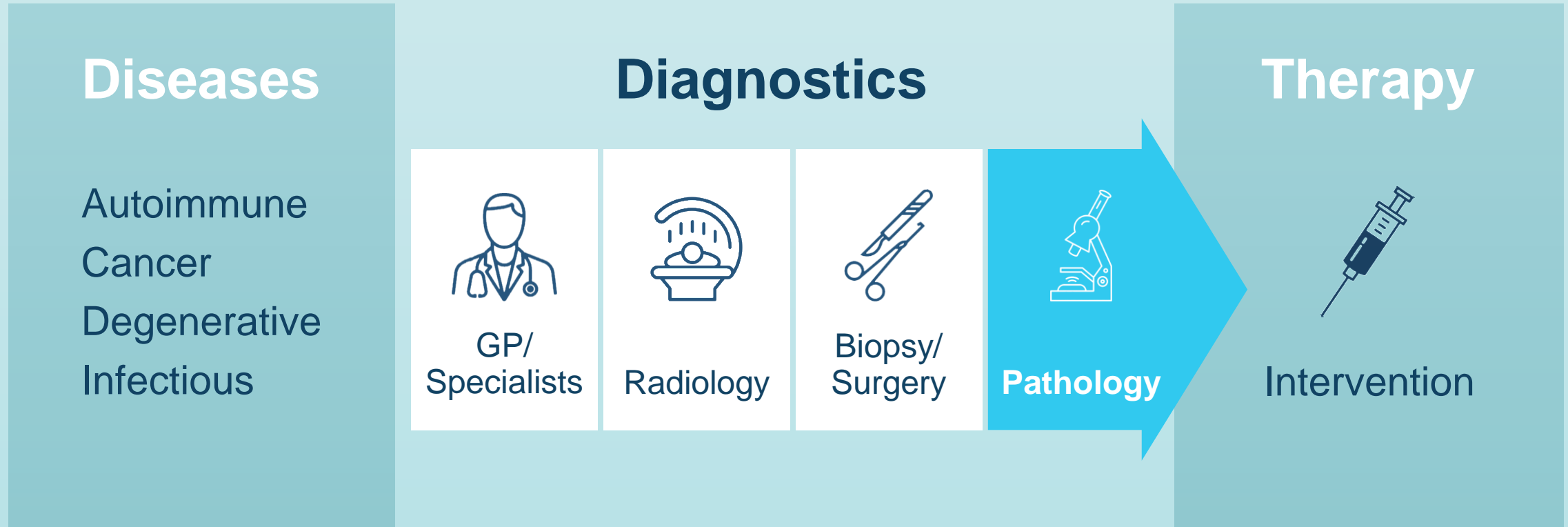


AI in cancer diagnostics: Application and validation

AI for Good Global Summit
Geneva, 29. 5. 2019

Frederick Klauschen
Charité and Berlin Institute of Health

Histopathology: Diagnose diseases, guide therapy decisions



350 Mio. diagnostic cases worldwide per year!

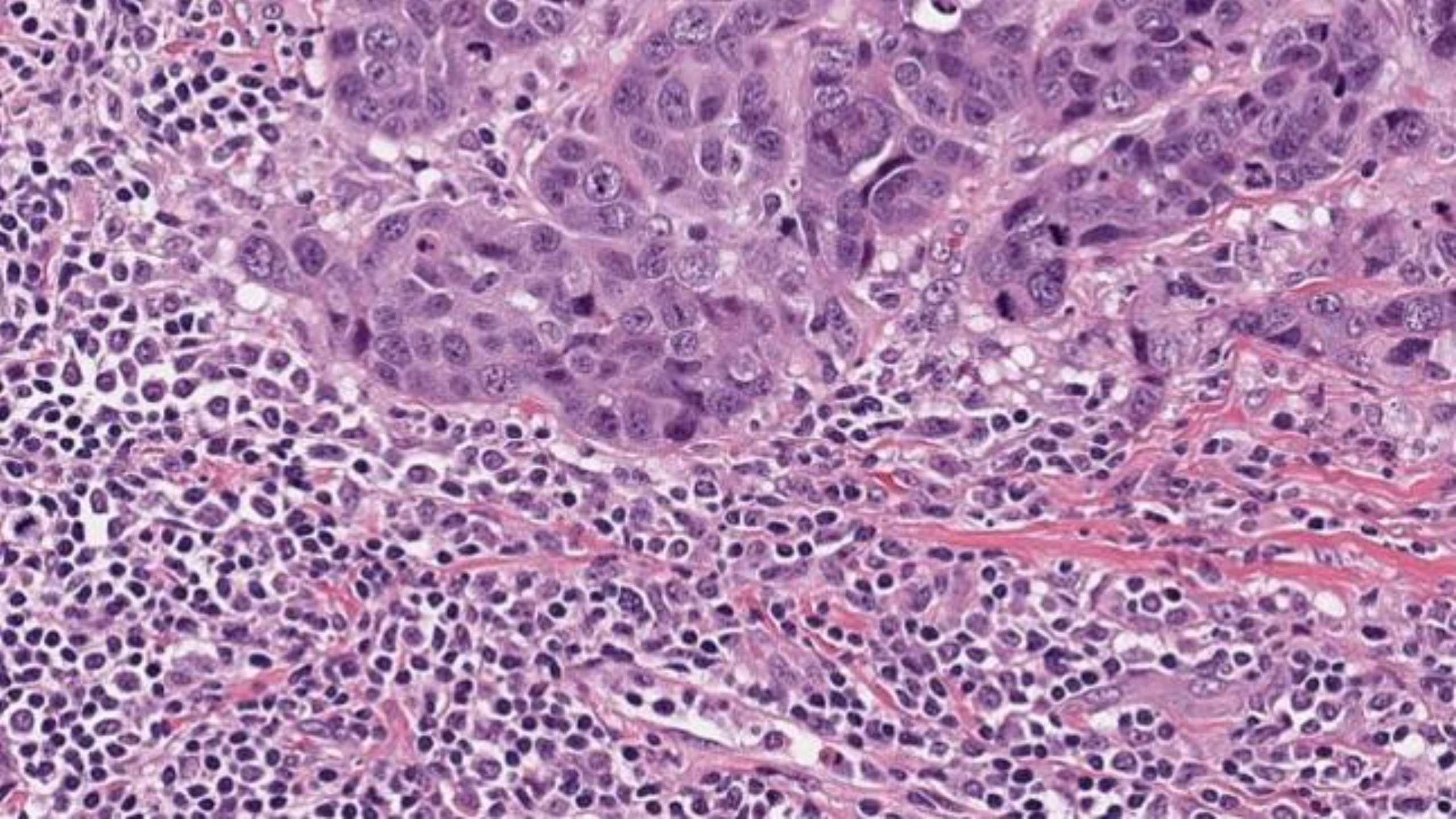


Histological slide

Microscopic diagnostics

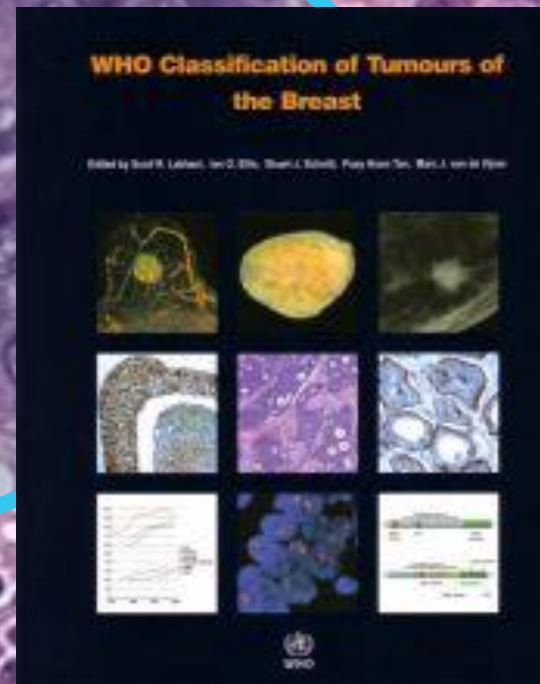


Manual evaluation



**Identify & classify
cancer!**

**Estimate
Immune cells!**



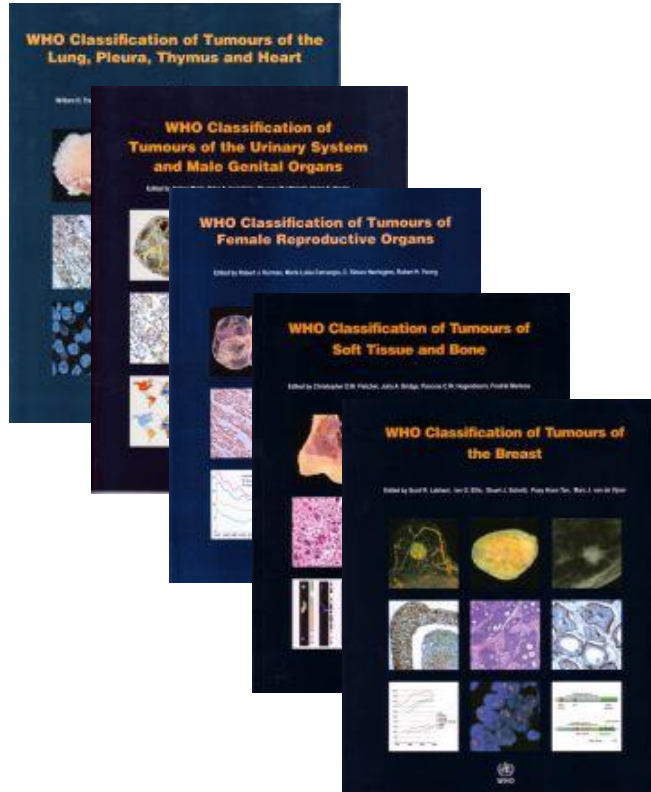
Re-classification of tumors based on molecular profiling

Multiplatform Analysis of 12 Cancer Types Reveals Molecular Classification within and across Tissues of Origin

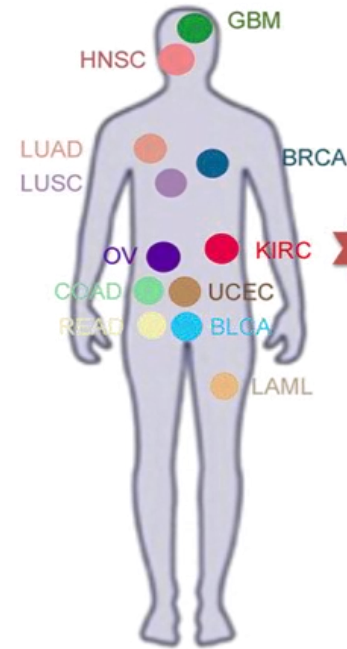
