

In situ RNA expression of CTCs and machine learning approaches for cell classification

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Overarching focus of our research group



- We hypothesize that the appearance of CTCs and ctDNA in blood is depending on
 - the histological architecture including



and the architecture differs in shedding and non-shedding areas of the tumor

Our overall objective is to elucidate which factors contribute to the dissemination of CTCs and ctDNA into blood

1-Page Graphical Abstract "Liquid biopsy in cancer"





- Implementation of European (CEN)- and International (ISO)- standardized workflows for liquid biospies (CTC + ctDNA)
- We thereby aim to implement CTCs and ctDNA in the clinical management of
 - Prostate Cancer (PC)-
 - Colorectal Cancer (CRC) and
 - Non-small Cell Lung Cancer- (NSCLC) Patients
- with sampling at multiple time points



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Prostate cancer - The aim





Figure 1. Swimmer plot indicating treatments that patients received, along with timing (and AR-V7 status) of CTC sampling, and whether or not PSA responses occurred during each therapy. Shaded boxes indicate failure to achieve PSA response; unshaded boxes indicate achievement of 50% PSA reduction on therapy. Percentage values indicate best PSA response. Daggers indicate deceased patient. Thirteen of 14 patients have previously been included in our prior publications, but only in the context of a single therapy.



Adapted from Nakazawa et al., Ann Oncol 2015

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Smart Biopsy CTC enrichment





1) Double negative selection

2) Size based gravity flow filtration

3) Cytospin of cell fraction for in situ padlock probe approach

- Depletion of blood cells
- Density gradient centrifugation

Kim et al., Analytical Biochemistry 2013 Lee et al., Oncology Letters 2017 <u>http://www.cytogenlab.com/</u>

Method: In Situ combined with Padlock Probes







In situ padlock technology = Visualisation of RNA transcripts

Modified from Larsson et al., 2010, Nature Methods Modified from Hofmann et al., MIMB 2019 Modified from Landegren., personal communication

Supervised machine learning - create training set



• Use drag and drop to sort cells into the classes

CPA/Classifier - DefaultDB_MyExpt_Cell_border_object.properties
File View Rules Nucleus cy5 cy7 cy3 atto425 atto488 texasred



Supervised machine learning - create training set



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6) Data visualization

Dimension reduction - t-SNE plot



Supervised machine learning count



Manual count

Comparison to manual analysis:

- cy3 negative = VCaP cell
- cy3 positive = PBMC

Cell count VCaP: No Cell: Negative= No RCP PBMC:

661 (cy3-) Not counted manualy 118 138 (cy3+ only)

Workflow for *in situ* padlock probe (PLP) analysis + machine learning classification

Multiplex in situ padlock probe assay

- CTC enriched patient samples (SmartBiopsy Cell Isolator)
- Visualize mRNA markers as fluorescent spots
- Panel of PBMC- and CTC-markers and VIM





Hmec

CellProfiler: image analysis

Signal detection, decoding, and feature extraction





CellProfiler Analyst:

supervised machine learning-based classification





Expert revision of candidate cells



Pro-apoptotic



El-Heliebi Amin **Do not post, unpublished data** Adapted from Sallinger, El-Heliebi et al., poster presentation ACTC 2021 Gyllborg et al., Nucleic Acid Research 2020 1 mm





- In situ padlock probes can be used to detect any expressed transcript
 - > = visualization of RNA
- + Can be quantified
- + Can be used instead of unspecific antibodies
- Very low expressed transcripts are challenging

Research Group El-Heliebi



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