Austrian Cohort Initiative Meeting

# Assessment of the Epigenome – Cohort Scenarios

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Austrian Cohort Initiative Meeting

# Assessment of the Epigenome – Cohort Scenarios

- Rationale for utilising DNA methylation for risk prediction
- ONA methylation and use of surrogate tissue
- Output: Potential cohort biobanks

Output Description (Cancer, Metabolic, Neurologic, etc)

### Unmet needs

## Unmet needs in disease prevention

### **CARDIOVASCULAR DISEASES**



Elevated blood pressure

**Risk factor** 

Risk prediction

SYS

DIA

### **OTHER DISEASES** (cancer, metabolic, neurologic, etc)

many

**Risk factors** 

**Risk prediction** 

???



**Preventive action** (primary/secondary)



### **Risk monitoring**

???

Risk monitoring



Cancer risk prediction

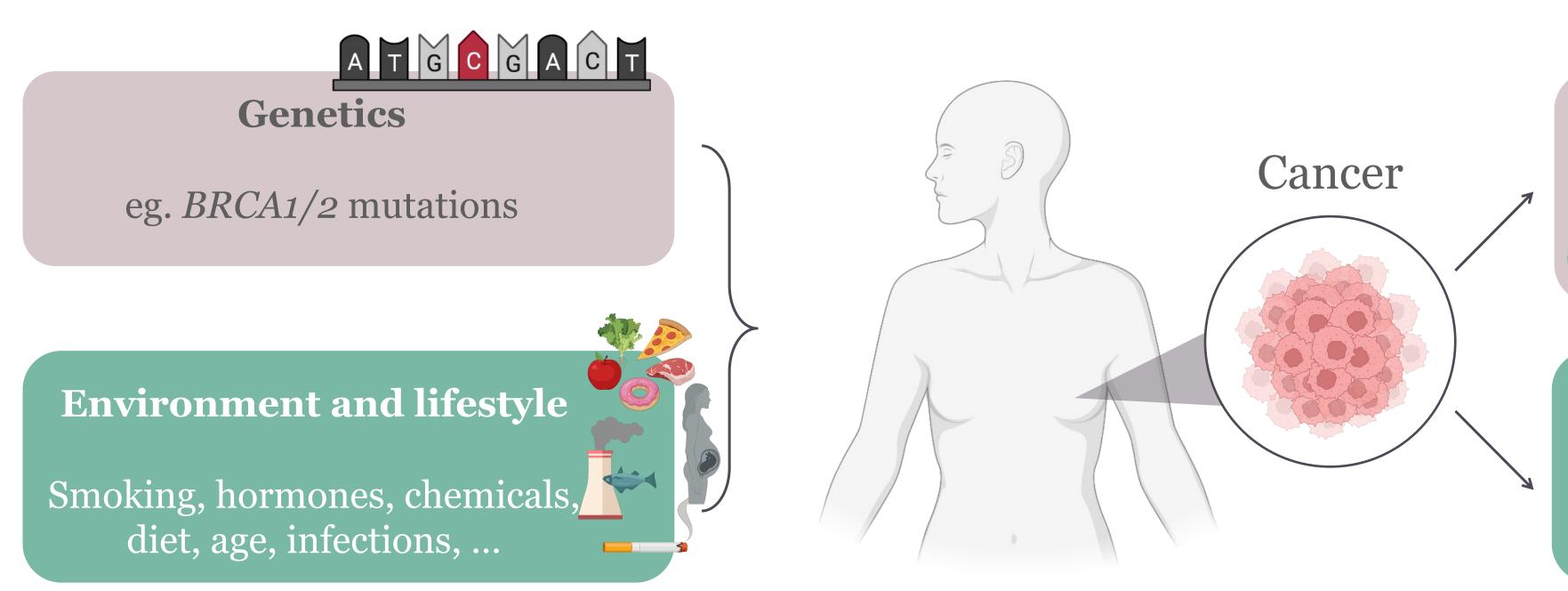
# Rationale for utilising **DNA** methylation for predicting disease risk – example: cancer







### Cancer risk prediction



## To enable cancer risk prediction, we need a tool that can account for both genetic and environmental factors

Genetic risk prediction based on mutation analysis or polygenic risk scores static, captures only proportion

**Epidemiological** risk prediction based on environmental factors does not account for interaction of genetics and environment

