

# Digital cytology- Applications in cervical cancer screening

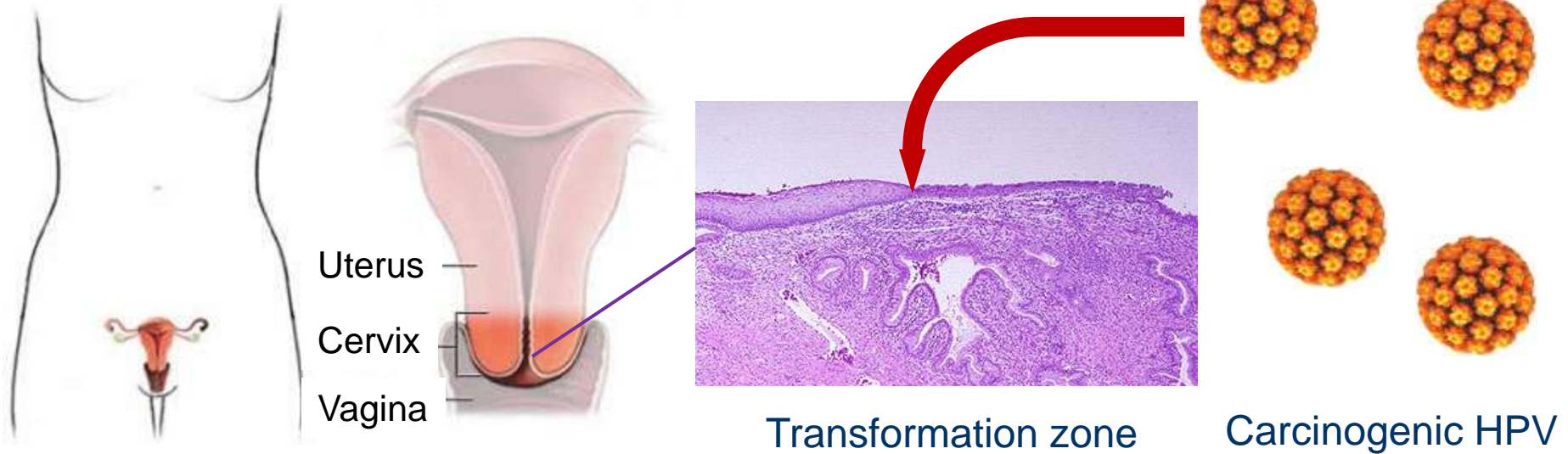
**PD Dr. med. Nicolas Wentzensen**

Hormonal and Reproductive Epidemiology Branch  
Division of Cancer Epidemiology and Genetics, NCI, NIH  
Bethesda, USA

# Statements

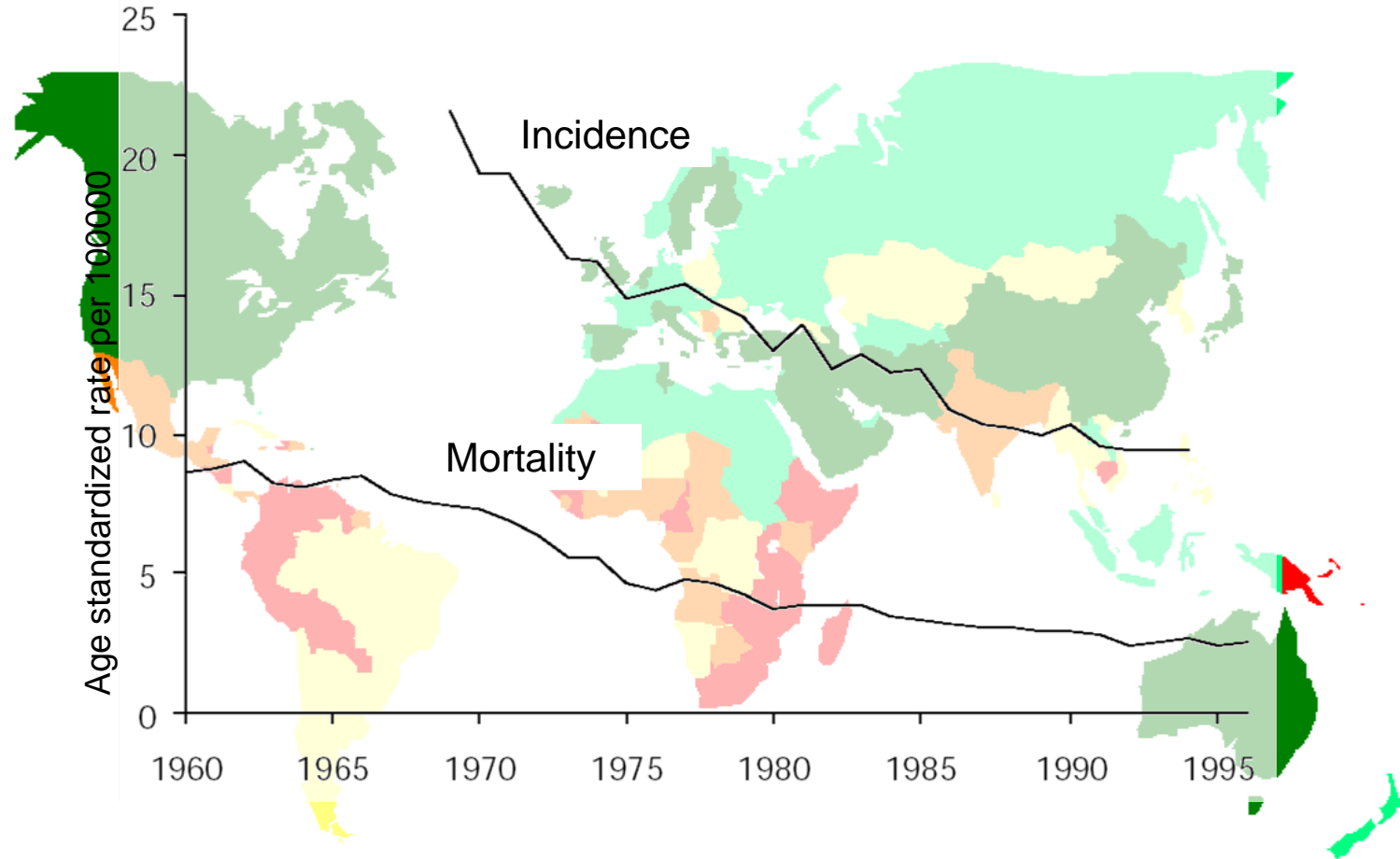
- I have no conflict of interest
- These are personal opinions and not official NCI statements

