



Mouse GBM: A Pilot Collection on NBIA

Sunny Jansen
Postdoctoral Fellow
Terry Van Dyke Lab
Mouse Cancer Genetics Program
NCI

Collaboration with Paul Mulhern and Brian Hughes

Introduction

Preclinical Goals



 Goal 1: Develop a molecular and imaging based classification for genetically engineered mouse models (GEMs) of glioblastoma multiforme (GBM)

 Goal 2: Use imaging to asses response of GBM GEMs to novel therapeutic regimens



Introduction

caBIG® Tools for Preclinical Data



NBIA

National Biomedical Imaging Archive

caArray

Gene expression microarrays



calntegrator

Integrate NBIA, clinical data (survival, IHC...), gene expression



Purpose



- Utilize NBIA and caIntegrator to correlate image-based features with 'clinical' data in GBM mouse models
 - Image-based features
 - Qualitative
 - Quantitative blood brain barrier permeability
 - 'Clinical' features
 - Genotype
 - Tumor histologic characteristics
 - IHC— e.g., CD31
- One time imaging in n=62 mouse patients



Animal models



 Intracranial injection of cell lines derived from several GEMs

Genotypes:

- Rb inactivation + Ras activation (TR)
- Rb inactivation + Ras activation + PTEN loss (TRP)
- Rb inactivation + Ras activation + EGFR loss (TRE)

