

# **AI-based fluorescent labeling for cell line development**

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Master thesis

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## **Erklärung**

Hiermit versichere ich, dass ich diese Master thesis selbstständig verfasst habe. Ich habe dazu keine anderen als die angegebenen Quellen und Hilfsmittel verwendet.

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## **Abstract**

Cell line development is an expensive and time-consuming process, however that is the most modern approach for producing the proteins needed in pharmaceuticals.



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# 1 Introduction

## 1.1 Motivation

## 1.2 Notation

# 2 Background

## 2.1 Biology

CLD Cell line fluorescence DIC vector CHO cell proteins transfection

Explain CLD process Its need for producing proteins Explain how cells are chosen and that it long Explain the role of fluorescence Explaining its pro and cons

Cell line development (CLD) is the process by which the cellular machinery is co-opted to manufacture therapeutic biologics or other proteins of interest. One can use different expression systems for cell line development: bacterial, plant-based, yeast, mammalian. (copy paste from <https://www.beckman.de/resources/product-applications/lead-optimization/cell-line-development>) Chinese hamster ovary (CHO) cells are the most popular mammalian cells used for protein production. (doi:10.1016/B978-0-08-100623-8.00007-4)

First step of CLD is the introduction of the gene of interest (GOI or a DNA vector) to CHO cells. This process is called a transfection. It is important to transfect a GOI into an optimal site of genome to secure a high protein expression over time during protein production, however practically transfection mostly results in a vector being inserted into a random site within a host cell genome. In case the gene was transfected in the inactive site of genome (and the majority of genome is not transcriptionally active) the cell will likely not express the gene. (doi:10.1016/B978-0-08-100623-8.00007-4) (doi:10.1016/j.coche.2018.08.002)

The second step is the selection of cell pools that have successful and stable gene integrations. The reason why not all of them are suitable for cloning is the following: during the transfection only 80% of cells receive the vector of GOI (doi:10.1016/B978-0-08-100623-8.00007-4), only the small percent of which actually integrate a vector into the genome and, as mentioned above, only a fraction of those cells are able to stably express a protein. (Reference needed). Such selection could be done with bulk sorting algorithm. (doi:10.1016/B978-0-08-100623-8.00007-4)

The third step in CLD is to clone the cells. The chosen stable pools of cells are phenotypically and genetically diverse - meaning they have different growth rates, metabolic profile and etc. This is not ideal for industrial production - all the cells used for protein production should be derived from the same clone ([25] here doi:10.1016/B978-0-08-100623-8.00007-4). In order to choose single best cells for further cloning one assesses several parameters like cell size, granularity, cell surface protein expression and etc. This can be done with Fluorescent Activated Cell Sorting (FACS) technology. (<https://doi.org/10.1517/14712598.4.11.1821>). Unfortunately fluorescence labeling is expensive and may ruin the cell due to its phototoxicity

(<https://doi.org/10.1371/journal.pone.0007497>). There is a limited number of available fluorescent channels in microscopes as well as such labels can also be inconsistent, depend a lot on reagent quality, and require many hours of lab work. Therefore there exists a need for fluorescent labeling in silico - without intervening into the cell.

Once the cells are cloned, phenotypical and genetical heterogeneity is reduced, the next step is to characterize clonally-derived cells based on the following criteria: cell size, growth rate, protein quality, titer, metabolites and etc. With this one can estimate clones productivity and titer. Such observations may take up to 90 days after which one can determine which cells are stable and therefore suitable for production. This is the last step of CLD process and consumes a lot of time and maintaining costs for feeding and cloning the cells. Predicting the stability of the cells directly from DIC images would reduce this time significantly allowing to escape this process completely.

## 2.2 ML

Background on Unet and ML in general Conrad (1997) hat ein Buch geschrieben. Es gibt auch andere Arbeiten (Patterson and Hennessy, 2004) die referenziert sind. In Abbildung 1 ist ein Sachverhalt dargestellt.

1 Autor: Conrad (1997) (Conrad, 1997)

2 Autoren: Liebeck and Conrad (2015) (Liebeck and Conrad, 2015)

3 Autoren: Liebeck et al. (2016) (Liebeck et al., 2016)

Online resource: ILSVRC2016 (2017)



Figure 1: Gallien zur Zeit Caesars

Jahr	Erster Consul	Zweiter Consul
1	C. Caesar	L. Aemilius Paullus
2	P. Vinicius	P. Alfenus Varus
3	L. Aelius Lamia	M. Servilius
4	Sex. Aelius Catus	C. Sentius Saturninus
5	L. Valerius Messalla	Cn. Cornelius Cinna
suff.	C. Vibius Postumus	C. Ateius Capito
6	M. Aemilius Lepidus	L. Arruntius

Table 1: Römische Konsulen

### 3 Prediction of cell organelles

## 4 Summary

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- Matthias Liebeck and Stefan Conrad (July 2015). "IWNLP: Inverse Wiktionary for Natural Language Processing". In: *Proceedings of the 53rd Annual Meeting of the Association for Computational Linguistics and the 7th International Joint Conference on Natural Language Processing (Volume 2: Short Papers)*. Association for Computational Linguistics, pp. 414–418.
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