

# METHOD AND SYSTEM FOR THE PRODUCTION OF RE-COMBINANT PROTEINS BY CELLS

## KEYWORDS

- > Micromolecular
- biology
- > Bioprocess engineering
- > Industrial
- biotechnology
- > Toxin-antitoxin
- systems
- > Genetic engineering

#### **Collaboration type**

Licence agreement Research cooperation

#### **IP Status**

Priority date: 2014/07/04 PCT publication number: W02016001447

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## **STATE OF THE ART**

Typically, production process of peptides or proteins of interest by host cells, or molecular cloning, requires the use of one or more replicon sequence(s), such as plasmids carrying one or more gene(s) encoding the protein(s) of interest. Cells in bioreactor usually present a high heterogeneity. It has been demonstrated that the growth of plasmid bearing cells, especially bacterial cells, is significantly reduced compared to plasmid free host cells. Indeed, peptide or protein production coming from gene of interest represents a significant burden on the metabolism of the cell. Therefore, it is necessary to select cells that keep the gene of interest. This selection is efficiently done by the use of a selective tool, such as antibiotic resistance gene or pair of poison and/or antidote genes. However, classical systems do not prevent frameshift from translation, truncated versions of the protein of interest... which could lead to a contamination by mutated proteins.

# THE INVENTION

The invention concerns a cassette sequence for the transformation of a host cell includes at least: a first nucleotide sequence encoding a pep¬tide or protein of interest to be produced by the host cell. The first se¬quence is linked to a second nucleotide sequence providing resistance to a toxin or encoding an antitoxin peptide to the toxin. The nucleo¬tide sequences are organized in such a way that production of the pep¬tide encoded by the second nucleotide sequence(s) is translationally coupled to production of the peptide encoded by the first nucleotide sequence. Thus, any modification of the protein of interest do not enable the expression of the antitoxin that leads to the death of the cell host. The quality and the yield of expression of the awaited product is then efficiently improved.

# **KEY ADVANTAGES OF THE TECHNOLOGY**

- > Efficient production of peptides or proteins of interests
- > High production and quality yield guaranties





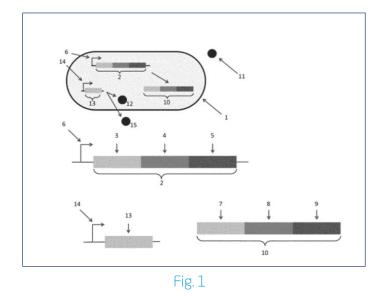


### TEAM

The laboratory specializes in the genetic analysis of Escherichia coli as well as its phages, plasmids and transposons. The aim of this research is to understand precisely how the poison/antidote systems work by investigating the targets of the poisons and by analyzing the molecular interactions between poisons and targets in order to develop new classes of antibiotics; by analyzing the molecular interactions between antidotes and ATP-dependent proteases; by analyzing the regulation mechanisms that trigger the programmed death of the bacteria caused by these systems. In addition, the laboratory is using the poison/antidote systems to develop innovative technological tools. Some of these tools are now used throughout the world in cloning experiments, i.e. the production of DNA fragments which are genetically identical.

### THE INVENTOR

LAURENCE VAN MELDEREN received her PhD in Molecular Biology in 1995 and is currently Professor at the Université Libre de Bruxelles (ULB). Her projects are centered on bacterial toxin-antitoxin systems and their impact on genome plasticity and evolution as well as bacterial physiology. Her group also focuses on molecular mechanisms involved in bacterial persistence and adaptation.



### **RELEVANT PUBLICATIONS**

> Messing up translation from the start: How AtaT inhibits translation initiation in E. coli, Van Melderen L., Jurenas D., Garcia-Pino A. (2018). E. coli. RNA Biol. 15(3):303-307.

> Novel toxins from type II toxin-antitoxin systems with acetyltransferase activity, Jurenas D., Garcia-Pino A., Van Melderen L. (2017). Plasmid. 93:30-35

