

Differences in Biomarker Expression in GBM Patients with Early Progression after Surgery

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Purpose

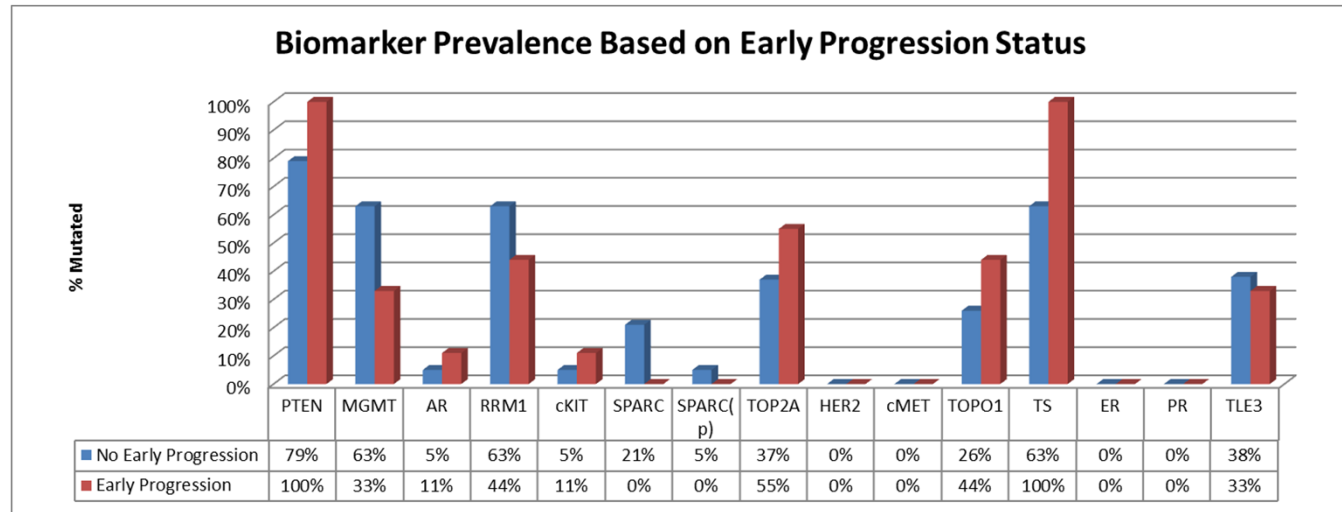
- Glioblastoma Multiforme (GBM) is locally aggressive
- GBM is the most common primary brain tumor in adults
- Surgery followed by temozolomide and radiotherapy are the standard of care
- Attempts have been made to increase radiation dose, pursue more aggressive chemotherapy, or add targeted therapy
- Biomarker expression may help individualize treatment decisions

Methods

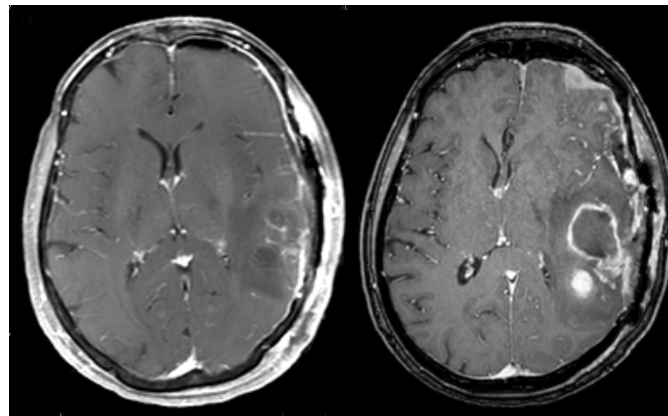
- 28 newly diagnosed GBM patients
- Treatment between January 2012 and May 2013
- Tumor profiling provided by Caris Life Sciences, Inc.
- Immunohistochemistry, FISH, CISH, MGMT promoter methylation and NextGen SEQ (Illumina TruSeq) were performed on formalin-fixed, paraffin-embedded tumor samples in a CLIA certified lab
- MRI imaging was performed on post-operative day 1 and at the time of CT simulation for radiation treatment planning
- Early Progression (EP) was defined as either new enhancing lesions or >25% increase in the T1 post-contrast enhancement.
- Two sided Fisher Exact Tests were used on contingency tables

Results

- Median time between MRI imaging 20 days
- 28.6% of patients had EP
- The genes more frequently expressed in EP group:
 - PTEN, p=NS
 - AR, p=NS
 - cKIT, p=NS
 - TOP2A, p=NS
 - TOPO1 p=NS
 - TS, p=0.06
- The genes more frequently expressed in the non-EP group
 - MGMT hypermethylation
 - RRM1
 - SPARC monoclonal
 - SPARC polyclonal



Example of Early Progression



Depicted above are A) the post-operative and B) Treatment planning MRI axial images revealing a new T1 post-contrast enhancing mass posterior to the resection cavity

Patient Characteristics			
Age at Diagnosis (years)		Location (n)	
Average	60	Frontal	18.00
Range	43-86	Parietal	7.00
Male (%)	26 (62)	Temporal	14.00
Female (%)	16 (38)	Occipital	1.00
Extent of Surgery		Basal Ganglia	3.00
Gross total (%)	27	Cerebellar	1.00
Subtotal (%)	73	Tumor Size (cc)	
Tumor Histology (n)		Average	35.9
Glioblastoma	39	Range	8.8-61.4
Gliosarcoma	2	Avg. FLAIR extent(cc)	110.2
Clear Cell Glioblastoma	1		

Conclusions

- There are different gene expression profiles in patients with EP
- This may lead to differing biology, aggressiveness and early treatment decisions
- This data will be confirmed in a validation set

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