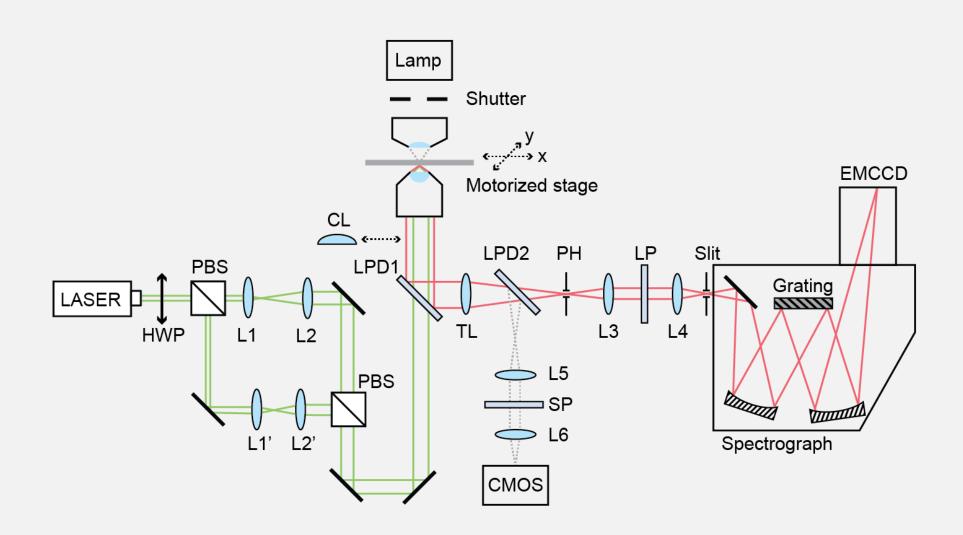
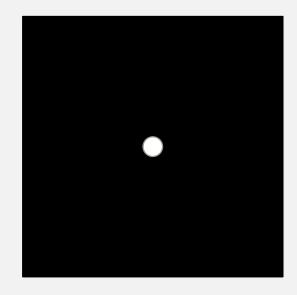
OPTICS SCHEMATICS

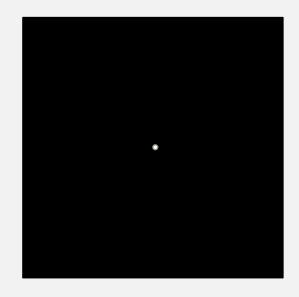


THREE ILLUMINATION PATTERNS



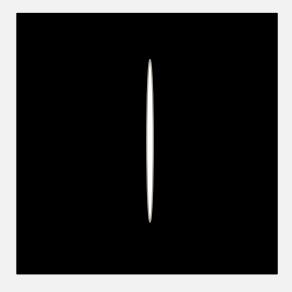
Single cell illumination

- ~8um
- Ideal for single cell measurement
- covers I/3 of a cell
- 300um pinhole