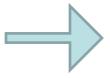


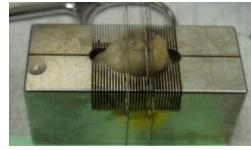
Brain Imaging Workflow











Four mice imaged at once

Philips Intera Achieva 3.0T MRI clinical scanner ~1hour imaging time

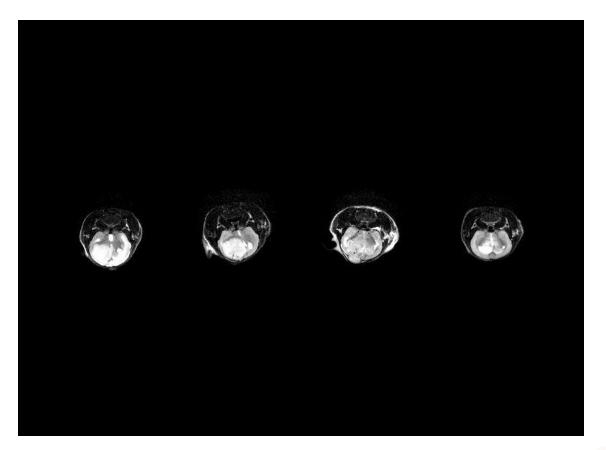
Post imaging euthanize and histology



MRI Pulse Sequences



T2 weighted pre contrast

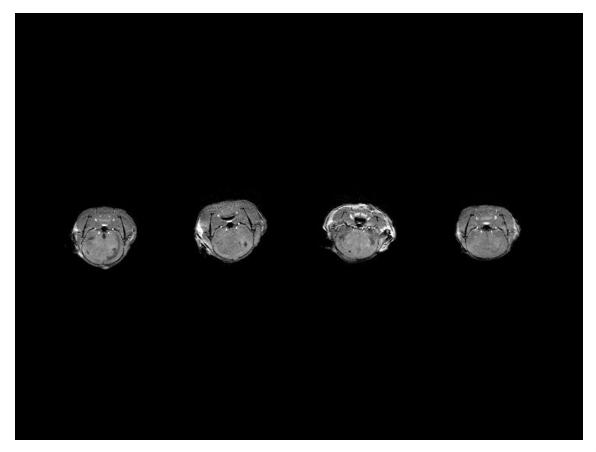




MRI Pulse Sequences



T1 weighted pre contrast

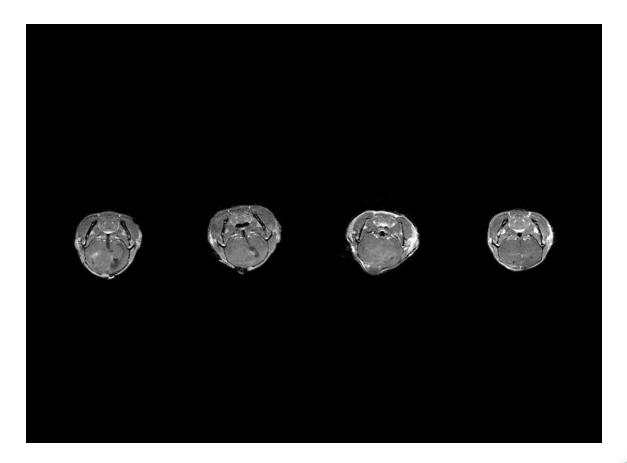




MRI Pulse Sequences



T1 weighted post contrast

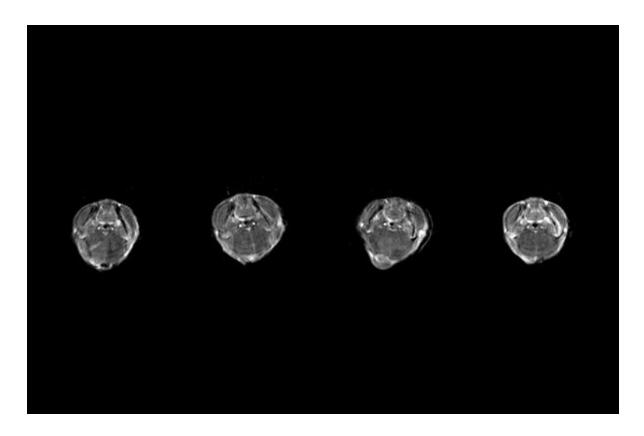




MRI Pulse Sequences



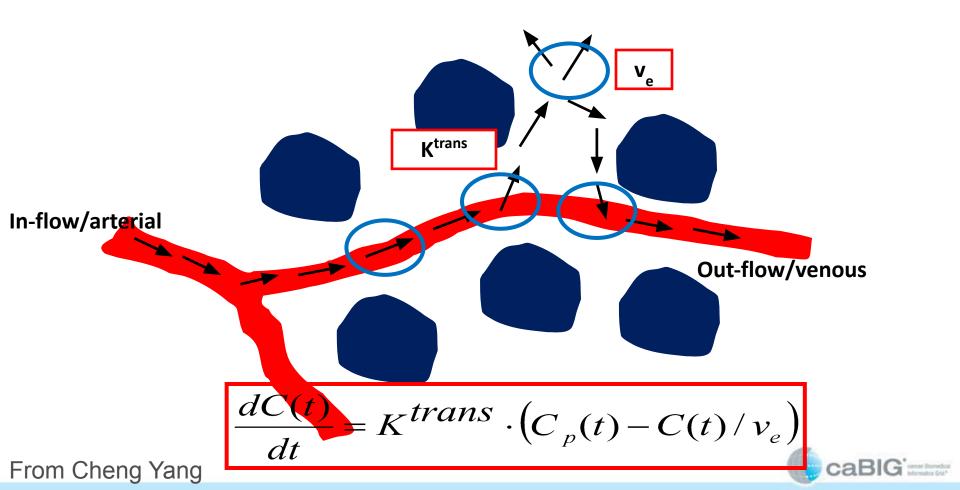
Dynamic contrast enhanced series



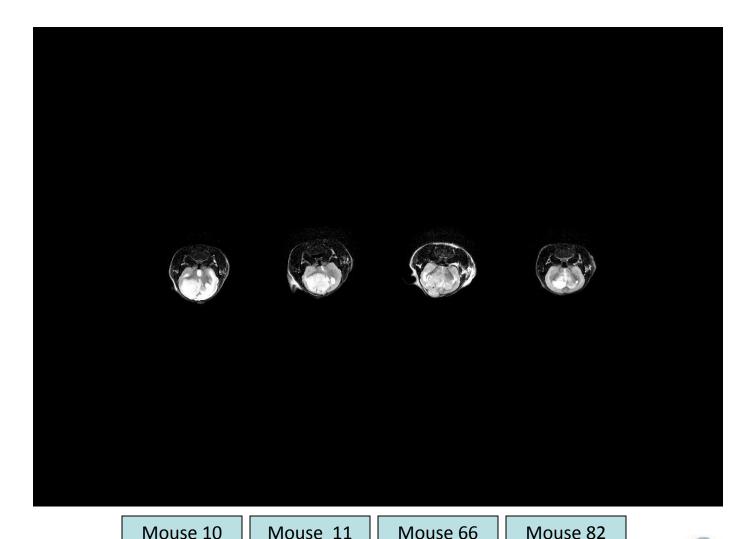


DCEMRI and two compartment model parameters

 Quantifiable information on the blood brain barrier permeability is extracted from the pattern of contrast agent uptake



Challenge: Cropping and splitting the image data



caBIG' unun Bernedent

Challenge: Cropping and splitting the metadata



DICOM header

```
(O10,0010) PN [SJ_DCE_10_11_66_82] # 18, 1 PatientsName
(0010,0020) LO [SJ_DCE_10_11_66_82] # 18, 1 PatientID
(0010,0030) DA [20100928] # 8, 1 PatientsBirthDate
(0010,0040) CS [M] # 2, 1 PatientSSex
```

