



s.dewit@utwente.nl

LB-250

AACR 2017

Liquid biopsy in advanced NSCLC: EpCAM+ and EpCAM- circulating tumor cells, tumor derived extracellular vesicles and cell-free circulating tumor DNA

Sanne de Wit¹, Menno Tamminga², Ellen Heitzer³, Joost F Swennenhuis¹, Ed Schuurin², Leonie L Zeune¹, Michael R Speicher³, T Jeroen N Hiltermann², Leon WMM Terstappen¹, Harry JM Groen²

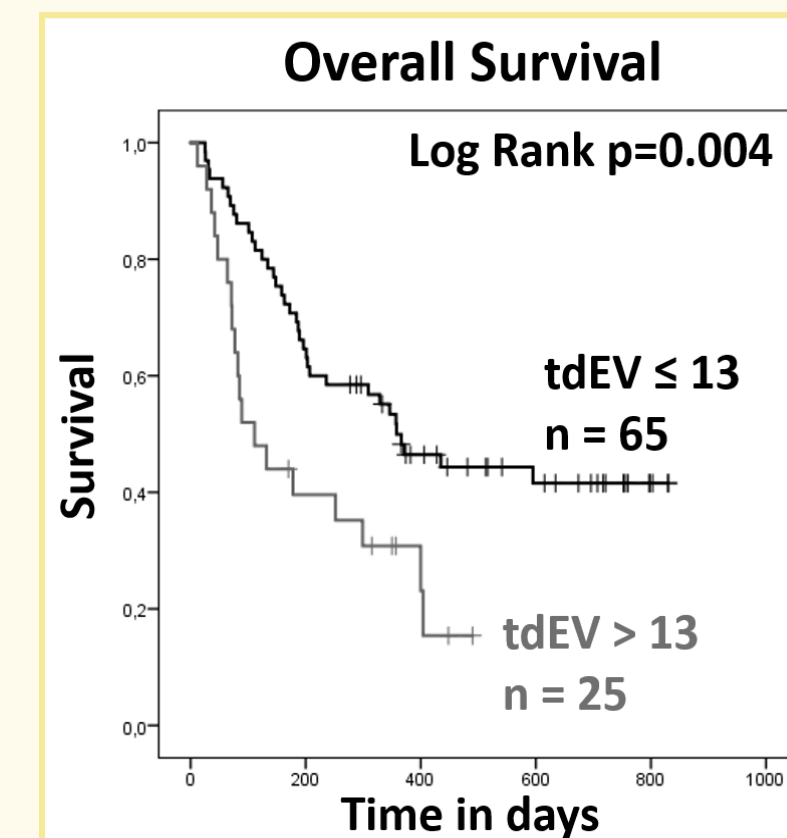
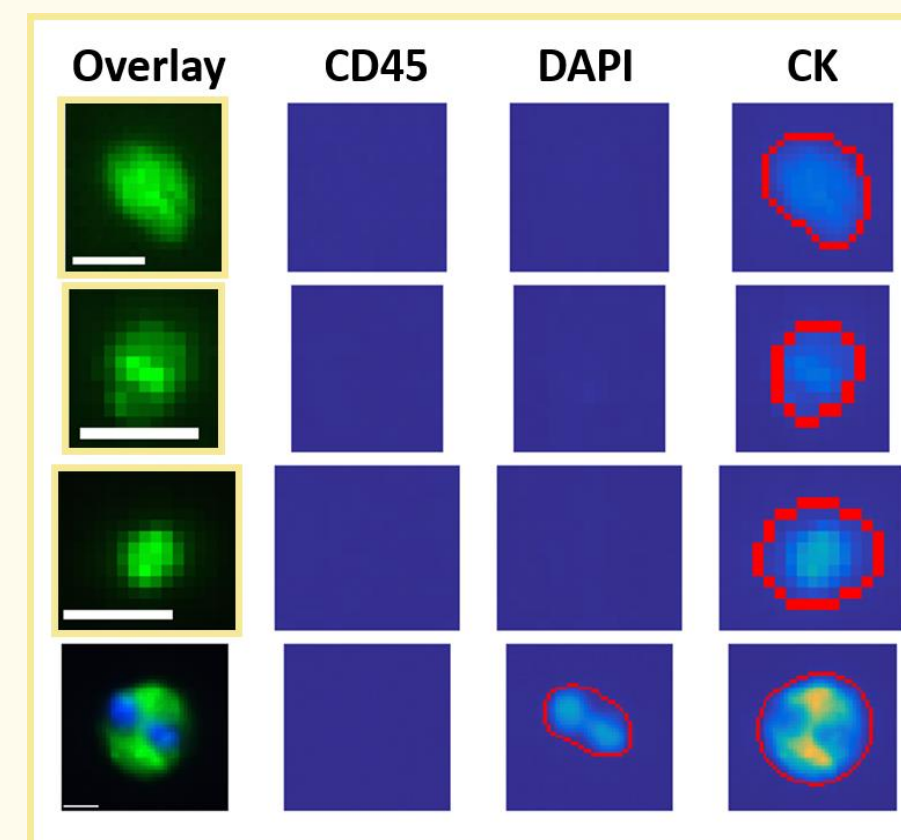
¹ Department of Medical Cell BioPhysics, MIRA institute, University of Twente, The Netherlands, ² Department of Pulmonology, University Medical Center Groningen, the Netherlands, ³ Institute of Human Genetics, Medical University of Graz, Austria