Cell

Cell 2014 158, 929-944.
Cell 2018.

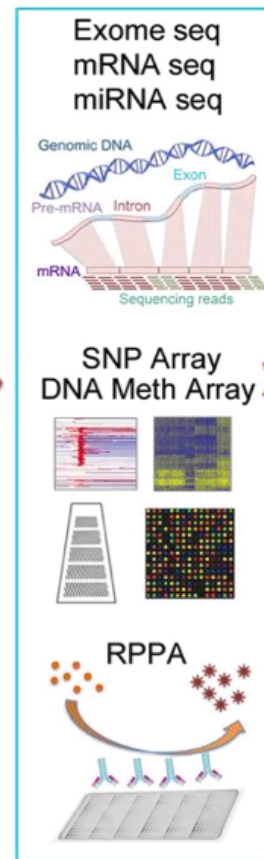
Katherine A. Hoadley,^{1,20} Christina Yau,^{2,20} Denise M. Wolf,^{3,20} Andrew D. Cherniack,^{4,20} David Tamborero,⁵ Sam Ng,⁶ Max D.M. Leiserson,⁷ Beifang Niu,⁸ Michael D. McLellan,⁹ Vladislav Uzunangelov,⁹ Jiashan Zhang,⁹ Cyriac Kandoth,⁸ Rehan Akbani,¹⁰ Hui Shen,^{11,22} Larsson Omberg,¹² Andy Chu,¹³ Adam A. Margolin,^{12,21} Laura J. van't Veer,³ Nuria Lopez-Bigas,^{5,14} Peter W. Laird,^{11,22} Benjamin J. Raphael,⁷ Li Ding,⁸ A. Gordon Robertson,¹³ Lauren A. Byers,¹⁰ Gordon B. Mills,¹⁰ John N. Weinstein,¹⁰ Carter Van Waes,¹⁸ Zhong Chen,¹⁹ Eric A. Collisson,¹⁵ The Cancer Genome Atlas Research Network, Christopher C. Benz,^{2,*} Charles M. Perou,^{1,16,17,*} and Joshua M. Stuart^{6,*}



