

16th
CONGRESS
Lung **ON**
CANCER

BARCELONA
27 / 28
NOVEMBER 2025

AI in digital pathology and multiplex immunohistochemistry (IHC)

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University Hospital Cologne

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... from a hemato-
oncologist point of
view!

CONFLICTO DE INTERESES

Grants or contracts from any entity (to institution)

Else-Kröner Forschungskolleg Cologne, German Cancer Aid, Dr. Rolf M. Schwiete Foundation, Amgen, AstraZeneca, Bristol-Myers Squibb, Janssen-Cilag, Novartis

Payment or honoraria (personal)

BeOne Medicines, Bristol-Myers Squibb, Takeda, Janssen-Cilag, Pfizer

Support for attending meetings and/or travel (personal)

Nationales Centrum für Tumorerkrankungen West, BeOne Medicines, Janssen-Cilag, Eli Lilly and Company, Takeda

AI vs human performance

Select AI Index technical performance benchmarks vs. human performance

Source: AI Index, 2025 | Chart: 2025 AI Index report

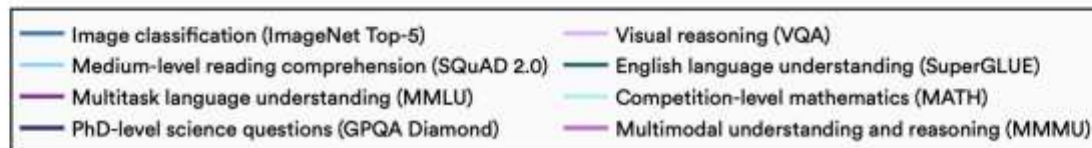
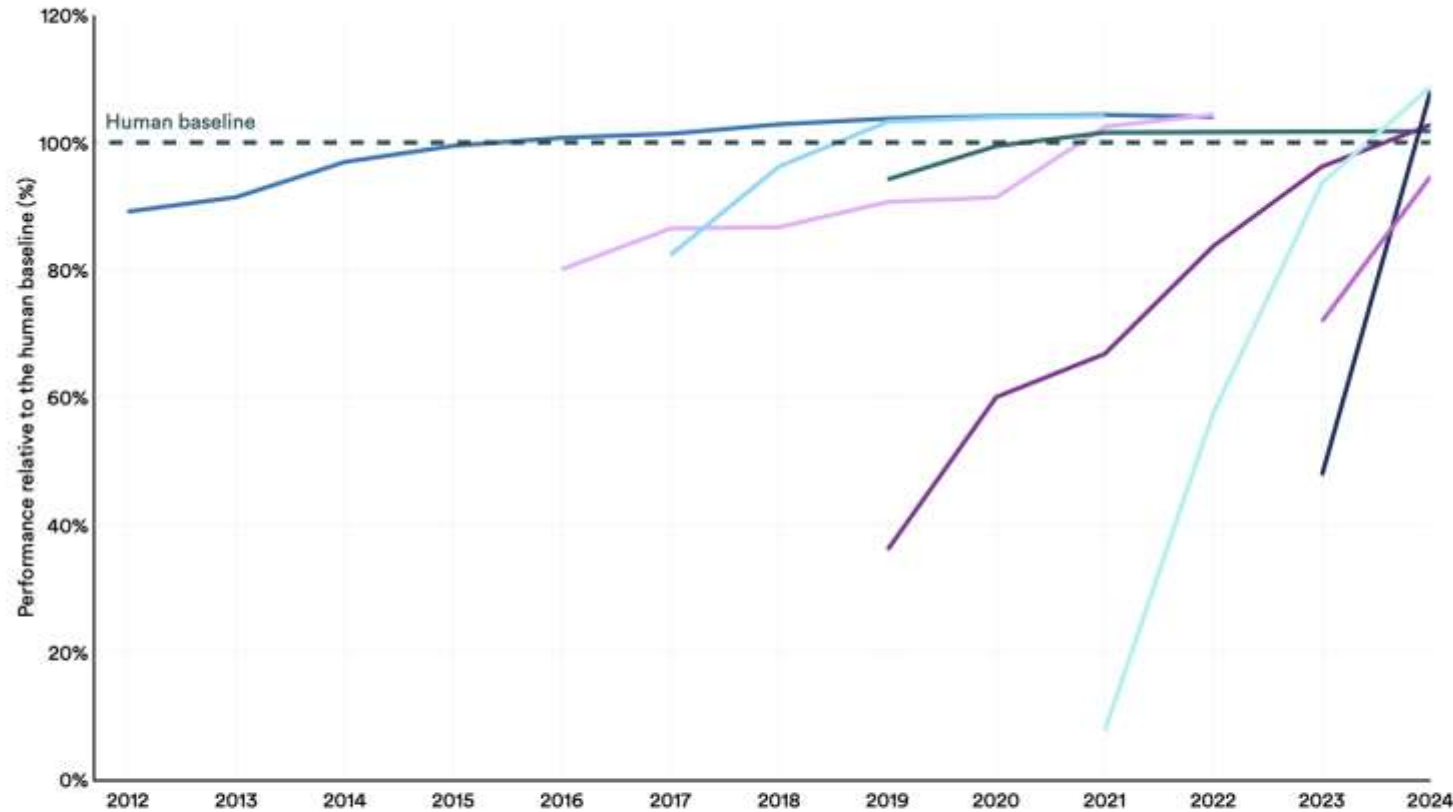


Figure 2.1.33²

AI is rapidly outperforming humans in diverse tasks

and the rate is increasing...

The human brain vs artificial neural networks (NN): a comparison of scale and efficiency

~80 billion neurons
~80 billion other cells
100 trillion synapses

>Exaflop computing
A billion-billion (1^{18})
operations/second for the
cost of **20 Watts**

The brain is
extraordinarily
energy-efficient

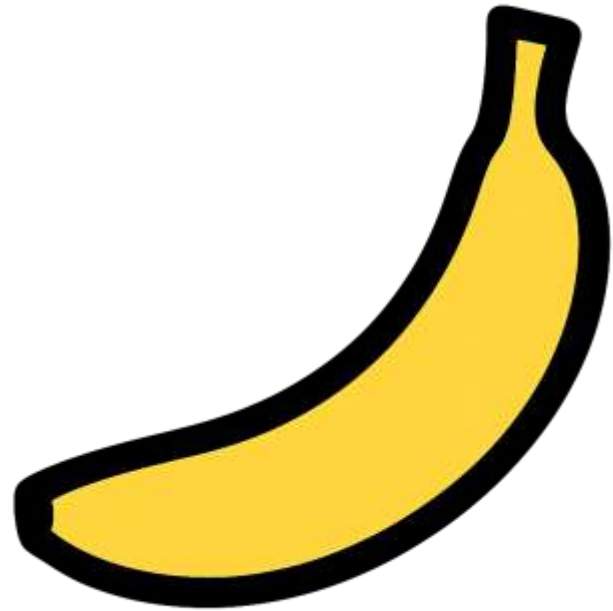


Artificial NNs
(node/unit of compute)

OpenAI - GPT4
Billions of parameters
(~synapses)

Millions of dollars to train

The first supercomputer to demonstrate
exaflop computing (Oak Ridge) needed
20 Megawatts to complete the compute



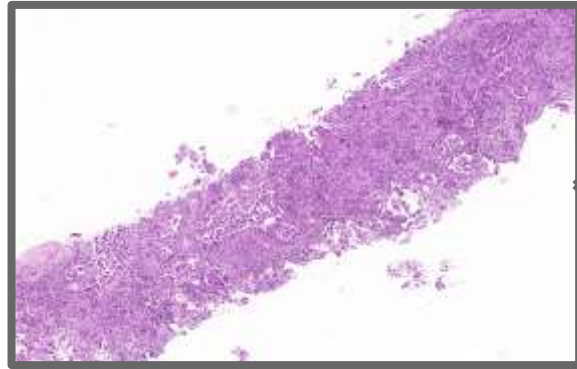
VS

**1,000,000
bananas
or
~120 T**

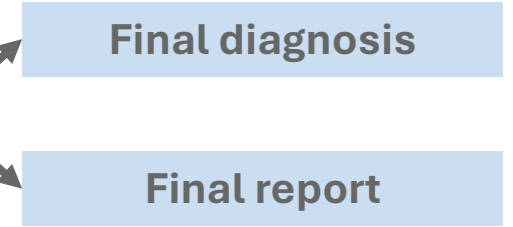
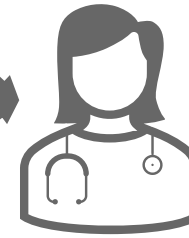
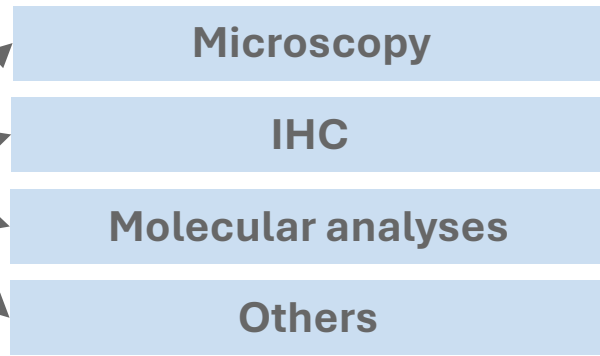
AI in digital pathology - ideally

