

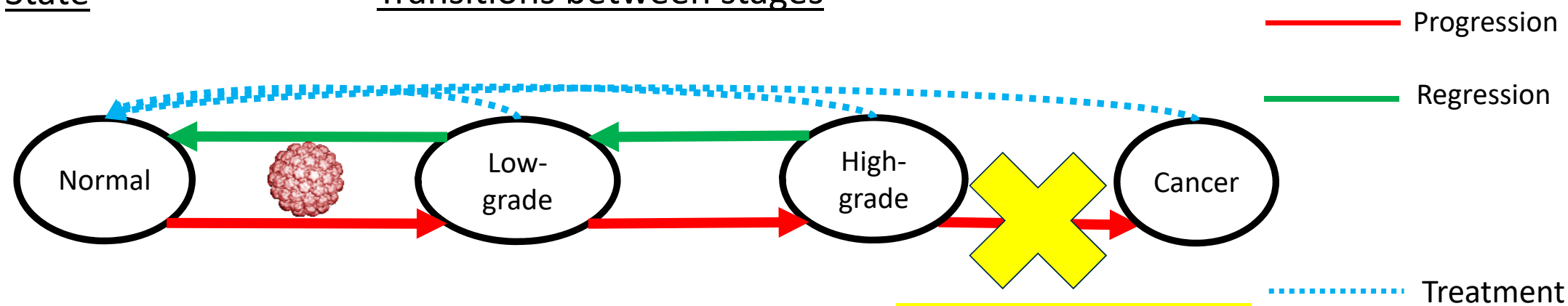
# Innovative methods in Cancer Screening

How Norway explores AI and uses the world's most powerful computer power to predict who has the largest risk of cervical cancer

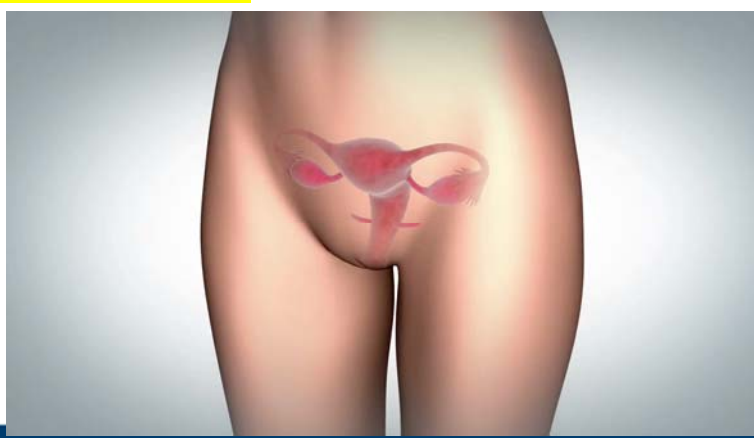
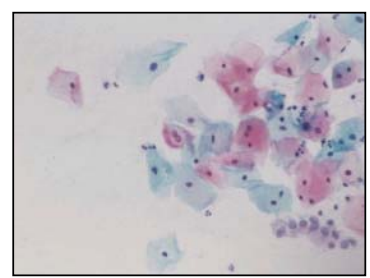
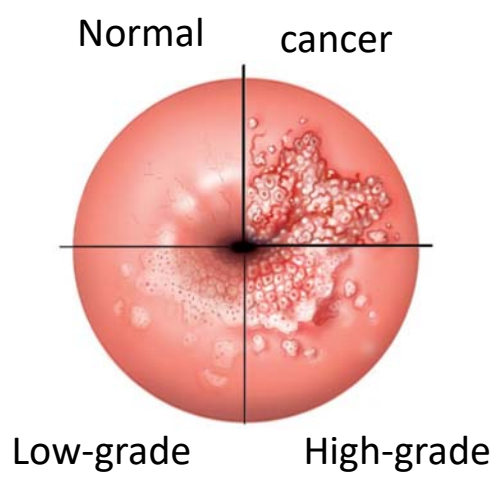
**Mari Nygård, MD, PhD**  
**Head Research Department, Cancer Registry of Norway**

State

Transitions between stages



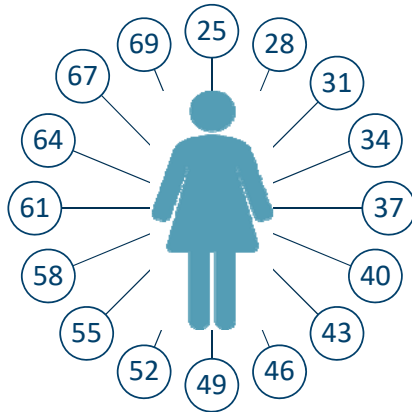
**Screening program**



# Mass-screening - a concept from 1960ies

## Screening

cytology exam for population without symptoms



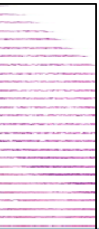
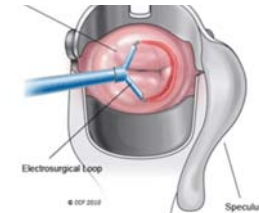
## Diagnostic confirmation and treatment



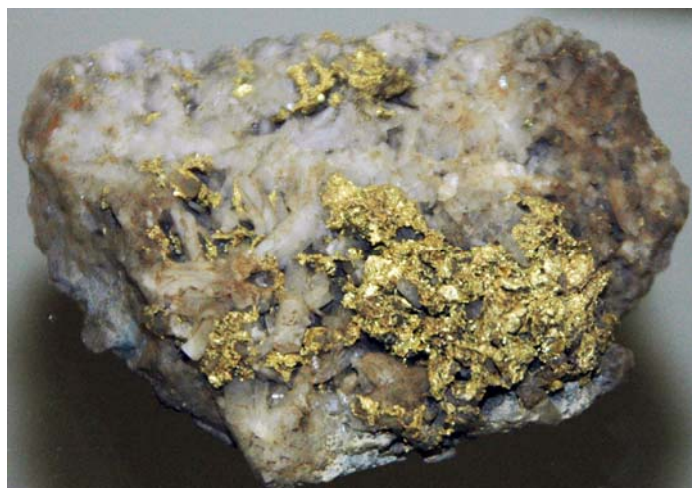
Punch biopsy  
forceps



Loop electrosurgical  
excision procedure



# Mass-screening program - a concept from 1960ies was implemented in 1995 in Norway



without symptoms

ID - women



**1.8 M women  
in the database**

date of the screening exam



**About 12.5 M exams  
Including date and results**

diagnose



ID- patient

date of the diagnostic exam

**Annually  
about 10,000 Punch biopsies  
About 6,000 treatments**

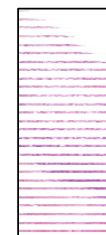
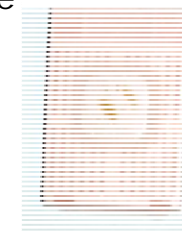
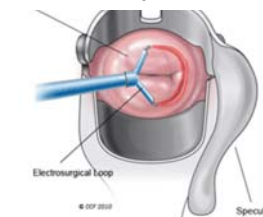