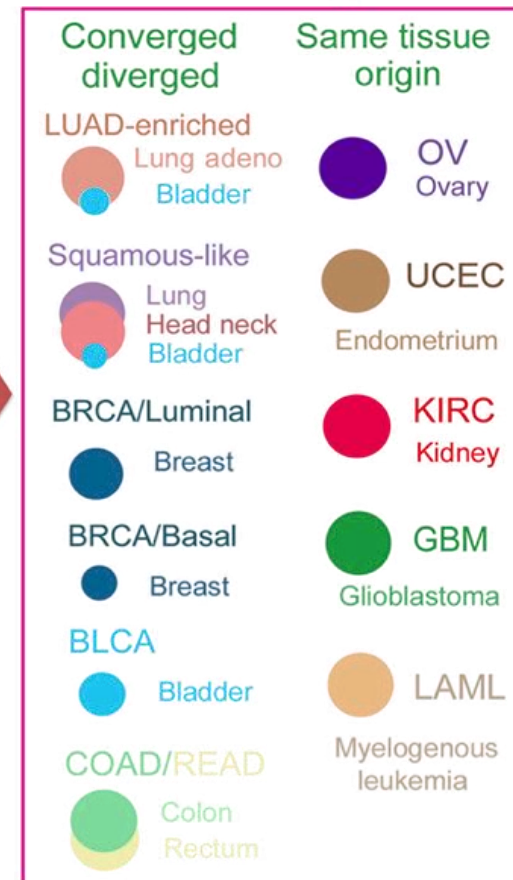
Pan-TCGA



Platforms



Reclassification of cancer types



Molecular Cancer Re-Classification

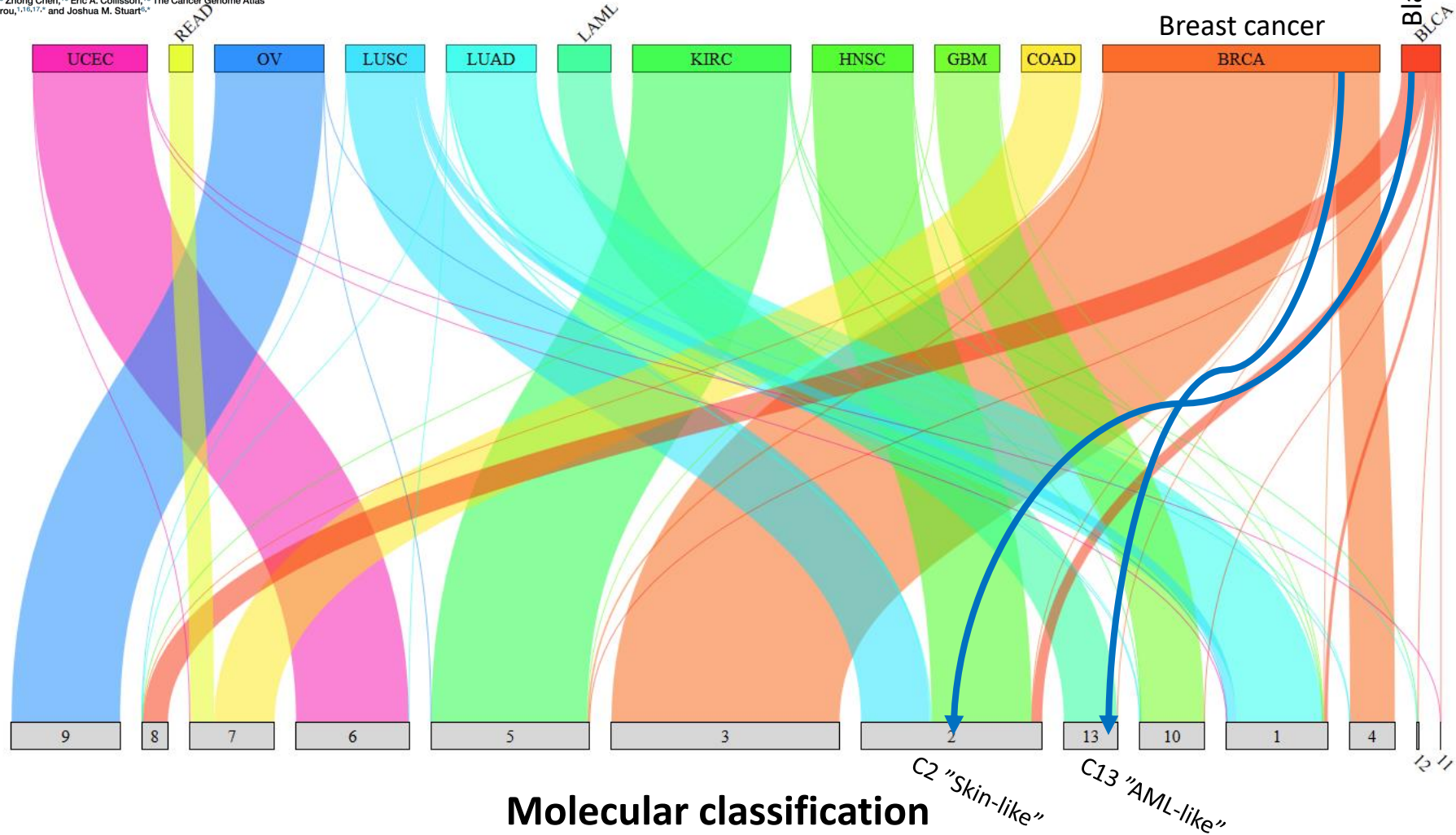
Multiplatform Analysis of 12 Cancer Types Reveals Molecular Classification within and across Tissues of Origin



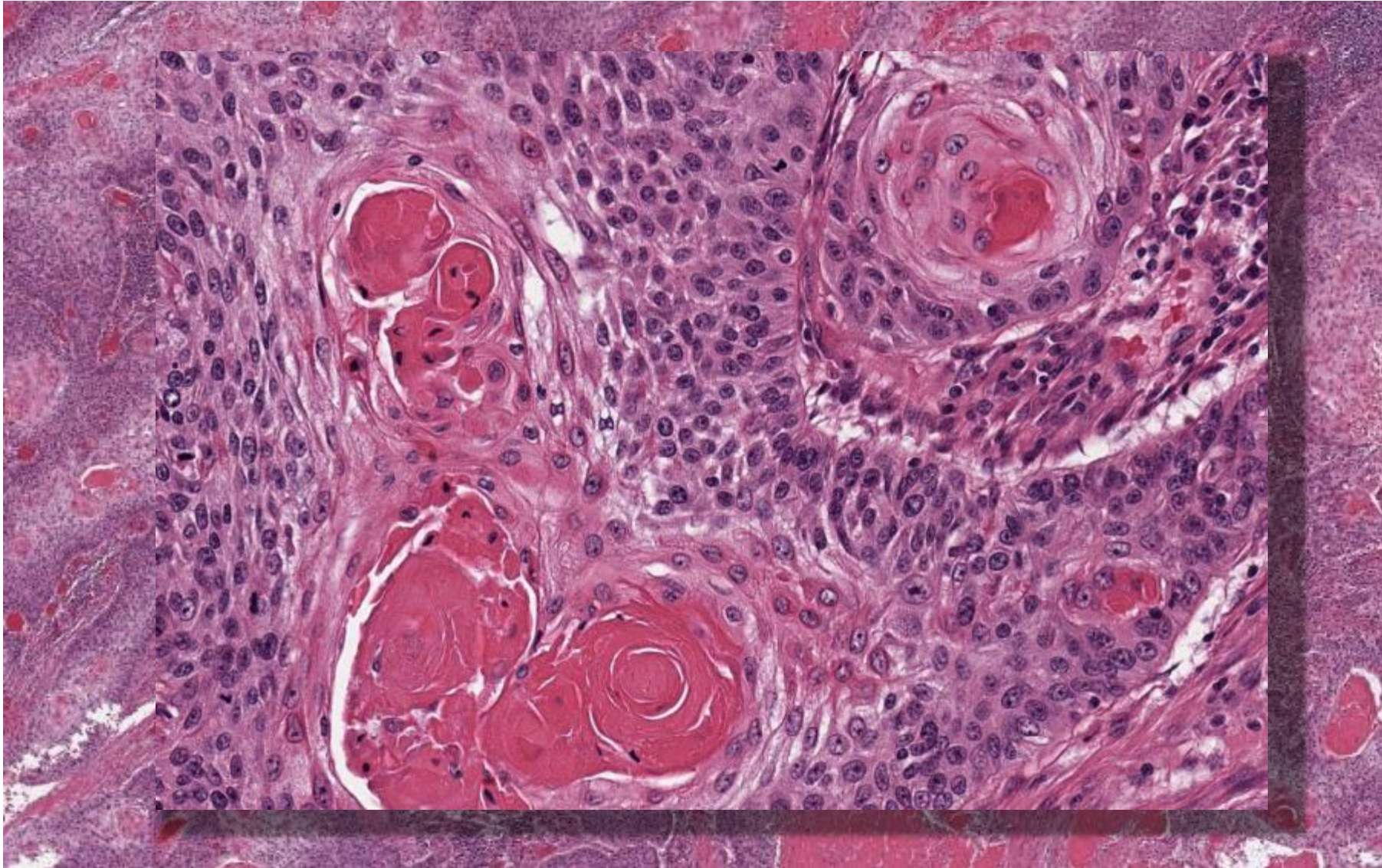
Cell Oct 2014 158, 929-944.