# Cervical cancer



- Second most frequent cancer in women worldwide
- Caused by persistent infections with carcinogenic human papillomaviruses (most importantly 16,18)

# Cervical cancer incidence and Pap based screening



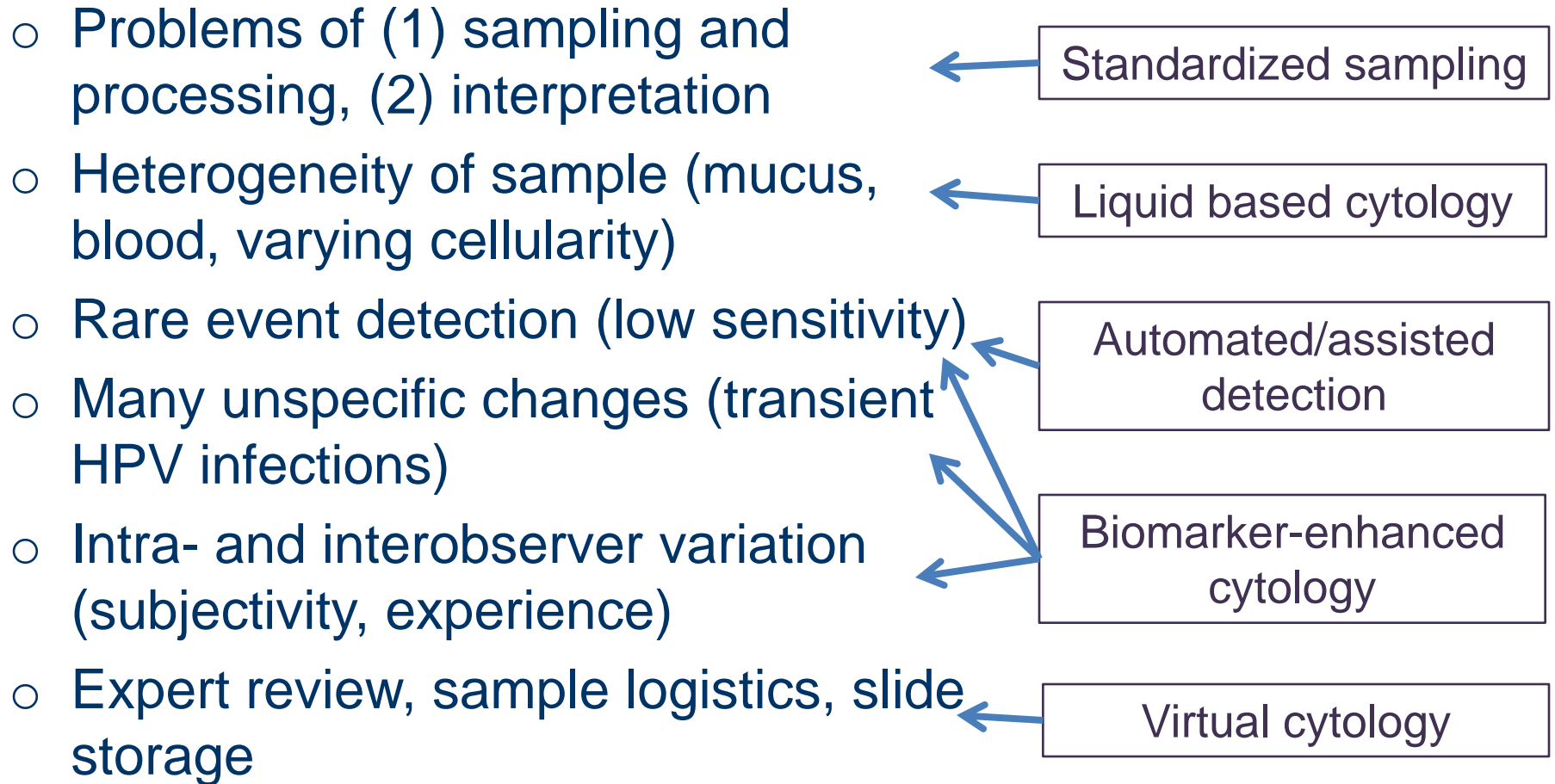
Reduction of cervical cancer incidence after introduction of screening (Canada)

# Characteristics of the Pap smear

- Introduced by Papanicolaou in the 1940s
- A smear is taken from the cervix and transferred to a glass slide, stained with several stains (nuclear, cytologic)
- Abnormal/precancerous cells are detected by morphology
- Presence of abnormal cells triggers further management and treatment



# Limitations of current cytological screening



Large epidemiologic/interventional studies are necessary to demonstrate the efficiency of these approaches