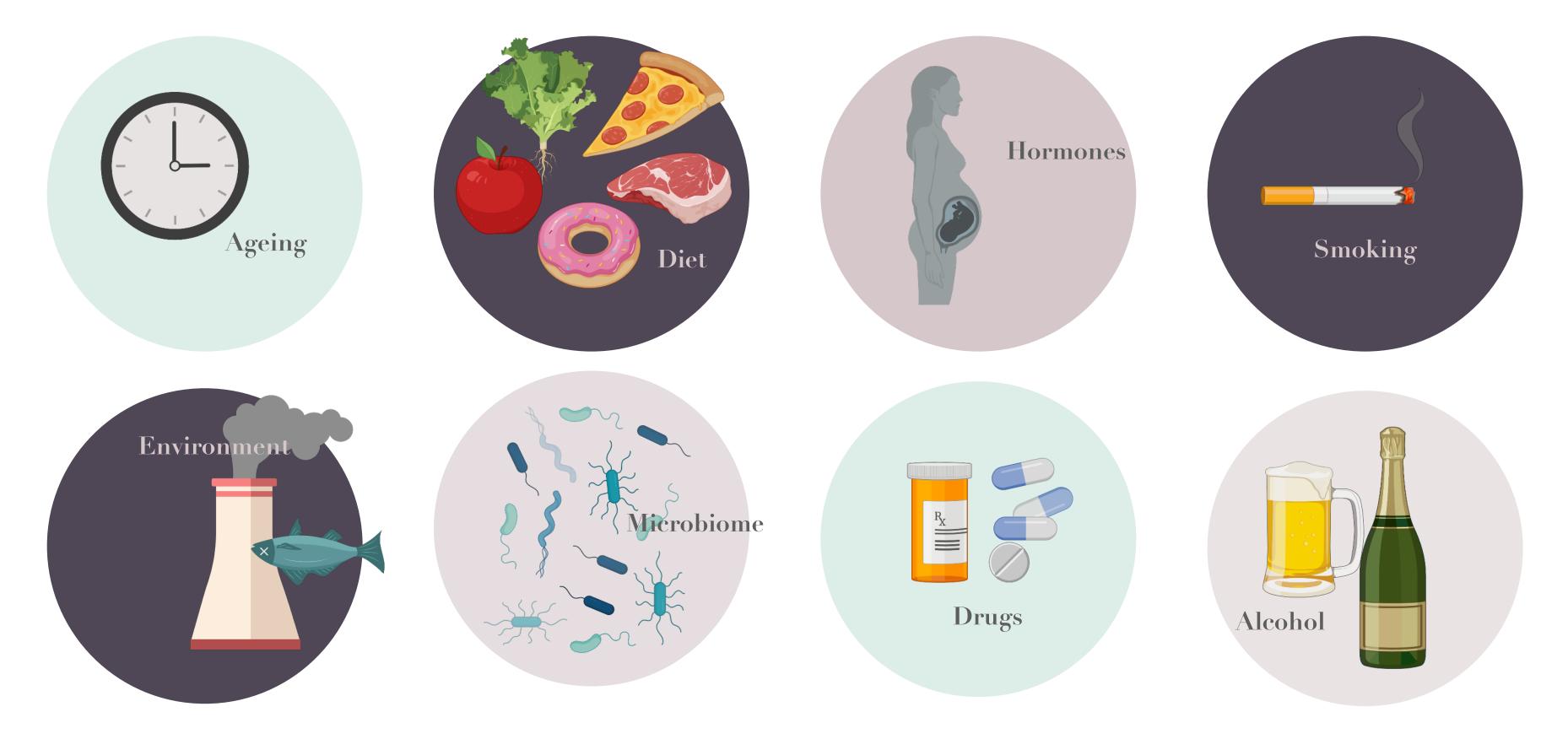






### **Epigenetics – DNA methylation**

## DNA sequence is determined at birth. *Epigenetics* can be influenced by external factors.





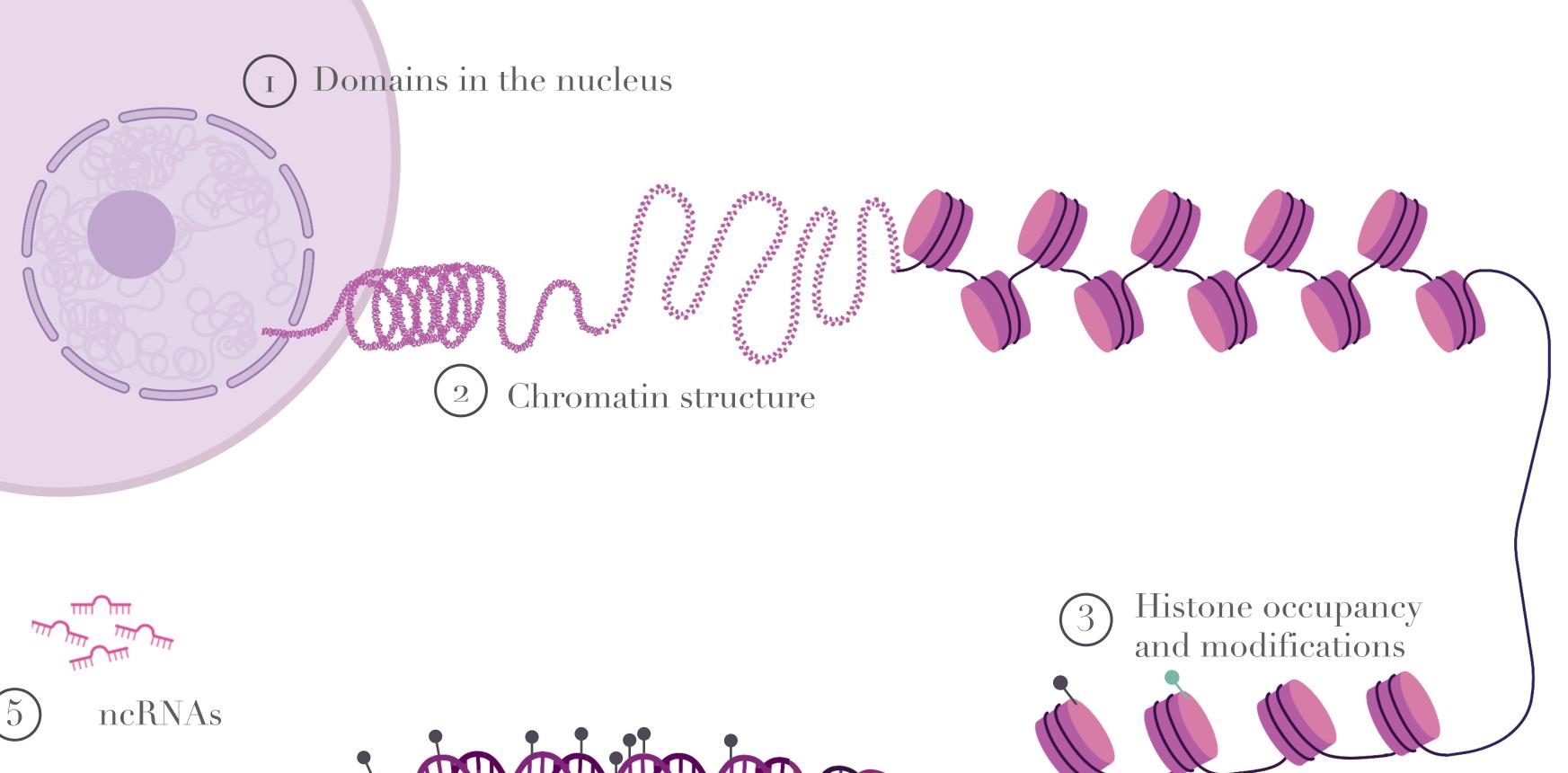
**Epigenetics – DNA methylation** 

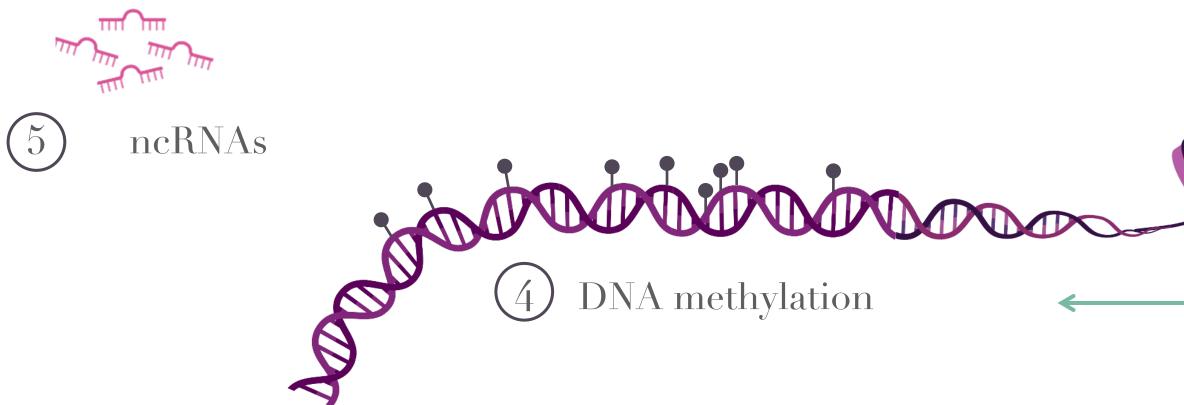
