

1. **Glioma Imaging Analysis pipeline based on Histological objects generation, resulting to histogram of Glioma images classification**
2. **A Genomic linkage of DSA-UI may validate this classification of gbm/lgg imaging use case, using mRNA gene expression.**

Pythonic Neuro Image Analysis pipeline to generate object detection

1. Generate random Tiles from selected GBM-SVS slide (grabTiles\_forTraining\_gbm.py), by doing
  - a). Simple Masking on lowResPILImage
  - b). Getting intended tiles using Girder calls on candygram server
2. Standard Colour deconvolution using HistomicsTK to generate Hematoxylin-stain, Eosin-stain images from the given tile
3. Generate histologic components or histologic objects by quantizing pixels in the above H-Stain and E-Stain channel into following groups (approximation)
  - hp - pixel value in h-channel
  - ep - pixel value in e-channel
  - havg - Average pixel values in h-channel
  - eavg - Average pixel values in e-channel
  - Label pixel p as purple if  $hp \leq havg$
  - pink (if  $hp > havg$  and  $ep \leq eavg$ )
  - white if  $hp > havg$  and  $ep > eavg$ 
    - a. Purple Pixels - Cell Nuclei material
    - b. Pink pixels - Stroma, Stomal cells' cytoplasms, mucin-poor epithelial cells's cytoplasms
    - c. White pixels - Lumina, mucin-rich epithelial cells's cytoplasms
4. Using circle fitting algorithm generate all the possible local histological objects.
 

[https://scipy-cookbook.readthedocs.io/items/Least\\_Squares\\_Circle.html](https://scipy-cookbook.readthedocs.io/items/Least_Squares_Circle.html)

  - a. Initially linear algebraic method will be used in implementing circle fitting algorithm

[https://dtcenter.org/met/users/docs/write\\_ups/circle\\_fit.pdf](https://dtcenter.org/met/users/docs/write_ups/circle_fit.pdf)

Based on algorithm complexity analysis, Jacobian method can be tried later
5. Using Bag of Words, get the frequency of these objects with the help of k-means and classify tissue images
 

(cancerous vs non-cancerous, gbm vs lgg) using SVM

Referenced publication (Local Object Patterns) has an accuracy 93.03% of representation and classification of pathology images of Colon tissue stained with H&E.

## Reference

### Image Analysis Pipeline :

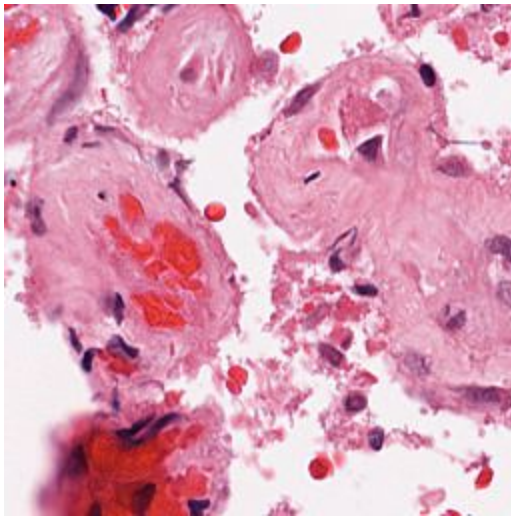
1. Local Object Patterns for the Representation and Classification of Colon Tissue Images, Gulden Olgun, Cenk Sokmensuer, and Cigdem Gunduz-Demir, IEEE JOURNAL OF BIOMEDICAL AND HEALTH INFORMATICS, VOL. 18, NO. 4, JULY 2014

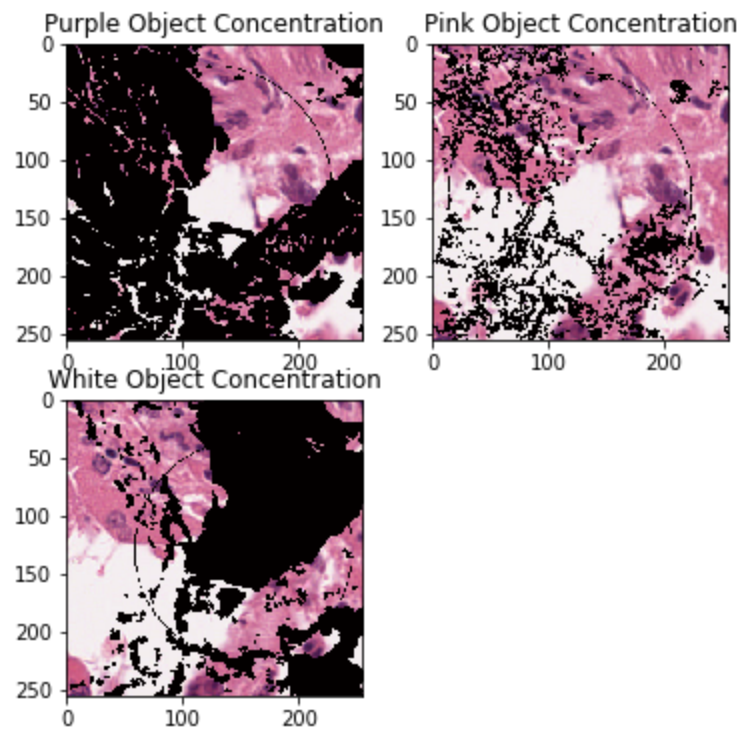
### Glioblastoma pathogenesis:

1. Discovery and validation of a glioblastoma co-expressed gene module, Leland J. Dunwoodie<sup>1</sup>, William L. Poehlman<sup>1</sup>, Stephen P. Ficklin<sup>2</sup> and Frank Alexander Feltus<sup>1</sup>

### Sample Tile:

/media/raj/Raj1\_5/10ktilles/train/gbm/TCGA-02-0001-01Z-00-DX2\_20x\_7744\_32416\_256x256.png





All the Purple, Pink and White Objects are marked in Black in the above images.

**Largest Object concentration, by applying k-means clustering and circle-fit algorithm**