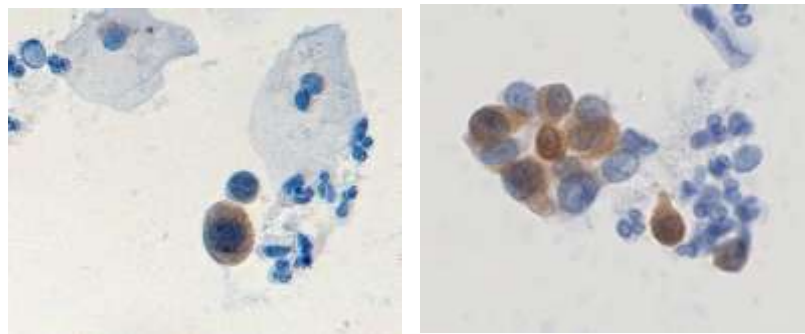
# Cytiga- a virtual cytology system



Virtual cytology



Biomarker enhanced cytology

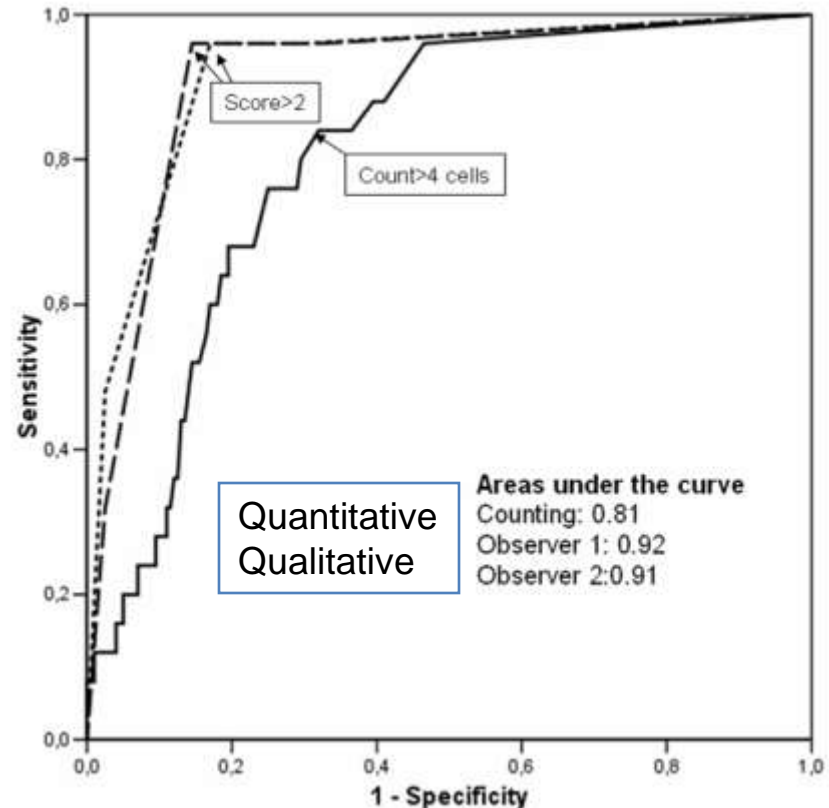
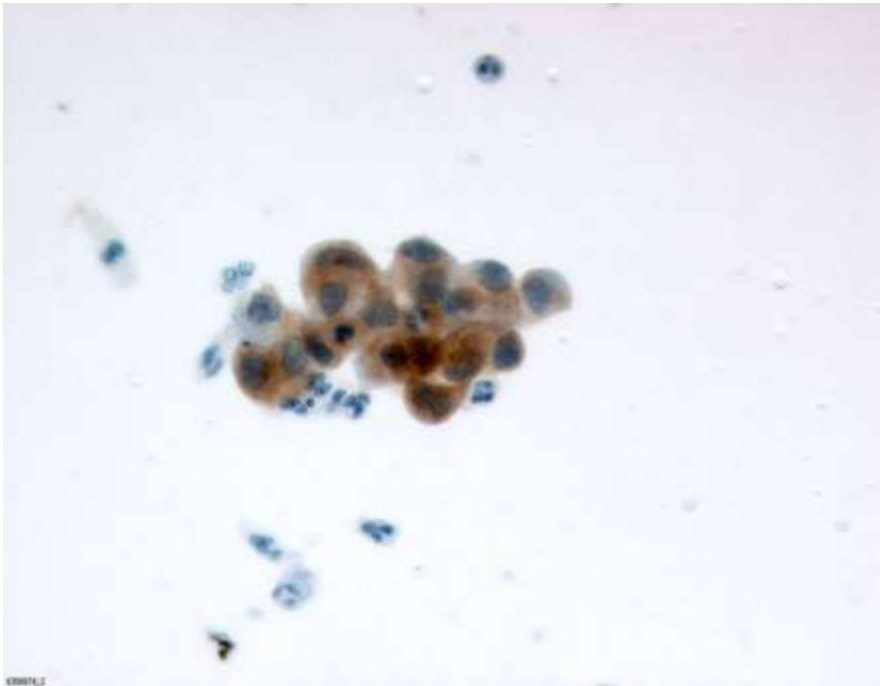


