**Notes for analyzing 386 well images from Laura:**

* CellProfiler is pretty weak,
  + Good for organizing data
  + Maybe good for pulling in images from Omero
  + Check out new version
* Matlab
  + Reading images and running imadjust gives a good starting point
  + Denoising with median filter doesn’t seem helpful
  + Edge detection + fill
    - Only useful for nuclei
    - ‘Canny’ and ‘log’ (Laplacian of Gaussian) both get nuclei that the other misses
      * Log or ‘zero cross’ seem equivalent and get the most nuclei
      * It would probably be best to combine these two
    - Check what CellProfiler does (it seems better).