



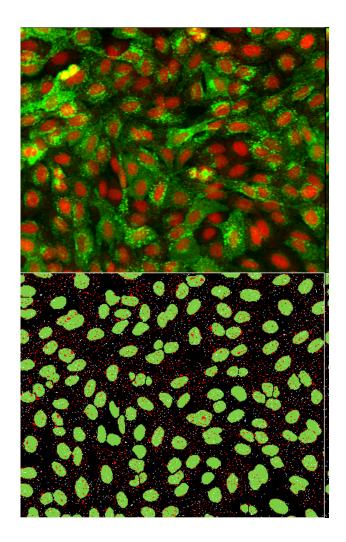
Together through life sciences.

#### MetaXpress® Software: Granularity Module

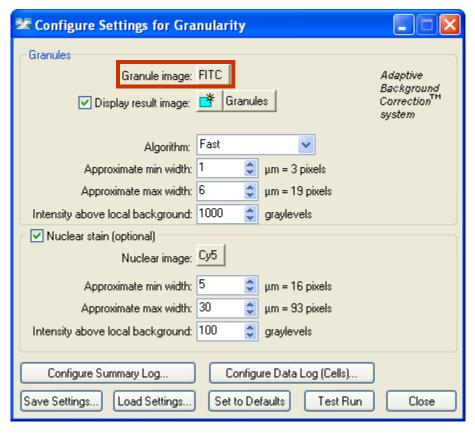


#### **Granularity Module Overview**

- The Granularity module can be used to analyze the number and intensity of granules (spots) per image and per cell
- This module does not require a nuclear wavelength.
- A nuclear stain (e.g. DAPI, Hoechst, or DRAQ5) is required to determine the number of granules per cell.

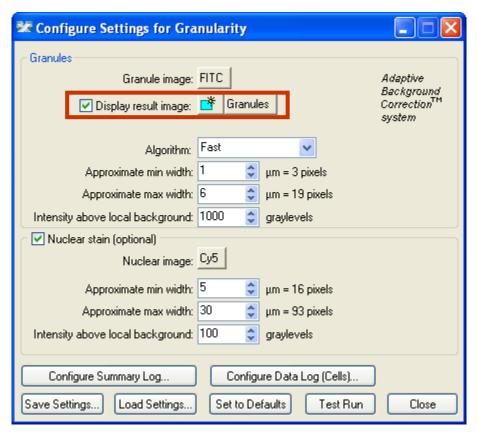




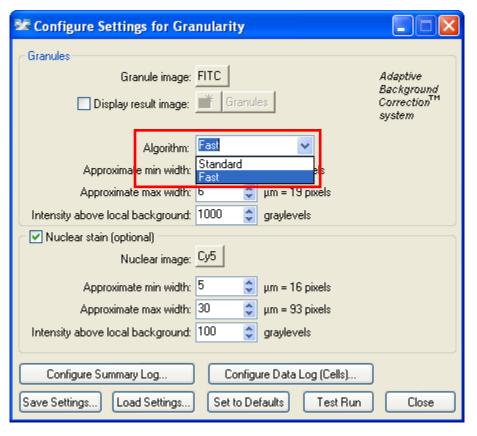


- Granule Image
- Select the image with granules here





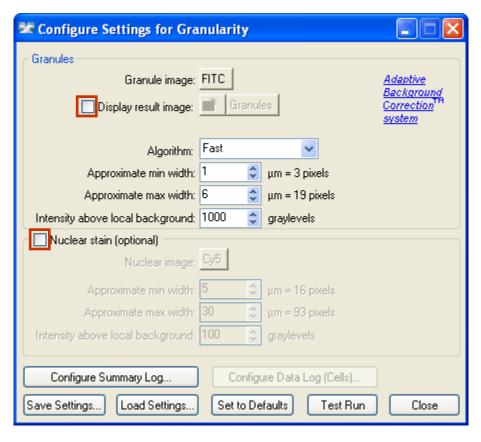
- Display result image
- Leave "Display result image" deselected (this is generally only used when journaling)