Conventional pathology vs AI-enhanced digital pathology

Conventional



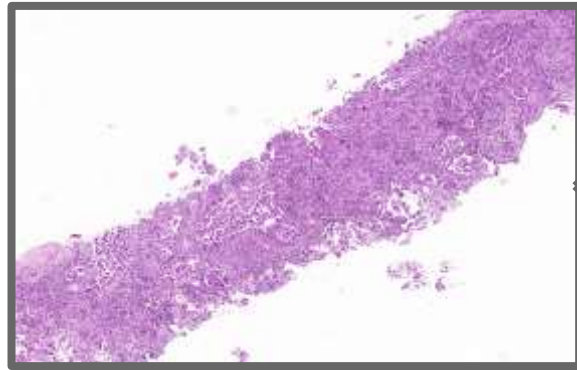
Tumor tissue and conventional HE slide



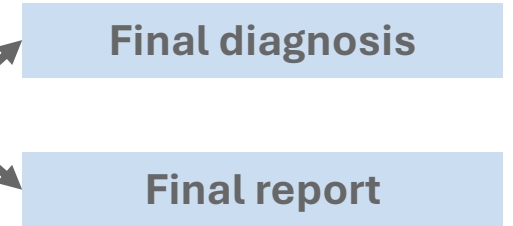
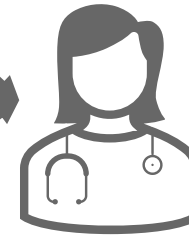
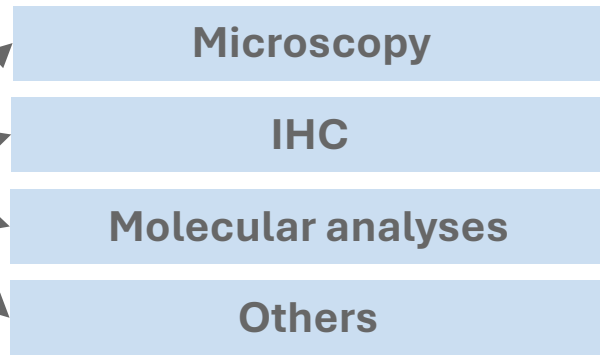
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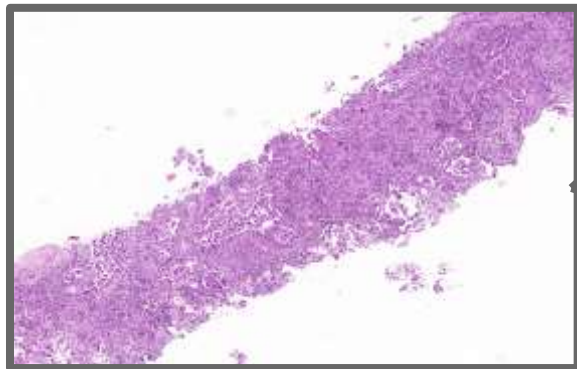
Conventional



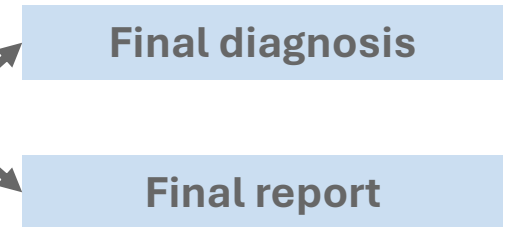
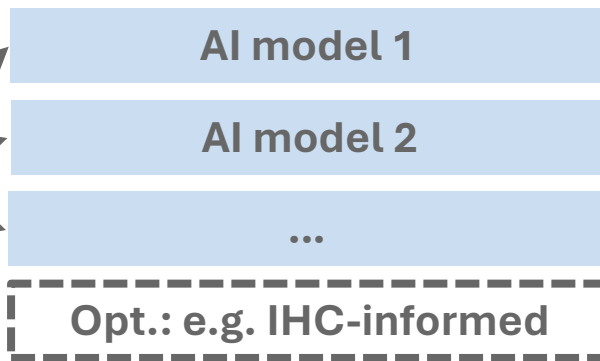
Tumor tissue and conventional HE slide



AI-enhanced

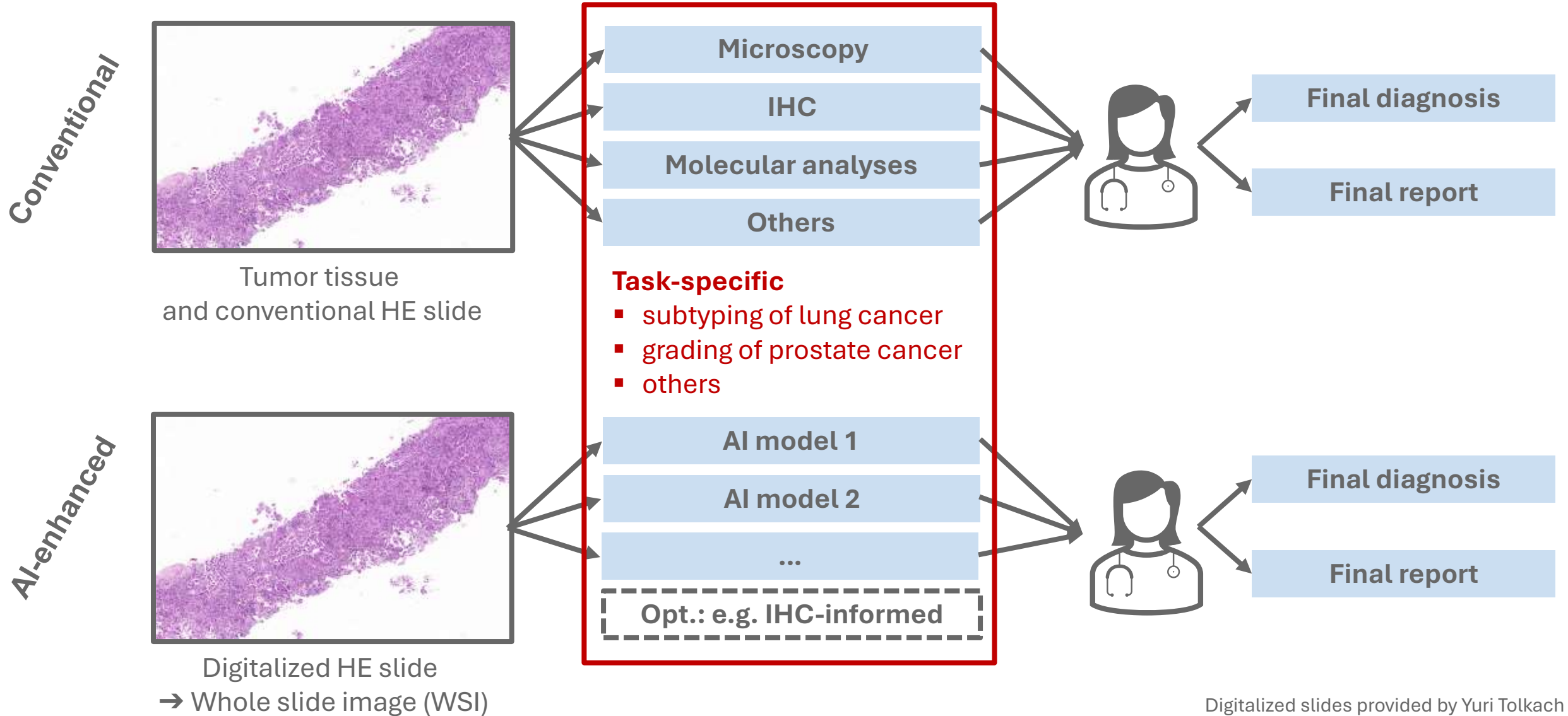


Digitalized HE slide
→ Whole slide image (WSI)