Punch biopsy  
forceps



date of the treatment

Loop electrosurgical  
excision procedure



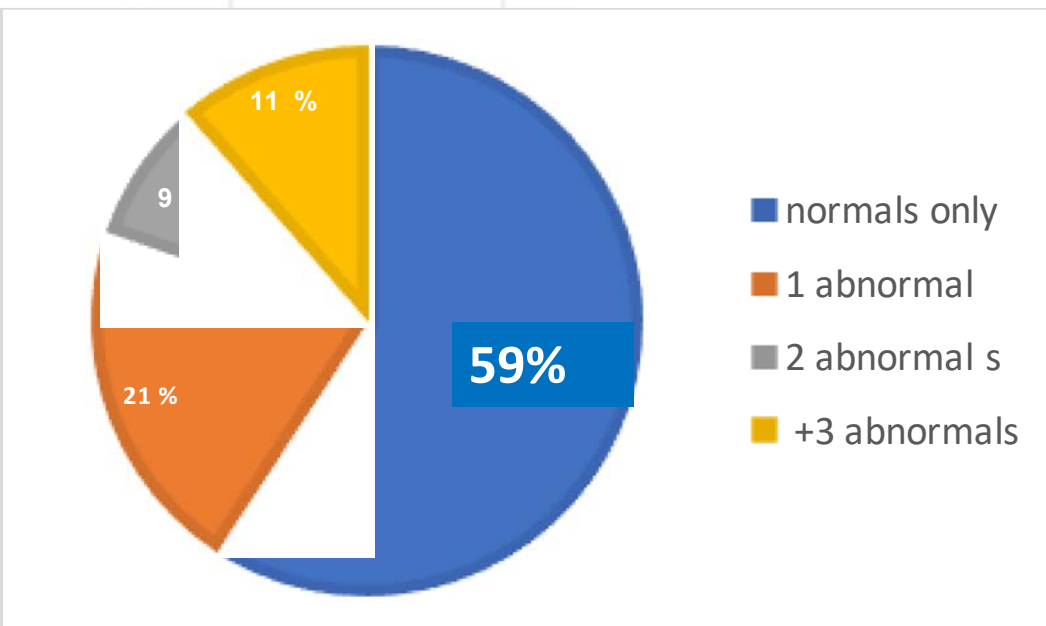
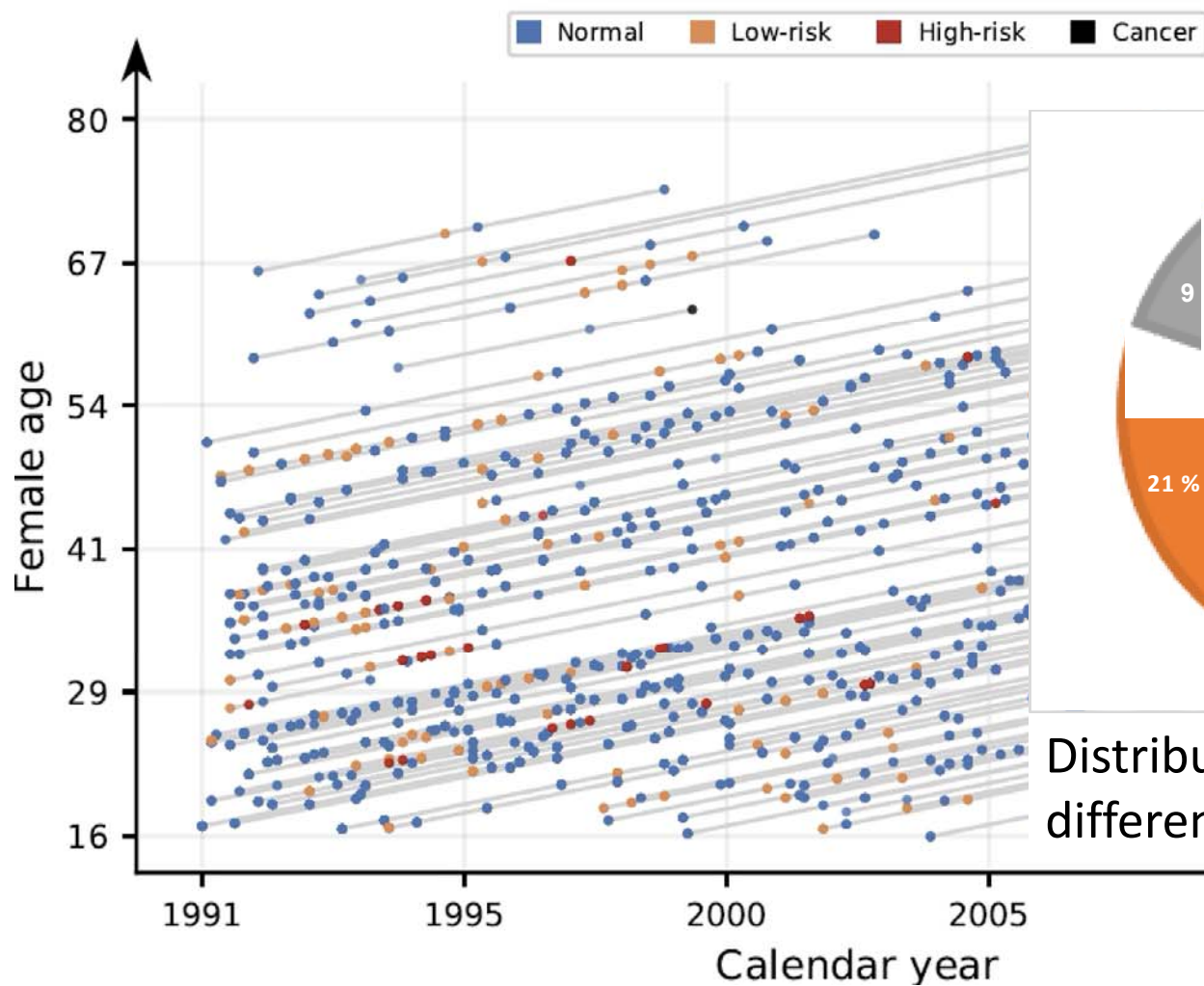
# Mass-screening program - a concept from 1960ies was implemented in 1992 in Norway

|   | ID - women                             | date of the screening exam                               | diagnose |
|---|--|--|----------|
| <u>Screening</u><br>cytology exam for population without symptoms | <b>1.8 M women<br/>in the database</b> | <b>About 12.5 M exams<br/>Including date and results</b> |          |