#### Algorithm

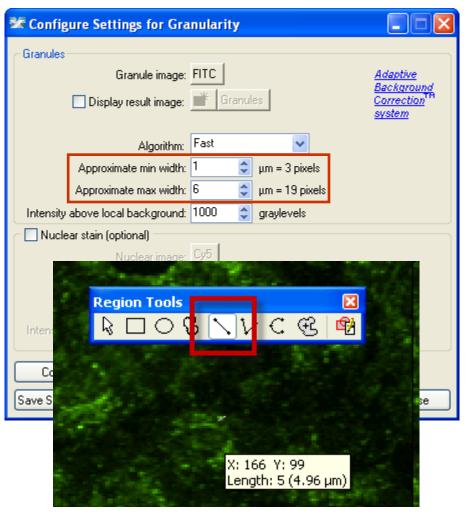
- This option is only available in MetaXpress software version 4.0 and higher and determines how quickly the analysis is performed.
- Fast algorithm can perform analysis up to twice as fast as Standard.
- Both algorithms produce similar but not identical results.





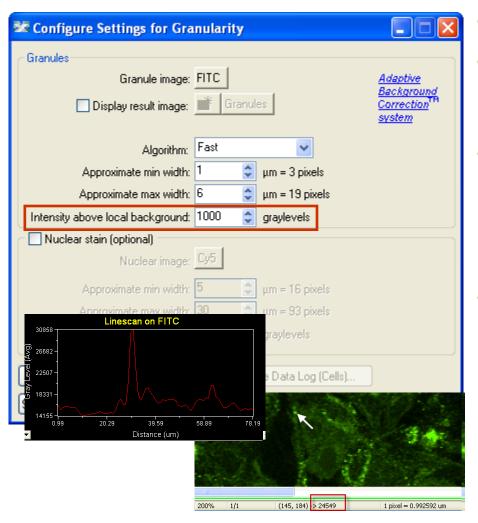
- Granules
- Deselect Nuclear stain





#### Granules

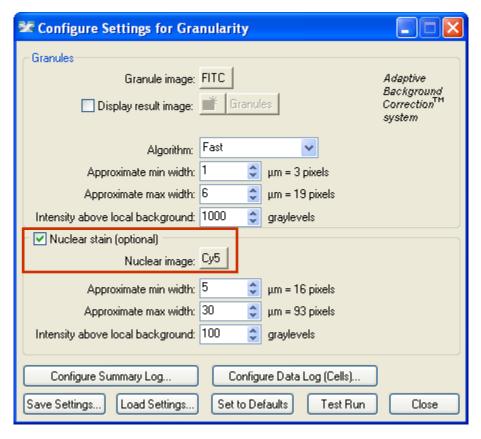
- Using the region tools measure the appropriate min (minimum) and max (maximum) width of qualifying granuless.
- Much smaller granules will be ignored
- Much larger granules will be split



#### Granules

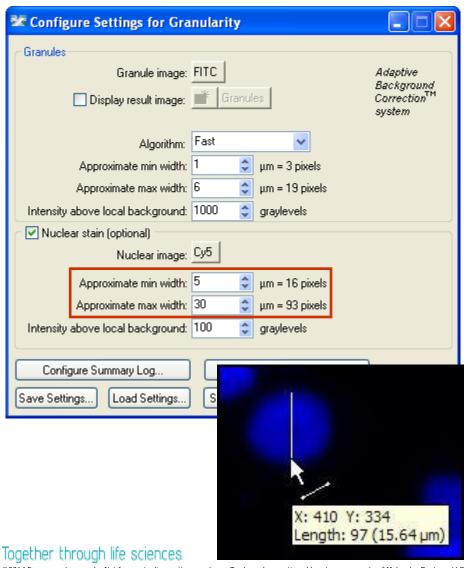
- The intensity above local background is used for finding the granules
- This value is a minimum and should be set slightly lower than the difference in intensity between a dim granules and its local background
- Draw a line across a cell into the background and use Measure > Linescan to determine intensity values; or simply mouse over the granules and the background and view the intensity values





- Nuclear Stain (not required)
- Select tick mark
- Select the wavelength for the nuclear image

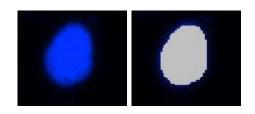




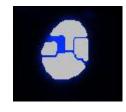
- **Nuclear Stain**
- **Set the Approximate min** width and Approximate max width for the range of nuclei that you want to detect
- The width is the short axis of a nucleus (in um)
- The region tools can be used to measure widths
- Much smaller cells will be ignored
- Much larger cells will be split