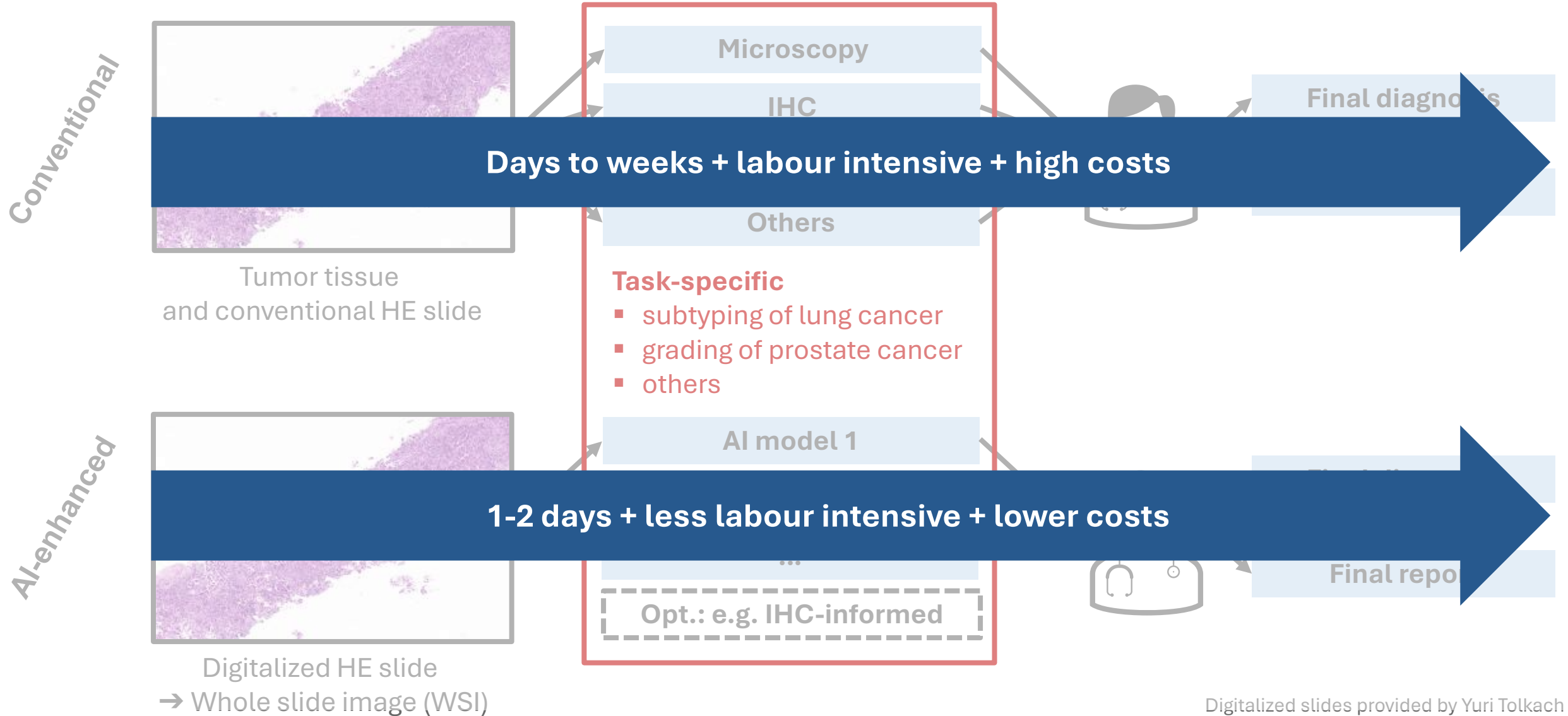
AI in digital pathology - ideally

Conventional pathology vs AI-enhanced digital pathology



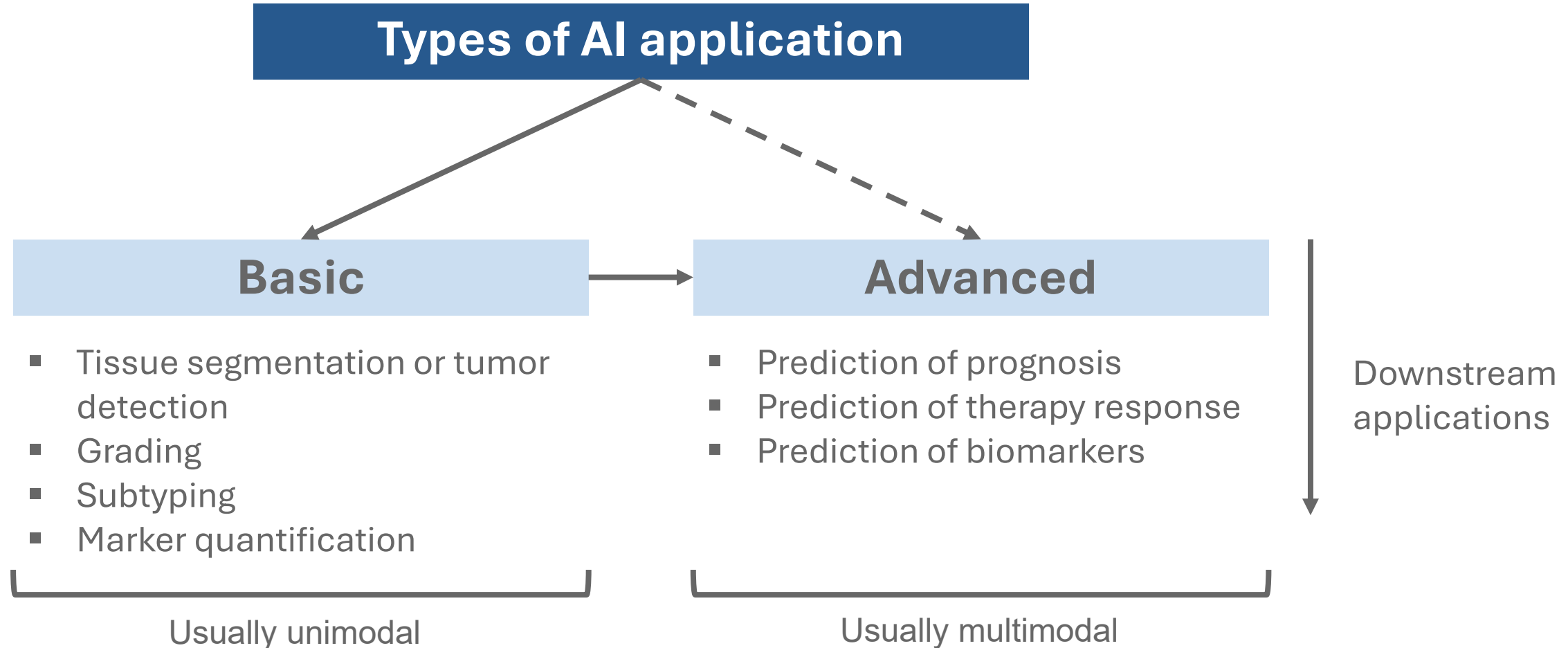
AI in digital pathology - ideally

Conventional pathology vs AI-enhanced digital pathology



AI applications in pathology and oncology

Basic to advanced = unimodal to multimodal

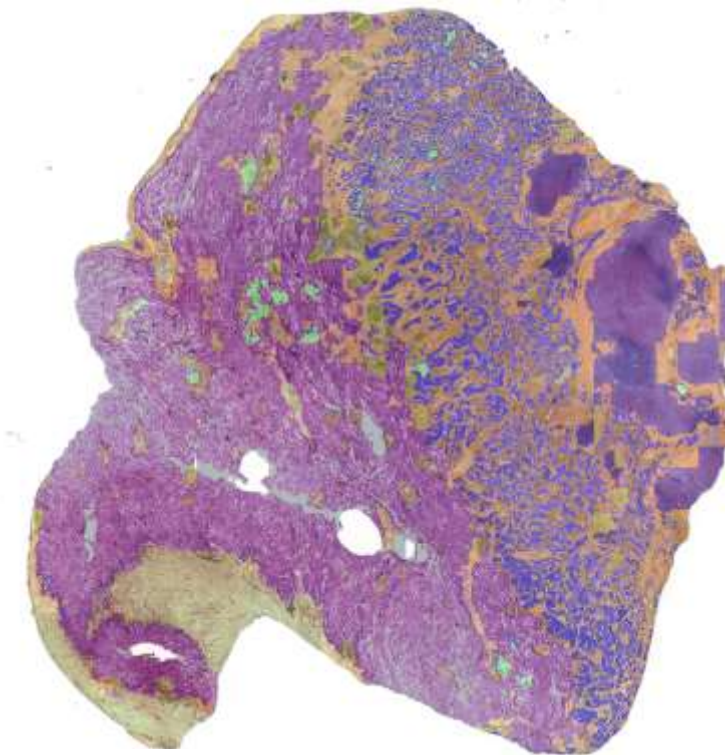
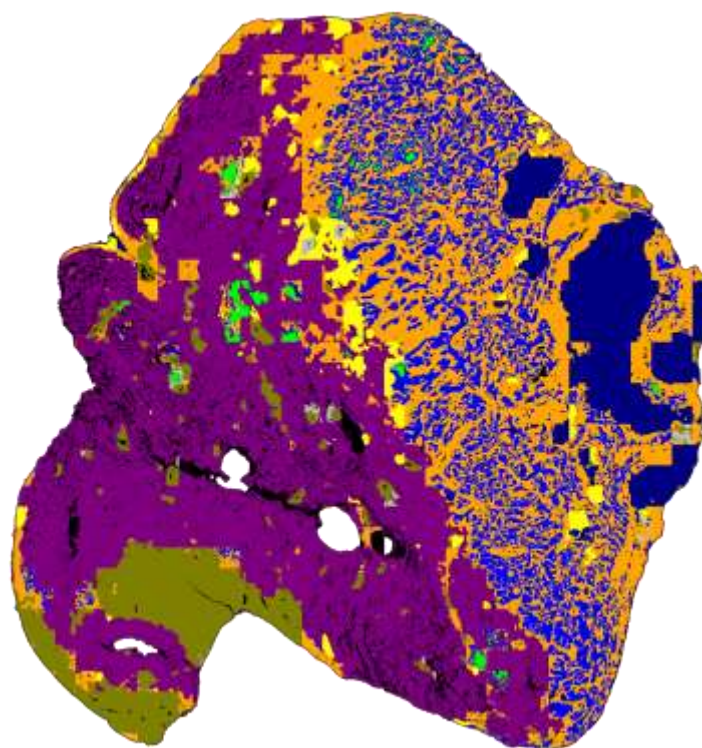
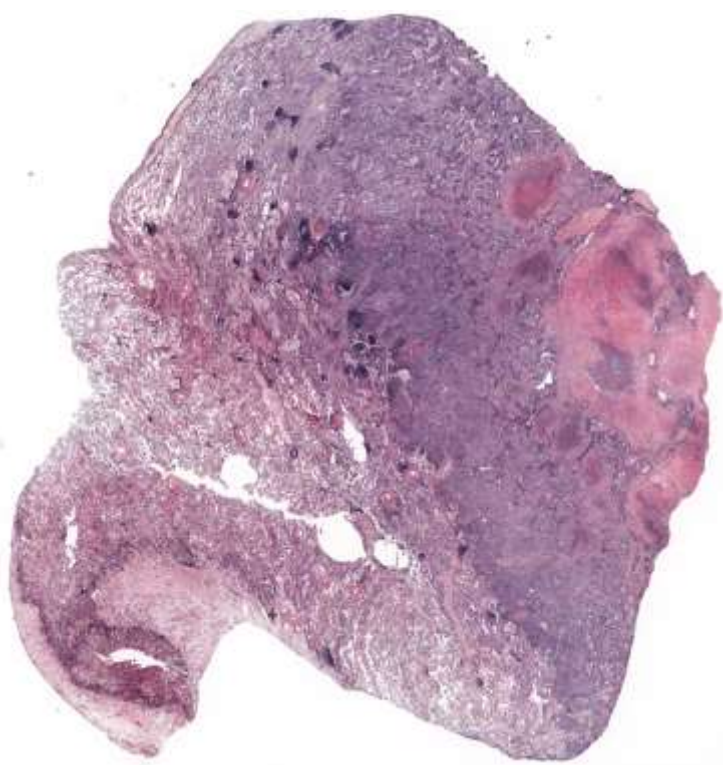