## DNA methylation





### **Epigenetics – DNA methylation**



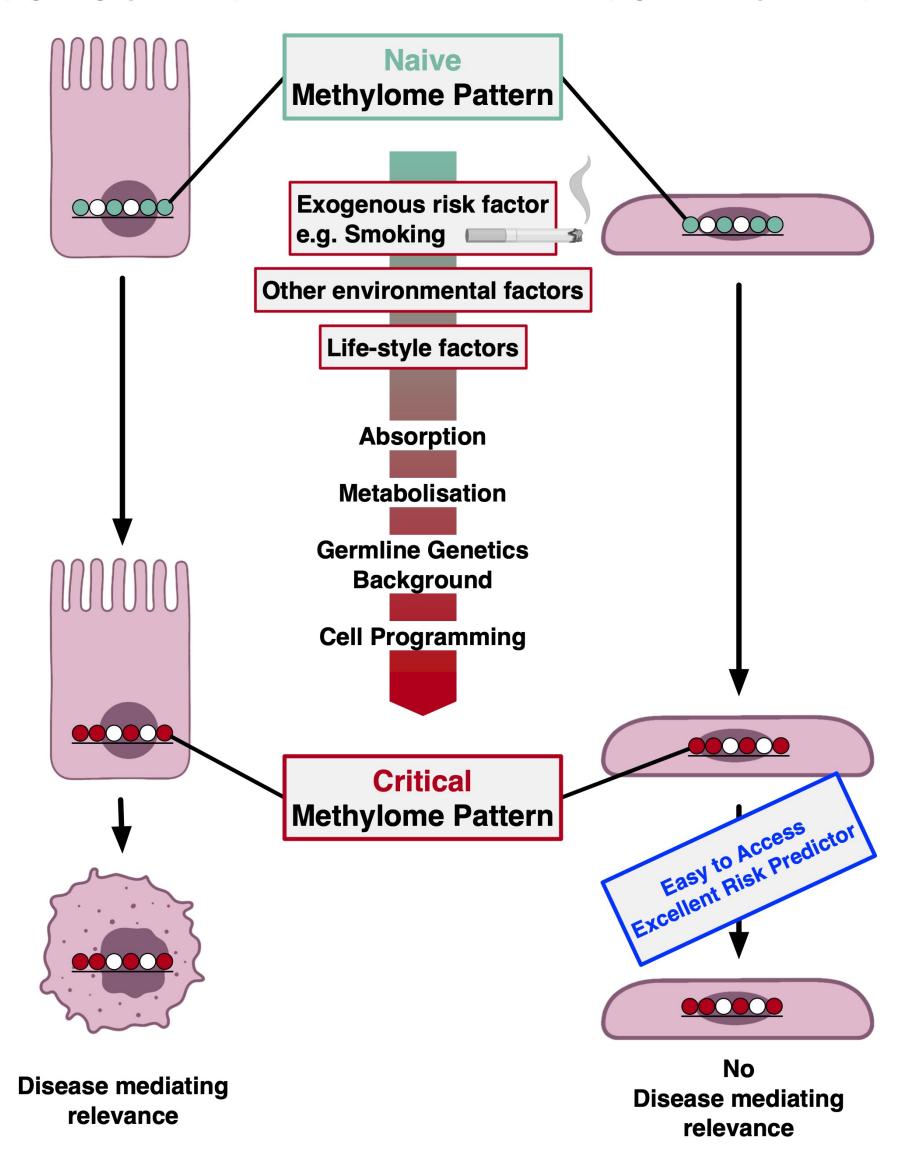


High-throughput methodologies available

### Surrogate tissue

### **Tissue at Risk**

(e.g. lung epithelium)



### Surrogate Tissue

(e.g. buccal epithelium)