#### **Effects of varying width settings**

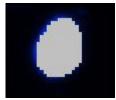


Min width too small: splits nuclei



Min width too large: omits smaller nuclei

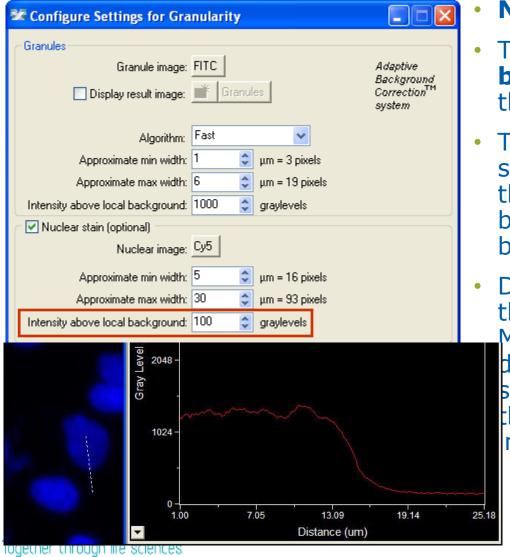
Max width too small: may shrink nuclear boundaries



Max width too large: may slightly enlarge nuclear boundaries

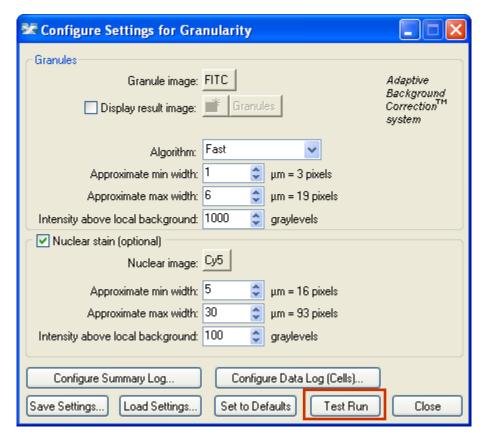






Nuclear Stain

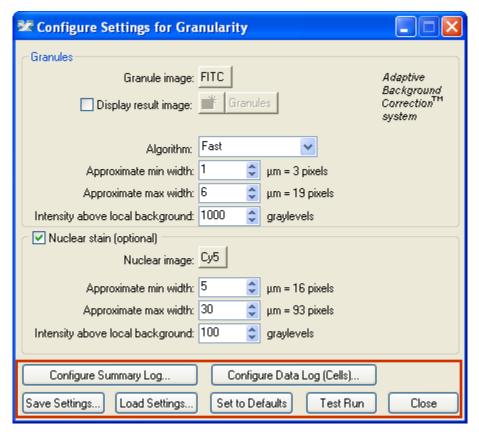
- The intensity above local background is used for finding the nuclei
- This value is a minimum and should be set slightly lower than the difference in intensity between a dim cell and its local background
  - Draw a line across a cell into the background and use
    Measure > Linescan to determine intensity values; or simply mouse over the cell and the background and view the ntensity values



- Select Test Run to view the cell segmentation
- Change settings if needed
- Save the settings



#### **Module Settings – General Settings**



- Configure Summary Log
   select site-by-site
   measurements
- Configure Data Log select cell-by-cell measurements
- Save Settings save analysis parameters to database
- Load Settings load saved analysis parameters
- Set to Defaults restore default analysis parameters
- Test Run test all settings together and display cell-bycell results for this site



## **Summary Data (site-by-site measurements)**

- Granules
- Granules Per Cell
- Total Granule Area
- Mean Granule Area
- Integrated Granule Intensity
- Average Granule Intensity
- Nuclei
- Total Nuclear Area
- Mean Nuclear Area
- Integrated Nuclear Intensity
- Average Nuclear Intensity
- Texture Index
- Cellular Texture Index
- Gradient Index
- Cellular Gradient Index
- Laplacian Index
- Cellular Laplacian Index

- Granules: Total number of granules
- Granules Per Cell: Total number of granules divided by the total number of nucleus
- Total Granule Area: The total area of the granules found in the image (in um<sup>2</sup>)
- Mean Granule Area: The total area of granules for all cells divided by the total number of nucleus (in um<sup>2</sup>)
- Integrated Granules
   Intensity: The total pixel intensity of the granules area
- Average Granules Intensity: The total pixel intensity of the granules area divided by the total number of nucleus



