

Software

Commercial

Microscopy

SymPhoTime64

A complete software suite for acquisition and analysis of time resolved fluorescence data. SymPhoTime features a powerful scripting language that allows customisation of existing analysis and even implementation of novel analysis techniques.

Phasor Analysis Phasor Analysis is currently not included in SymPhoTime64. However, you can use Globals developed by the Laboratory for Fluorescence Dynamics to analyse FLIM data via the Phasor approach.

Spectroscopy

EasyTau

EasyTau is the systems software of PicoQuants time resolved spectrometer FluoTime 300. EasyTau features a scripting language to automate more complex experiments. Examples of EasyTau scripts can be found on github: https://github.com/PicoQuant/EasyTau-Scripts

FluoFit

Decay curve analysis software

Open Source

Please note that these are external projects for which PicoQuant takes no responsibility and provides no support. In case of problems please contact the respective authors. Also note that such projects may not be maintained continuously. In particular, please check if they use current versions of file formats and/or programming libraries before you invest in larger development work based on such projects.

If you develop improved or new code, please consider contributing your code back to the community.

The compilation below is probably not complete. If you are the maintainer of a software that deals with TCSPC data please edit this page and add your software listing. Alternatively you can also use https://support.picoquant.com to contact PicoQuant support to submit your software project.

Data Analysis

Complete packages

• **PAM** is a software package for quantitative analysis of fluorescence microscopy and spectroscopy data, with a focus on experiments using pulsed interleaved excitation.http://pam.readthedocs.io

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- Open Tools for Solution-Based Single-Molecule Fluorescence Spectroscopy
 - **FRETBursts**: software for burst analysis of freely-diffusing single-molecule FRET (smFRET) measurements including μs-ALEX. FRETBursts Homepage and paper.
 - **PyBroMo** simulator for single molecule FRET experiments of freely diffusing particles.
 - Researchgate group Open Computational Tools for Single-Molecule Spectroscopy
- PyCorrFit is a general-purpose FCS evaluation software that, amongst other formats, supports the established Zeiss ConfoCor3 ~.fcs file format. PyCorrFit comes with several built-in model functions, covering a wide range of applications in standard confocal FCS. In addition, it contains equations dealing with different excitation geometries like total internal reflection (TIR).
- FLIM-FRET analyzer: open source software for automation of lifetime-based FRET analysis. This is a Microsoft Windows application originally developed in C#. The main goal of this application is to provide the FLIM community a user-friendly application.
- Fluortools.Fluorescence Decay and Anisotropy Analysis.
- PyCorrelate:Pycorrelate computes fast and accurate cross-correlation over arbitrary time lags. Cross-correlations can be calculated on "uniformly-sampled" signals or on "point-processes", such as photon timestamps. Pycorrelate allows computing cross-correlation at log-spaced lags covering several orders of magnitude. This type of cross-correlation is commonly used in physics or biophysics for techniques such as fluorescence correlation spectroscopy (FCS) or dynamic light scattering (DLS).
- QuickFit 3.0: QuickFit 3 is our open-source data evaluation software for FCS and imagingFCS (imFCS) measurement. Actually QuickFit 3 itself is a project manager and all functionality is added as plugins. A set of tested plugins is supplied together with the software.
- FCS Point Correlator FCS Bulk Correlation Software accepting pt3 and ptu files.
- FLIMFit. FLIMfit is an open source package for rapid analysis of large FLIM datasets.
- Extensible Time-tag Analyzer. ETA, the extensible time tag analyzer, is an event driven programming language with graphical user interface for analyzing, plotting, and fitting of time tagged data. More info.

Libraries/scripts/readers

- Demo Code for reading PicoQuants PTU File format
- Reading PicoQuants PTU File formats to MatLab 100x faster than the Demo Code
- libpicoquant: Tools for reading PicoQuant file formats: https://github.com/tsbischof/libpicoquant
- **Open Microscopy's BioFormat** supports PicoQuant bin files (pre-histogrammed Image Data File) https://www.openmicroscopy.org/site/support/bio-formats5/formats/picoquant-bin.html
- ImageJ data readers-
 - Import Filter for pt3 files:
 - http://imagejdocu.tudor.lu/doku.php?id=plugin:inputoutput:picoquant .pt3 image reader:start
 - Import of ptu/pt3 image files (+intensity/average lifetime stacks): https://github.com/ekatrukha/PTU Reader
- A Python library to read PTU files
 - From the authors: "The main purpose of the library is to compute g² from the photon arrival times as fast as possible, as our files often go into multi GB sizes. With a good SSD we can parse >3GB/s. It can also compute intensity time traces and post select photons based on that."
- https://github.com/QuantumPhotonicsLab/readPTU Probably same as above
- ptuT3toT2 a program that converts ptu files from picoquant from T3 mode to T2 mode
- **Photon-Tools** Python utilities for working with photon timestamp data from fluorescence spectroscopy experiments https://github.com/bgamari/photon-tools

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- **Photon Correlation** A suite of command-line tools for performing calculations on photon arrival data, as produced by hardware such as the PicoQuant PicoHarp and HydraHarp: https://github.com/tsbischof/photon_correlation
- dspFRET: Matlab code for analyzing TTTR data from PicoQuant. Typical application is analysis of single pair FRET measurements with pulsed interleaved excitation (PIE-FRET). https://github.com/RobinsonLab/dspFRET
- exciton diffusion (Matlab)
- **FoCuS-point**: FoCuS-point utilises advanced time-correlated single-photon counting (TCSPC) correlation algorithms along with time-gated filtering and innovative data visualisation. FoCuS Homepage and paper
- TCSPC curve fitting for phd files (Python)

Data Acquisition

- **qudi**: A modular laboratory experiment management suite which supports PicoHarp300. https://github.com/Ulm-IQO/qudi
- **pyPL**: wxpython integration of software and hardware for micro-photoluminescence measurements: WinSpec, PicoHarp, and Thorlabs/APT piezos and stepper motors. https://github.com/kaseyrussell/pyPL
- PicoHarp Multiscan with Prior Stage: https://bitbucket.org/ywsong2/uky_ptu_convert

Demos about Controlling PicoQuant Devices

Sepia II and Solea

https://github.com/PicoQuant/PQLaserDrv

TimeHarp 260

• https://github.com/PicoQuant/TH260-Demos

PicoHarp 300

- PicoHarp 300 TCSPC system hardware plug-in for ScopeFoundry (Python)
- Python binding, Tango device server and Taurus gui to access the PicoHarp300 photon counter
- https://github.com/PicoQuant/PH300-v2.x-Demos
- https://github.com/PicoQuant/PH300-v3.x-Demos

HydraHarp 400

- Control HydraHarp400 (python)
- https://github.com/PicoQuant/HH400-v1.x-Demos
- https://github.com/PicoQuant/HH400-v3.x-Demos

Other

PicoHarp Library in Python

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https://github.com/srgblnch/PicoHarp300

GitHub repositories mentioning PicoQuant and supporting the PicoQuant t3r data file format: https://github.com/search?&q=picoquant+OR+picoharp+OR+hydraharp+OR+TimeHarp

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