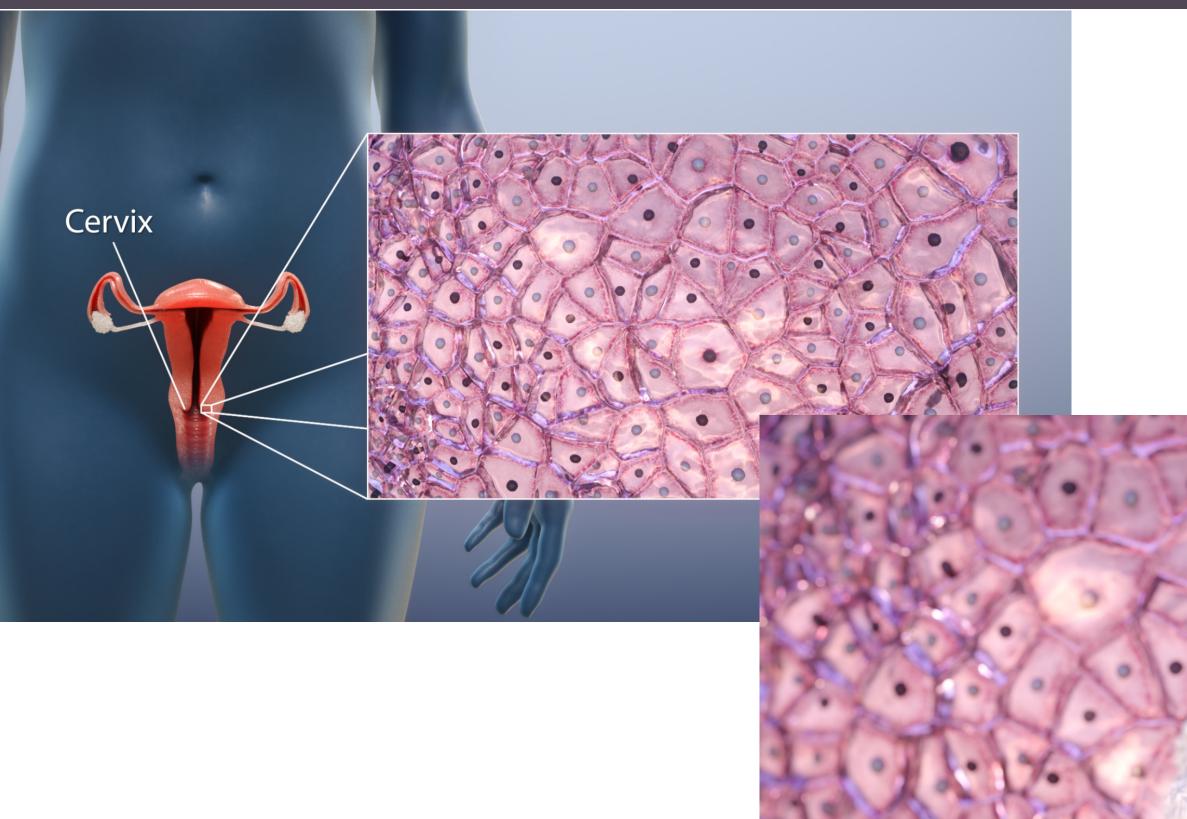
Needs of surrogate tissue

# How does an ideal surrogate tissue look like?

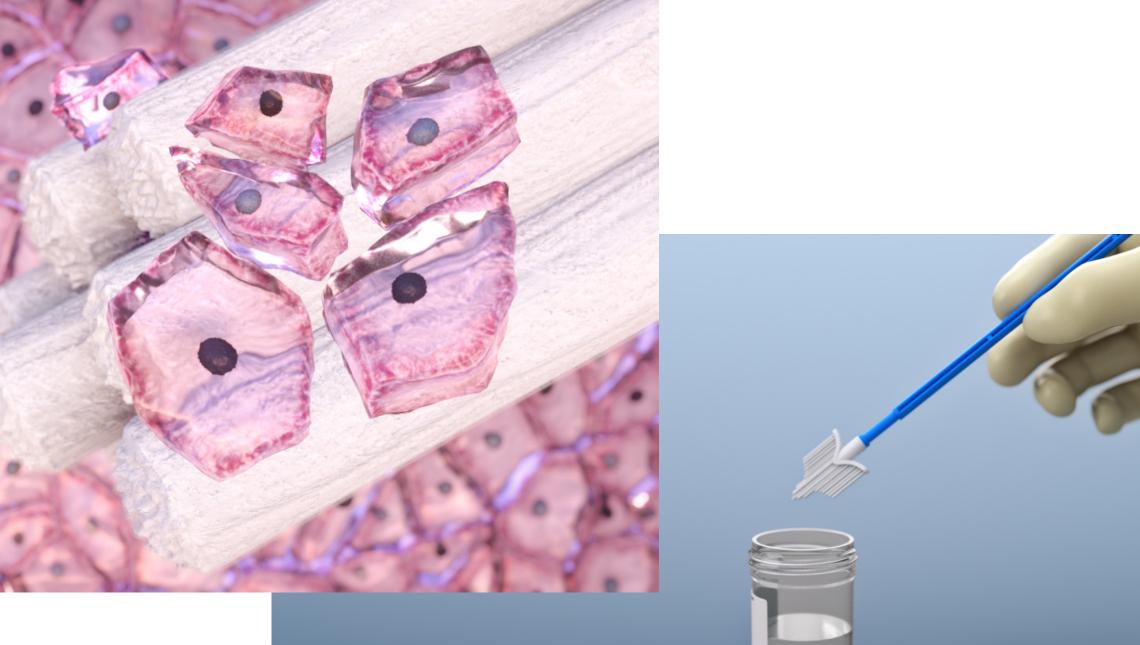
- Easy accessible (ideally self-collected)
- Capturing specific disease pathways  $\bigcirc$
- Variability of features reflective of tissue at risk
- Similar/same embryological origin



## Surrogate tissue – easy accessible









### Surrogate tissue – ideally self-sampling is feasible



Wash your hands before usage.

Remove the Evalyn Brush from the packaging. Do not throw the packaging away, as it is necessary for sending the Evalyn Brush to the laboratory after usage.



Press the sides of the pink cap with your thumb and index finger to remove the pink cap from the Evalyn Brush. Ensure that you do not touch the white brush of the Evalyn Brush with





Place the packaging containing the Evalyn Brush into the plastic bag provided and seal it.





fold the transparent end to insure the white brush does not extend again. Place the pink cap eack on the Evalyn Brush using our thumb and index finger. You will hear a click when it is properly in place.

Obtain the sample whilst in a standing position. Assume a comfortable stance (e.g. as if you were about to insert a tampon).



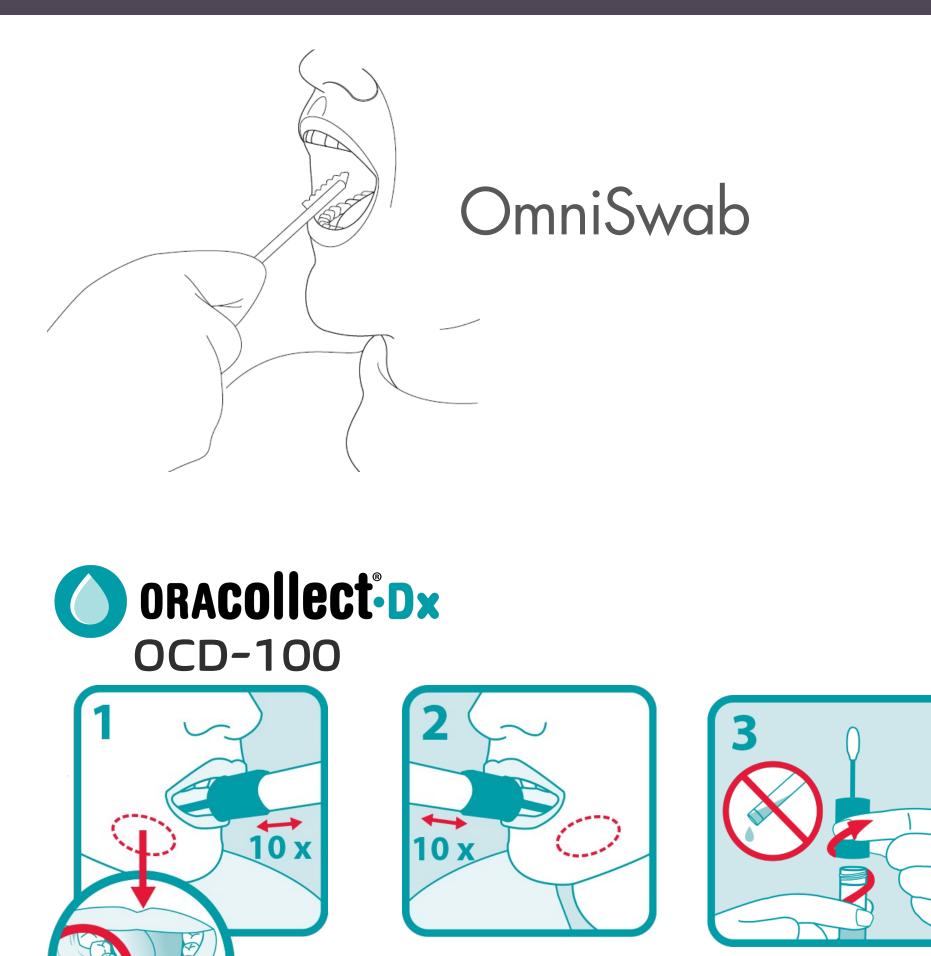
send the plastic bag containing the Evalyn Brush together with other required information.