Basic: Tissue segmentation

A multi-institutional model by the University Hospital Cologne

A supervised (explainable) model to segment tissue of digitalized HE-stained whole slide images (WSI)

→ **Trained in 345 annotated WSI and validated in 4,097 WSI**



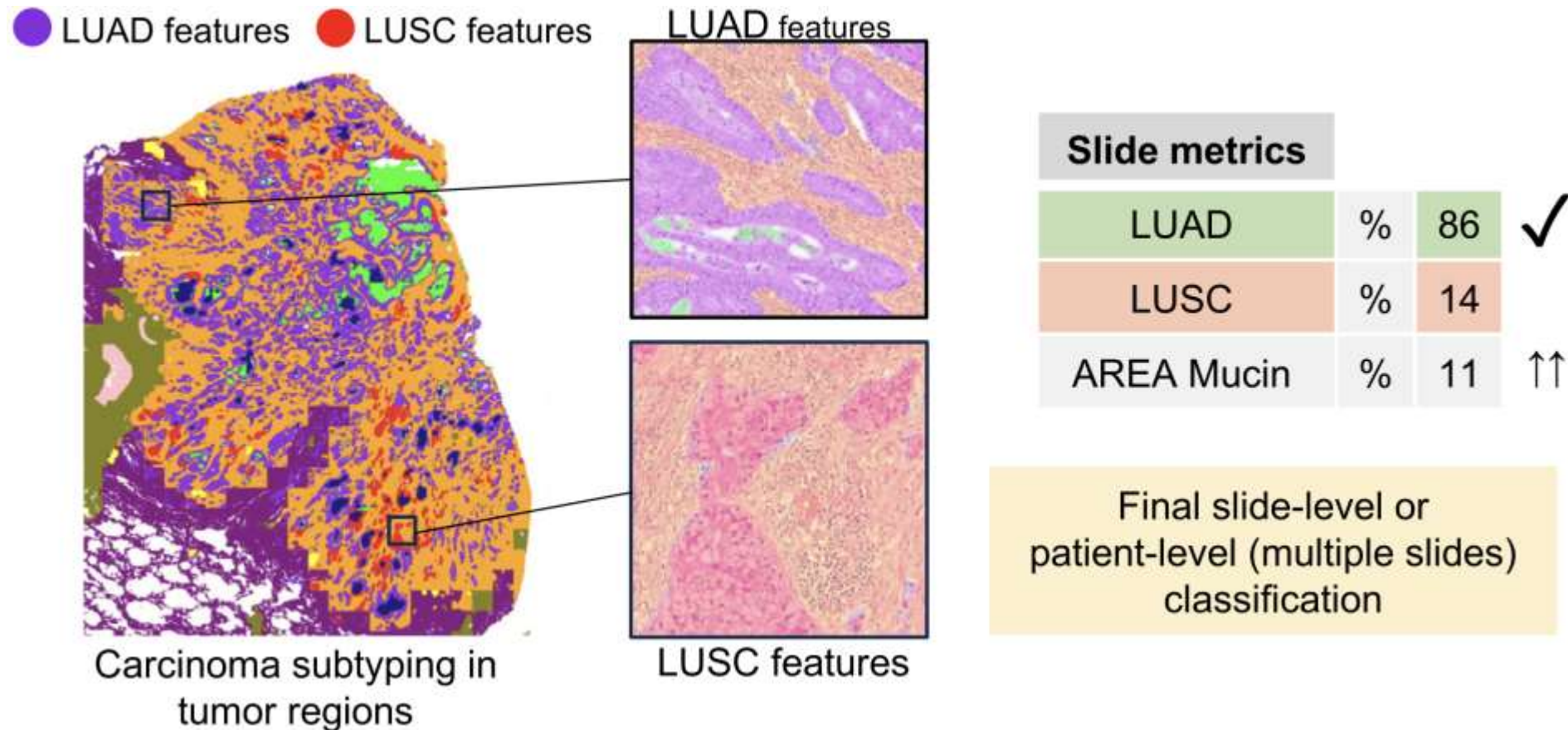
- TUMOR ● TU_STROMA ● NECROSIS ● MUCIN
- TLS ● LUNG_BENIGN ● STROMA ● BRONCH
- BLOOD ● GLAND_PERIBR ● CARTIL ● BACK

Kindly provided by Yuri Tolkach

Basic: Histologic subtyping

A multi-institutional model by the University Hospital Cologne

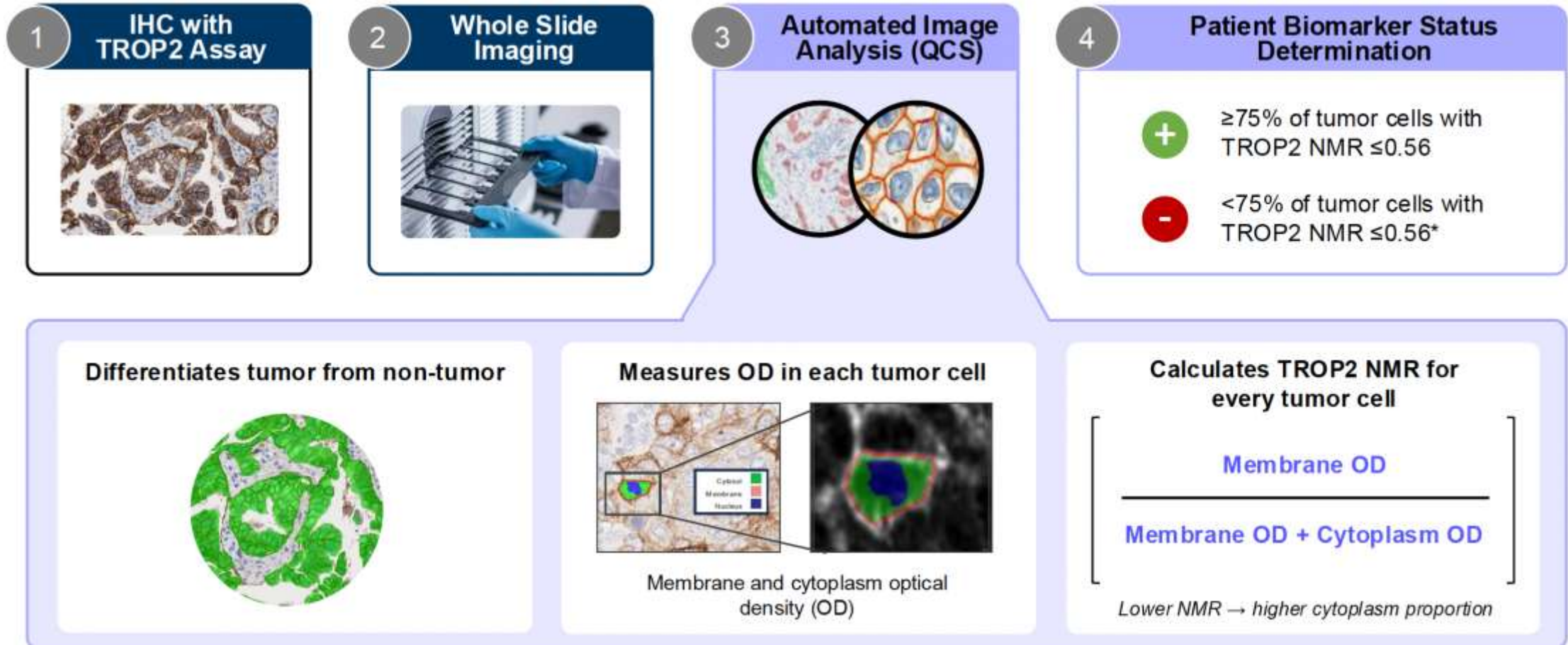
Histologic subtyping (LUAD vs LUSC) with a high overall accuracy of 0.929 to 0.978