Automated/assisted detection



# The biomarker example: p16 ICC

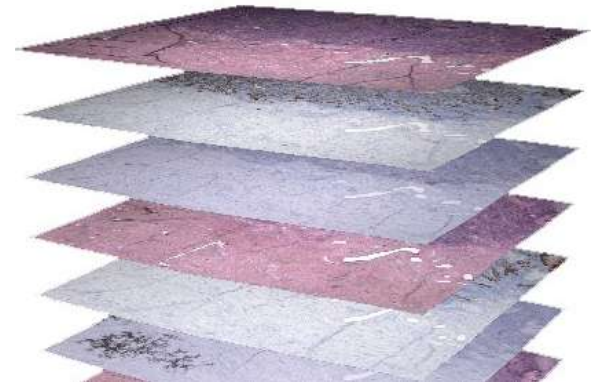
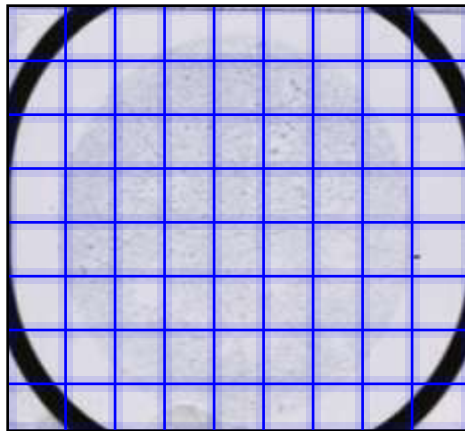
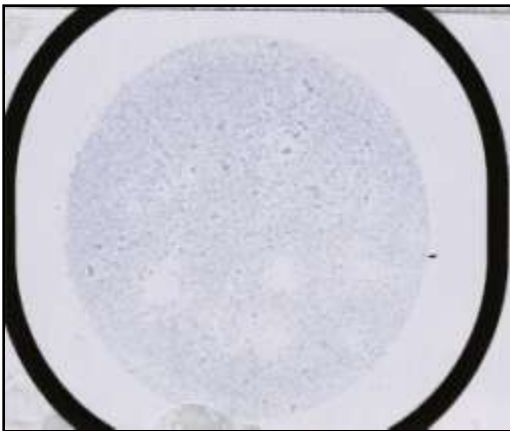
- Direct link to HPV carcinogenesis: high sensitivity and specificity for detecting cervical precancers
- Improved rare event detection (locator function)
- Qualitative (nuclear morphology) and quantitative (cell counting) evaluation possible





# Virtual cytology

- Slide is scanned, image is stored in tiles
- Image can be accessed via internet through a web browser plugin, only the active area is transmitted
- Z-stack option: Multiple images are superimposed to analyze 3D cells



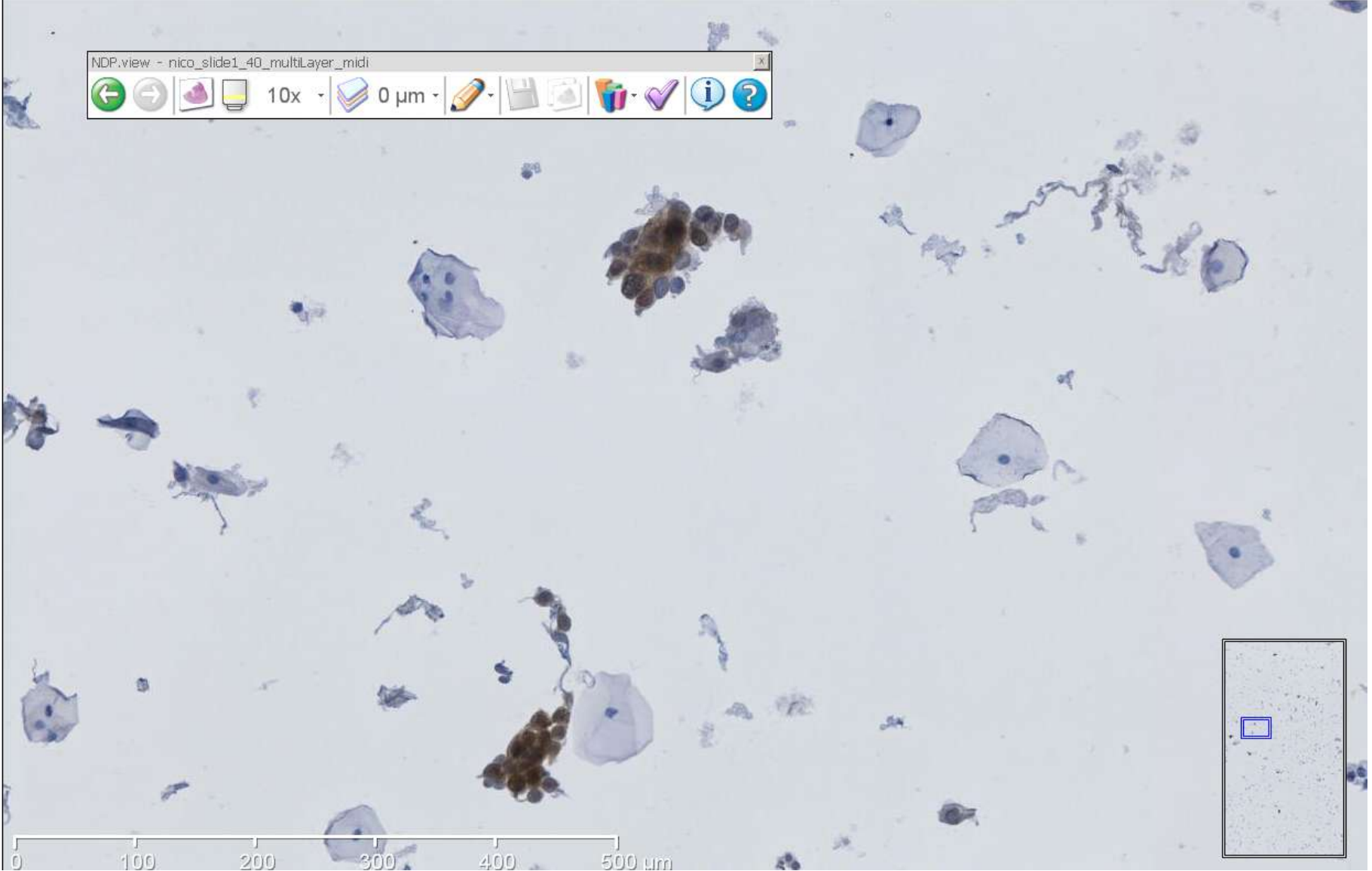
# Using a virtual slide



NDP.view - nico\_slide1\_40\_multiLayer\_midi



Navigation and editing icons: back, forward, home, zoom, 10x, 0 μm, pencil, eraser, selection, checkmark, info, help.