DEFINITION tdEV are EpCAM+, DAPI-, cytokeratin+, CD45-, slightly round, surface < 150 μm^2 , perimeter > 4 μm .

METHOD Analysis of the CellSearch cartridge (n=90) with open source program ACCEPT after processing 7.5 mL blood for detection of CTC. Cut-off value >13 is based on mean+1SD of 127 healthy controls (HC).

CONCLUSION Presence of tdEV is significantly associated with poor overall survival.

Patients (n=90)	HC
tdEV ≤ 13	72% 12%
tdEV > 13	28% 88%
Mean	21.6 6.7 (± 6.3)
Min	0 0
Max	381 37
Median	7 5



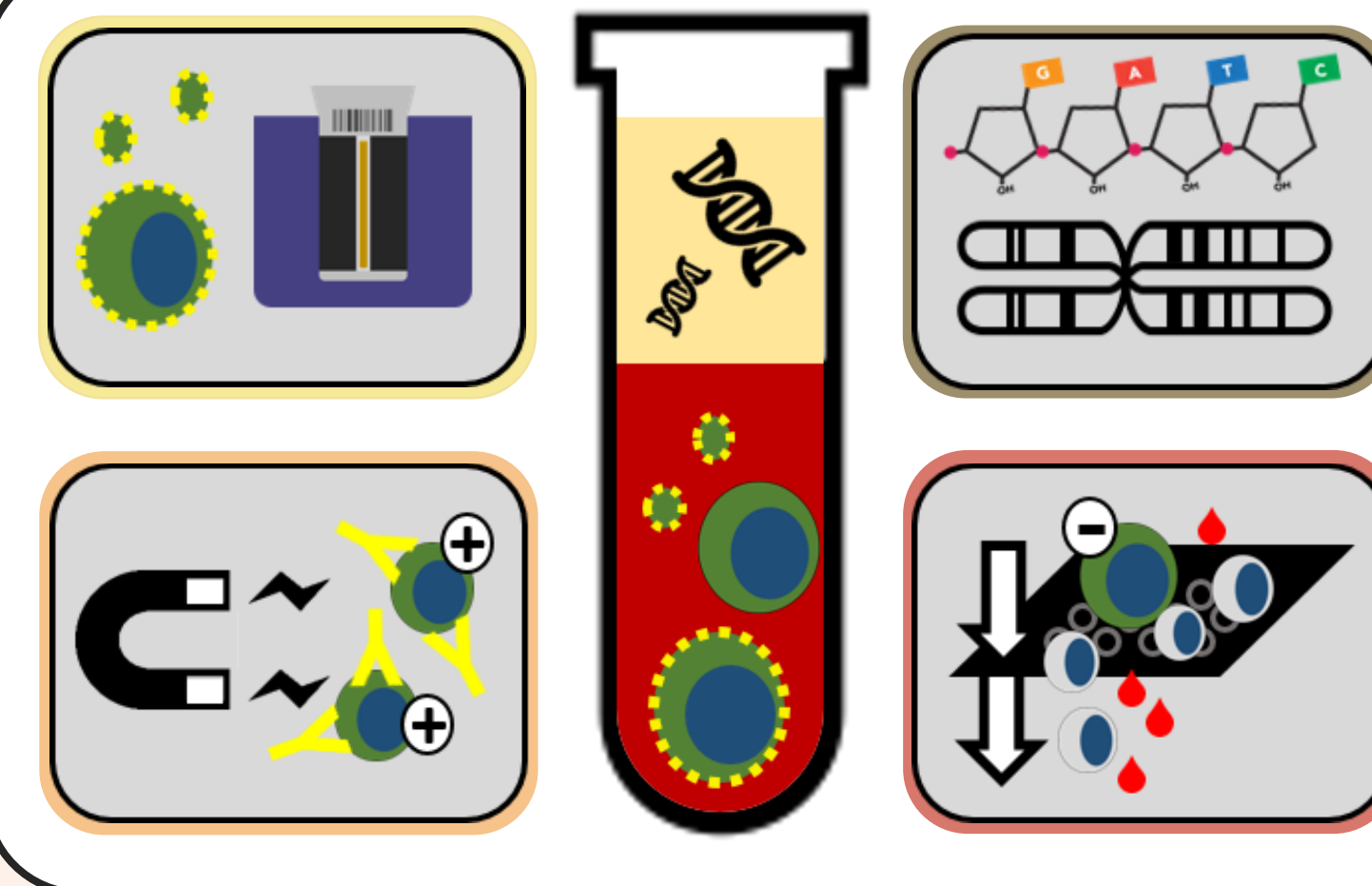
WHY The need for a liquid biopsy in non-small cell lung cancer (NSCLC) patients is rapidly increasing as more targeted therapies become available.

WHAT Four biomarkers are explored for their potential to represent a liquid biopsy.

WHO In metastatic NSCLC patients before treatment we investigate the biomarkers in relation with overall survival.

HOW In just one 7.5 mL CellSave tube of blood.

