



# Custom Libraries Illumina preparation and Shipping Guidelines

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## SERVICE CONDITIONS

*Please read carefully!*

The samples must meet the quality and quantity criteria and the shipping conditions, as indicated in the “Sample Requirements” and “Shipping” sections below. In case such specifications are not available the Customer shall enquire before sending any material. If storing and shipping conditions or sample specifications do not meet the required standards, IGA Technology Services may ask for an additional processing fee and the standard turnaround time might be delayed. Please note that the experimental setup and processing will be based on the information reported in the Sample Spreadsheet (**see Documents/Guidelines section in <https://igatechnology.com/>**). Keep in mind that absorbance-based methods (e.g., Nanodrop) might largely overestimate the DNA quantity. Clients will be informed upon QC if samples are non-compliant and must decide whether to proceed or submit replacements. The proceeding with non-compliant samples absolves IGA Technology Services of liability for experiment failures (inadequate library or low sequencing due to poor sample quantity and/or quality) and allows it to charge the full-service fee.

### **Contacts**

All requests for information about the service must only be sent via email ([custom\\_libraries\\_illumina@igatechnology.com](mailto:custom_libraries_illumina@igatechnology.com)) and will be dealt with within a maximum of two working days from receipt.

In case problems arise during the sample processing, the Customer will be contacted directly to agree on possible solutions. Notifications will be sent to the contact person indicated in the Sample Spreadsheet at the sample reception and delivery.

For sample return, data storage and delivery refer to general Terms and Conditions.



## LIBRARY REQUIREMENTS

Pseudonymization.

***Please refer to general Terms and Conditions.***

### Libraries specifications

- Diluent: 10mM Tris-HCl pH 8.5 or others commercial DNA resuspension buffer.
- Concentration range: 5ng/uL – 50ng/uL (by fluorimetry)

Keep in mind that absorbance-based methods (*e.g.*, Nanodrop) might largely **overestimate** the DNA quantity.

- Minimum volume: 25uL

**Obligatory!** Indicate in the Library sheet of the Sample Spreadsheet:

- Library preparation kit
- Index sequences for demultiplexing and name in case of commercial kit usage
- When using a homemade library construct it is mandatory to compile the full Adapter sequences used

## SHIPPING

**For batches of  $\leq 24$  samples**, 1.5 mL or 2 mL tubes can be accepted. They must be sealed with parafilm. The tubes must have on the vial top, a clear and permanent sign (or a thin label) with a **progressive number** corresponding to information specified in the Sample Spreadsheet.

**For batches of  $> 24$  samples**, samples are only accepted in 96-well plate, sealed with adhesive/heat-sealed aluminum foil or multiwell strip caps. Each plate must be labeled with a plate identifier indicated on the Sample Spreadsheet.

**IMPORTANT!!!The 0,5 mL and 0.2 mL tubes as well as strips are not accepted!**



For more details, please follow “Shipping and Packaging Guidelines” PDF file available at <https://igatechnology.com/igatech/documents/>

It is **MANDATORY to send us the compiled Sample Spreadsheet**, both with the shipped parcel and via e-mail. To properly track and safeguard your samples, send us the Tracking Number via e-mail (see Above).

**Shipping Address:**

**IGA Technology Services srl**

**via Jacopo Linussio 51**

**33100 Udine Z.I.U.**

**Italy**