

# Report

## Clone selection report of cell samples containing vector

Prepared for:	Company name
	Company address
Customer name:	Name
	Position within a company
	email
Internal project number:	XXX
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## Goal

In this study, # transgenic SAMPLE with the vector XXX sequence were analyzed.

The aim of this analysis was to:

1. Determine the presence of sequence variants in the integrated vector sequence.
2. To determine sister clones.

An overview of the TLA technology and technical details of the performed analyses is provided in the manual "[Introduction to the terminology and methods used in transgene & integration site TLA analyses & ddPCR\\_v3](#)".

## Summary

Sample	Sequence variant present in integrated vector sequence	If Yes,				Similar to other clones
		Annotation	Position	mutation	%	

## QC information

### Sample and Study details

Sample receipt date  
 Condition of sample at receipt  
 Start date in the lab  
 Sequencing run  
 Date data analysis  
 Deviations from the protocol  
 TLApp version:

### Study Personnel

Lab technician  
 Data Analyst  
 QC Analysis and Report



### Quality control

The results are independently verified and reviewed and are an accurate and complete representation of the study. The scope of accreditation for ISO/IEC 17025:2017, accredited by the Dutch Accreditation Council RvA, Registration number L671, entails all analytical services including: determination of the integrity of the transgene vector sequence; determination of the vector integration site(s), next generation sequencing (NGS) and bio-informatic data analysis.

Scientific approval

Date

Signature