Katherine A. Hoadley,^{1,20} Christina Yau,^{2,20} Denise M. Wolf,^{3,20} Andrew D. Cherniack,^{4,20} David Tamborero,⁵ Sam Ng,⁶ Max D.M. Leiserson,⁷ Bifang Niu,⁸ Michael D. McLellan,⁹ Vladislav Uzunangelov,⁵ Jiashan Zhang,⁹ Cyriac Kandath,⁹ Rehan Akbari,¹⁰ Hui Shen,^{11,22} Larsson Omberg,¹² Andy Chu,¹³ Adam A. Margolin,^{12,21} Laura J. van't Veer,³ Nuria Lopez-Bigas,^{5,14} Peter W. Laird,^{11,22} Benjamin J. Raphael,⁷ Li Ding,⁸ A. Gordon Robertson,¹³ Lauren A. Byers,¹⁰ Gordon B. Mills,¹⁰ John N. Weinstein,¹⁰ Carter Van Waes,¹⁰ Zhong Chen,¹⁰ Eric A. Collisson,¹⁰ The Cancer Genome Atlas Research Network, Christopher C. Benz,²⁴ Charles M. Perou,^{1,15,17,*} and Joshua M. Stuart^{4,*}

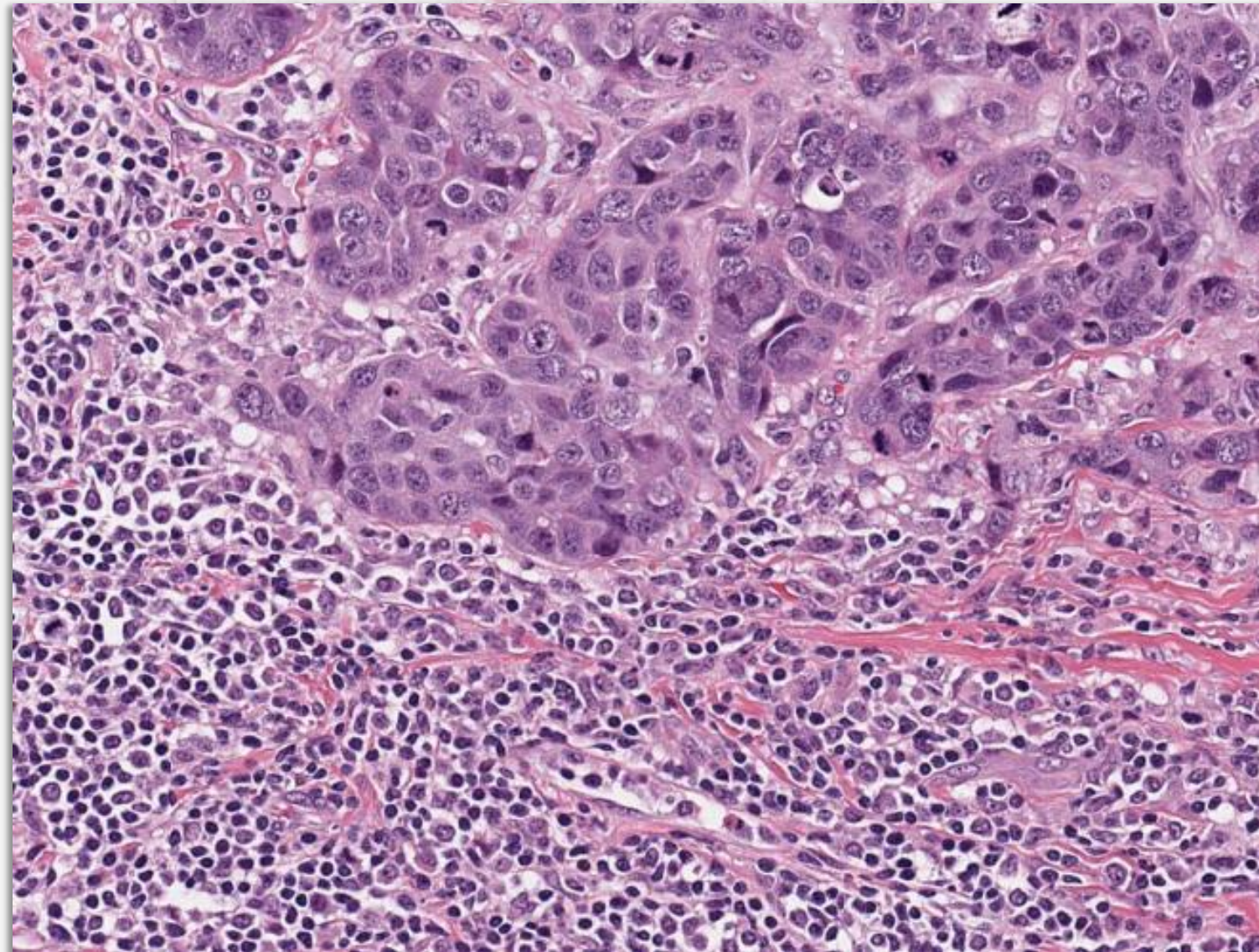
WHO organ-based classification



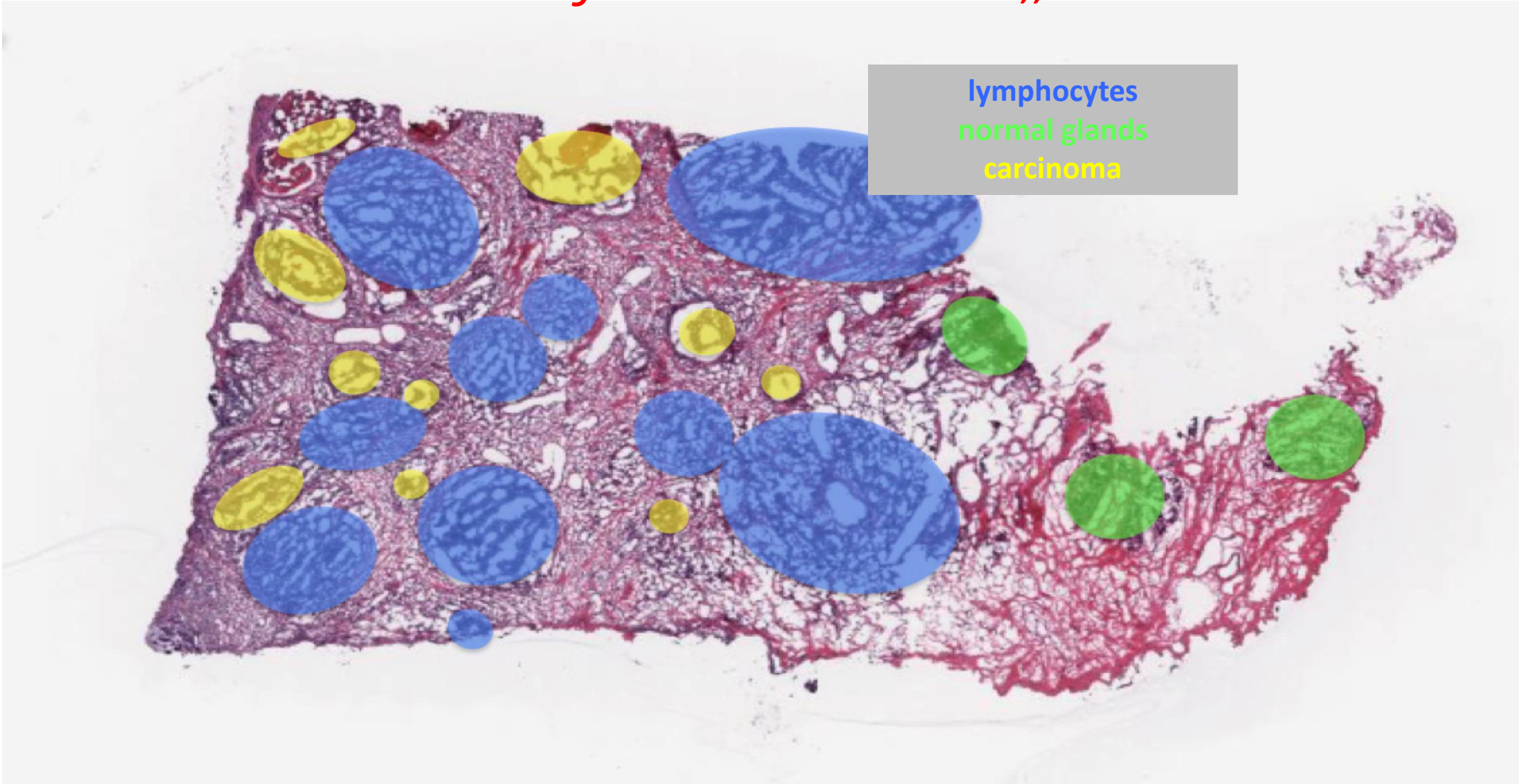
Bladder Ca *re-classified as C2 „Skin-like“*



Breast Ca *re-classified as* C13 – „Leukemia-like“



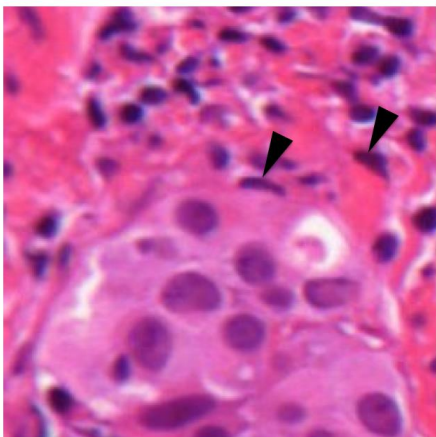
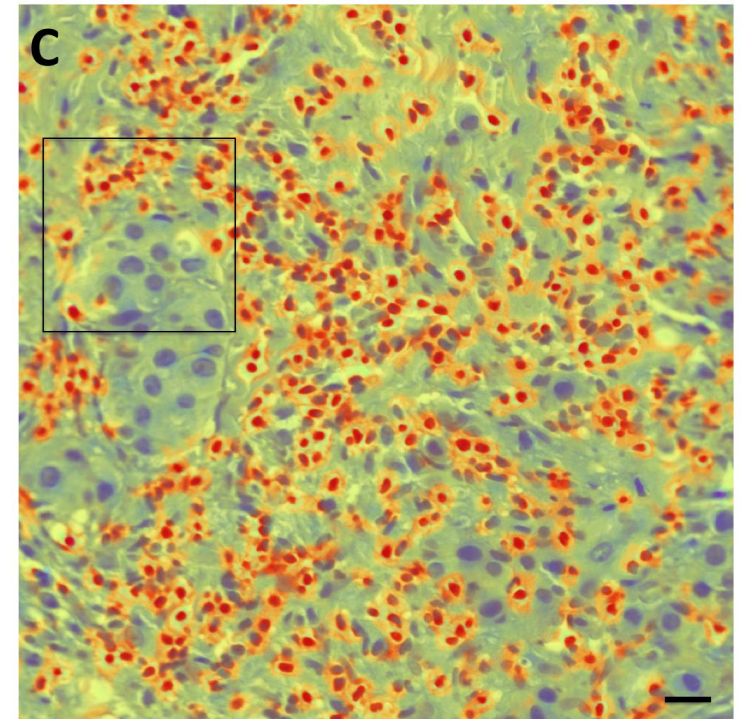
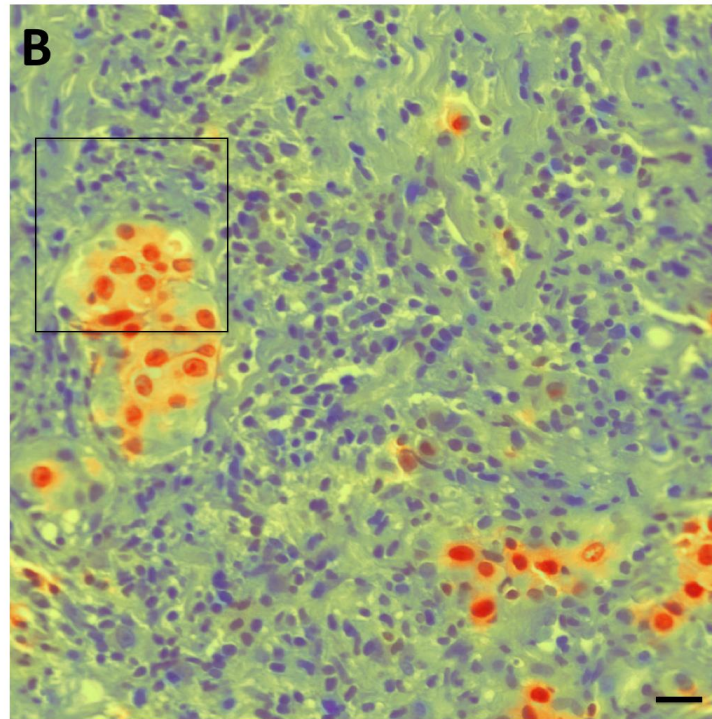
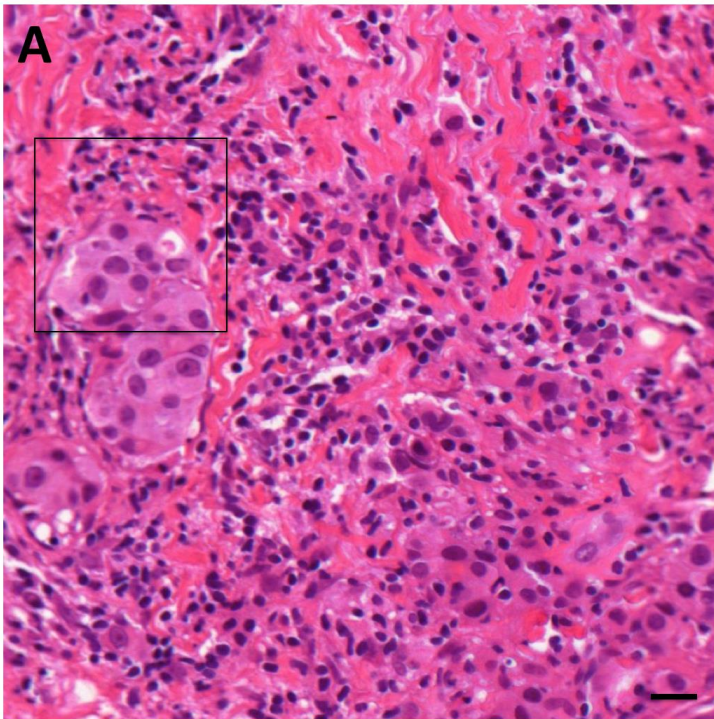
Breast Ca *re-classified* as C13 – „Leukemia-like“