90% of test are normal, 2% suggest disease, 5% suggest that “something is going on», 3 % are technical failures

## Diagnostic confirmation and treatment

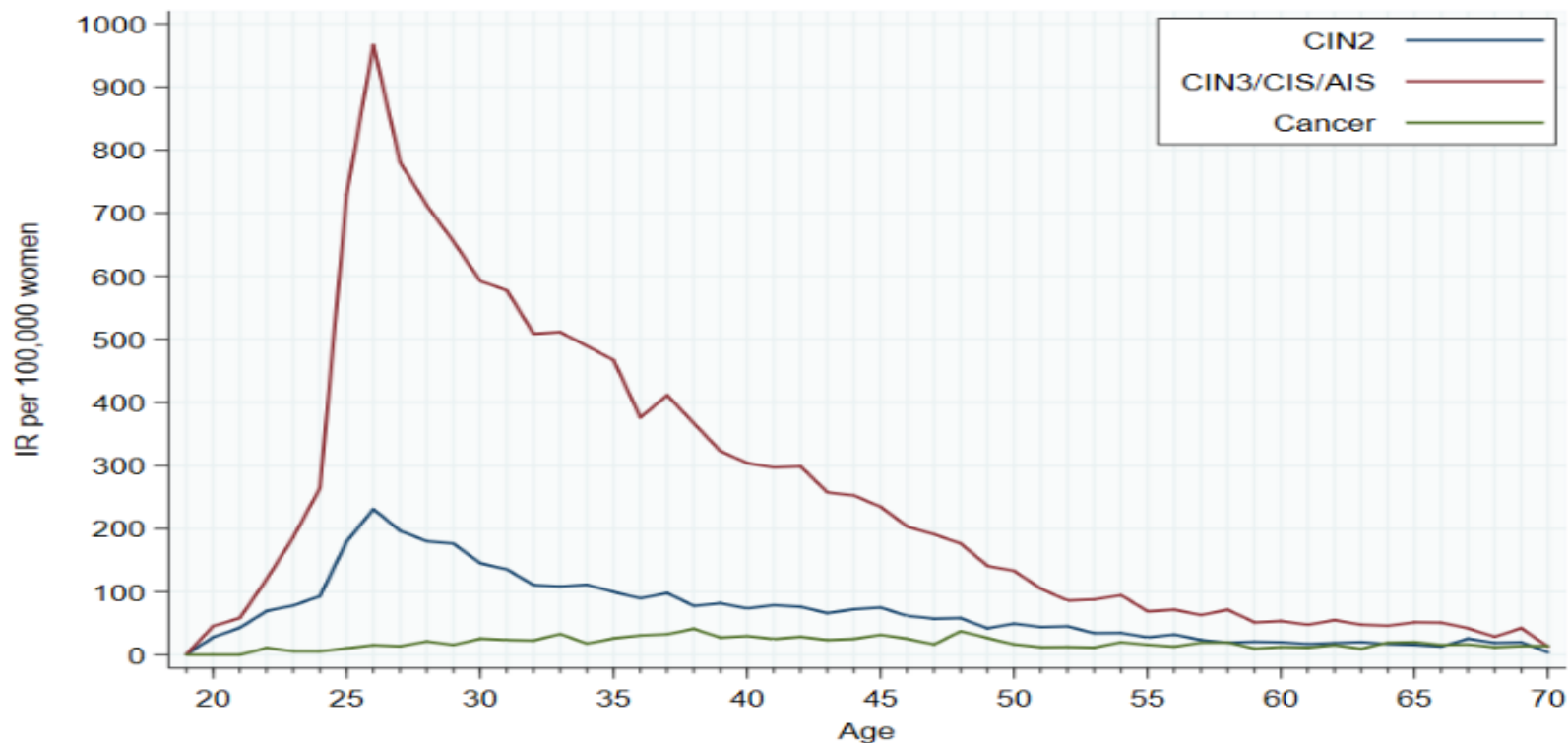
|  |                                   |
|--|-----------------------------------|
| <b>Annually<br/>about 10,000 Punch biopsies<br/>About 6,000 treatments</b> | <b>Including date and results</b> |
|--|-----------------------------------|



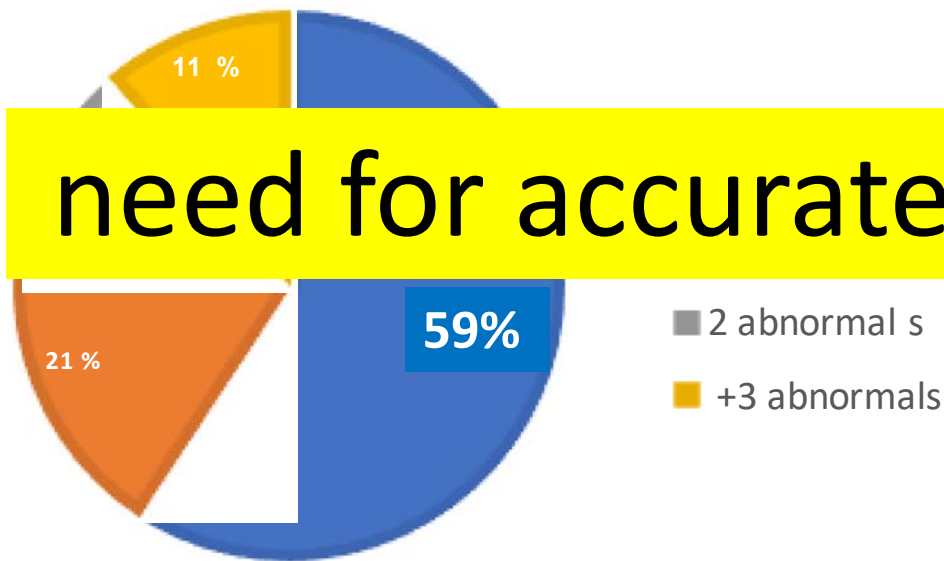
Distribution of 25-69 yrs old women with different screening results



