longest (2008-2012) and has shown significant increases over period of collection
- Completeness of WHO grade varied significantly by histology over time
 - Highest in gliomas



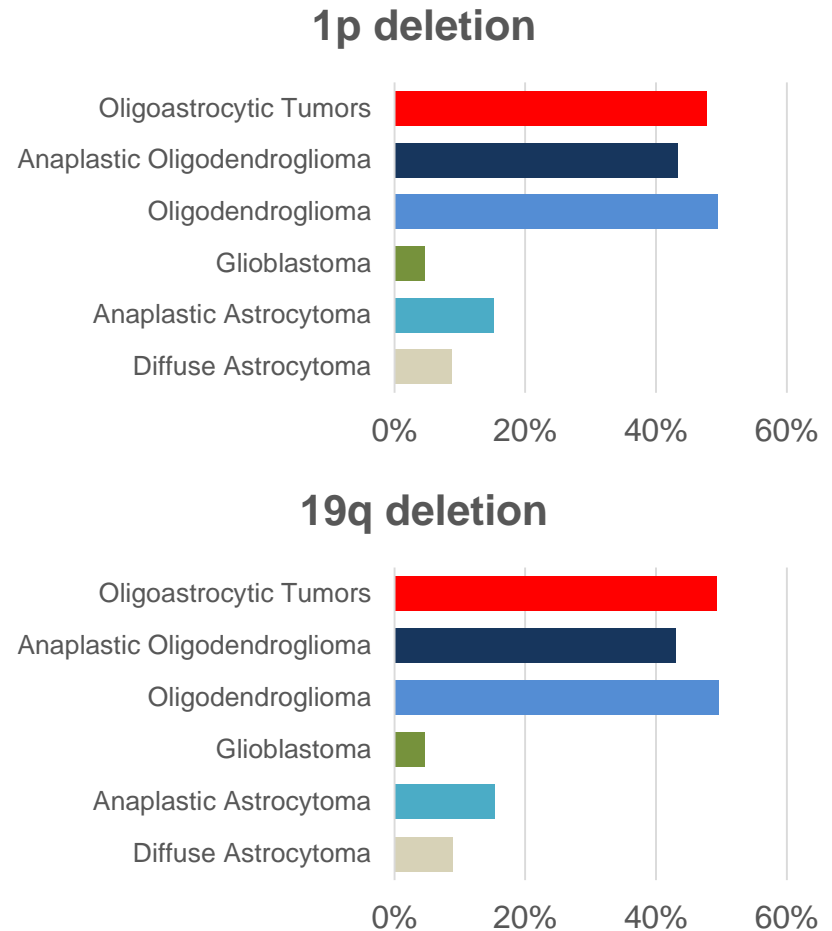
MGMT Methylation by Histology

- Completeness of MGMT methylation information varied significantly by histology
- Completeness was highest in histologies where factor is most clinically relevant
 - Glioblastoma



1p/19q Codeletion by Histology

- Completeness of 1p/19 codeletion status varied significantly by histology
- Completeness was highest in histologies where factor is most clinically relevant
 - Oligodendroglioma
 - Oligoastrocytoma



New WHO classification schema

- 2016 revision of WHO central nervous system tumor classification incorporates molecular markers
 - *IDH1/2* mutation
 - 1p/19q codeletion
- Cancer diagnosis is **increasingly molecular**, and it is **critical** that these data are incorporated in cancer registries

Diffuse astrocytic and oligodendroglial tumours	
Diffuse astrocytoma, IDH-mutant	9400/3
Gemistocytic astrocytoma, IDH-mutant	9411/3
<i>Diffuse astrocytoma, IDH-wildtype</i>	9400/3
Diffuse astrocytoma, NOS	9400/3
Anaplastic astrocytoma, IDH-mutant	9401/3
<i>Anaplastic astrocytoma, IDH-wildtype</i>	9401/3
Anaplastic astrocytoma, NOS	9401/3
Glioblastoma, IDH-wildtype	9440/3
Giant cell glioblastoma	9441/3
Gliosarcoma	9442/3
<i>Epithelioid glioblastoma</i>	9440/3
Glioblastoma, IDH-mutant	9445/3*
Glioblastoma, NOS	9440/3
Diffuse midline glioma, H3 K27M-mutant	9385/3*
Oligodendroglioma, IDH-mutant and 1p/19q-codeleted	9450/3
Oligodendroglioma, NOS	9450/3
Anaplastic oligodendroglioma, IDH-mutant and 1p/19q-codeleted	9451/3
<i>Anaplastic oligodendroglioma, NOS</i>	9451/3
<i>Oligoastrocytoma, NOS</i>	9382/3
<i>Anaplastic oligoastrocytoma, NOS</i>	9382/3

Louis et al, 2016



Molecular Markers and Cancer Registration

- Population-based cancer surveillance data are a **critical resource** for measuring the impact of cancer on a population
- Availability of complete and accurate molecular markers by histology is **essential** for measuring clinical impact on a population
 - IDH1/2 mutation
 - 1p/19q codeletion



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