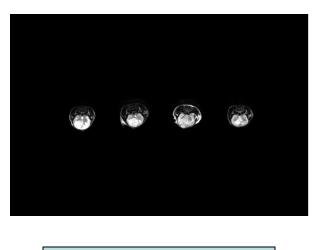
```
(0018,9089) FD 0\0\0
(0020,000d) UI [1.3.46.670589.11.17169.5.0.4584.2011021511013196216] # 52, 1 StudyInstanceUID
(0020,000e) UI [1.3.46.670589.11.17169.5.0.4564.2011021511582801965] # 52, 1 SeriesInstanceUID
(0020,0010) SH [350910091] # 10, 1 StudyID
```

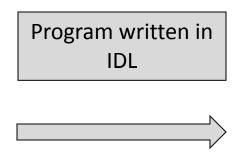
```
(0028,0010) US 1024 # 2, 1 Rows
(0028,0011) US 1024 # 2, 1 Columns
```

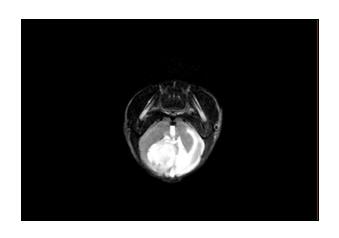


Cropping and splitting the image data









10 11 66 82

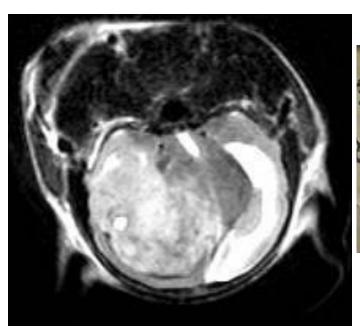
Mouse 10

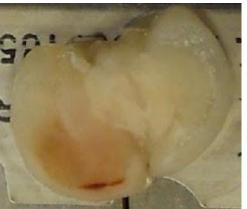
Advantage: Image arrays 16 times smaller

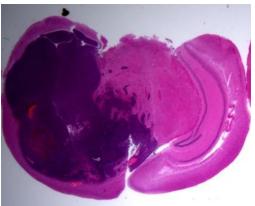


Cropping and splitting the image data











Cropping and splitting the metadata



DICOM header

```
(0010,0010) PN
                  [SJ_DCE_10_11_66_82]
                                                                       18, 1 PatientsName
                  [SJ_DCE_10_11_66_82]
(0010,0020) LO
                                                                       18, 1 PatientID
(0010,0030) DA
                  [20100928]
                                                                        8, 1 PatientsBirthDate
(0010,0040) CS
                                                                         1 PatientsSex
(0020,000d) UI [1.3.46.670589.11.17169.5.0.4584.2011021511013196216] # 52, 1 StudyInstanceUID (0020,000e) UI [1.3.46.670589.11.17169.5.0.4564.2011021511582801965] # 52, 1 SeriesInstanceUID
                                                                         # 10, 1 StudyID
(0020,0010) SH [350910091]
(0028,0010) US 1024
                                                                                2, 1 Rows
(0028,0011) US 1024
                                                                                2. 1 Columns
```



Program written in IDL

```
[TVD_GBM_IC2_11311010]
 0010,0010) PN
                                                            20, 1 PatientsName
(0010,0020) LO
               TVD_GBM_IC2_11311010]
                                                            20, 1 PatientID
(0010,0030) DA [20100928]
                                                             8, 1 PatientsBirthDate
(0010,0040) CS [F]
                                                             1 PatientsSex
(0020,000d) UI
               [1.3.46.670589.11.17169.5.0.4584.2011021511013196216.1] #
                                                                           54, 1 StudyInstanceUID
               [1.3.46.670589.11.17169.5.0.4564.2011021511582801965.1] #
                                                                           54, 1 SeriesInstanceUID
(0020,000e)
(0028,0010) US 256
                                                             2, 1 Rows
                                                             2, 1 Columns
(0028,0011)
           US 256
```