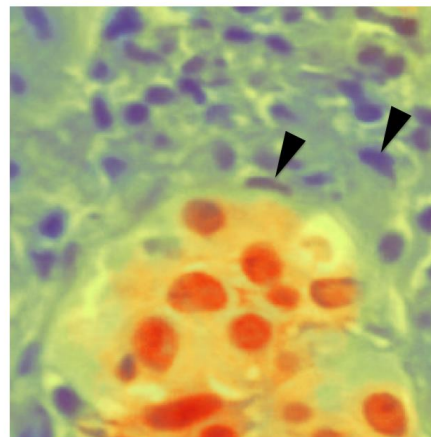
Incorrect classification because of

- incomplete training data
- imprecise evaluation of histopathology.

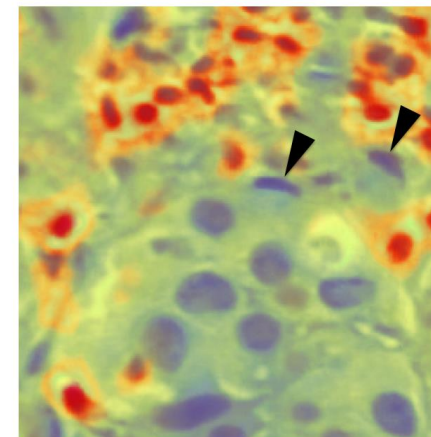
AI-assisted histopathology



original



cancer



TiLs

Binder et al.

Artificial Intelligence in Diagnostics:

PATIENTS IN DANGER?

Need for validation and
benchmarking of AI in medicine!

ITU/WHO Focus Group AI for Health

Topic Group Histopathology



- Topic group dedicated to benchmarking AI approaches in histopathology
- First use case: Detection of breast cancer cells and tumor-infiltrating lymphocytes
- Define what should be annotated and how
- Define criteria for benchmarking
- Provide server infrastructure to perform benchmarking

Annotation of the histopathology images

Specifications:

- Digitized histological slides in standard staining
- Comprehensive tissue component annotations:

cancer tissue

multiple subtypes

focus on NST (no-special-type) and invasive-lobular breast cancer

normal tissue

normal breast gland and duct epithelium

connective tissue (fibers, cells)