LIQUID



BIOPSY

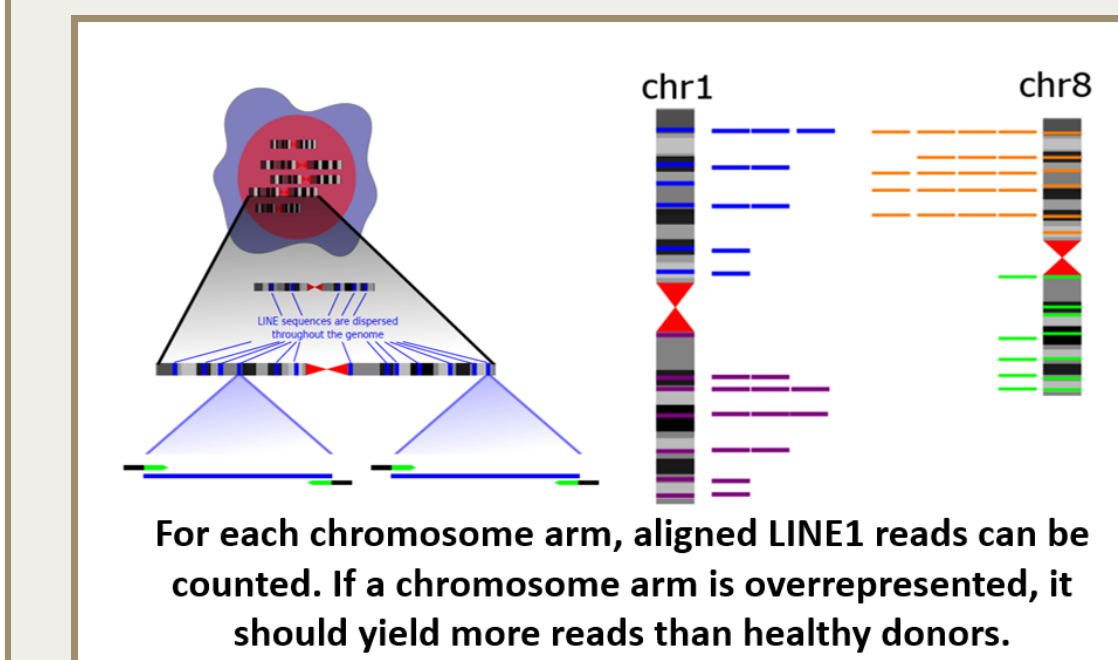
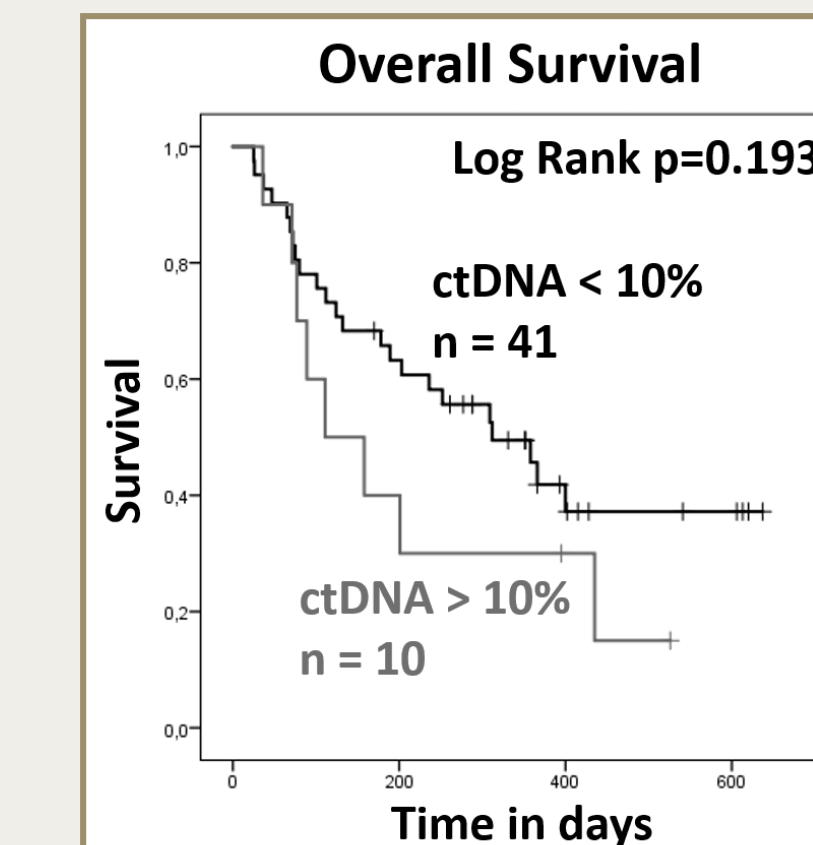
CONCLUSIONS

tdEV showed the strongest association with overall survival. Addition of any combination of the biomarkers did not increase this association. Remaining question is what the efficiency is to extract treatment relevant information from these biomarkers.

Patients (n=50)	
All 4 biomarkers positive	6%
3 biomarkers positive	12%
2 biomarkers positive	20%
1 biomarker positive	58%
Correlation EpCAM+ CTC	p=0.019
Correlation EpCAM- CTC	p=0.571
Correlation tdEV	p=0.134
Correlation ctDNA	p=0.082

DEFINITION DNA present in plasma originating from the tumor

METHOD Plasma was collected from the CellSave tube and ctDNA concentration was measured with the mFAST-SeqS approach (n=51). This approach relies on the amplification of uniquely mappable *LINE1*-sequences across the genome and can be used as a general measure of aneuploidy in a plasma sample. Detection limit of ctDNA concentration is $\geq 10\%$ mutant alleles.