PSF illumination

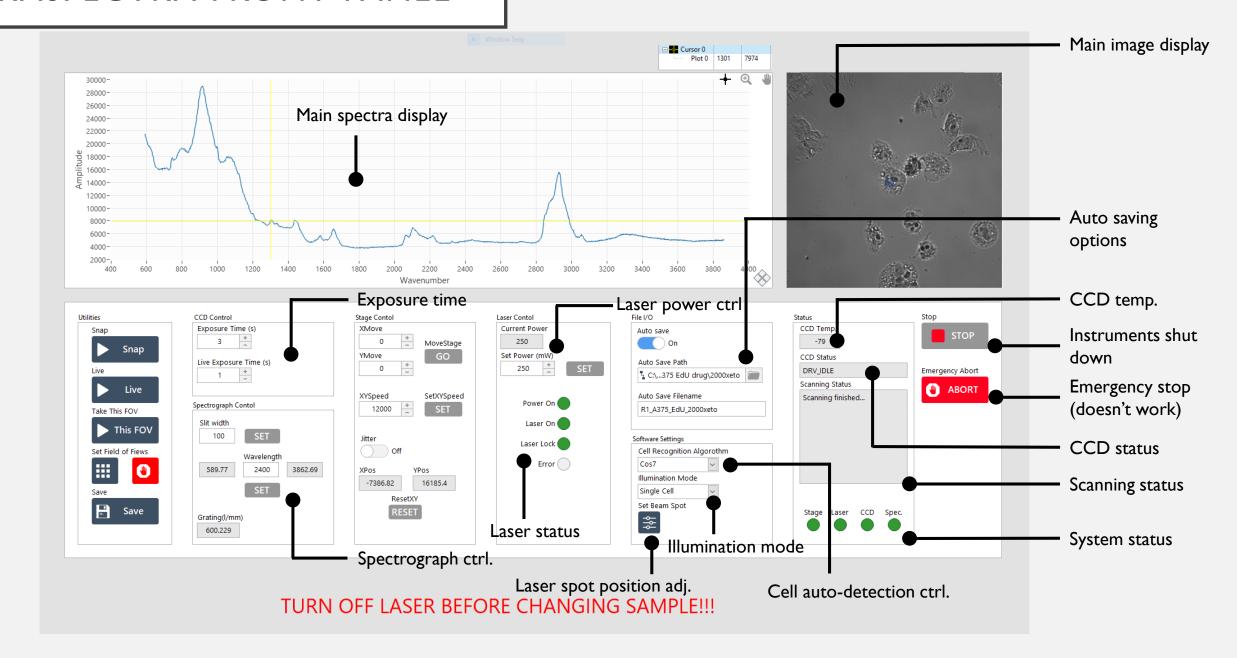
- ~300nm
- Ideal for scanning



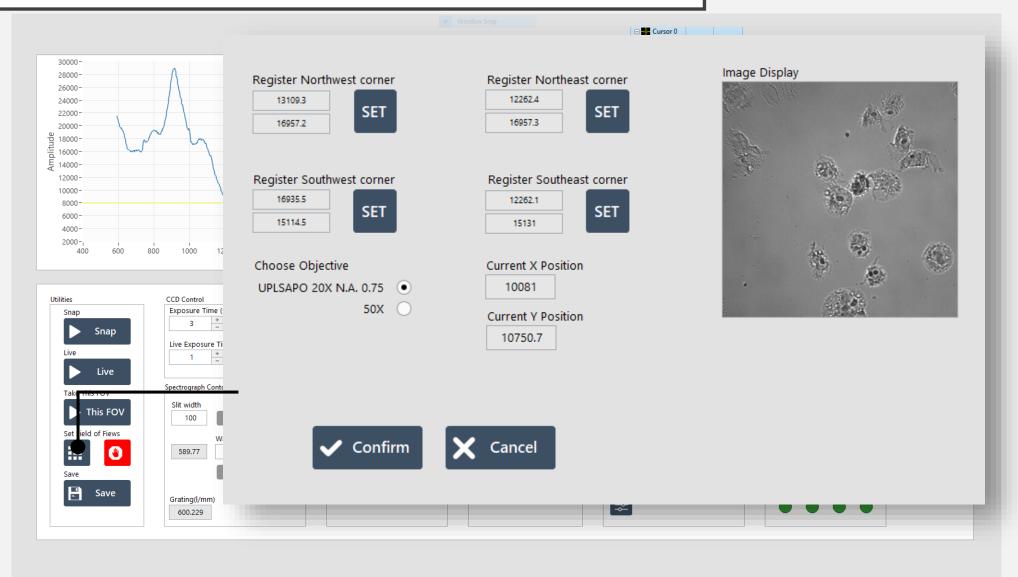
Linear illumination

- ~8 by 80um
- Ideal for bulk measurement
- Optically 'averages' multiple cells
- Without pinhole