Basic: Biomarker quantification

Quantitative continuous scoring (QCS) to quantify TROP2 expression

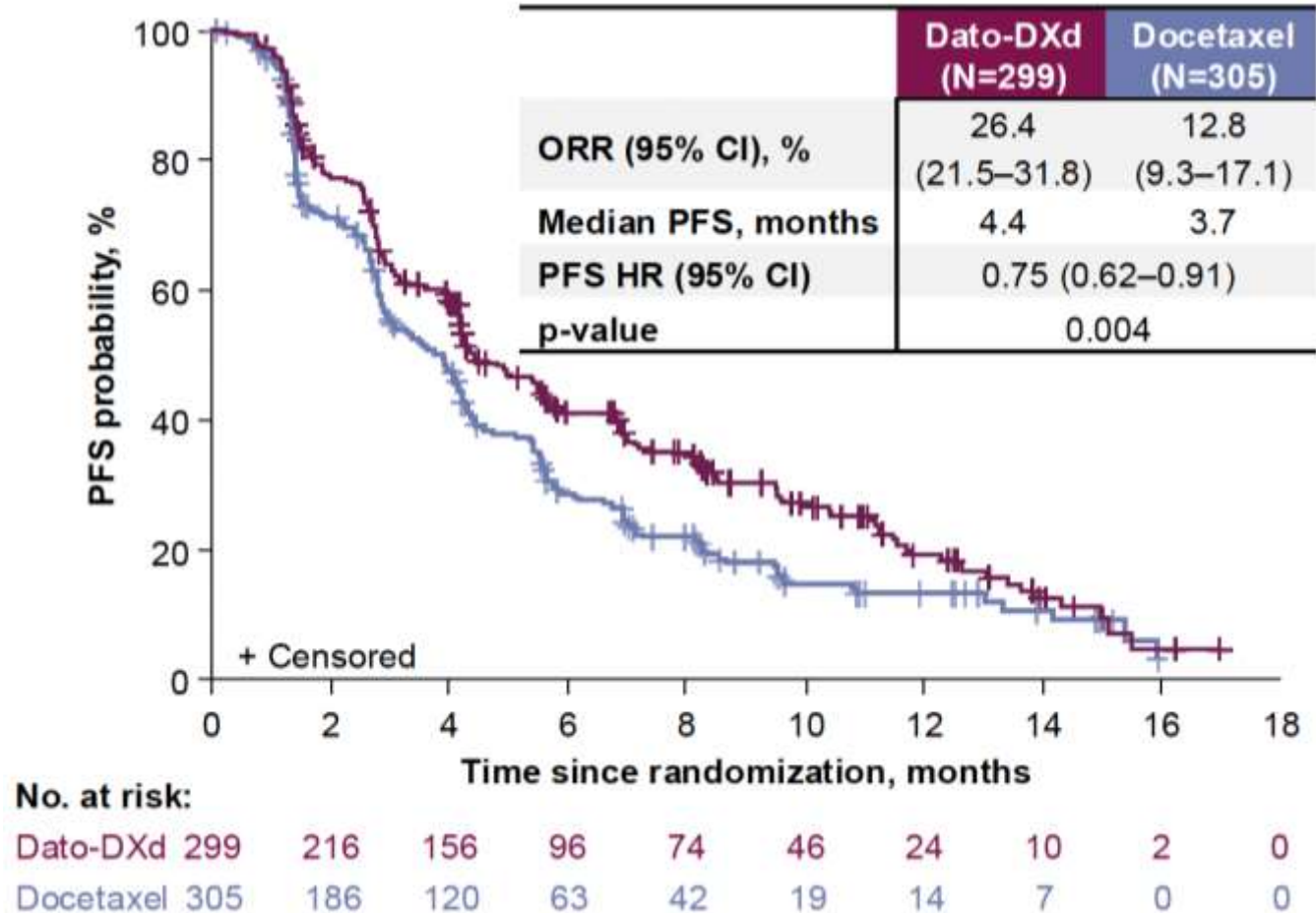
A supervised (explainable) model to determine the normalized membrane ratio (NMR) of surface markers



Basic: Biomarker quantification

Results from the TROPION-Lung01 study (NCT04656652)

PFS and ORR by BICR of the TROP2 ADC datopotamab-deruxtecan vs docetaxel



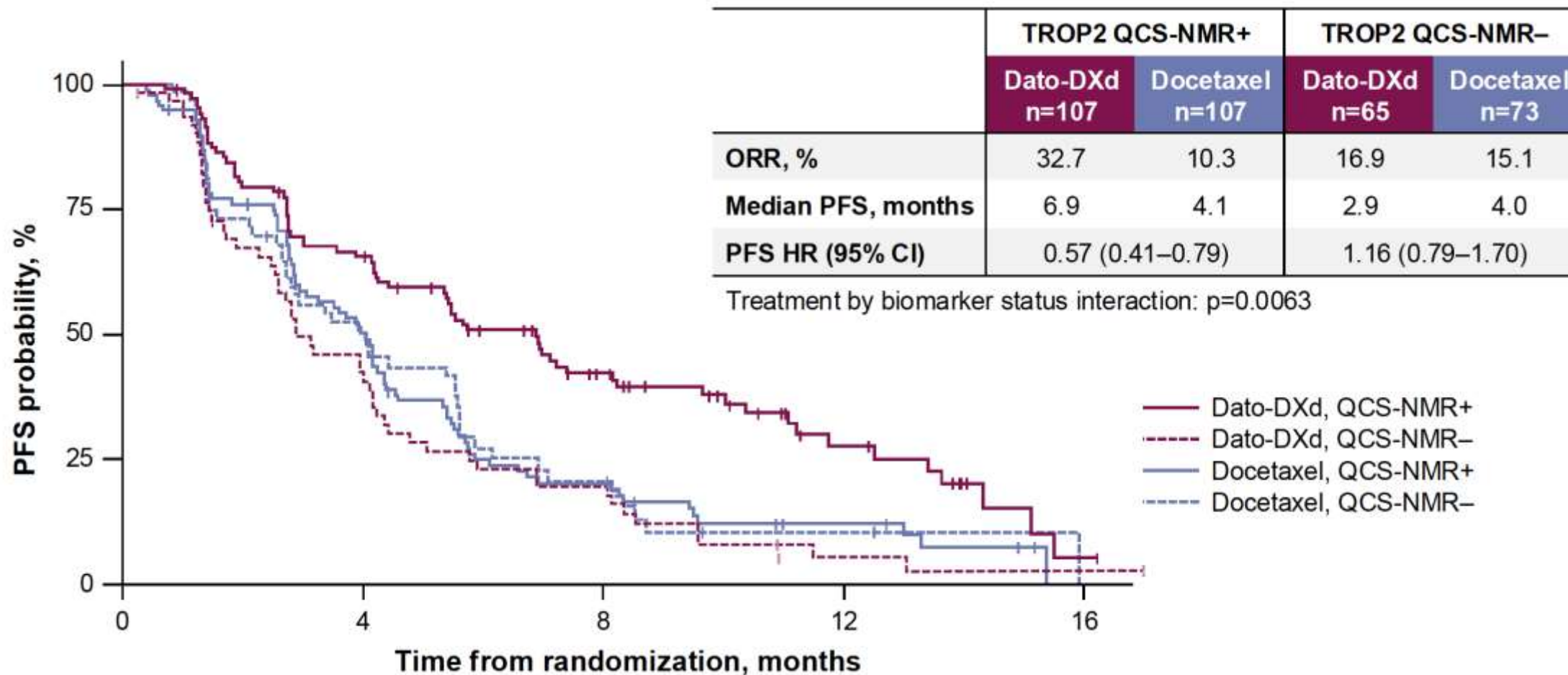
- No difference in OS observed!
 - HR, 0.94 (0.78-1.14)¹
- Conventional read out of TROP2 expression did not correlate efficacy²

Left, Garassino et al. WCLC 2024 – PL02.11
 1. Ahn et al. J Clin Oncol 2024; doi.org/10.1200/JCO-24-01544; 2. Shimizu T, et al. J Clin Oncol 2023; doi: 10.1200/JCO.23.00059.

Basic: Biomarker quantification

Results from the TROPION-Lung01 study (NCT04656652)

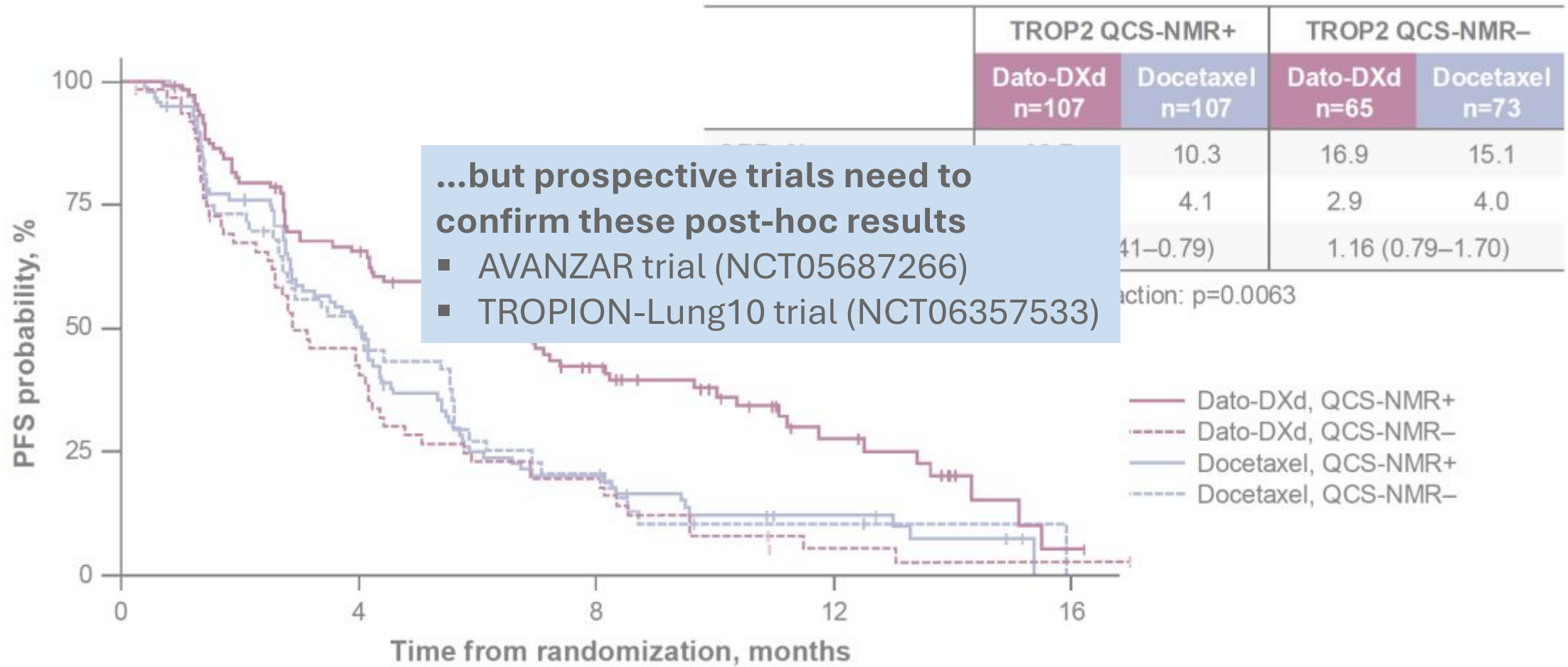
PFS and ORR by BICR in patients stratified by TROP2 NMR (N=352)