Mouse GBM Pilot Collection Deployment to CBIIT instance of NBIA

Demo: Mouse GBM on NBIA

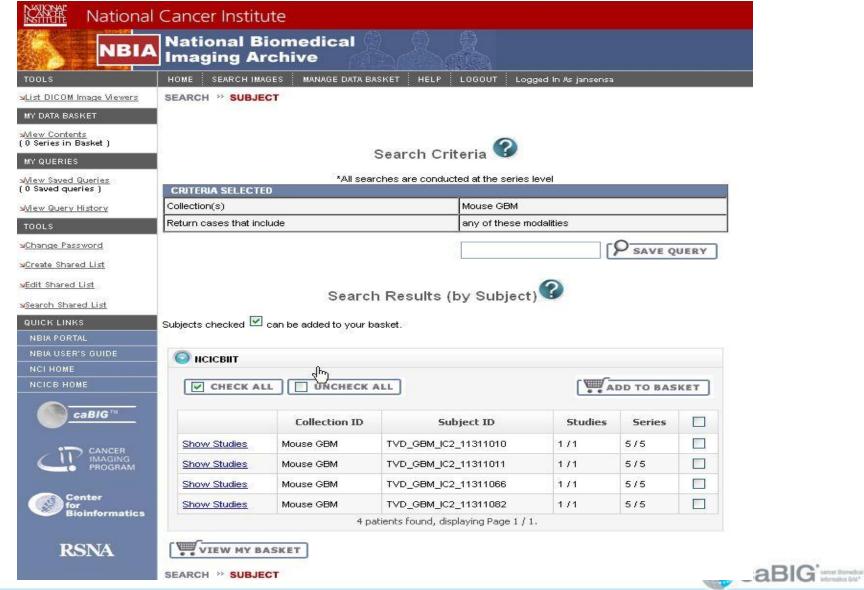




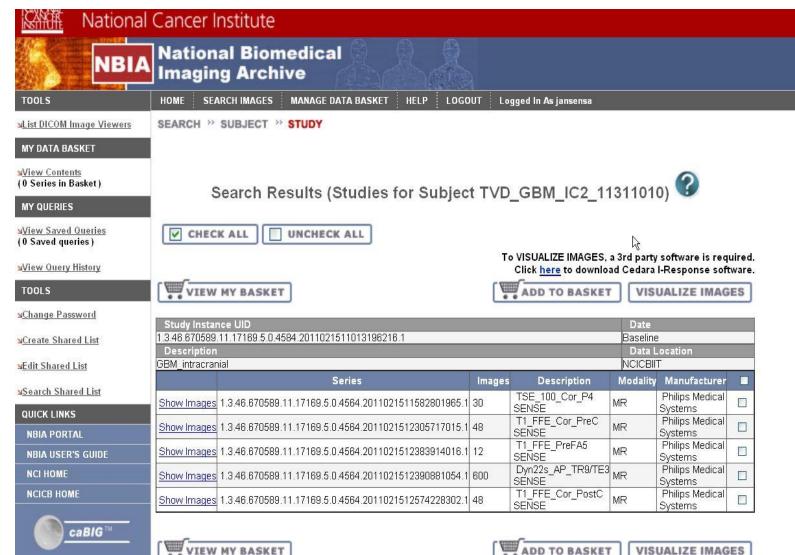
Available on NBIA (mm/dd/yyyy) Series Includes Annotations W Annotated Non-Annotated Subject ID(s)	Collection(s) Collection Descriptions		Selected Mouse GBM ELECT ALL EMOVE ALL
Annotations Mon-Annotated Non-Annotated			То:
Subject ID(s)		Annotated Non-Annotated	
	Subject ID(s)		I
			SUBMIT [





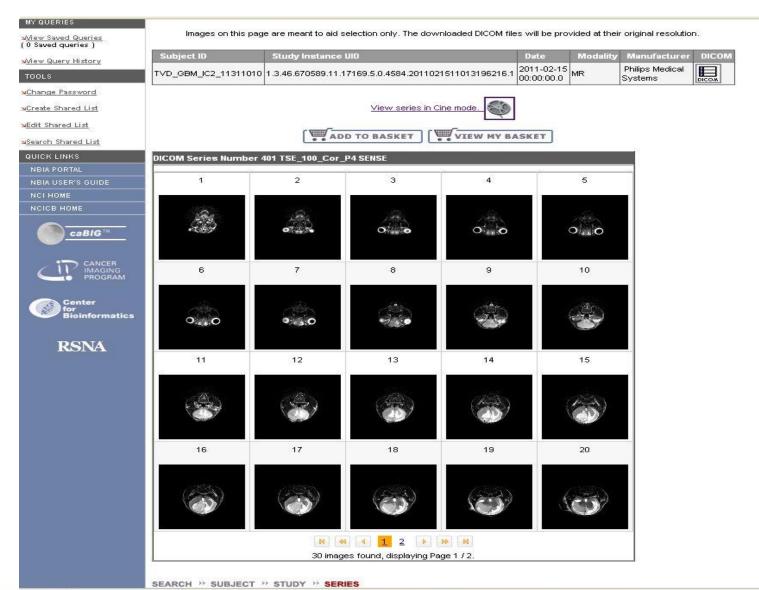












Summary



- Progress to date: Successfully deployed individual mouse GBM MRI datasets to NBIA
- Next steps
 - Set up own instance of NBIA
 - Qualitative and quantitative image analysis
 - AIM features
 - K^{trans}, v_e
 - Correlation of image-based features with genotype, histology, pathology
 - caIntegrator



Mouse GBM Pilot Collection Deployment to CBIIT instance of calnt

Demo: Mouse GBM on calntegrator



Summary



Future directions

- caArray
- Center for Advanced Preclinical Research (CAPR)
- Mouse Mammary pilot collection and image-based prognostic markers