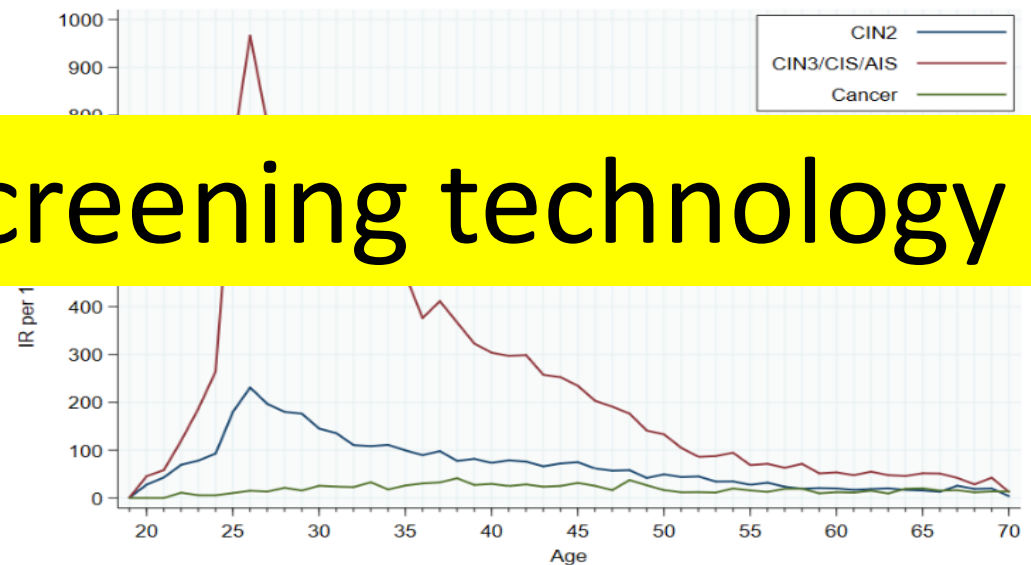
# Cervical pre-cancers and cancers by age in 2014-2016 Norway



# Harms of the mass-screening program: over-screening and over-treatment



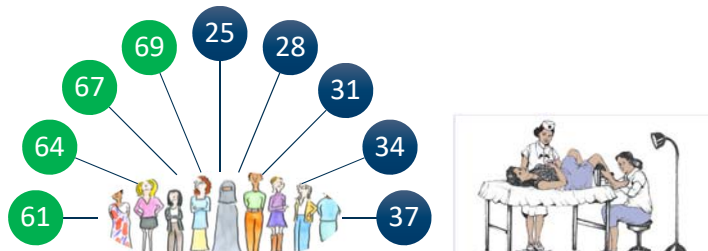
Distribution of 25-69 yrs old women with different screening results



Cervical pre-cancers and cancers by age in 2014-2016 Norway



# Mass-screening program is not reaching to all women

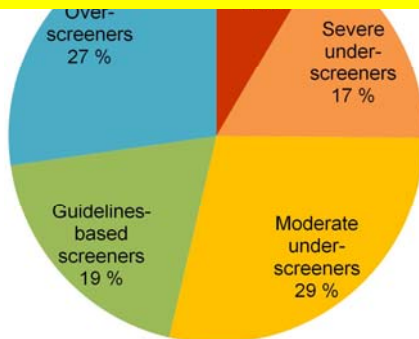


Proportion of those with one screening test in suggested time interval in Norway by age and year:

