

CellReporterXpress[®] Software Guide

Quick Start guide



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Support Resources

- Help button within CellReporterXpress[®] Software
- Support and Knowledge Base: <u>http://mdc.custhelp.com</u>
- Email Technical Support: <u>support@moldev.com</u> (US) <u>techsupport.eu@moldev.com</u> (EU)
- Telephone Technical Support: 800-635-5577 (US) or +44 118 944 8000 (EU), select options for Technical Support → Cellular Imaging Products →ImageXpress Products





Purpose

This document provides a step-by-step review of CellReporterXpress to:

- Running a saved acquisition protocol
- Creating a new acquisition protocol with on-the-fly analysis
- Viewing Images





Turn on the system and open the software

1. Turn on the System.



- 1. ImageXpress Pico power
- 2. Computer and Monitor
- 3. Log into the computer

2. Start the software.



Double click on the CellReporterXpress icon on the desktop

-or-

Open Chrome (or Safari) browser and type the web address (localhost:8080 for attached desktop)





MOLECULAR DEVICES

LOG	IN
	Login
	Password
1	Remember me
	LOG IN

Enter your LOGIN and PASSWORD, then select LOG IN.

NOTE On most computers attached to your organization's network, this will be your network user name and password.

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Load and run a saved protocol



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Welcome to CellReporterXpress by Molecular Devices



ACQUISITION

Start new experiments by acquiring images from a device.



EXPERIMENTS

Review experiments and run analysis.



STATUS

Monitor acquisition and analysis progress on all devices on the network.



CONFIGURATION

Manage labware and stain libraries, register devices and services



DEVICE

Manage devices



Select the Acquisition panel.



US (Toll-free) 1-800-635-5577 US (Toll-free) 0-800-635-5577 UK (Freephone) 00800-665-32860

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Find the card for your saved Settings and click on the **Run** o icon on the bottom right corner of the card.





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STEPS

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Run Protocol Experiment Name * I am reusing an existing acquisition protocol. Barcode Experiment Description Validation Acquisition Parameters Analysis Parameters 1. Enter an **Experiment Name** and, 655090 optionally, a description. \checkmark Objective x10 0.4 x 0.3mm \bigcirc Exposure (?) Acquisition region 1 (?) Selected Wells 28 ? Analysis regions 1 \bigcirc Selected measurements 11 \bigcirc Analysis services \bigcirc **Device** IXN-AWS-FAS 172.31.30.81 (\mathbf{X}) Device Temp Storage D:\

Free: 4.42 GB

IXN-AWS-FAS C:\ProgramData\Molecular Devices\MD.LocationService\Data

Free: 10.36 GB

2. On the right side of the screen, click the **Run Experiment** button to start the acquisition.



 (\mathbf{X})

Data Storage

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Expected: 8.50 GB

Expected: 11.34 GB

Home Status							Ð	8.1	G
IN PROGRESS	FAILED	SUCCEEDED							
Name	Owner	Start Time		Acquisition Status	Image Processing S	itatus	Analysis Status	Cancel	
TL-GFP Transfection Effici 10X	iency tim	ı.baranowski@moldev.com	Aug 2, 2017 15:06	Aug 2, 2017 15:20	Succeeded	Succeeded		Succeeded	
					Device desktop-pjpbh3o Addresses 10.133.18.75,169.254.255.254 Progress N/A	Service amsm Progress N/A	rhgc8kv1	Service amsn Progress 56/§	ivhgc8kv1 56

Acquisition progress as shown in the Monitor panel







Configure new acquisition and analysis



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DEVICES







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Acquisition De	evice
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Click on		Online 🔵 🕁			
Device Name Serial Number	AMSNVL-4T7SXF2 16ABC521C370_8091	Connected on Version	amsnvl-4t7sxf2 2.0.6648.20242	IP	169.254.139.233 169.254.241.106
Device Model	IX Pico	Free space	54.35 GB	MAC	14ABC521C374 025041000001

STEPS

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~ \$}		Select your Objective from the list	l	4x 4.13mm x 3.46mm PL FLUOTAR 4x/0.13 objective		<u> </u>	
ē				10x 1.65mm x 1.39mm HC PL FLUOTAR 10x/0.32 objective			4x
				20x 0.83mm x 0.69mm HC PL FLUOTAR 20x/0.40 objective			•
f (x)				40x 0.41mm x 0.35mm HC PL FLUOTAR L 40x/0.60 CORR objectiv	e		
	•	스 A1		63x 0.26mm x 0.22mm HC PL FLUOTAR L 63x/0.70 CORR objectiv	9]	
Ō							
		* ÷ *					
		2.63 mm					



















If the image is not already in focus, automatically adjust focus position by clicking on the **Auto** button **Focus offset** box. Alternatively, manual adjust focus by clicking the arrows or typing an offset value into the field.







