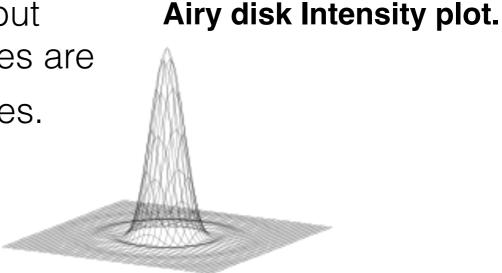
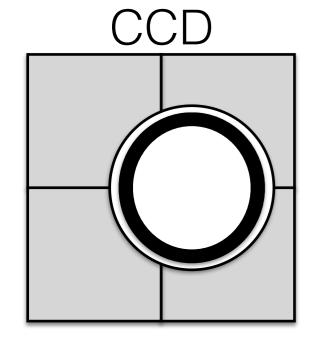
# Imperial College London

## **Single Molecule Localisation Microscopy and OMERO**

Ian Munro

Imaging is fundamentally diffraction-limited but more information available by knowing all sources are fluorescent molecules - i.e. small point sources.





The location of a single point-source can be

determined to sub-pixel resolution.

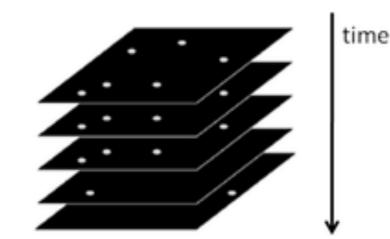
## Label sample with fluorophores, crucially ensure that