65% 25-39 yrs old

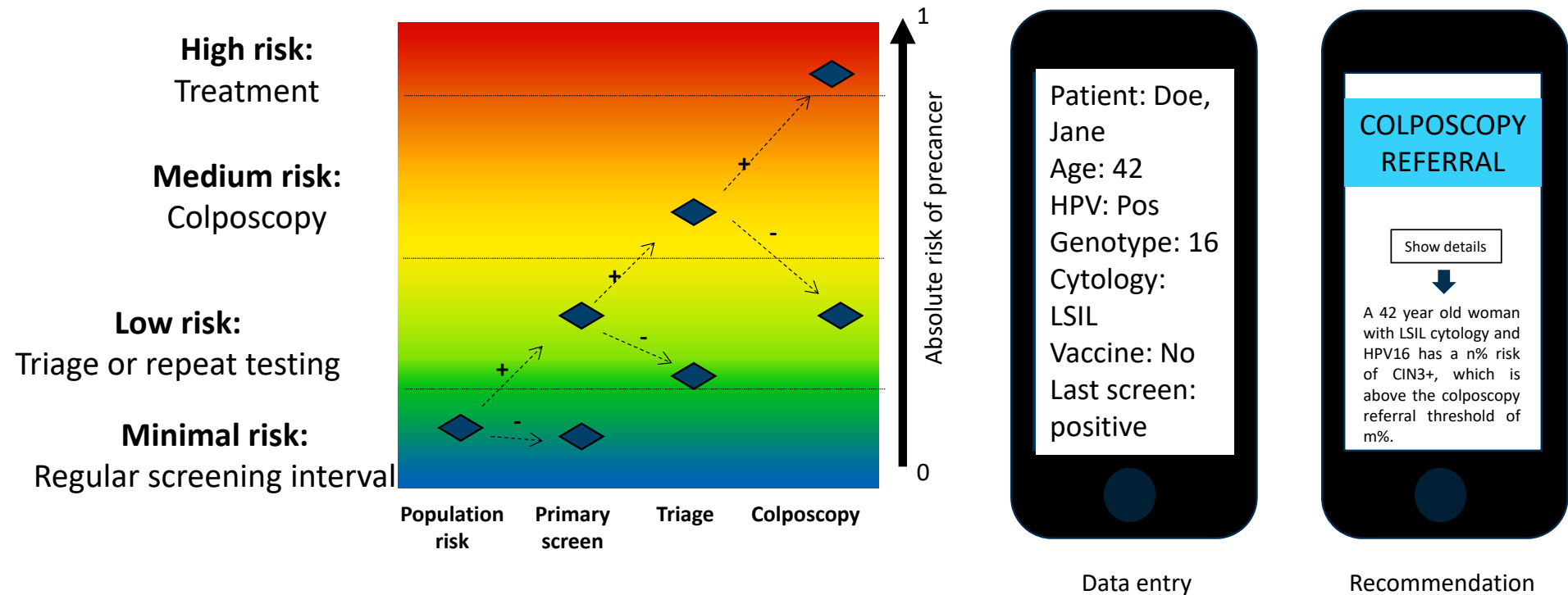
71% 40-54 yrs old

Screening should reach to those who need it most



Cancer incidence is highest among suboptimally screened

# Harmonizing Screening According To Risk



**Future Issues: Integration of Important Modifiers of Cervical Cancer Risk ( personalized screening)** Screening History, Integration of new biomarkers e.g., genotyping, p16 Ki67, mRNA, Methylation, HPV Vaccination

# *An HPC Application to Population-Level Cancer Screening Data*

HPC and Data Intensive Resources at LLNL

Personal level data from the Cancer Registry of Norway



# Cancer

## Registry of Norway

INSTITUTE OF POPULATION-  
BASED CANCER RESEARCH



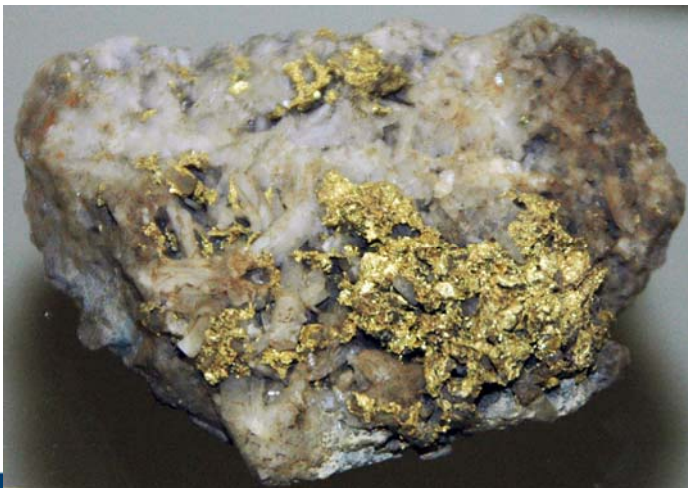
LLNL-PRES-753007

This work was performed under the auspices of the U.S. Department of Energy by Lawrence Livermore National Laboratory under Contract DE-AC52-07NA27344. Lawrence Livermore National Security, LLC

# Aim

... to make cancer prevention programs more flexible, scalable and sustainable by use of the personalized algorithms.

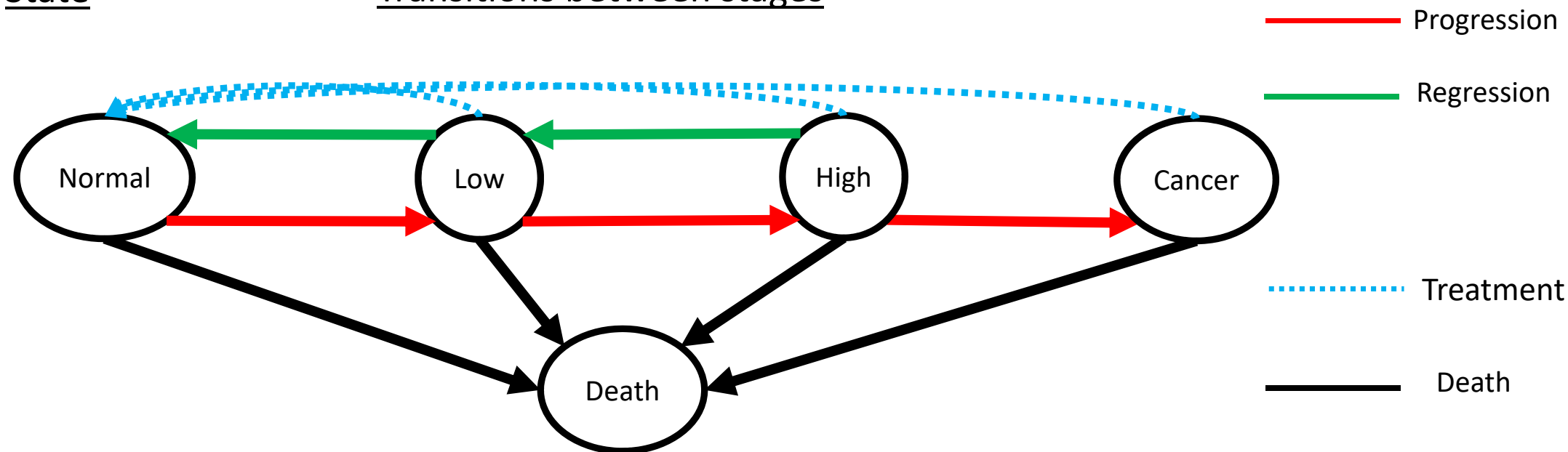
If proven effective, our project will introduce a paradigm shift from mass-screening to a personalized prevention.