2. Repeat this step for each wavelength. By selecting the next underlined wavelength tab.









gion Selection to Acquire 1. Select the Region Selection to Acquire icon 4.13 mm \$ \$ 3.46 mm \bigcirc Site # 1 Selected Actual to capture Field Of View Actual Coverage 41.97% Covered Area 14.27 mm² Storage per Well 61.18 MB

? FROM CENTER AUTO ٠ Target area in % 33.01 > I 100% MANUAL # of X sites < || > # of Y sites || >

[→

2. Select the region to acquire images from in each well. The default setting is one image in the center of the well. To adjust the region increase the number of sites by selecting the arrows or typing in the field.

Alternatively, customize regions to acquire by manual adding regions.











wavelength of the

25

target object.

Manual

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Reset

of interest by clicking on the objects within the image.

Re-select the flashing Test Analysis icon to test new settings. Adjust settings as needed.





G	> Acquisition >	Cell Count Walk Through					?	ки о	∖ [→
STEPS STEPS	Acquisition >	Cell Count Walk Through	Nuclei Intensity	Click the Measurem select the summary a measurements to rep	ents icon to and cell bort.	MEASUREMENTS Summary Measurements Select / Deselect All Cell Count Cell Total Intensity Cell Average Intensity Cell Average Integrated Intensity Cell Average Integrated Intensity Total Area Average Area	Cell Measurements Recommended 128 44575.1 348.243 2.43831e+7 190493 8195.47 64.0271		
		DAPI 🔻	Min Width Max Width Manual	5 30 Auto Reset					
localhost/in	aex.ntml#/								





습	> Acquisition	Cell Count Walk Through			0	ки О	[→
STEPS	+ . 0 of 26.22	- O ::		SAVE AN Analysis New ar Settings	ALYSIS Settings: nalysis template avatar: Use captured pic ck to upload Save	× ture	
		Target	Intensity 100 Min Width 5 Max Width 30 Manual Auto Reset	Click	on the Save Analys the analysis settings	is icon	to





ON O Analyze Entire Acquisition Area



Acquisition Area 8.55 mm² Analysis Area 8.55 mm²

The entire image will be analyzed by default.

Region Selection to Analyze

To customize the region of analysis, toggle the **Analyze Entire Acquisition Area** button to the "OFF" position.

Add regions for analysis by adding a new region or resizing by dragging the blue corners of the region box.



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<mark>ک</mark> > STEPS	Acquisition > cell 2. Type for you	e a name (require r new acquisition	ed) and description (option n protocol.	al)	◎ 23 ペ [→
	Protocol Name * acquisition Protocol Description New self-guided plate acquisit	on and analysis.			
• • • • •	Acquisition Parameters A	nalysis Parameters 24x16	384 Greiner 781091		
	Stains Objective	DAPI, FITC x4	4.13 x 3.46mm		
1. Cl	ick on the Save P Selected Wells	42			
	Device	AMSNVL-4T75XF2 Click the Save P	169.254.139.233, 169.254.241.106		





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STEPS			2. Type in a	a name and des	scription (optional) of you	ır experim	ent.				
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		$\overline{\bigcirc}$	<u>Objective</u>	x4	4.13 x 3.46mm						
<u> </u>		\bigcirc	Focus and Exposure								
Ē		\bigcirc	Acquisition regions	1							
		1. S	elect the Run	Protocol icon.							
		$\langle \! \rangle$	Selected measurements	11							





Review mages



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localhost/index.html#/







T1 T1 \sim Thumbnail View 12 13 18 19 1 2 3 4 5 6 7 8 9 10 11 14 15 16 17 2 А Double click on any thumbnail to see the В corresponding full resolution image. С D Е F G н Cellular















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