* Load all HDAB images
* Set deconvolution stains consistently across all
* Create full image annotation
* Train a pixel classifier
  + Create a training image composed of multiple representative sections, for classes tumour, necrosis, and ignore
* Save pixel classifier
* Load pixel classifier, create objects on full image
  + Create objects, new object type = annotation, minimum hole size 20 000, split objects, delete existing objects, create objects for ignored classes
* Run **reclassify small objects to best neighbour.groovy**
  + Sets class of small objects to tumour
  + (need to implement) merges all annotations of same class together
* Perform manual annotation corrections as needed
* Add intensity features
  + Preferred pixel size can be the resolution of Visium. Also do another measurement calculation at whole image res to compare values
  + Include hematoxylin, DAB, OD sum, mean, stdev, median
* Cell detection