fatty tissue, bone tissue, nerves

blood and lymphatic vessels

immune system

Lymphocytes, plasma cells

Granulocytes, monocytes/macrophages

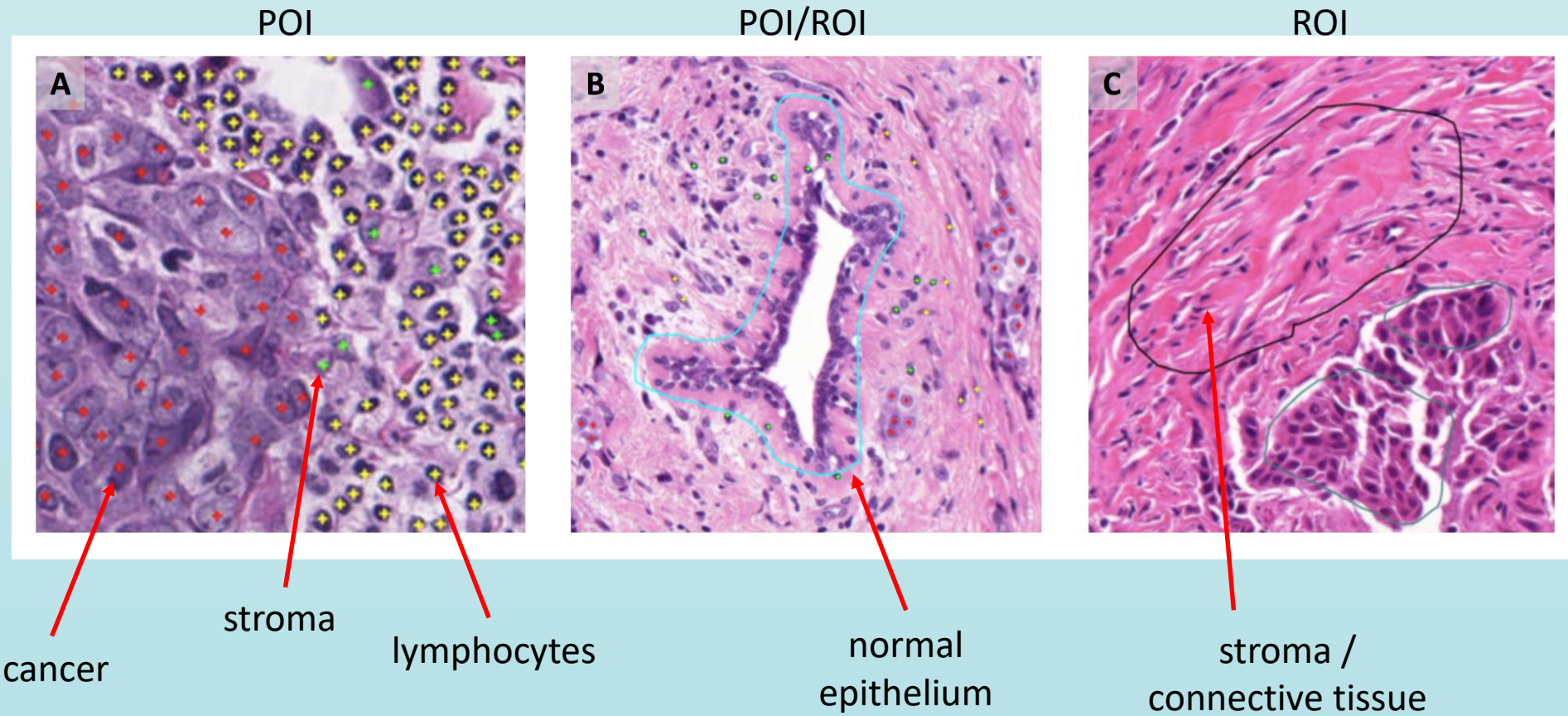
necrotic tissue

artifacts

Background

- Positive and negative annotations

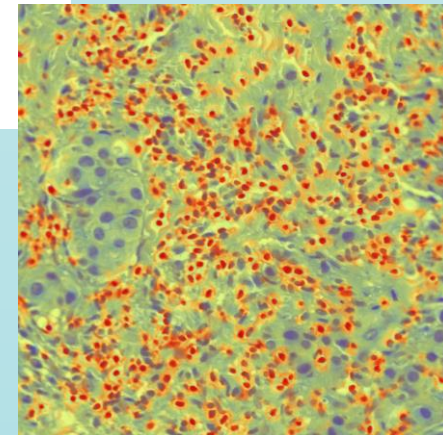
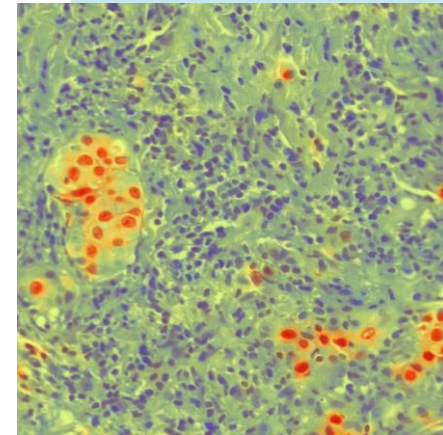
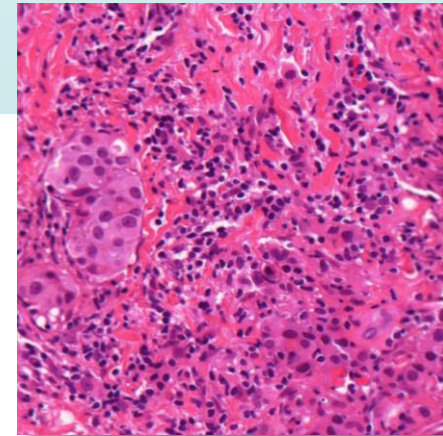
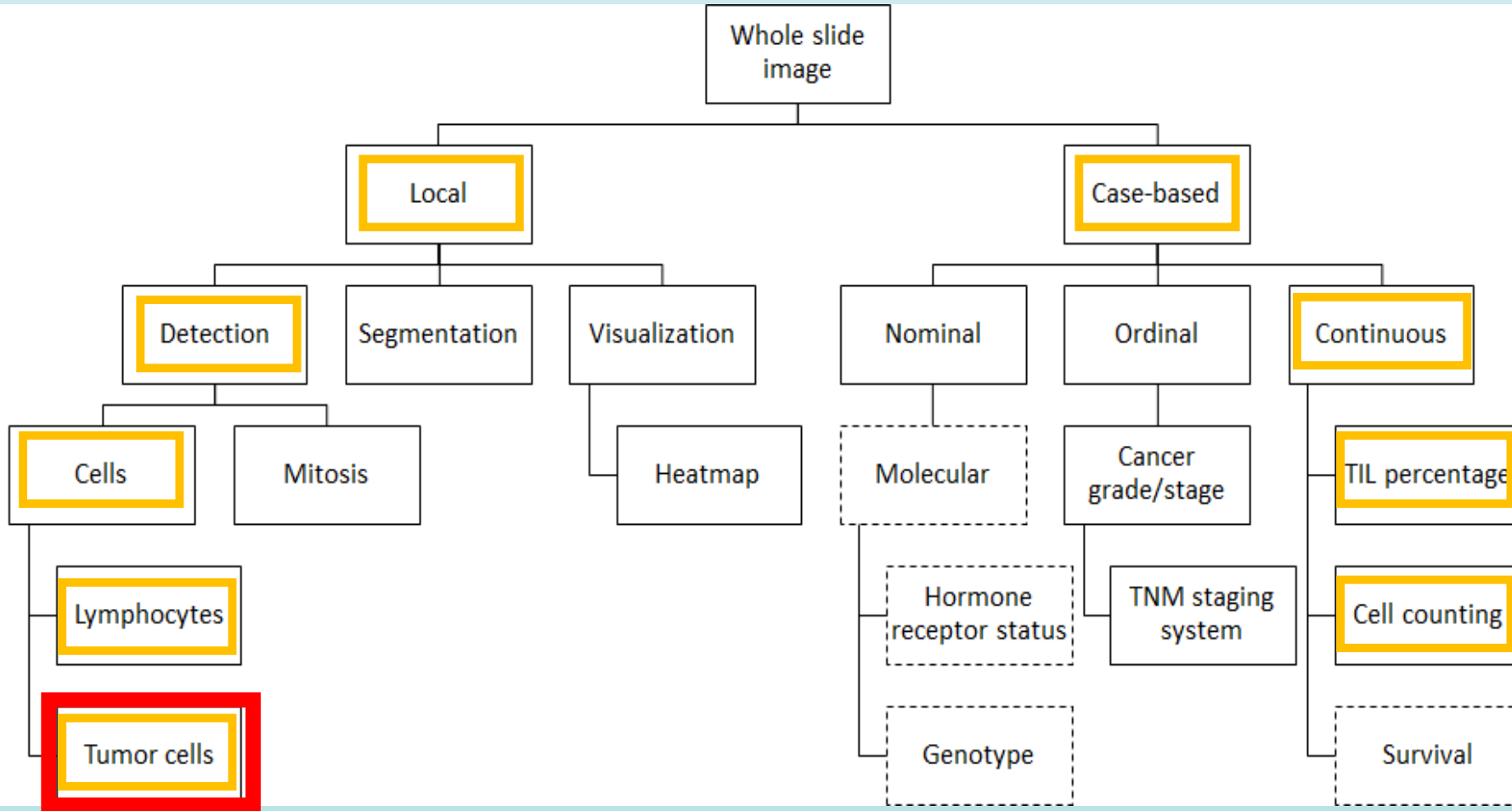
Annotation of the histopathology images