# Hidden Markov Model - joint probability model

State

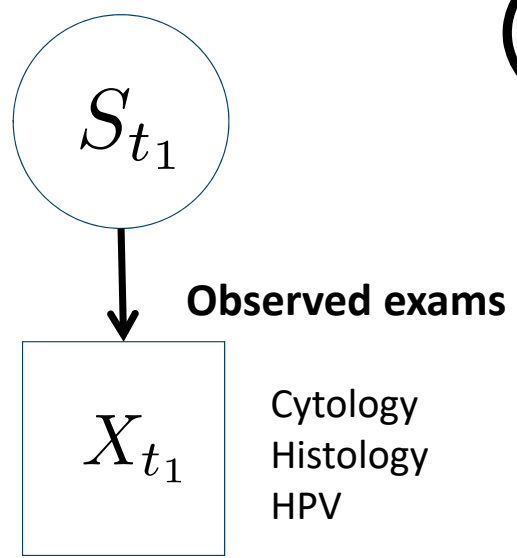
Transitions between stages



Model Screenings as Noisy Observations with Possible Misclassification

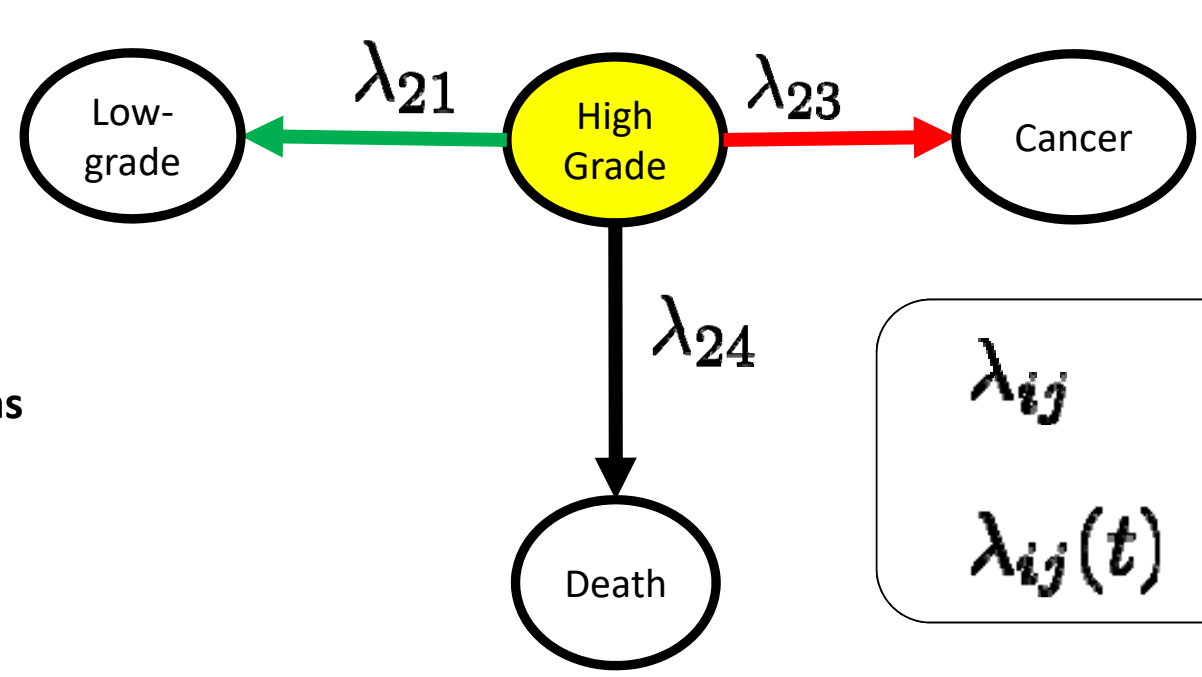
# Continuous-Time Markov Process for Cervical Cancer

disease state  
always **unobserved**



- States
- 1 Low Grade
  - 2 High Grade
  - 3 Cancer
  - 4 Death

- Transitions
- Progression
  - Regression
  - Death



Transition intensities determine how likely it is to transfer out of a current stage and into another.

$\lambda_{ij}$ *Age-ignorant*

$\lambda_{ij}(t)$ *Age-dependent*

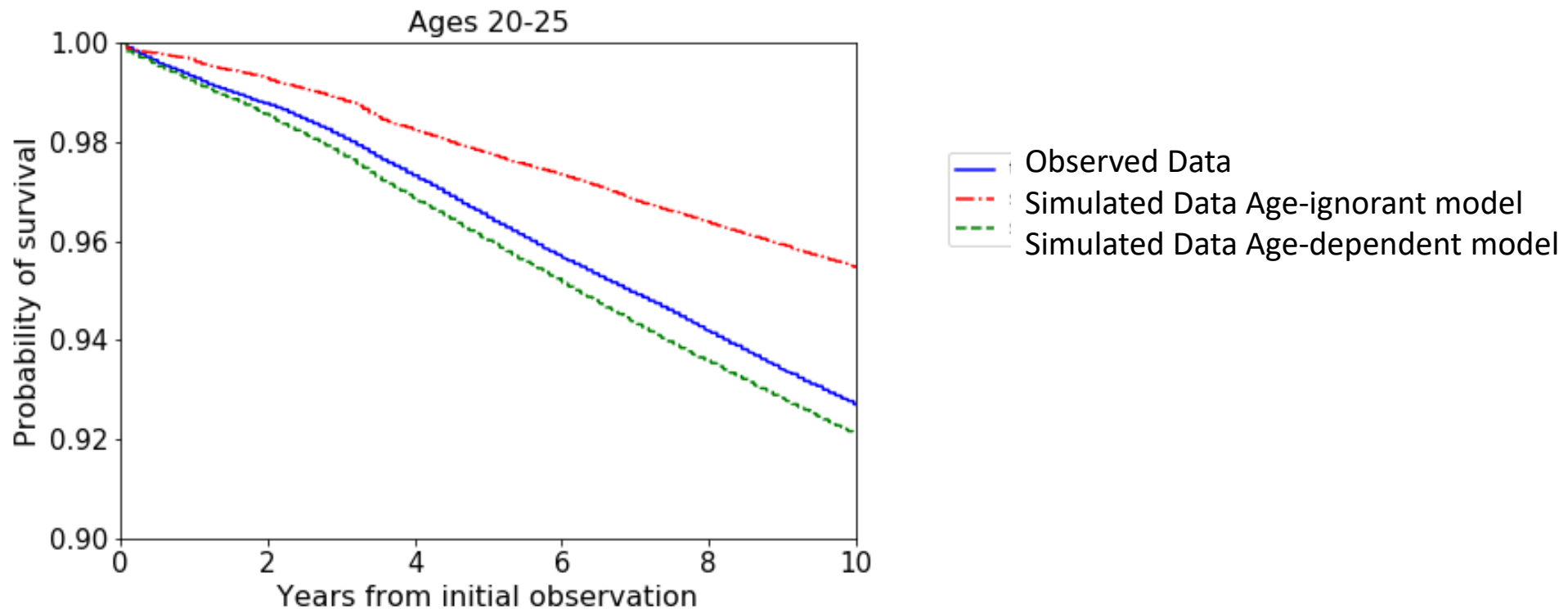
# HPC and Data Intensive Resources at LLNL

- **Catalyst, Quartz (Intel Ivy Bridge)**
  - 7,776 cores across 324 nodes with 24 cores per node; 96,768 cores across 2,634 nodes with 36 cores per node
- **Inference Algorithm**
  - Implemented a modified EM algorithm in parallel to compute Maximum Likelihood Estimates (MLEs) of all model parameters.
- **Simulations and Model Fit**
  - Simulated synthetic population data from learned models  
*Age-ignorant*  
*age-dependent model*
  - Compute Kaplan-Meier estimators for different outcomes comparatively for observed and synthetic data to assess goodness-of-fit.