0 100 200 300 400 500 μm



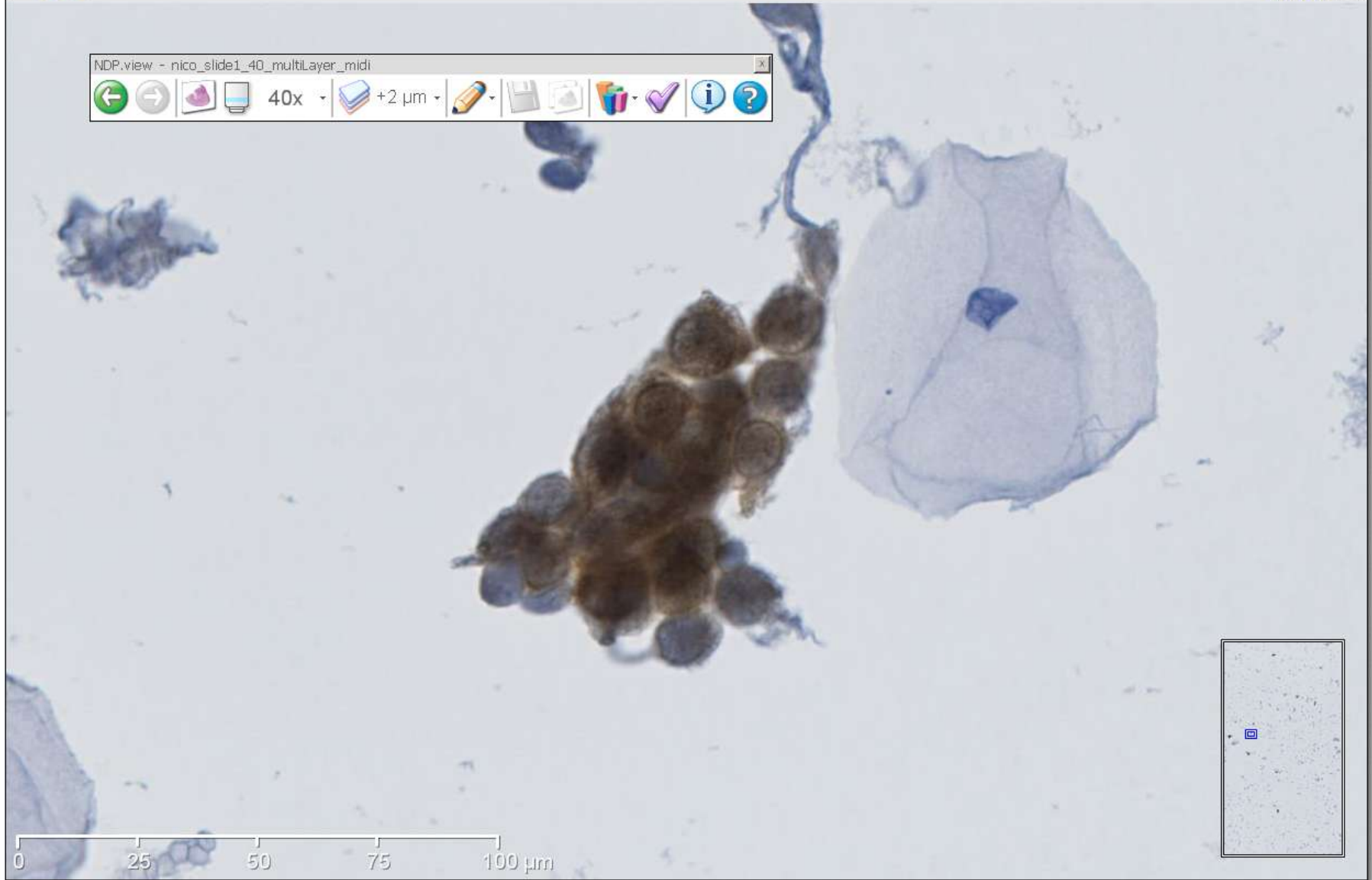
# nico\_slide1\_40\_multiLayer\_midi

Welcome Dr. Nicolas Wentzensen  
(Sign out)

NDP.view - nico\_slide1\_40\_multiLayer\_midi

Navigation icons: Back, Forward, Home, Stop, Refresh, Print, Mail, New Folder, Favorites, Home

40x | +2  $\mu$ m | [Pencil icon] | [Save icon] | [Print icon] | [Share icon] | [Heart icon] | [Info icon] | [Help icon]



0 25 50 75 100  $\mu$ m





# nico\_slide1\_40\_multiLayer\_midi

Welcome Dr. Nicolas Wentzensen  
(Sign out)

NDP.view - nico\_slide1\_40\_multiLayer\_midi

Navigation icons: Back, Forward, Home, Print, Mail, New Window, Copy, Paste, Undo, Redo, Erase, Fill, Stroke, Stroke Width, Stroke Color, Stroke Style, Stroke Dash, Stroke Opacity, Stroke Weight, Stroke Color, Stroke Style, Stroke Dash, Stroke Opacity, Stroke Weight, Info, Help

40x | 0  $\mu\text{m}$



0 25 50 75 100  $\mu\text{m}$





# nico\_slide1\_40\_multiLayer\_midi

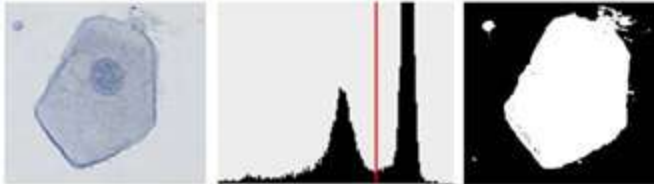
Welcome Dr. Nicolas Wentzensen  
(Sign out)