### Surrogate tissue – ideally self-sampling is feasible



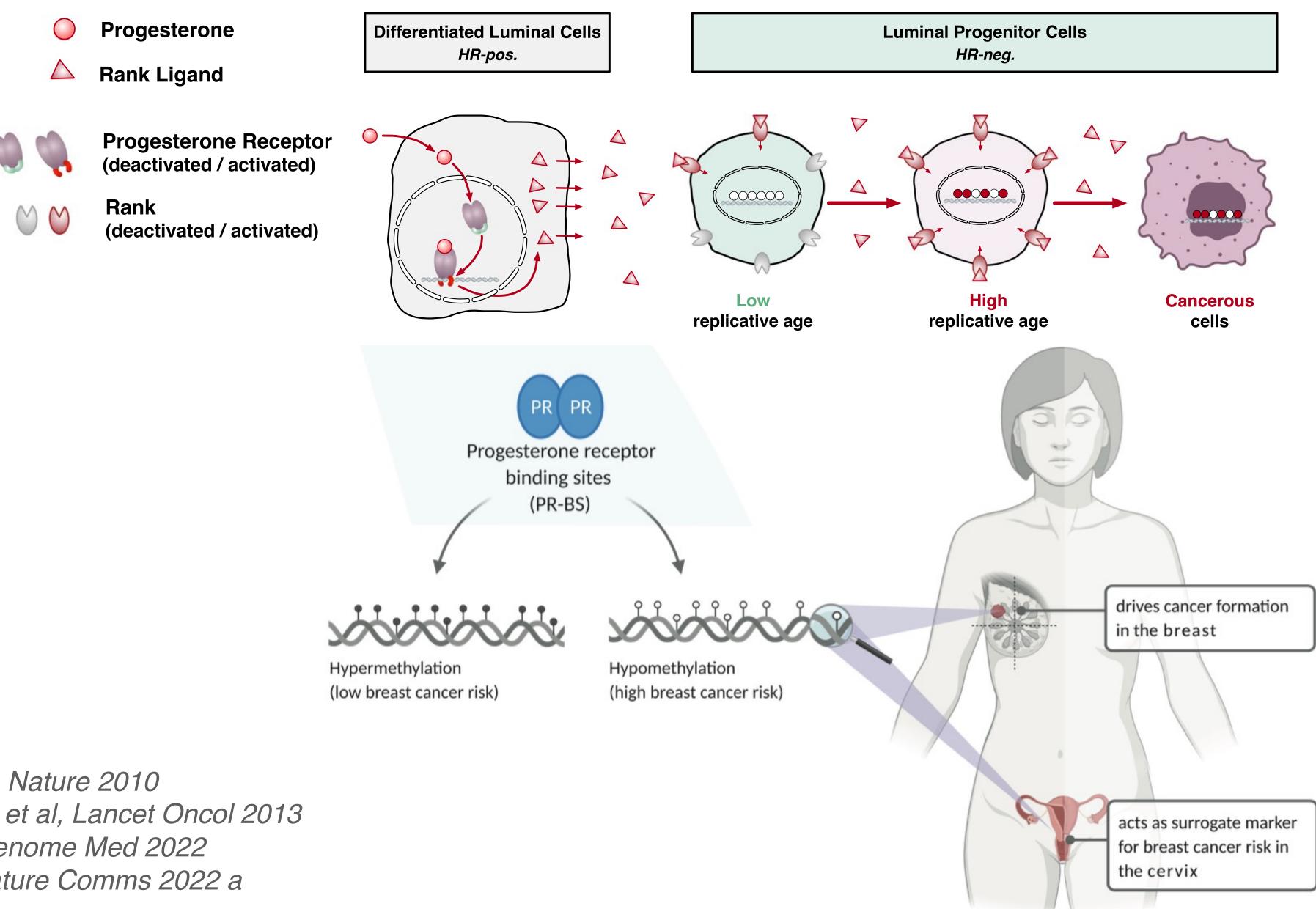




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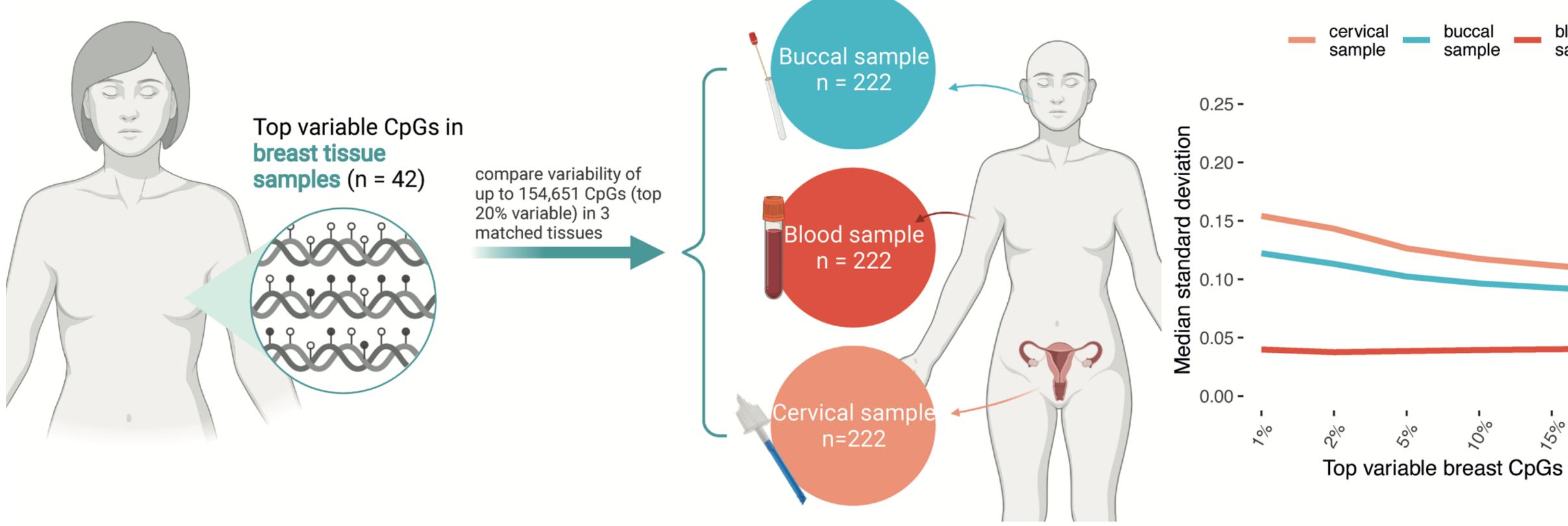