# Simulated Data vs Observed Data transition from normal to high-risk



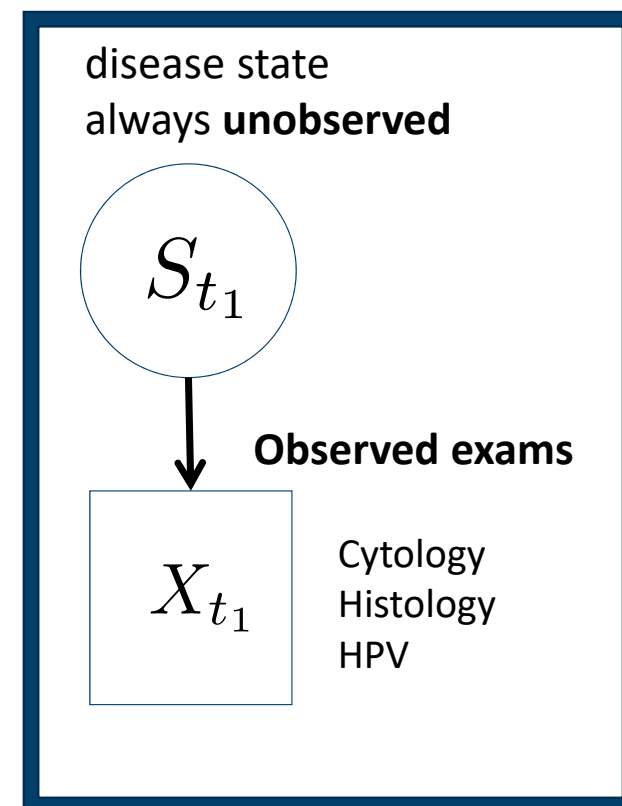
# Learning likelihoods

- Probability of cytology/HPV result conditioned on hidden state

|                 | <u>Test result</u>     |                       |
|-----------------|------------------------|-----------------------|
|                 | <u>normal cytology</u> | <u>hrHPV positive</u> |
| Normal state    | 99%                    | 5.7%                  |
| Low-risk state  | ~0%                    | 48.9%                 |
| High-risk state | 12%                    | 92.4%                 |
| Cancer          | 2.5%                   | 98.8%                 |



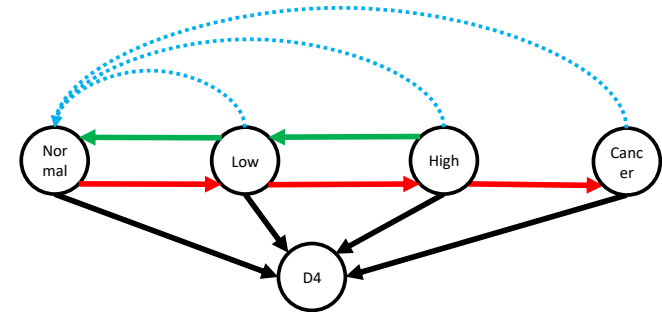
HPV test is more sensitive



# Learning likelihoods for transmission intensities

- Progression

Low-risk → High-risk 0.4% to 4.2%



# Learning likelihoods for transmission intensities

- Progression

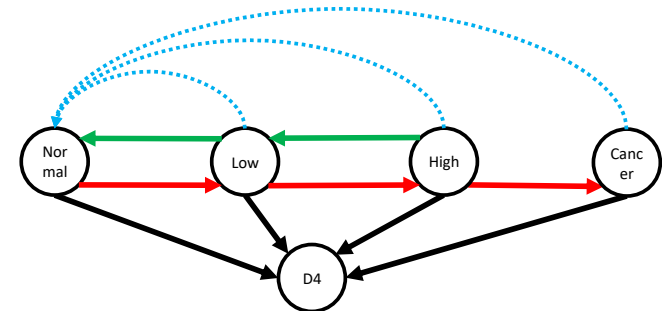
| Age group | 20-24 | 35-39 | 60+ |
|-----------|-------|-------|-----|
|-----------|-------|-------|-----|

Low-risk → High-risk 0.4% to 4.2%

2.5%

3.5%

3.4%



# Learning likelihoods for transmission intensities

## Regression

Age group

20-24

35-39

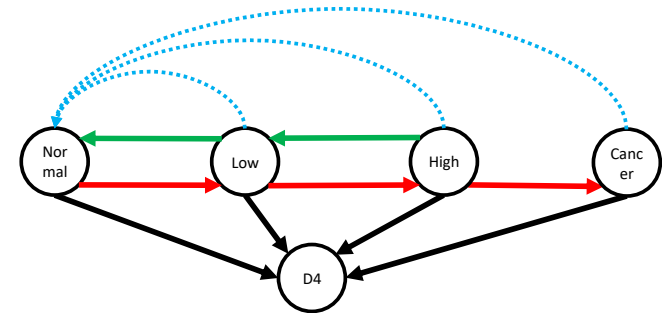
60+

High-risk → Low-risk 5% to 22%

10.9%

6.7%

5.8%



# WORK IN PROGRESS

# Predict Future Disease States

- Compute probabilities for future disease states based on a patient's past test results.