NDP.view - nico\_slide1\_40\_multiLayer\_midi

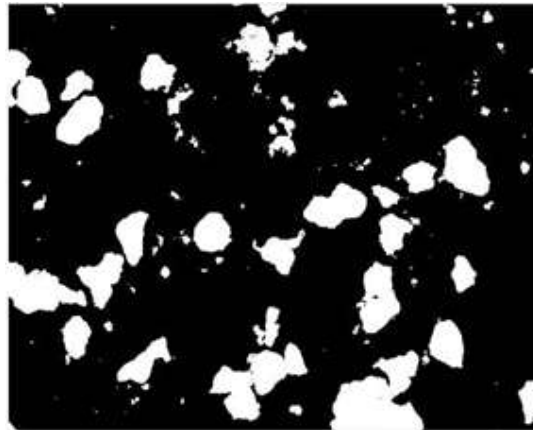
Navigation icons: Back, Forward, Home, Print, 40x, -2 μm, Eraser, Save, Print, Layers, Checkmark, Info, Help



# Cell detection



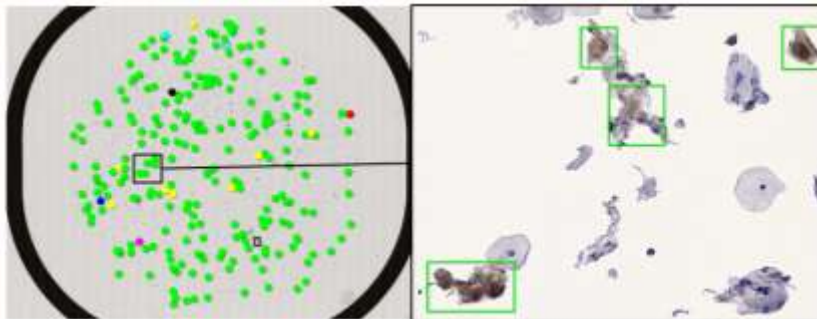
Set color-based cutoff  
between cells and background