## Surrogate tissue – capturing specific disease pathways



Schramek et al, Nature 2010 Widschwendter et al, Lancet Oncol 2013 Bartlett et al, Genome Med 2022 Barrett et al, Nature Comms 2022 a

### Surrogate tissue – reflective of variability in tissue at risk

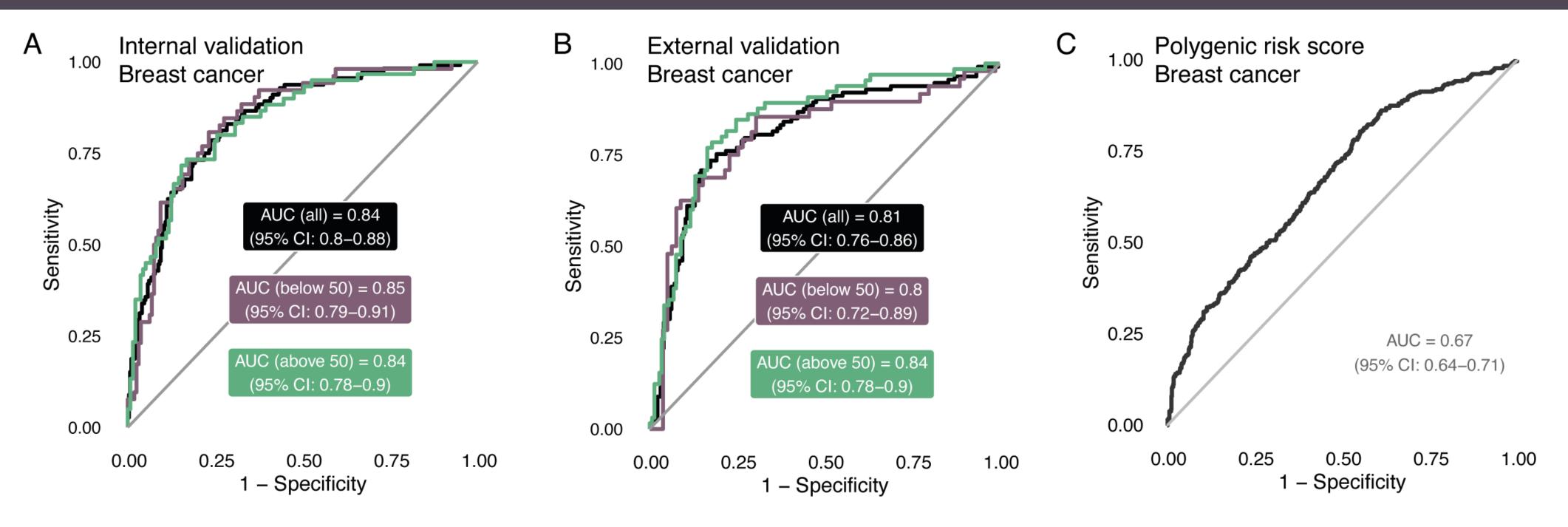








### **Breast Cancer risk prediction – WID-BC**



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Quantile	Control	Cancer	OR (unadjusted)	OR (adjusted)	Risk group	Controls	Cases	OR (unadjusted)	OR (adjusted)
Internal validation					Internal validation				
(-1.53, -0.58)	75	2	1.00 (reference)	1.00 (reference)	low PRS <sub>313</sub> , low WID™-BC	109	6	1.00 (reference)	-
(-0.58, -0.28)	74	5	2.42 (0.48,19.25)	2.29 (0.45, 17.15)	high PRS <sub>313</sub> , low WID™-BC	70	9	2.3 (0.78-7.3)	2.94 (0.95, 9.94)
(-0.28, 0.07)	74	17	8.01 (2.17,56.31)	8.47 (2.23, 55.81)	low PRS <sub>313</sub> , high WID™-BC	54	25	8.2 (3.3-23)	10.17 (3.68, 33.75)
(0.07, 1.62)	74	88	41.11 (12.33,274.77)	41.73 (12.2, 262.62)	high PRS <sub>₃13</sub> , high WID™-BC	47	67	25 (11-69)	26.05 (11.15, 72.13)
External validation					External validation				
(-1.53, -0.58)	58	8	1.00 (reference)	1.00 (reference)	low PRS <sub>313</sub> , low WID™-BC	82	9	1.00 (reference)	-
(-0.58, -0.28)	69	8	0.84 (0.29,2.46)	0.89 (0.3, 2.67)	high PRS <sub>313</sub> , Iow WID™-BC	57	13	2.1 (0.82-5.4)	2.26 (0.89, 5.96)
(-0.28, 0.07)	50	14	2 (0.78,5.46)	2.57 (0.95, 7.51)	low PRS <sub>313</sub> , high WID™-BC	41	29	6.3 (2.8-15)	10.59 (3.97, 32.71)
(0.07, 1.62)	48	83	12.19 (5.62,29.86)	15.67 (6.59, 42.38)	high PRS <sub>₃13</sub> , high WID™-BC	30	60	18 (8.1-42)	18.35 (7.71, 49.24)

Barrett et al, Nature Comms 2022 a

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### Surrogate tissues

## Various issues to consider

- Tissue specificity
- Storage requirements
- Age dependence
- Hormonal (cycle) exposure
- Circadian rhythm