$$P(S_t^{\text{new}} | \mathbf{X}_{t_1:t_k}^{\text{new}}), \quad t > t_k$$

*When should a patient be tested next?*

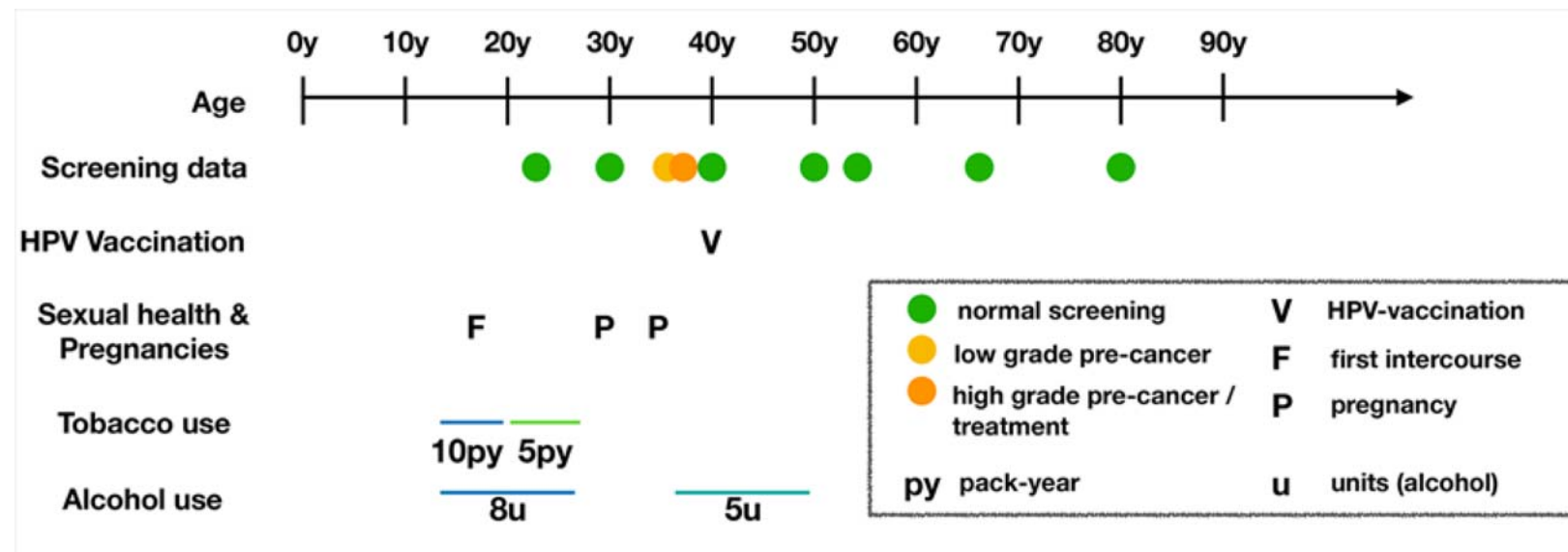
## ***Idea***

*Given a learned model, choose the minimum time interval that results in a positive diagnosis based on a Bayes optimal decision rule.*



# Increasing complexity: multi-modality datasets

- Detailed information on HPV types impacting the disease differently
- HPV vaccination status: age at vaccination, vaccine type
- Other possible life-style determinants (biomarkers)



# 2019 - expanding the group with PhD and Post-Doc



Lawrence Livermore National Laboratory



The Research Council  
of Norway

Data-driven Framework for Personalised Cancer Screening



EEA Financial Mechanism 2014-2021 Baltic Research  
Programme

simulamet



Karolinska  
Institutet

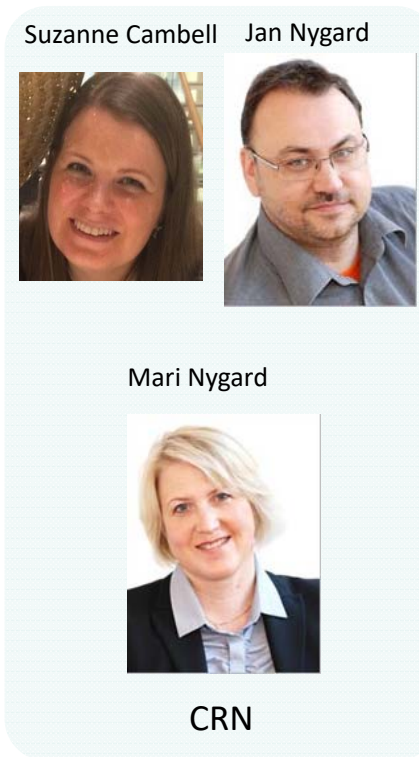
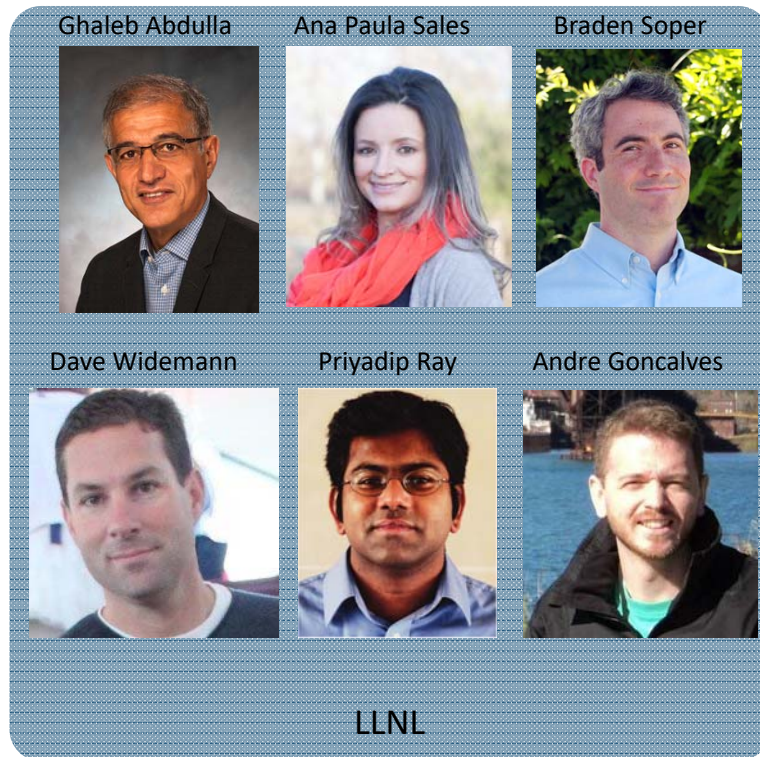


UNIVERSITY  
OF TARTU



Lawrence Livermore National Laboratory

# Our Team



- Supported

- Giske Ursin (CRN)
- Jim Brase (LLNL)
- Rob Sharpe (LLNL)
- Jason Paragas (LLNL)



## Collaborators

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Mari Nygard, Cancer Registry of Norway

Jan Nygard, Cancer Registry of Norway

Hang Deng, Rutgers University