Apply cutoff on slide



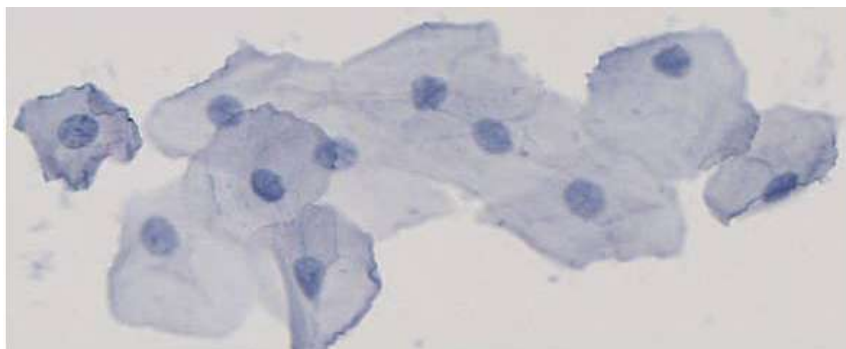
Small object removal



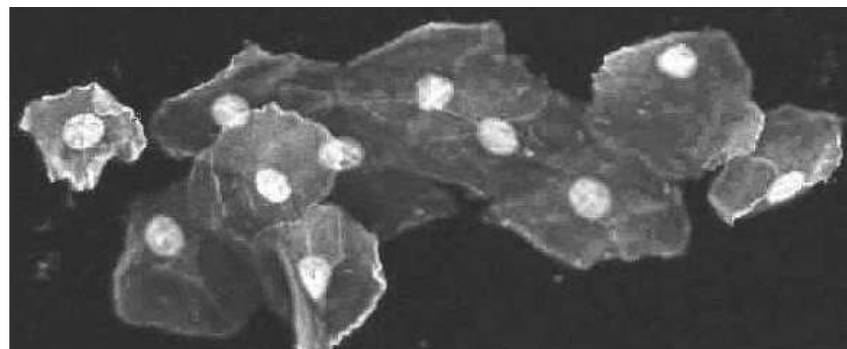
Overlay annotations on  
original image



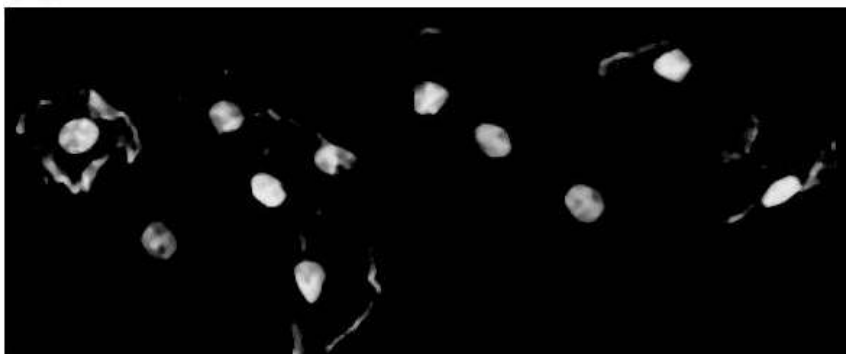
# Nucleus detection



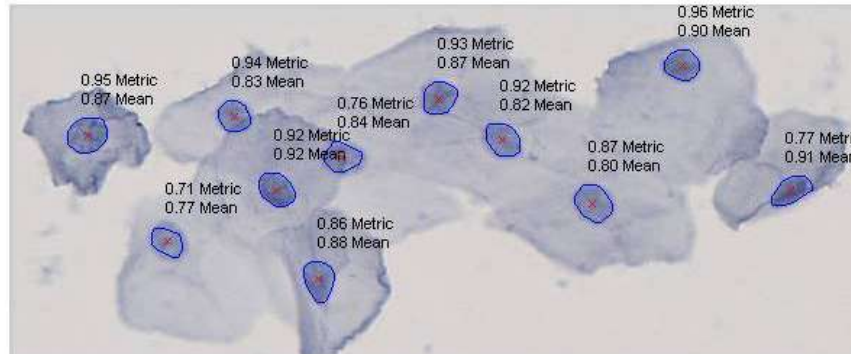
Original image



Probability function for nuclear color



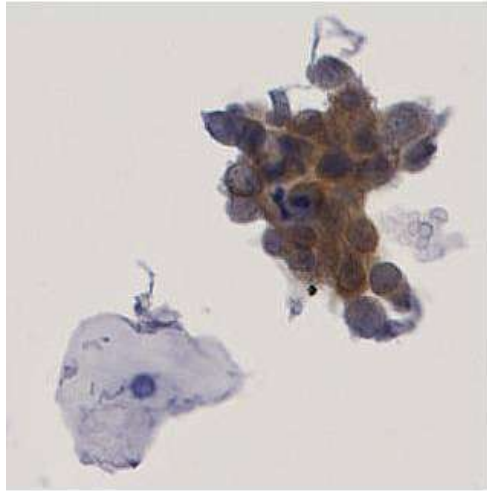
Subtraction of cytoplasm



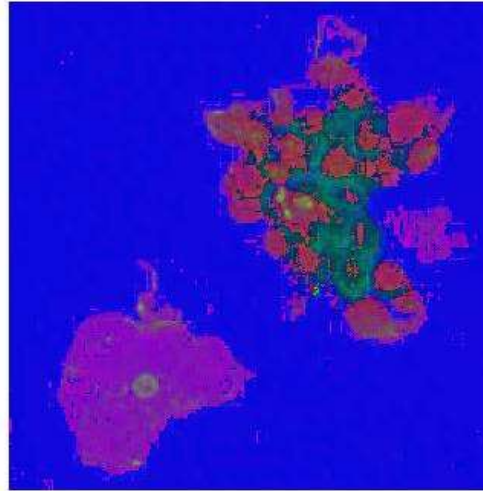
Nuclear annotation on original image  
(probability and roundness)



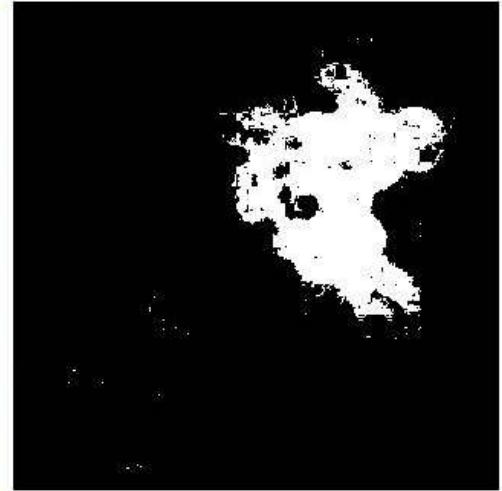
# Biomarker detection



Original image



HSV color space



Applying threshold for  
HSV color of p16 stain

# Detection of p16 - stained cells

		Man		
		positive	negative	total
Machine	positive	98	22	120
	negative	11	~50,000	50,011
	total	109	50,022	50,131

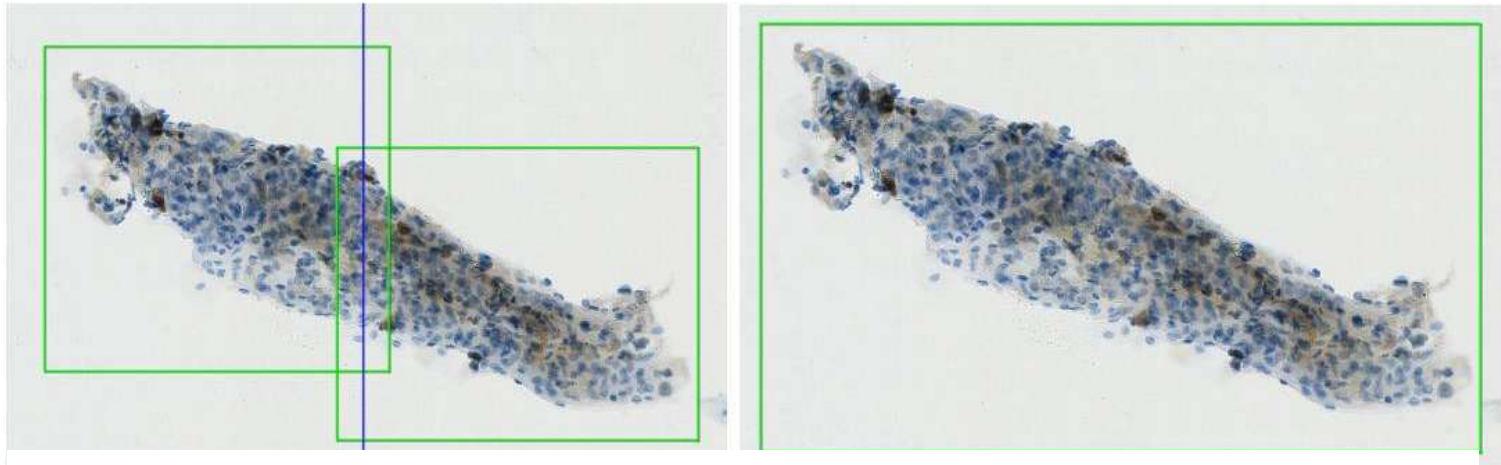


McNemar's test: Not significant

Man: 109/125 (87%)  
Machine: 120/125 (96%)

- High agreement between automated and manual detection of p16 positive cells
- Few discrepancies related to missed cells in manual evaluation and faint staining / artifacts in automated evaluation

# Challenges and artifacts



large object spanning multiple tiles



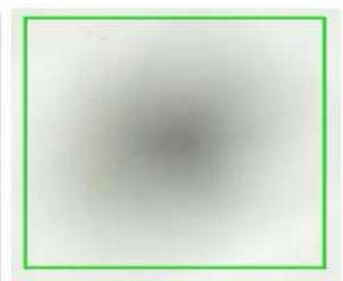
air



unfocused cell



dirt/precipitate



dust

# Cell annotation

Cy@TIGA  
CYTOLOGY @ TISSUE IMAGING AND ANALYSIS CENTER

fk\_1

(Back) (Help) (Sign out)