CONCLUSION ctDNA concentration did not significantly correlate to overall survival, but might be reached by increasing the number of patients.

Patients (n=51)	
ctDNA < 10%	74%
ctDNA > 10%	26%
Min	0.8
Max	66.7
Median	2.3

Tumor Derived Extracellular Vesicles

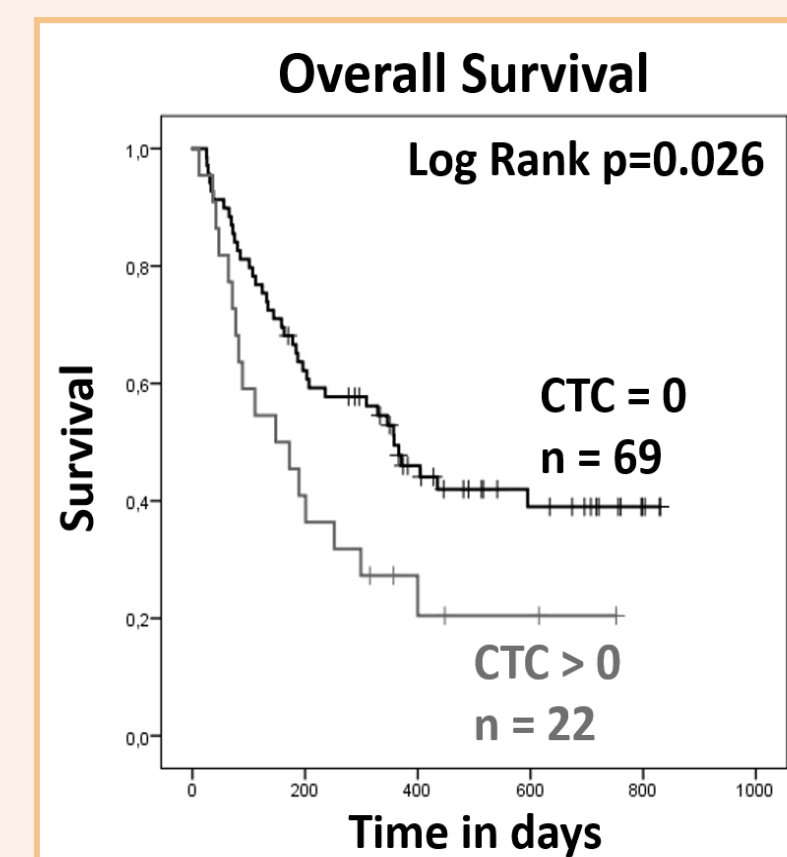
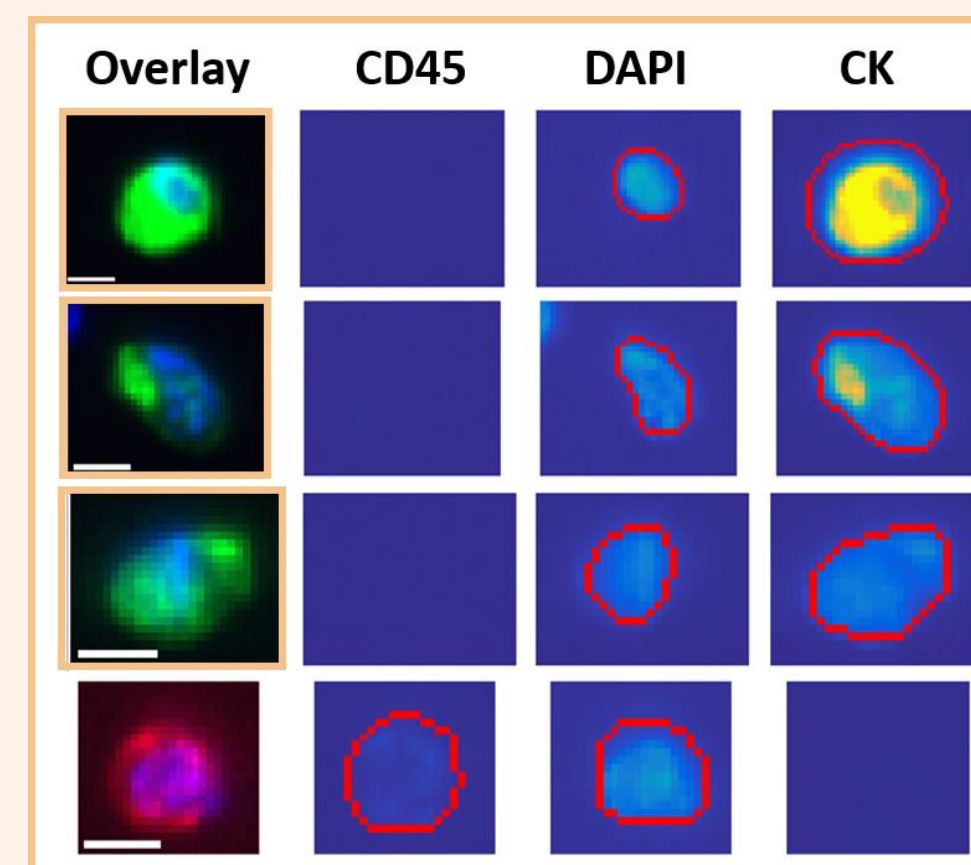
EpCAM+ Circulating Tumor Cells