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- Texture Index
- Cellular Texture Index
- Gradient Index
- Cellular Gradient Index
- Laplacian Index
- Cellular Laplacian Index

- Nuclei: Total number of nuclei (cell count)
- Total Nuclear Area: The total area of the nucleus for all cells found in the image (in um<sup>2</sup>)
- Mean Nuclear Area: The average area of nucleus for all cells found in the image (in um<sup>2</sup>)
- Integrated Nuclear Intensity:
   The total pixel intensity of the nuclear stain over the nuclear area
- Average Nuclear Intensity: The total pixel intensity of the nuclear stain over the nuclear area, divided by the total number of cells



#### **Summary Data (site-by-site measurements)**

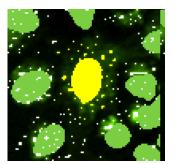
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- 🗸 Cellular Laplacian Index

- Texture Index: Standard deviation of intensity values in the image.
- Cellular Texture Index: Cell-by-cell standard deviation of intensity values near the nuclei. (Requires use of nuclear stain).
- Gradient Index: A texture-dependent measurement that reflects the amount of local intensity contrast. Measures the difference between the maximum and minimum intensity within a local neighborhood.
- Cellular Gradient Index: Cell-by-cell Gradient Index measured near the nuclei. (Requires use of nuclear stain).
- Laplacian Index: Similar to the morphological gradient, however this morphological measurement reflects fluctuations in the gradient.
- **Cellular Laplacian Index:** Cell-by-cell Laplacian Index measured near the nuclei. (Requires use of nuclear stain).



### Cell Data (cell-by-cell measurements)

- Cell: Assigned Label #
- Cell: Granule Count
- Cell: Granule Total Area
- Cell: Granule Integrated Intensity
- Cell: Granule Average Intensity
- Cell: Nuclear Total Area
- Cell: Nuclear Integrated Intensity
- Cell: Nuclear Average Intensity
- Cell: Texture Index.
- Cell: Gradient Index.
- 🗸 Cell: Laplacian Index



Highlighted cells shows assigned granules to cell

- Cell: Assigned Label # Cell label number (1 through total cell number)
- Cell: Granules Count: Number of granules detected for a specific cell. (Note: a granules is assigned to its nearest nucleus)
- Cell: Granules Total Area: Area covered by all the granules assigned to a specific cell in um2
- Cell: Granules Integrated Intensity: The total pixel intensity of the granules assigned to a specific cell
- Cell: Granules Average Intensity:
   The total pixel intensity of the granules assigned to a specific cell divided by the number of granules assigned to a specific cell



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- Cell: Granule Count
- Cell: Granule Total Area
- Cell: Granule Integrated Intensity
- Cell: Granule Average Intensity
- Cell: Nuclear Total Area
- Cell: Nuclear Integrated Intensity
- Cell: Nuclear Average Intensity
- Cell: Texture Index
- Cell: Gradient Index
- Cell: Laplacian Index

- Cell: Nuclear Total Area Total square microns of the nucleus
- Cell: Nuclear Integrated
   Intensity Total pixel
   intensity of the nuclear stain
   in the nucleus
- Cell: Nuclear Average
   Intensity Average pixel
   intensity of the nuclear stain
   in the nucleus



### Cell Data (cell-by-cell measurements)

- Cell: Assigned Label #
- Cell: Granule Count
- Cell: Granule Total Area
- Cell: Granule Integrated Intensity
- Cell: Granule Average Intensity
- Cell: Nuclear Total Area
- Cell: Nuclear Integrated Intensity
- Cell: Nuclear Average Intensity
- Cell: Texture Index
- Cell: Gradient Index
- Cell: Laplacian Index

- Cell: Texture Index: Standard deviation of intensity values of a cell
- Cell: Gradient Index: A texturedependent measurement that reflects the amount of local intensity contrast. Measures the difference between the maximum and minimum intensity within a local neighborhood of a cell
- Cell: Laplacian Index: Similar to the morphological gradient, however this morphological measurement reflects fluctuations in the gradient of a cell





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