Basic: Biomarker quantification

Results from the TROPION-Lung01 study (NCT04656652)

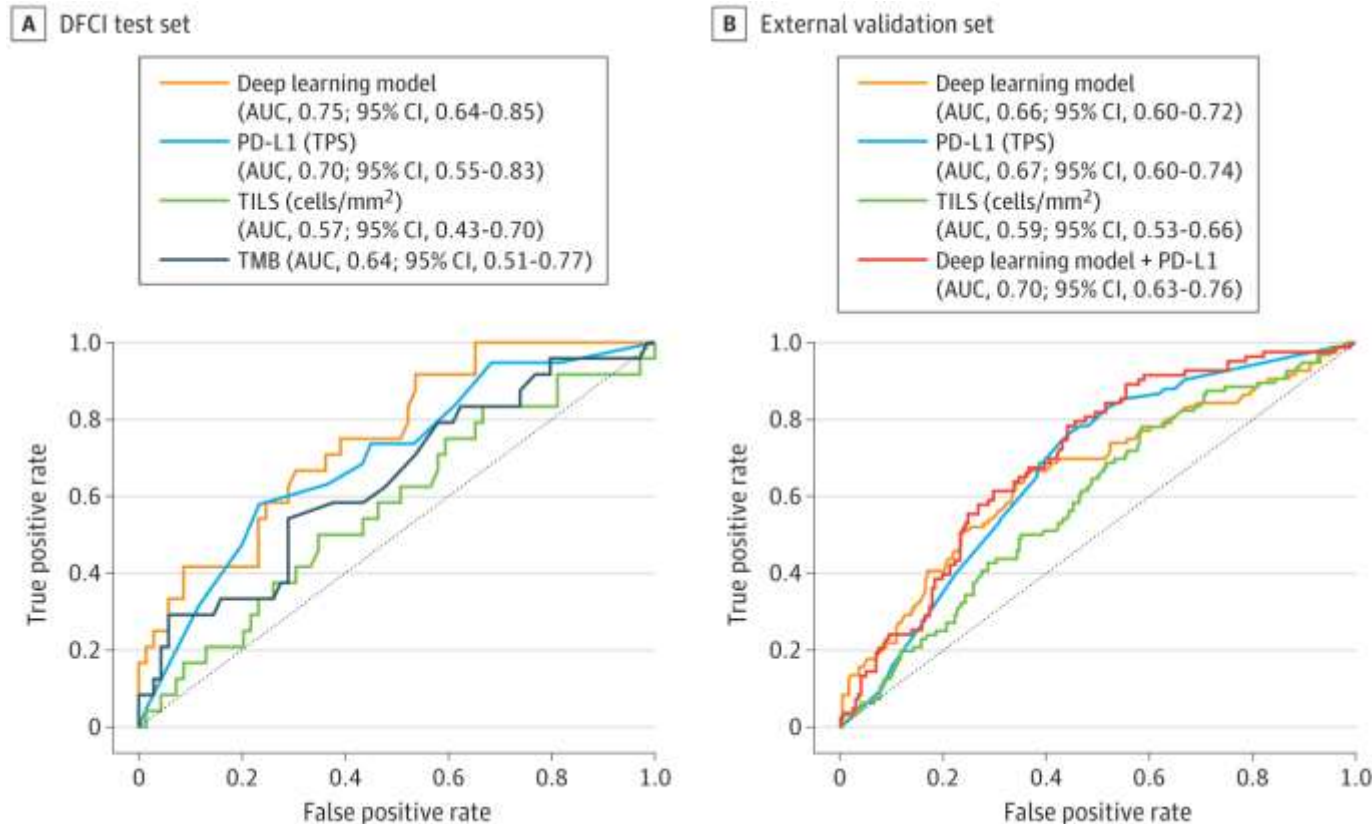
PFS and ORR by BICR in patients stratified by TROP2 NMR (N=352)



Advanced: Prediction of treatment response

A multi-institutional model from the Dana-Faber Cancer Institute

Deep-IO: A supervised (explainable) stratification model to predict response (responders vs non-responders) to immune checkpoint inhibitors (ICIs) monotherapy in NSCLC



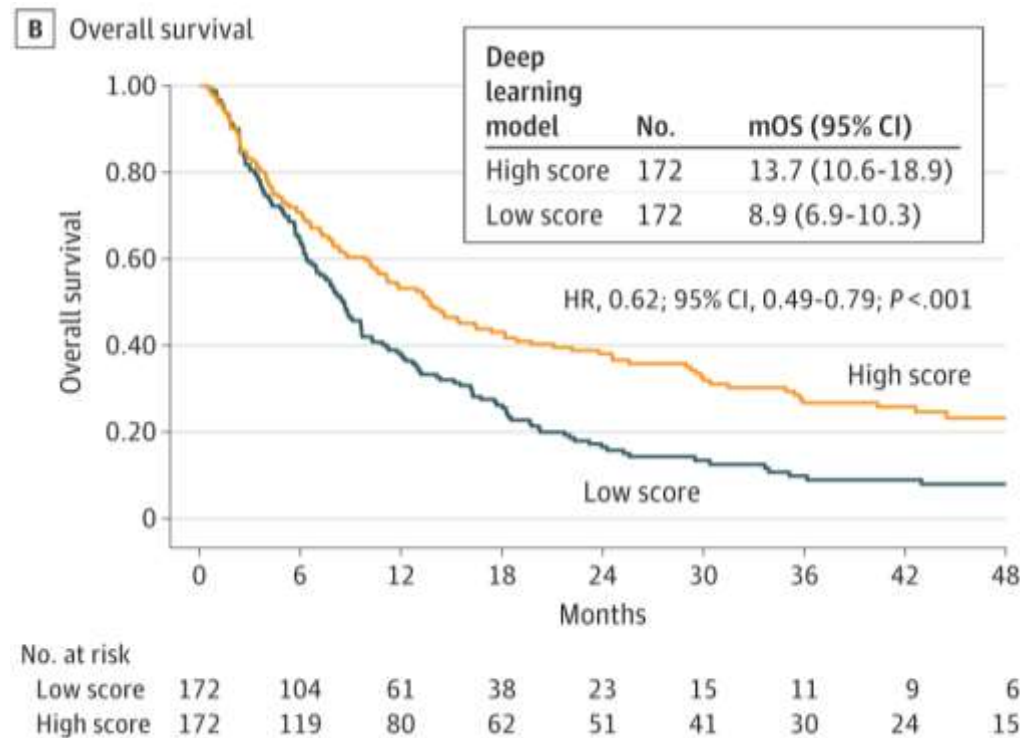
Performance compared to other predicting markers

- Diverging results between test and validation cohort (robustness and applicability?)
- Overall performance <0.8 (ll ideal threshold) = 0.66 – 0.75

Advanced: Prediction of treatment response

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Ready for decision making?

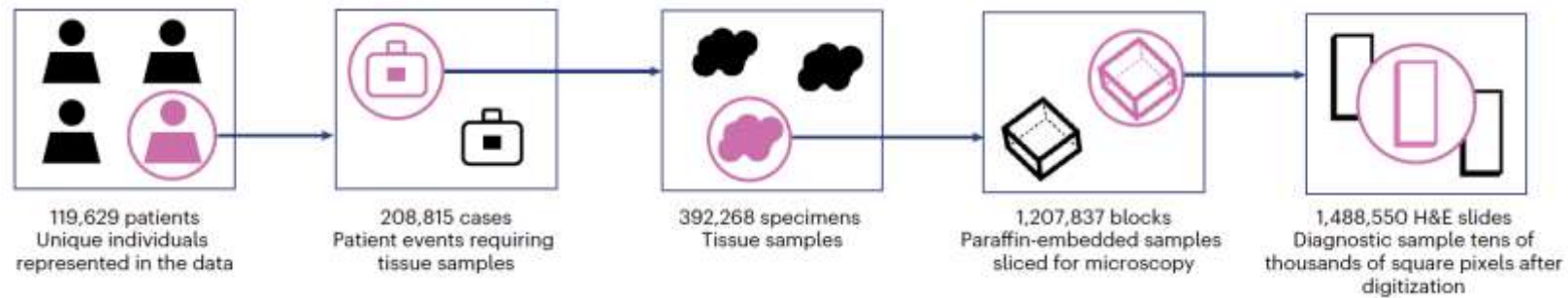
- No!
- Interesting results, which may compete with PD-L1 IHC!
- But, prospective randomized trials are necessary!