DEFINITION CTC are EpCAM+, DAPI+, cytokeratin+, CD45-, round, >4 μm in size, DAPI-CK overlay >50%.

METHOD 7.5 mL blood measured with CellSearch for detection of CTC by immunomagnetic selection (n=91 and HC n=39).

CONCLUSION Presence of EpCAM+ CTC is significantly associated with poor overall survival.

Patients (n=91)	HC
CTC = 0	76% 97%
CTC ≥ 1	24% 3%
CTC ≥ 3	13% 0%
CTC ≥ 5	5% 0%
Min	0 0
Max	186 1
Median	0 0

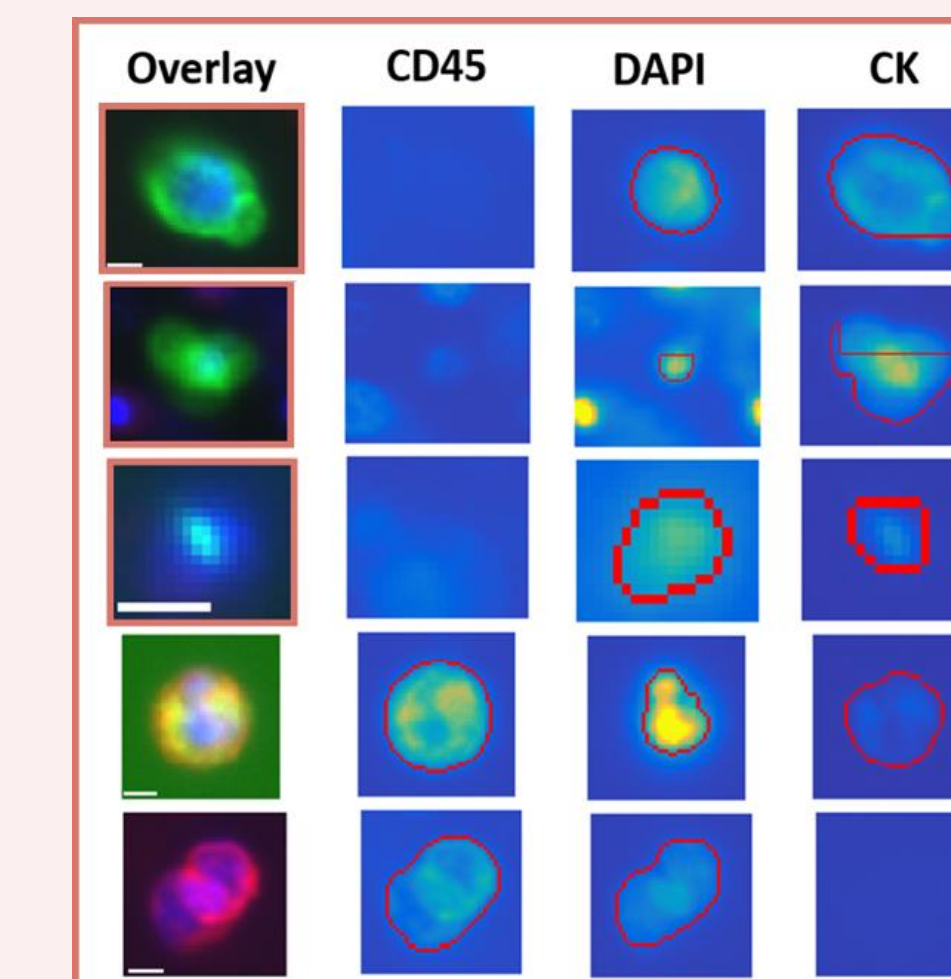
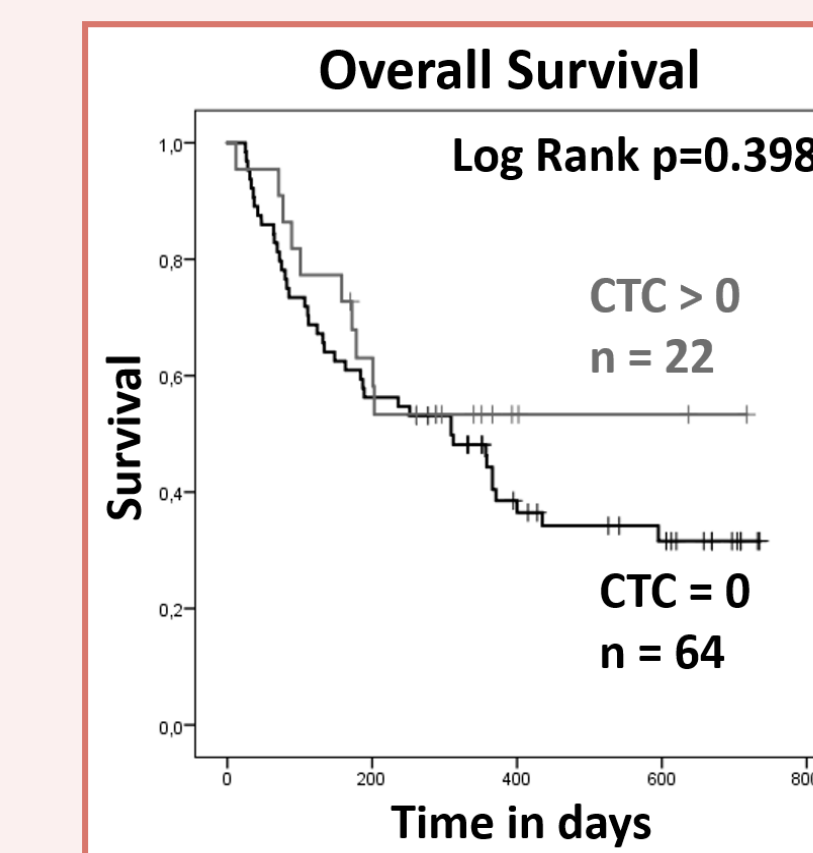


Circulating Tumor DNA

EpCAM- Circulating Tumor Cells

DEFINITION CTC are EpCAM-, DAPI+, cytokeratin+, CD45-, DAPI-CK overlay.

METHOD Blood discarded by CellSearch after immuno-magnetic isolation was filtered through 5 μm pores and stained with a CK-antibody cocktail (n=86). HC (n=27) spiked with ~ 300 EpCAM- NSCLC cell line NCI-H1650 cells (1.4×10^2 EpCAM antigens and size 12 μm): mean recovery = 31% [min 11-max 350].



CONCLUSION Blocking of the filter influences CTC recovery. Presence of EpCAM- CTC are not correlated with overall survival.

Patients (n=86)	HC
CTC = 0	76% 76%
CTC ≥ 1	24% 24%
CTC ≥ 3	13% 6%
CTC ≥ 5	5% 6%
Min	0 0
Max	186 12
Median	0 0