**First program to be executed is**

writeImage.m

**input dataset** - D:\subjectsHandled\B.E-Report\2017-18\VIII-Sem-Project\Kirupa-Keerthana-kavya\Project\ALLDataSet\ALL\_IDB1

**output** - D:\subjectsHandled\B.E-Report\2017-18\VIII-Sem-Project\Kirupa-Keerthana-kavya\Project\Biomarkers for the Identification of Acute Leukemia (April '18)\Code\Udata

MKmeans.m

==========

- calls MKmeanscl.m

MKmeanscl.m

==========

- calls DoWatershed.m

**Deep Learning**

**UpdatedKeras.py –**entire deeplearning process. Final output we will get

VisualizeFeatures.py – same as UpdatedKeras.py. It contains additional codings to display the intermediate values in each layer

**Total layers – 3 excluding fully connected**

**In each layer 3 sublayers- convolution,activation, maxpooling**

**Totally – 9 layers numbered from 0 to 8**