OS of ICI monotherapy in NCLC stratified by Deep-IO score
Cut-off: median score

Current developments: Foundation models

One fits all models?

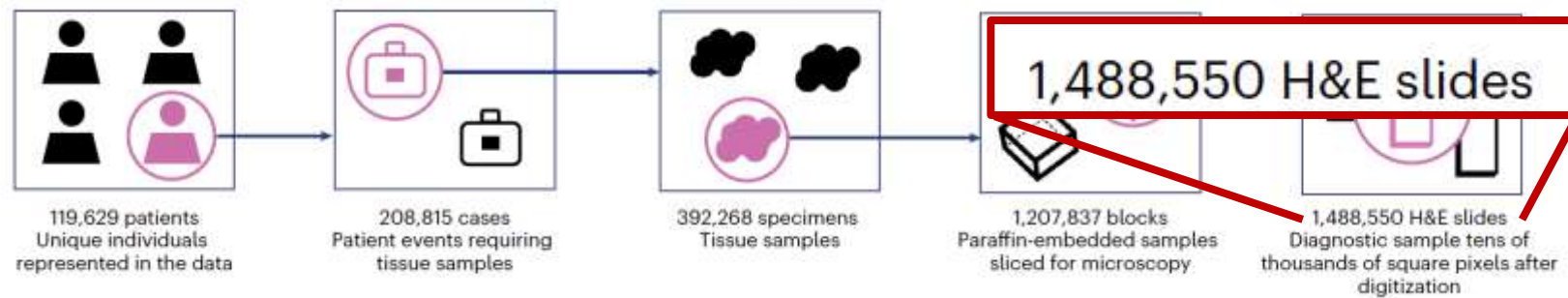
Virchow – A pan-cancer, multi-purpose foundation model



Current developments: Foundation models

One fits all models?

Virchow – A pan-cancer, multi-purpose foundation model

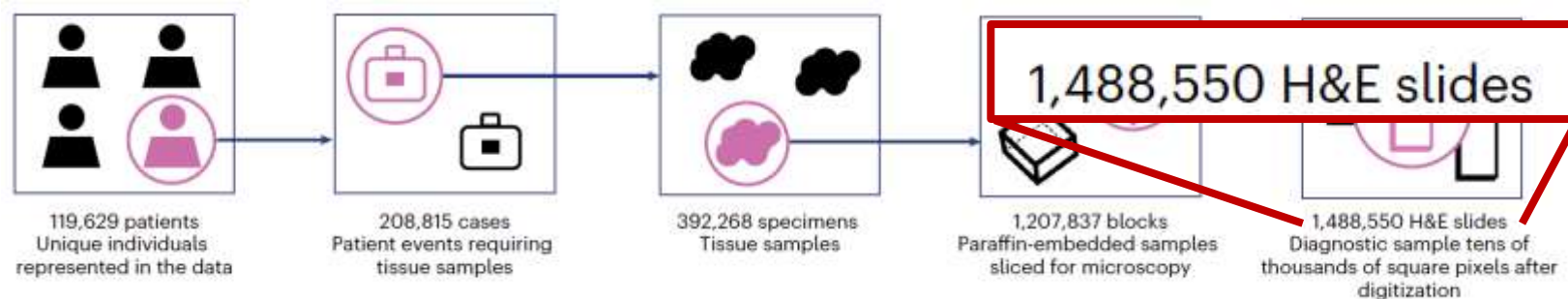


Learning model
Self-supervised
Unlabelled data

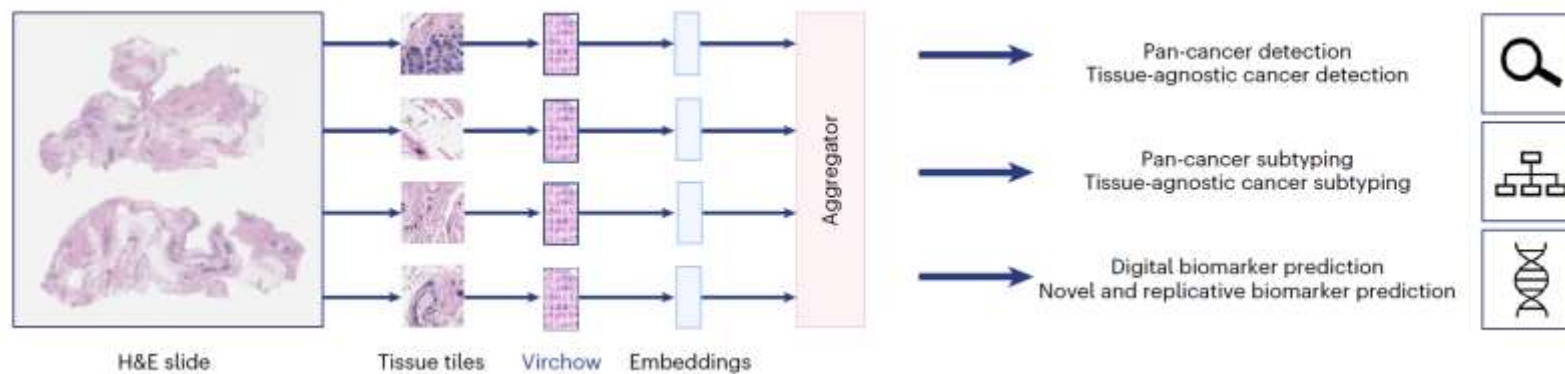
Current developments: Foundation models

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Virchow – A pan-cancer, multi-purpose foundation model



Learning model
Self-supervised (SSL)
Unlabelled data

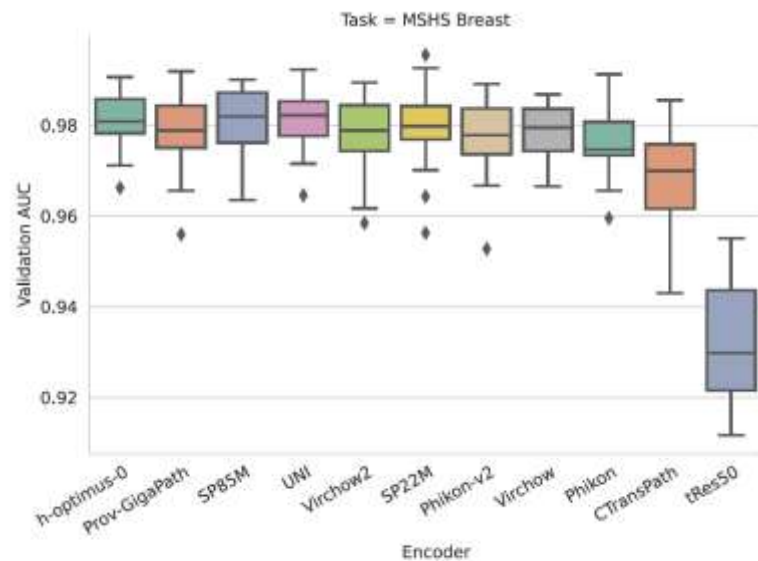


Tasks
Multi-purpose
Multi-task
Multi-modal

Current developments: Foundation models

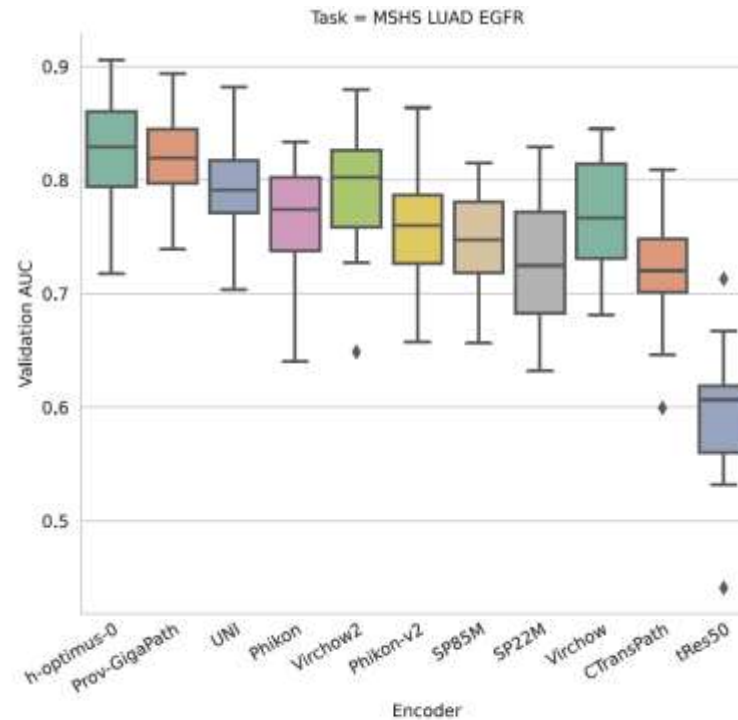
One fits all models?