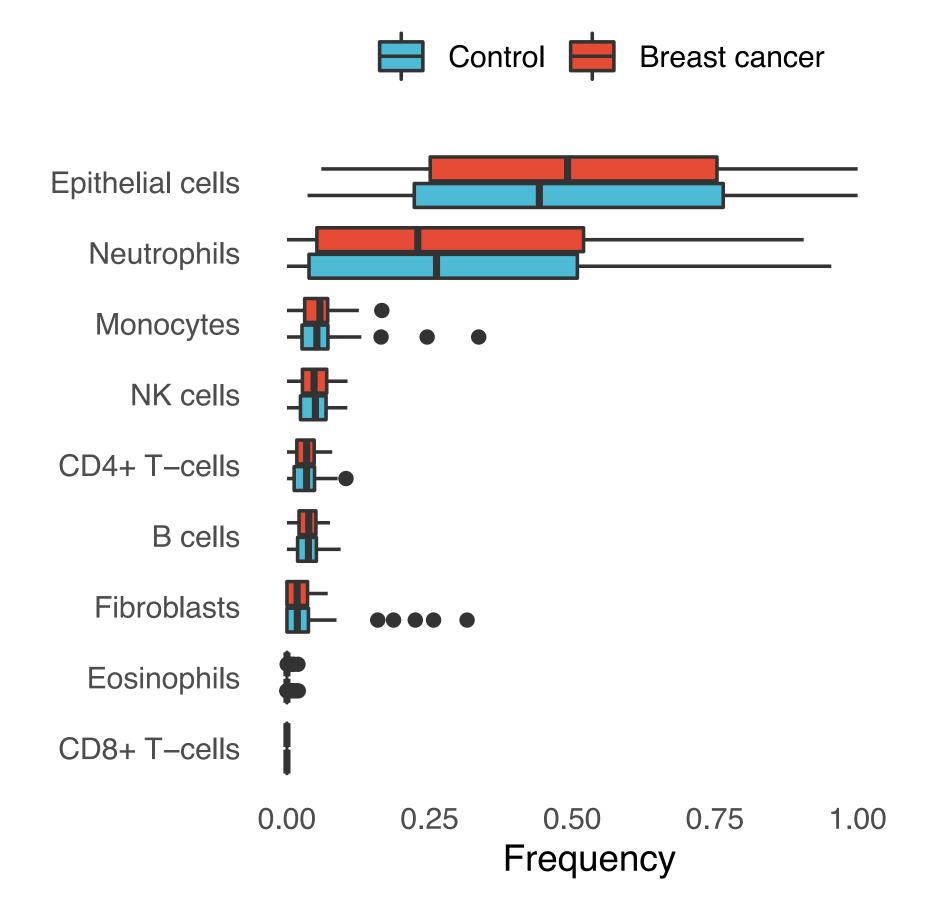




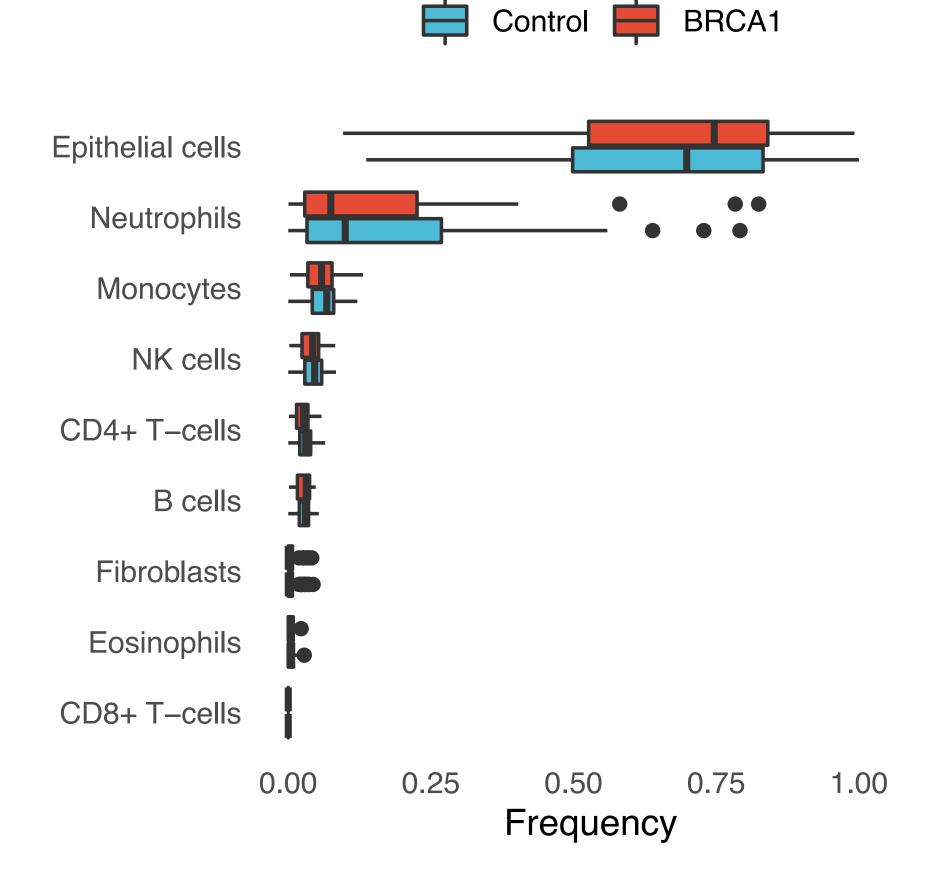


## Assessment of proportion of cell types utilising DNAme<sup>n</sup>in heterogenous samples

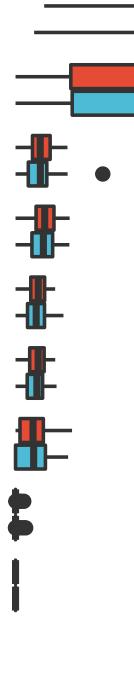








Barrett et al, Nature Comms 2022 a





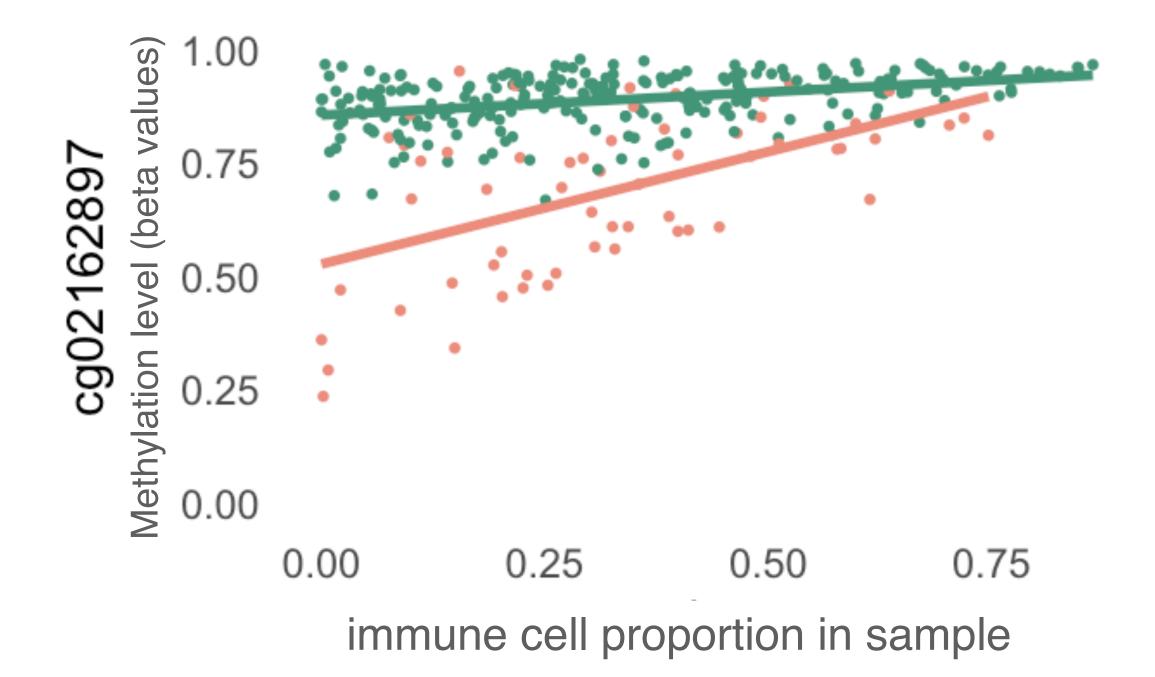
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### Tissue dependent recording of environmental exposure