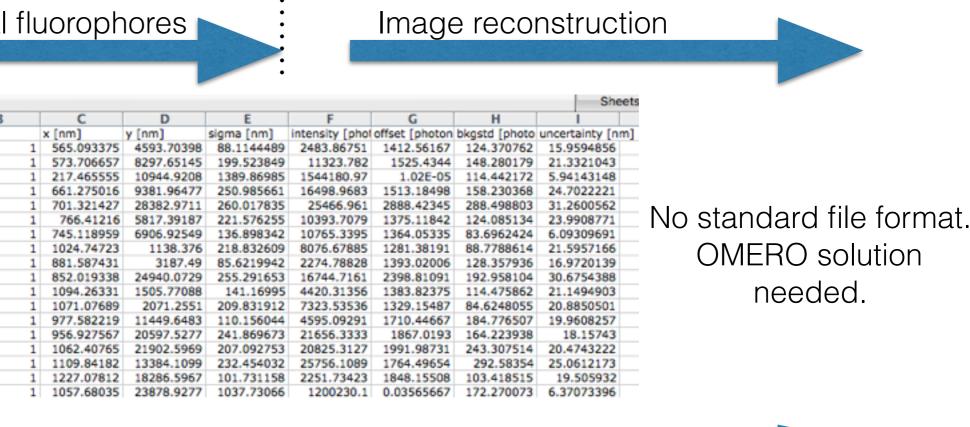
most are switched off PALM, STORM etc





# **Processing Pipeline**





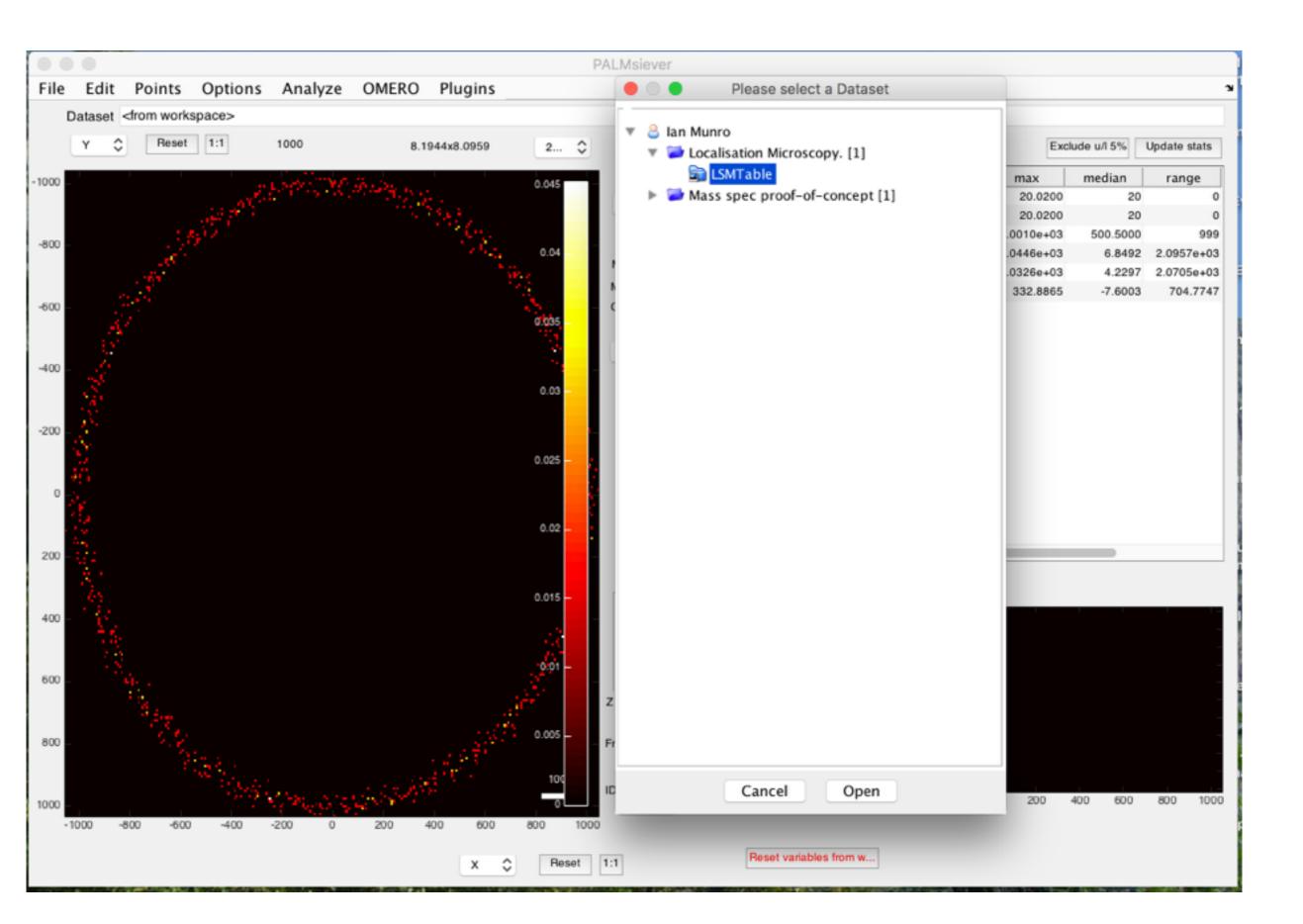
PALMsiever. Matlab

#### Localisation of individual fluorophores

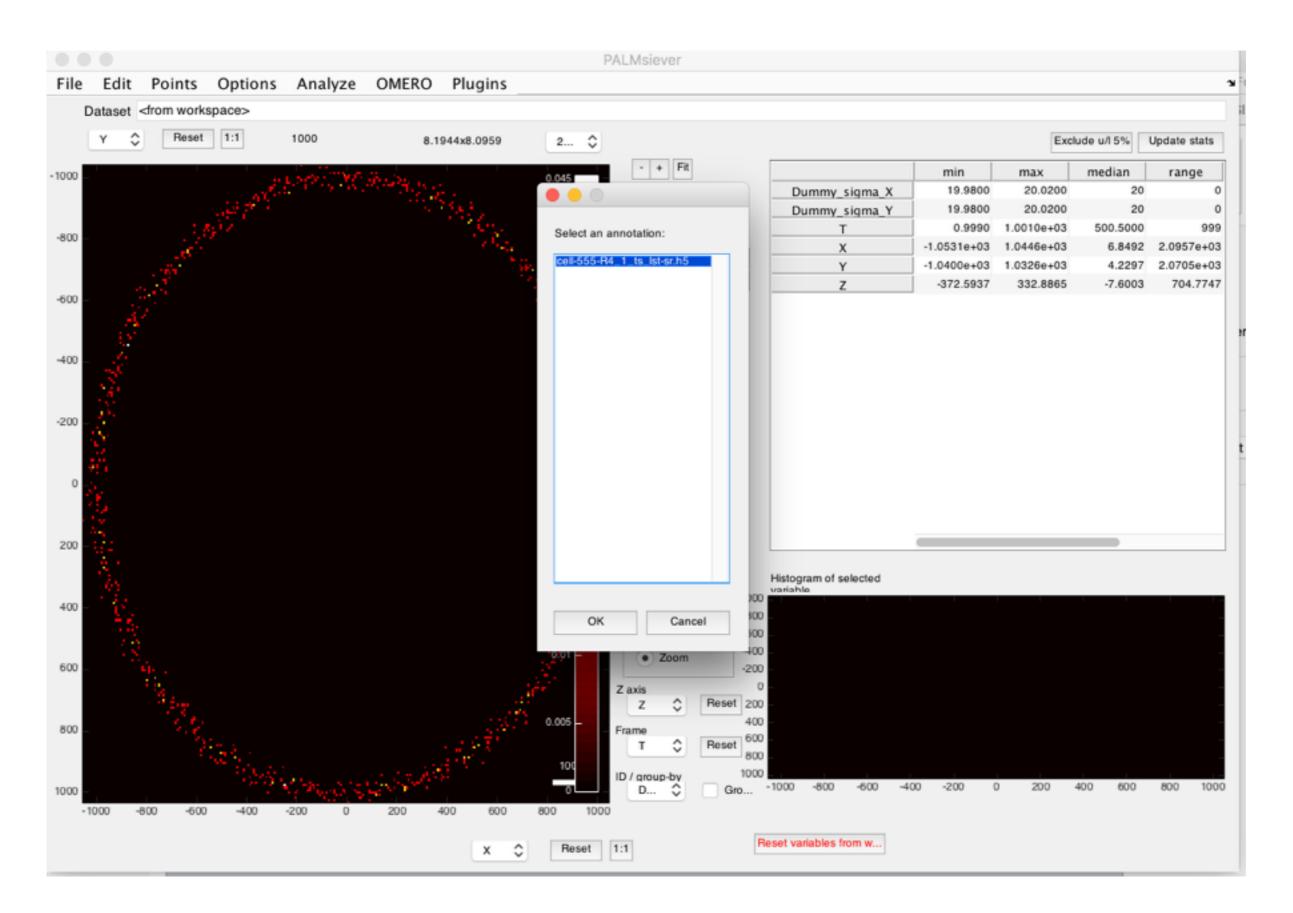
frame

ThundarSTOPM Jova	Imagai plugin	
ThunderSTORM. Java,	imagej plugin	-

#### OMERO menu added to PALMsiever allows user to choose a dataset



#### Then choose a Table



### Super resolved image

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OMERO menu added to ThunderSTORM allows user to choose a dataset

	(Fiji Is Just) ImageJ
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*Oval*, elliptical or brush selectio	ns (right click to switch)
	Please select a Dataset
	🔻 🚨 Ian Munro
	<ul> <li>Television Microscopy. [1]</li> </ul>
	🗊 LSMTable
	Mass spec proof-of-concept [1]
	Cancel Open

#### N.B. same UI as PALMsiever

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	6	1	766.412	5817.392	221	10393.708	1375.118	124.085	23.991
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	8			1138.376			1281.382	88.779	21.596
	9			3187.49			1393.02	128.358	16.972
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	11			1505.771			1383.824	114.476	21.149
	12			2071.255			1329.155	84.625	20.885
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## **Future work**

### **UI** improvements.

UI was to-hand and easy but too many Imagej-OMERO interfaces already. Clean disconnect on closing Fiji needed. Currently Imagej OMERO login separate from ThunderSTORM. Select Table in a single step rather that as a 2-step process. More extensive OMERO interface required for PALMsiever?

#### **Other File formats.**

Handling in Bio-Formats required? possible to re-use ThunderSTORM import code to generate OMERO Tables?