Annotations  
← 52 →

Go to annotation Nr:  Go

Scoring

- Score 1
- Score 2
- Score 3
- Score 4
- Artifact

Given score:

Comments  
no comment

Submit

Move direction

0 50 100 150 200  $\mu\text{m}$

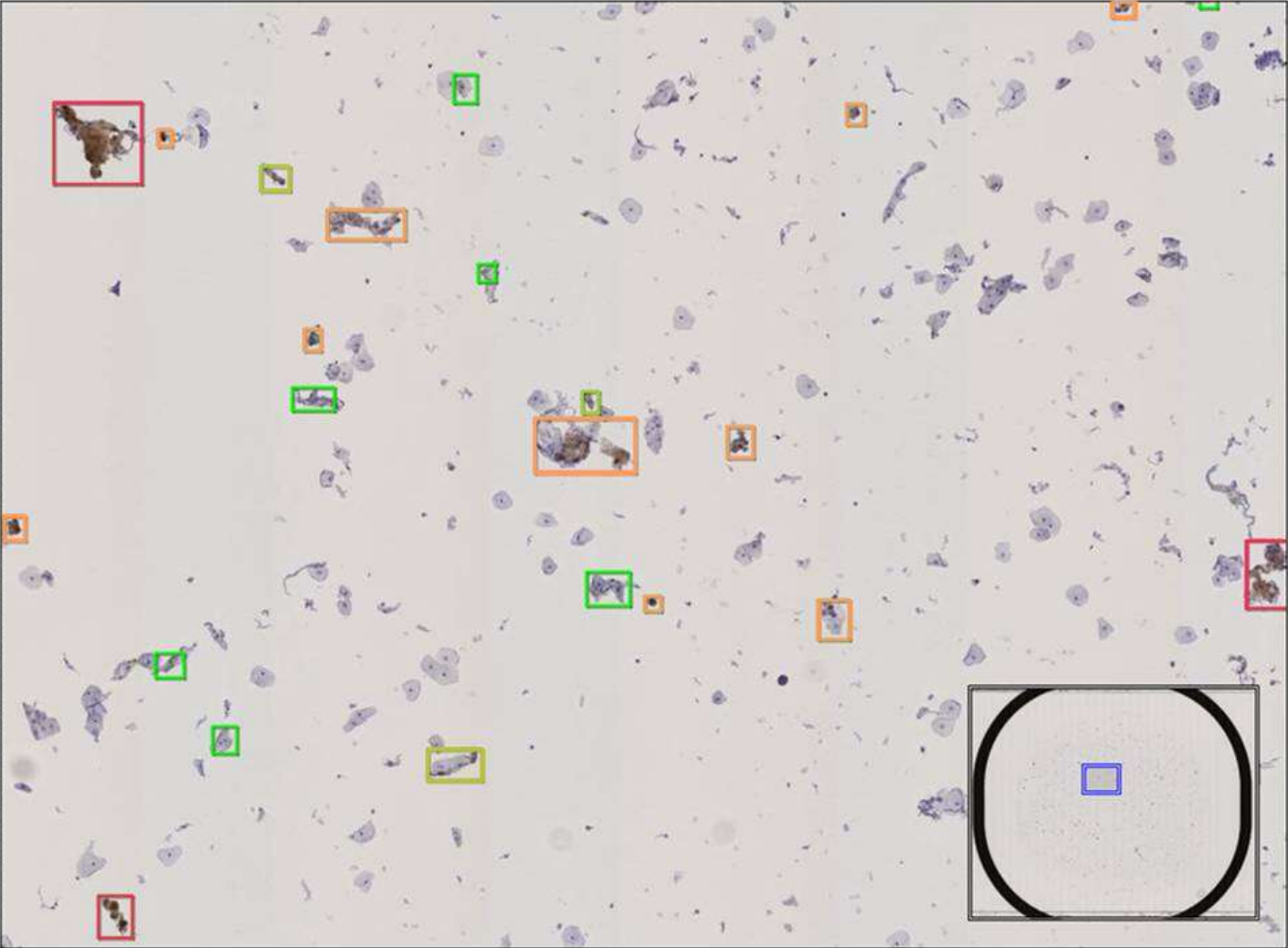
Total amount of cells: 72512. Total amount of cluster: 67.

- Cell detection
- Cell counting
- Nucleus detection
- Nuclear measures
- Biomarker detection
- Flexible annotation system for open and blinded review

# Database for slide evaluation

**Cy@TIGA** CYTOLOGY & TISSUE IMAGING AND ANALYSIS CENTER

first (Back) (Sign out)



Annotations

← 0 →

Go to annotation Nr:

Scoring:

- Score 1
- Score 2
- Score 3
- Score 4
- Skip
- Artifact

Comments:

Move direction

⊖ + - ← → ↑ ↓


Show / Hide Annotations

Image Controls:

Export Image

Name:

Path:



# Next steps

- Fine-tuning of biomarker detection
- Expand automated evaluation
- Analyze large series of p16 stained slides, e.g. from NCI cervical cancer screening studies
- Challenges for cytology:
  - New QC regulations (case loads)
  - HPV vaccination against 16, 18
  - Primary HPV testing
- Many opportunities for biomarker enhanced virtual cytology
- **Goal: Create a system for biomarker research and for routine cytology use**



# Team



UniversitätsKlinikum Heidelberg



Niels Grabe  
**Bernd Lahrmann**  
Thora Pommerencke



Magnus von Knebel Doeberitz  
Miriam Reuschenbach  
Denise Eschenbach  
Nina Hensch

NATIONAL  
CANCER  
INSTITUTE

Nicolas Wentzensen