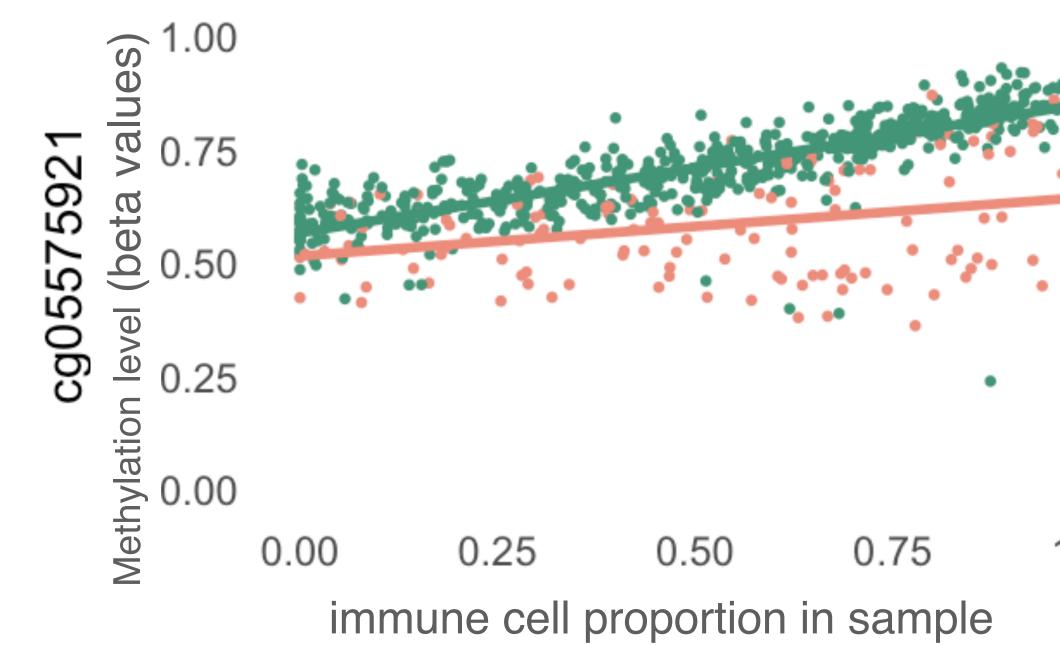


Control Smoker

**Epithelial effect** 



### *Immune cell effect*



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1.00



### Austrian Cohorts

# SUGGESTIONS for AUSTRIAN COHORTS







### Austrian Cohorts

- Capturing factors determining health and disease
- Three different albeit complementary settings spanning the entire life-course
- Uniqueness (compared to existing international efforts)
  - Longitudinal and repetitive sampling
  - Self-sampling
  - Including non-blood (i.e., epithelial)







### Austrian Cohorts - Initiative 1: In utero exposure ("Austrian Birth Cohort")

- Only a few dedicated centres across Austria (E, W, S, N) • Invitation of pregnant women (via office gynaecologists) who are likely to give birth in the relevant
- centres
- Collecting samples (blood, faeces, saliva, vaginal swab, buccal sample assessing metaboloms/toxins, microbiomes, etc.) and data/questionnaires throughout pregnancy and documenting the course of pregnancy
- Collecting placental and umbilical blood at birth and isolate:
  - Placental cells (fibroblasts, endothelial cells, macrophages, trophoblastic cells)
  - CD34 cells from umbilical blood
- Readout from foetal/placental cells are:
  - DNAmut
  - DNAme
  - Other omics
  - In a subsetset: Organoids and possibly plasticity of cells (i.e., proportion of successful induction of pluripotency from placental fibroblasts, etc.)
- Link of exposure/data collected throughout pregnancy with the readout data

 Long-term follow up data of children low numbers

**Challenges:** Standardised procedures, laborious and time-consuming procedures, expected relative







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- Women attending Mammography Screening Centres
- Dedicated App to obtain epidemiological data
- Link to registries
- Electronic consenting
- Collection of four specimens at the outset:
  - Cervical self-sample (Evalyn brush  $\rightarrow$  Thinprep)
  - Healthcare professional sample (Cervex  $\rightarrow$  Thinprep)
  - Buccal self-sample (HRC-100)
  - Urine
- Repeat collection (3 specimens) every year (approached by post, self-sampling only)
- Collection of epidemiological (volunteer-reported) every 6 months
- Access to imaging data (i.e., breast density)
- Initiative expanding to other European countries (SWE, IT)
- Core purpose is to develop/validate risk predictive algorithms **Challenges:** To standardise procedures, to secure buy in from screening centres

- Dedicated campaign in specific Austrian cities, villages, communities, companies
- Women and men > 18 years of age (no upper age limit)
- Providing buccal self-sample every 6 months (HRC-100; sent by post)
- Dedicated App to obtain epidemiological data (volunteer-reported) every 6 months
- Link to registries
- Electronic consenting
- associated with disease risk (i.e., cancer, metabolic, cardiovascular, neurodegeneration) **Challenges:** To standardise procedures, to secure buy in from screening centres

• Core purpose is to identify factors that accelerate or decelerate the ageing process (e.g. telomere lengths, DNAmut like CHIP, DNAme, etc) and to assess whether the slope of the ageing curve (i.e., as assessed by longitudinal/repetitive measurements) and its relation to chronological age is





### Example:

the ageing process (Initiative 3)

Assess whether molecular data that indicate disease risk (e.g., for BC; evidence obtained in Initiative 2) are being initiated in utero by specific exposures (e.g., Bisphenol as assessed in Initiative 1) and whether molecular features which are driven by in-utero exposure are involved in











### Austrian Cohorts

## DISCUSSION