Challenges – systematic analysis of performance



Example for detection

Breast cancer detection performance



Example for prediction

EGFR mutation prediction performance

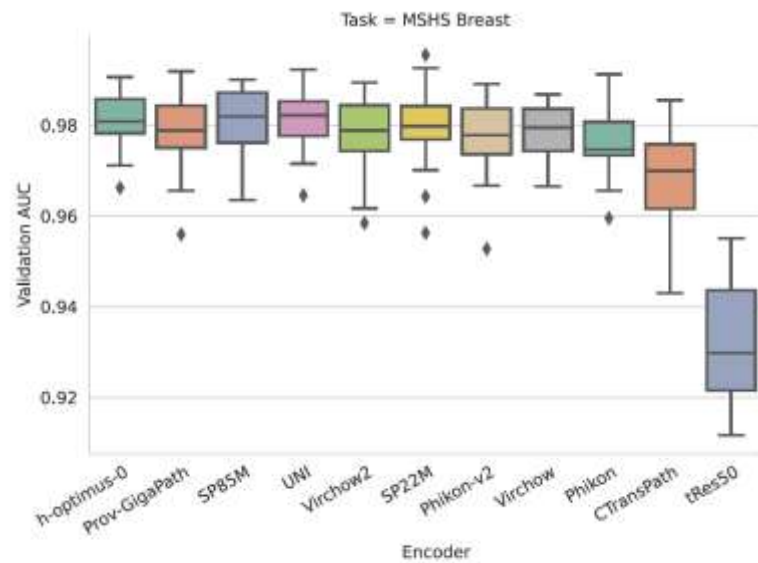
Key findings

- Overall good performance in cancer detection
- Heterogenous performance in biomarker prediction
- No strong correlation between size of pre-training data set and performance – improvement with more samples?

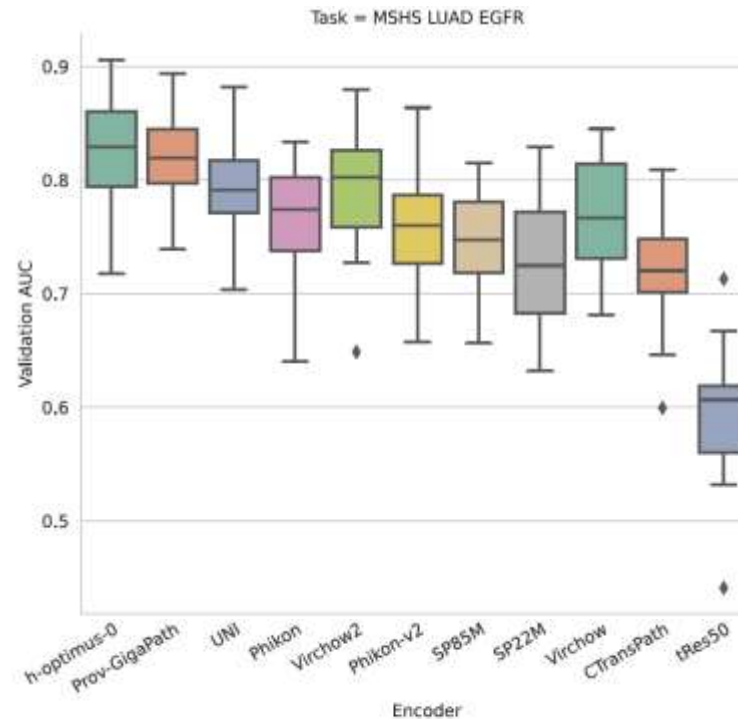
Current developments: Foundation models

One fits all models?

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Example for detection
Breast cancer detection performance



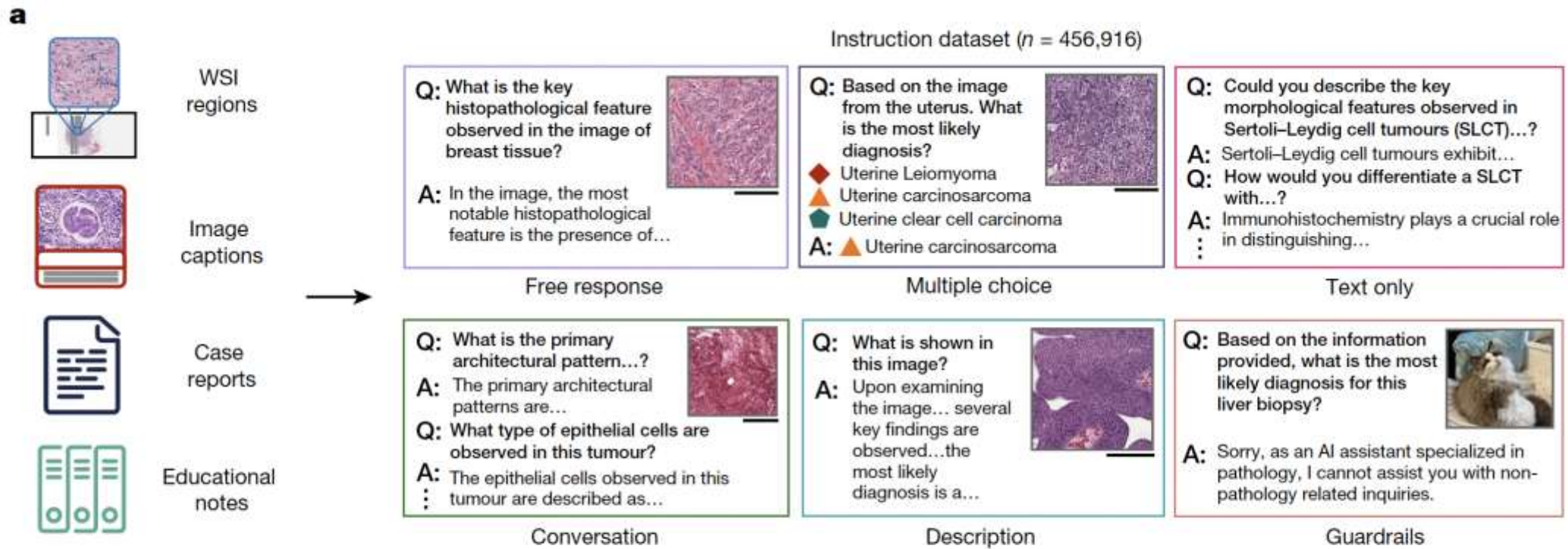
Example for prediction
EGFR mutation prediction performance

Would I base a clinical decision on the AI-based mutation prediction?

- **No – not yet**
- **Refinement need**
- **Including other modules like clinical characteristics and radiomics**

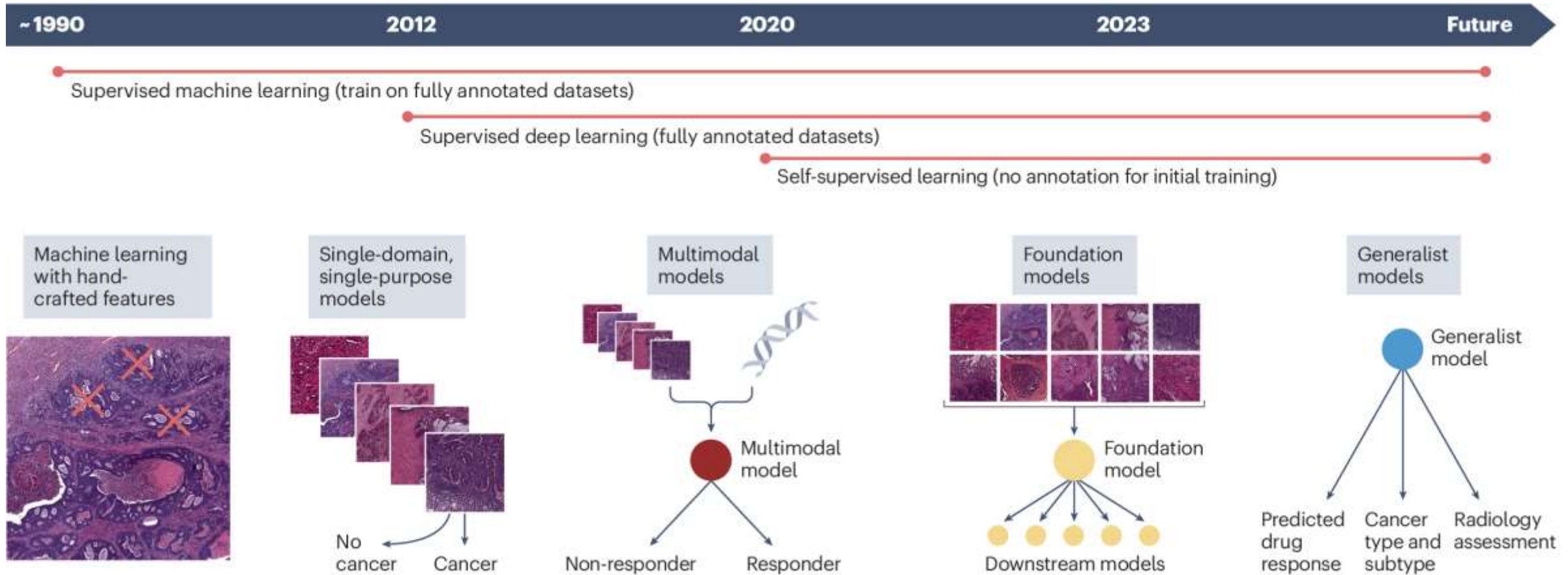
Outlooks: AI co-pilots

Integrating foundation models and specialized LLMs = Multimodal LLM



AI development – summary

From single purpose models to foundation models and beyond



Challenges and hurdles

Why is AI in digital pathology still not routine?

- Task-dependent performance and reliability
 - Currently no approved one-fits-all foundation model
- Generalizability, reproducibility and external validity
 - Clinical evidence gap
 - Lack of prospective trials
 - Lack of clinical meaningful robustness of prediction models
 - Biopsies still represent major challenges due to low tissue quality
- Reimbursement
 - Uncertain and different national reimbursement strategies
- Complex regulatory processes (IVDR)
 - However around 50 CE-IVDR tools are available in the EU, only 2 in the US¹
- Workflow integration and infrastructure
 - High implementation costs and uncertain reimbursement
 - Uneven global implementation

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THANK YOU

