



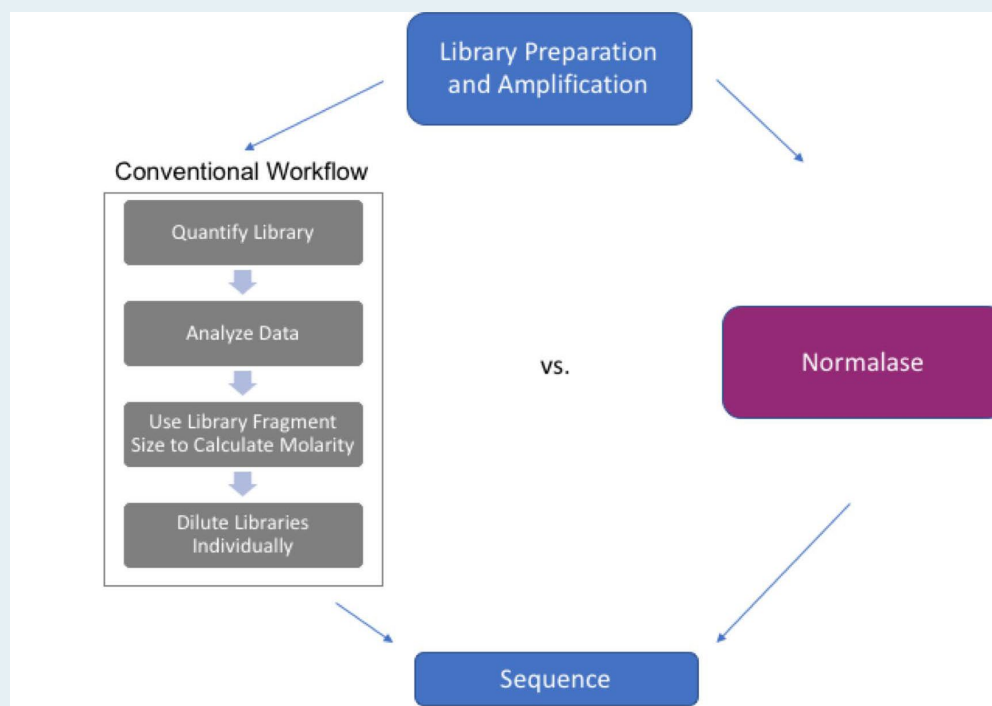
# Swift Normalase Kit for Libraries on Illumina® Platforms

## Radically Streamlining NGS Library Quantification and Normalization

**Saves time and increases throughput:** Uniform sample processing with fewer handling steps to generate balanced library representation in multiplexed sequencing

**Reduces variability to save on sequencing costs:** Improved library balancing allows higher multiplexing per run, predictable read depth, reducing the need to resequence.

**Flexible design for many workflows:** Compatible with diverse library preparation methods and sample types to produce more evenly balanced sequence data



The Swift Normalase Kit offers a novel enzymatic library normalization technology that is augmented by the power to consolidate library normalization and pooling for loading on Illumina® sequencing platforms. The Normalase workflow eliminates the need for library quantification and concentration adjustment prior to library pooling, resulting in optimal cluster density and library balance. The Swift normalization method

can easily be integrated into standard protocols to improve turnaround time loading and accuracy for NGS laboratories.

[Read more](#)

## Upcoming Conferences and Presentations

There will be a strong **TATAA** presence at both Bioscience 2018 ( 7th Nov, Uppsala) and at the 4Bio Summit (27-28th November, Rotterdam) in the forthcoming month.

### Bioscience 2018, 7th Nov, Uppsala

Of particular interest to some **TATAA** customers will be Lukas Paul's (Lexogen) presentation 'Using time-resolved gene expression profiling to identify direct transcriptional targets' at Bioscience, 2018.

Click on the Bioscience 2018 tab for full conference details.

[Bioscience 2018](#)

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### 4Bio Summit, 27-28th Nov, Rotterdam

The Postillion Convention Centre in Rotterdam will play host to the 4Bio Summit between 27th and 28th Nov, 2018. Here you can join two senior members of **TATAA's** team and participate in their satellite courses.



Filip Stern will be hosting a post-conference workshop on NGS library construction and quality control.

The course gives an introduction to Next Generation Sequencing (NGS) and its many applications. The course will focus on considerations for the NGS experiment design, the different sequencing platforms, quality control of samples, library preparation techniques, and quantification of libraries for sequencing.

A pre-conference workshop on expression biomarkers will be hosted by Mikael Kubista.

This course is aimed at a range of personnel from Quality Control managers to laboratory personnel and will focus on how to measure, validate and quality control the workflows in compliance with recent guidelines.

To register for these workshops and to see all our upcoming courses please click on the following tab:

[Registration & Courses](#)

## FREE GenEX Tutorial



A fantastic free to use tutorial on GenEx is now available on-line.

Click on the tab below to gain access to this informative tutorial.

[Free GenEx Tutorial](#)

## News

[Press release – TATAA Biocenter and Arbor Biosciences Announce Strategic Partnership](#)

[Want to learn about Molecular microbiological identification and typing?](#)

Dr Mikael Kubistas webinar "Normalization of qPCR data"

Webinar "AR-V7 Key in Tx of mCRPC" by Andrew Armstrong, MD, presented at ASCO

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