Provision of test and benchmarking images

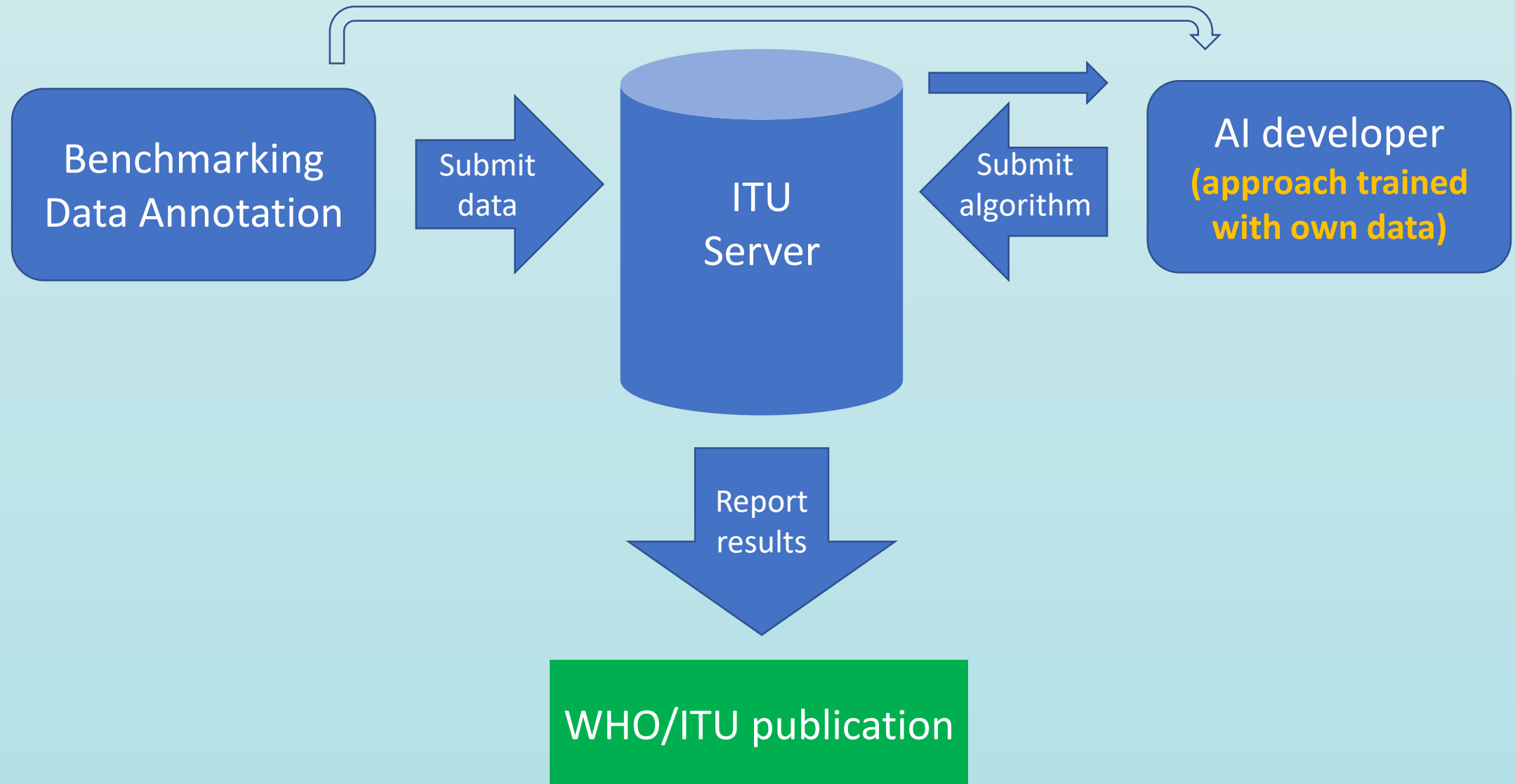
- image data 3000x3000 at 400x
- consensus annotations by two pathologists
- 4 exemplary images made available with test annotations to provide overview of data
- 86 annotated images not public, available only for benchmarking on WHO/ITU servers with
- 258k annotations

3 Benchmarking

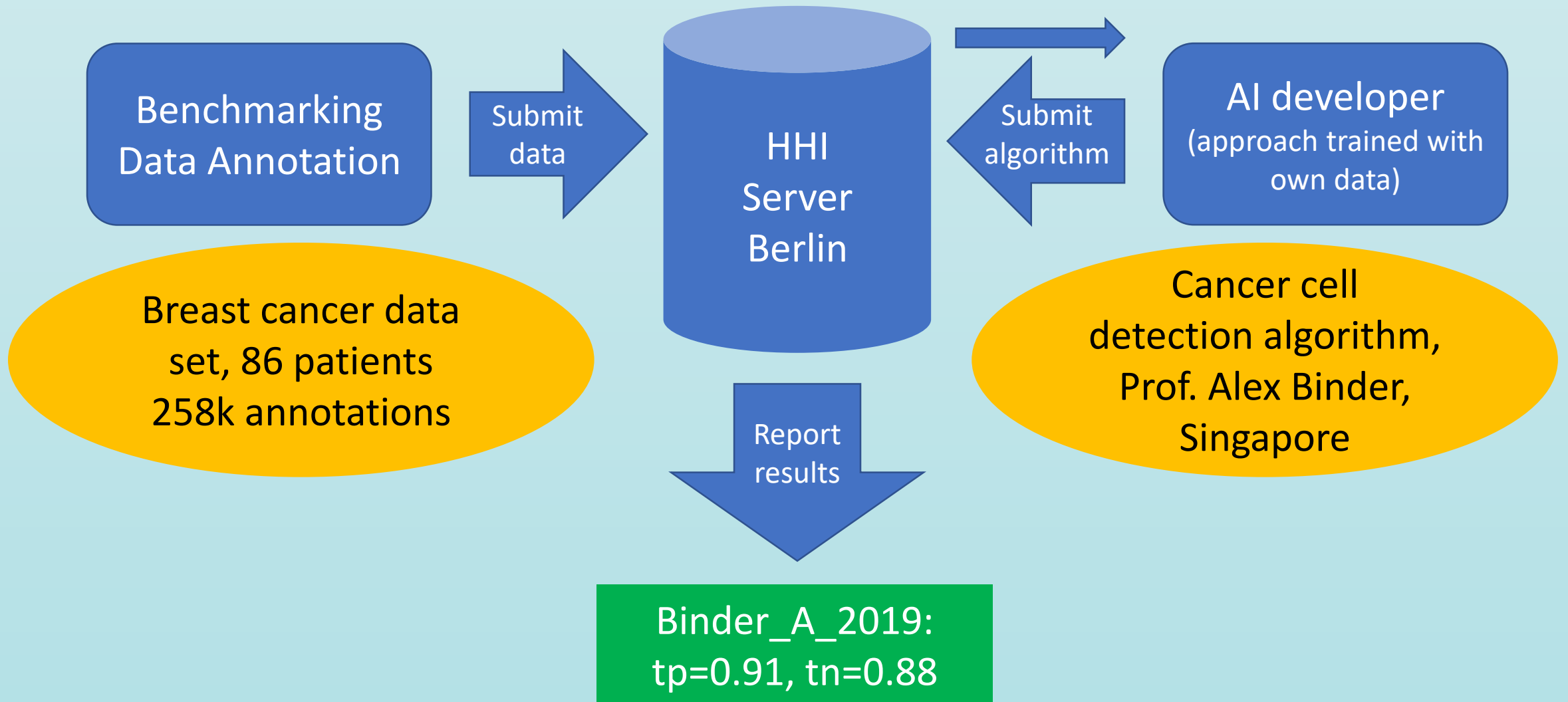


true positive, true negative, precision

Benchmarking pipeline



Benchmarking breast cancer cell detection



Prof. Dr. Alexander Binder
Singapore University of Technology and Design (SUTD)
Singapore

Dr. Markus Wenzel, Dr. David Neumann, Prof. Dr. Thomas Wiegand
Fraunhofer Heinrich-Hertz-Institute, Berlin & WHO/ITU Focus
Group AI4HEALTH

Dr. Alexander Arnold, Dr. Bruno Sinn, Prof. Dr. Frederick Klauschen
Institute of Pathology, Charité Universitätsmedizin Berlin
Berlin Institute of Health