RASPECTRA FRONT PANEL

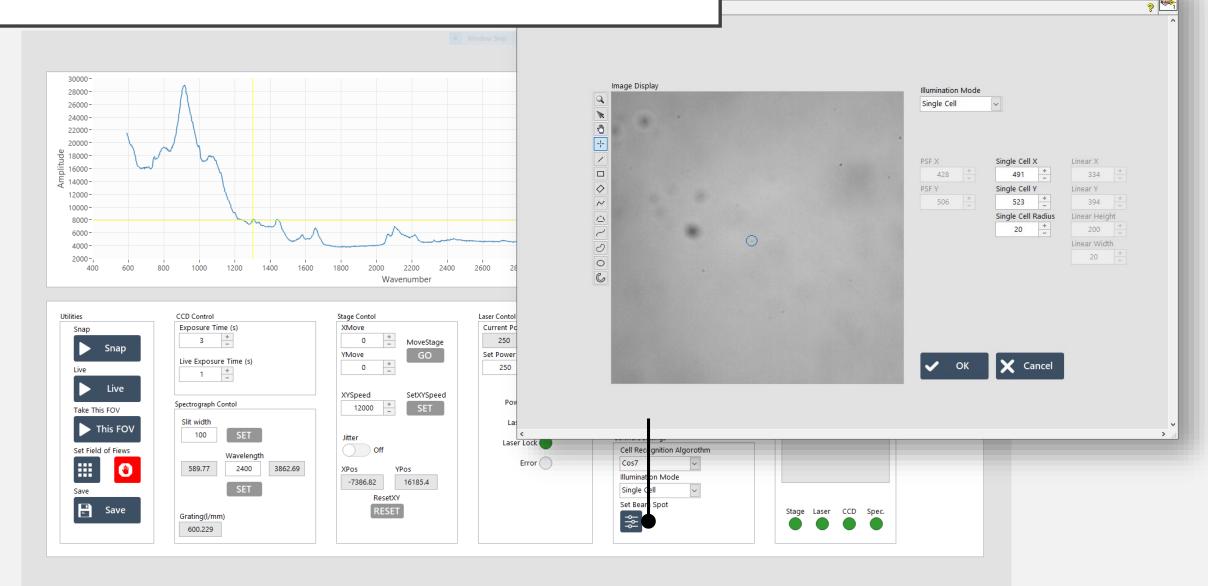


RASPECTRA SETTING SCANNING RANGE



TURN OFF LASER BEFORE CHANGING SAMPLE!!!

RASPECTRA SETTING BEAM POSITION



TURN OFF LASER BEFORE CHANGING SAMPLE!!!

SPECR 3.0 CLASS

```
Class specr
Class specr
                                                  -methods
    -properties
         .data
                                                      %% data management
                                                      .specr()
                           %wavenumber
              .wavenum
                                                      .importdata()
              .spc
                           %spectral data
                                                      %% plots
              .bgcor
                           %fitted bkgrd
                                                      .plot()
              .removed
                           %removed data
                                                      .plotMean()
         .label
                                                      %% preprocessing
              .propertyName
                                                      .removeBg()
              .propertyValue
                                                      .removeOutliner()
              .filename
                                                      .removeLowSNR()
              .cellType
                                                      .trim()
              .probes
                                                      .normalize()
              .treatment
                                                      .calibrate()
              .history
                                                      .delete()
         .stats
                                                      %% utilities
              .spcMean
                                                      .setLabel()
              .spcStd
                                                      .getMean()
              .spcResolution
                                                      .getSNR()
              .spcSNR
                                                      .getNum()
              .cellNum
```

SPECR 3.0 FUNCTIONS

Auxiliary functions:

```
%% core functions
spcinitialize() %initialize properties fields
                %select data with GUI
spcuiimport()
                %open files with GUI
spcuiopen()
spcsort()
                %sort spectra
                %find ID using regexp
spcgetid()
spcisfield()
                %if input is a valid field
spcfindregion() %find index given wavenumber
                %basic function for normalization
spcnormalize()
                %basic function for trimming
spctrim()
spcknneval()
                %evaluate accuracy with knn model
%% data management
                %combine two specr objects
spcmerge()
spcmergedata() %combine two specr objects
spcsplitdata() %split data for training and testing
spcstack
                %stack arrays
%% bkgrd correction
                %fit baseline with various model
spcbgsub()
%% plots
                %plot spectral data
spcplot()
spcmarkpeaks() %mark common peaks on the plot
spcploterrorellipse() %mark 2D error bar
%% calibration
                %fit peaks with various model
spcpksfit()
spccalibstd()
              %calibrate with standard samples
spcfindstdpks() %detect peaks with parameters
```