



## Call for tenders

### ***Bring your Own Samples: bulk RNAseq workshop - 2026***

In the frame of its RARE TRIPS (*Training Research Interdisciplinary Platforms*), the MarMaRa institute is opening a workshop call to support preliminary data in genomics and encourage collaborative research projects within the Rare Diseases community. The **“Bring your own samples: bulk RNA seq workshop”** aims to provide the opportunity to participate to a step-by-step **training on bulk RNA sequencing methods**.

- 1 week **Workshop BYOS-bulk RNAseq (1-8 samples)\***
- 1-2 sessions call / year (spring & fall) upon demand and applications
- [BulkRNA seq poly A stranded protocol](#) by Illumina sequencer (see details in the application form)

#### **Who can apply?**

- This workshop is open to **PhD students and early-career researchers new to RNA-seq**
- Involvement of **Master 2 students** is encouraged.
- **Exploratory and collaborative projects are encouraged**
- *When applicable, the name of the rare disease(s) addressed by the project should be clearly indicated.*

#### **Calendar steps:**

- Application: **January 16 – February 18, 2026**
- Notification of selected projects: by February 27, 2026
- Validation of the RNAs samples (quality control): by March 15, 2026
- BYOS-bulkRNAseq workshop: **April 13 -17, 2026, TGML, Luminy campus (in person)**
- *Bulk RNA seq: preprocessing analysis* – MarMaRa PhD program remote training, May 2026 (1 day, to be confirmed)

\*The workshop will be funded by the **project TIGER, RARE TRIPS**, through **MarMaRa Institute, A\*Midex** and **France 2030 plan**.

#### **General eligibility criteria:**

- **Each team may submit only one application per year to this call.**
- For TAGC members (not organized into teams), a maximum of four applications are accepted per call session.
- If two applications are submitted by the same team within the same call, the second-ranked applications will be subjected to the budget available.

## Ethicals aspects

By submitting an application, each participant acknowledges and agrees that exploitable results outputs cannot be guaranteed. Consequently, the RARE TRIPS / TGML platform assumes no responsibility for the samples or the resulting data.

## Specific eligibility criteria:

- All MarMaRa's members are eligible: researchers (PhD student, post docs, AHU, CR, MCU, DR, PU, IR)
- **To be eligible, sample qualities need to be assessed by the hosting platform prior mid-March 2026.**
- To be eligible, the applicant must commit to present their project and/or its results to MarMaRa's scientific specific events or annual symposium in the following academic year. As stated in all MarMaRa's document, laureates must participate to at least 2 institute's activities<sup>1</sup>.

## Evaluation rule:

The RARE TRIPS committee will review all applications and contact the project leader shortly after the deadline regarding sample quality (pre-mRNA quality control) to assess feasibility of the project for the current session. If quality control is insufficient or if sample biases are detected, applicants will be encouraged to resubmit their application in a future session.

## Documents required:

- **Application form** (document to download [here](#))

**Application must be submitted as a single PDF file** ("MarMaRa\_AO\_BYOS\_bulkRNAseq\_[Name of project leader]\_[Laboratory – team]\_Janv26") **in this [AMUBOX folder](#).**

## Terms of engagement:

- All scientific publications and communications which have benefited of the financial support of MarMaRa must mention the following sentence:

*"This work received support from the French government under the France 2030 investment plan, as part of the Initiative d'Excellence d'Aix-Marseille Université - A\*MIDEX (AMX-19-IET-007) through the Marseille Maladies Rares Institute (MarMaRa). High throughput sequencing was performed at the TGML Platform, supported by grants from Inserm, GIS IBISA, Aix-Marseille Université, and ANR-10-INBS-0009-10."*

<sup>1</sup> Annual symposium, JIMR, seminars, PhD program, WG CIVIS, WG 3M...

- Publications must be deposited in the A\*Midex HAL AMU collection, the University's open archive.

**Deadline 18/02/2026**