

# Sequencing report

## Table of contents






<b>1 Project</b>	<b>1</b>
<b>2 Sample AAV_pCMV_k-Firefly_WPRE_SV40pA</b>	<b>2</b>
2.1 Overview	2
2.2 Visual representation of the sequencing results	3

## 1 Project

Name	pTransgeneReporter_V2
Customer	[REDACTED]
Internal ID	pTransgeneReporter_V2
Quote ID	pTransgeneReporter_V2
Genetic Engineering Specialist	Pauline Vieugué

## 2 Sample **AAV\_pCMV\_k-Firefly\_WPRE\_SV40pA**

### 2.1 Overview

Name	AAV_pCMV_k-Firefly_WPRE_SV40pA
Type	vector (circular) 
NGS reference	
Concentration (ng/μL)	1279
Expected sequence length	5448 bp
Estimated sequencing depth	1777 X
Verified sequence	5305 base pairs (3 fragments)  <i>download available on your account</i>
Mycoplasma contamination	 <0.1%
Comparison vs expected sequence	 100% validated

 **Comments from the Genetic Engineering Specialist**

“ ... ”

## 2.2 Visual representation of the sequencing results

