

# Establishment of ctDNA as a biomarker in melanoma follow-up

Project Start: January 1<sup>st</sup>, 2026

Enrollment: at least 200 patients

Duration: 1 year

## Objective

- Evaluate the value of circulating tumor DNA (ctDNA) by as a biomarker for detection and relapse
- Replace or complement other monitoring methods (e.g., S100)
- Establish a standardized approach for ctDNA assessment

## Project Design

**Type:** multi-center EADO–EUMelaReg project

### Site inclusion criteria:

- BioRad QX200™ or QX600™ infrastructure
- Accredited medical laboratory according to local standards e.g., ISO15189
- Access to EUMelareg (can be granted for this specific project only)

### Patient inclusion criteria:

- High-risk resectable cutaneous melanoma, AJCC stage IIIB or higher
- Confirmed mutations that are covered with Oncobit™ PM (BRAfV600D/E/E2/K/R or NRASQ61K/L/R)
- Signed consent to provide routine clinical data (according to local ethical standards)

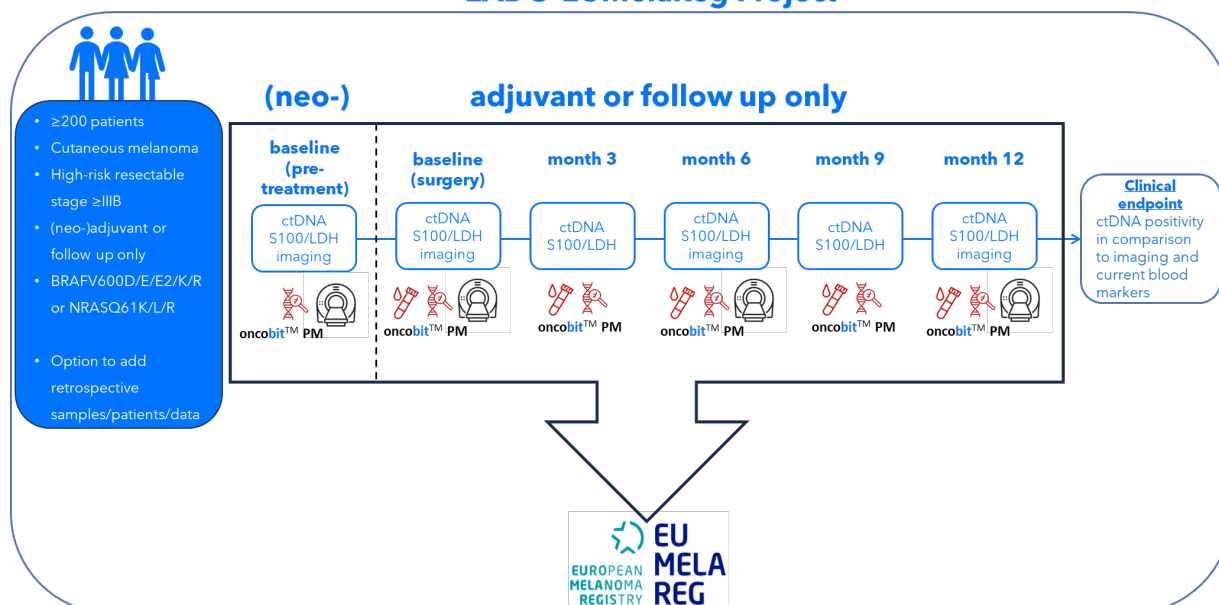
### Sampling time points (as part of clinical routine):

- Quarterly plasma collection for ctDNA analysis (or more frequently if patients are seen according to local standards and if feasible to do within the local routine clinical setting)
- S100/LDH and other markers if performed according to local standards
- Imaging according to local standards every 6 months
- Collection of clinical information including treatment information

### Clinical Endpoints:

- % ctDNA positive in comparison to imaging, S100, LDH or other markers

## EADO-EUMelaReg Project



## Funding

### Oncobit Contribution

Oncobit will provide centralized support for the ctDNA assessment of the project. Specifically, Oncobit will fund:

- **Oncobit™ PM kits and cloud-based analysis software:** Provided together as a **CE-IVDR certified system**
- **Digital PCR kits**, including positive controls, are optimized and designed for mutation-specific detection
- The **software platform** enables secure upload, standardized analysis of digital PCR data, and automated reporting with mutation allele frequency (MAF), variant calls, and quality metrics

### Participating Site Contributions

Each participating clinical site is responsible for the operational aspects of patient management and sample processing. Specifically, sites will fund:

- **Sample collection:** Blood draw and plasma separation.
- **DNA extraction:** Using site-validated methods suitable for circulating free DNA (cfDNA).
- **Execution of digital PCR:** Running Oncobit™ PM assays on BioRad QX200™ or QX600™ platforms.
- **Clinical data collection and management:** Data regarding clinical routine needs to be added to the EuMelaReg database (treatment documentation, imaging results, blood biomarker results). Sites are responsible for internal project protocol/IRB approval if needed according to their local standards.

## Contact

For further information regarding this clinical project or interest in participation, please contact [Prof. Dr. Reinhard Dummer](#) or [Dr. Claudia Scheckel](#), or complete the [feasibility form](#).