

Overview

We use automated Dye-Terminator Sequencing (Sanger Sequencing) to provide long high quality reads from the most various DNA templates. In more than 10 years we have provided service to support hundreds of basic and translational research projects. In addition, we participate in the diagnostic activity of the CGT lab.

Service

We perform standard sequencing as well as sequencing projects. We do fragment analysis for the most different applications. We assist in primer design and in data analysis.

Innovation

We routinely revise existing protocols to increase the quality of our work. In addition, we develop new techniques according to researcher's needs.

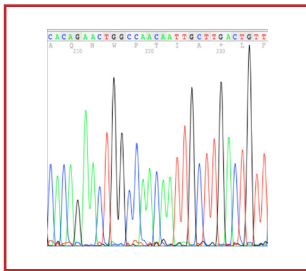
Quality

We provide read length of 600-800 bases and Universal Primers free of charge. We have a rapid turn-over time (1-3 working days) and automated data delivery in .abi format.

Our laboratory is UNI EN ISO 9001:2008 certified (n. IT256850).



SEQUENCING



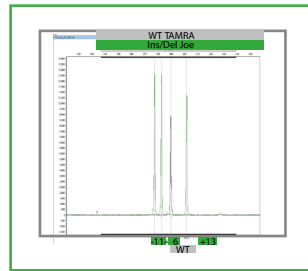
Sequencing in single tubes and 96 well plate format

Primer Walking

Re-sequencing

Variant Analysis (SNP & Mutations)

FRAGMENT ANALYSIS



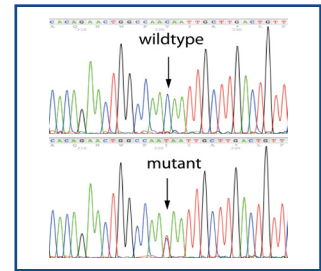
Microsatellite analysis

Cell line verification

CrispR/Cas9 genotyping

MLPA

DATA ANALYSIS



Data Analysis for Re-Sequencing and Primer Walking projects

Data Analysis for Cell line ID and CrispR/Cas9 genotyping

Assistance in Trouble Shooting for Standard Sequencing