Real-world data at scale: How machine learning can enable learning from all patients

Discussion Leader: Lotte Steuten, PhD

Discussants:
Maarten Ijzerman, PhD
Corey Benedum, PhD
Natalia Kunst, PhD

Panelists



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Moderator
Deputy Chief Executive,
Office of Health Economics



Maarten Ijzerman, PhD

Discussant
Dean, Professor,
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Erasmus School of Health Policy
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DiscussantQuantitative Scientist,
Machine Learning
Flatiron Health



Natalia Kunst, PhD

Discussant
Associate Professor,
Department of Health
Management and Health
Economics,
University of Oslo

Agenda

- 1. The Challenge of Data Linkage to Analyse Complete Episodes of Care
- 2. Machine Learning Extraction and RWD Generation at Scale
- 3. Improving Medical Decision Making: Evidence And Uncertainty Consideration

Erasmus School of Health Policy & Management



The Challenge of Data Linkage to Analyse Complete Episodes of Care

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Ezafus,

Acknowledgement



Fanny Franchini, Karen Trapani, Sallie Pearson

"Predicting the Population Health Impact of New Cancer Treatments"

Medical Research Future Fund, Australia

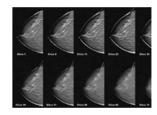


10 Mb



radiography

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digital mammography

100 Mb



CT scan

6,000 Mb



digital pathology

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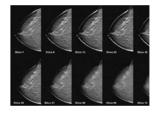
whole-genome sequencing

10 Mb



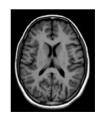
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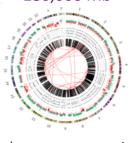
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whole-genome sequencing

US\$ 8 billion



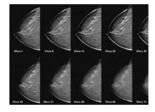
global market (2026) for healthcare data storage

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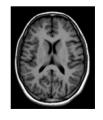
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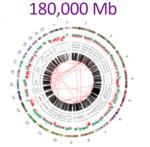


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digital pathology



whole-genome sequencing

US\$ 8 billion



global market (2026) for healthcare data storage

US\$ 35 billion



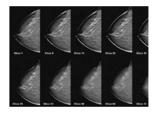
global market (2028) for electronic health records

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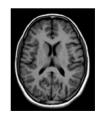
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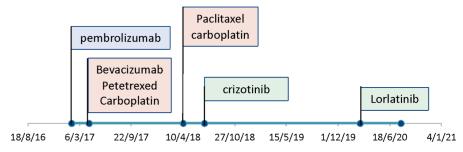
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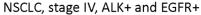
US\$ 56 billion

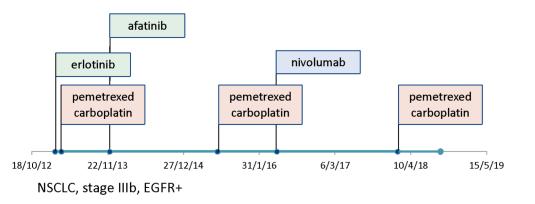


global market (2025) for checkpoint inhibitors

Listing new treatments: ALK+ NSCLC

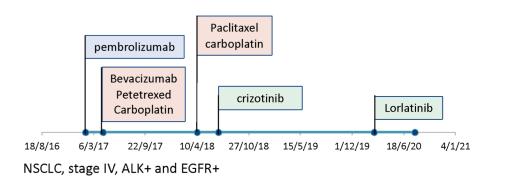


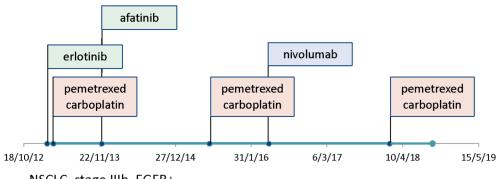


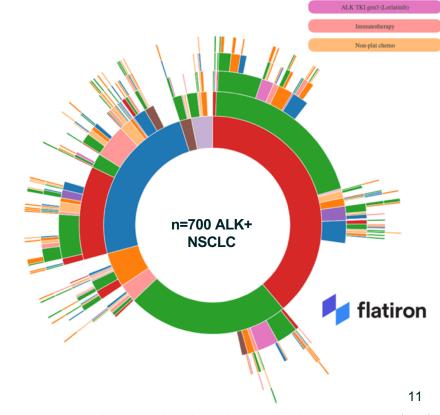


(zafus

Listing new treatments: ALK+ NSCLC







Plat chemo

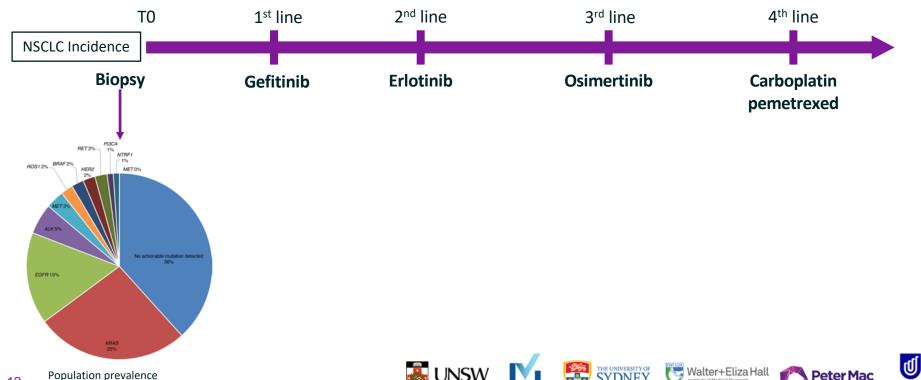
ALK TKI gen1 (Crizotinib)

ALK TKI gen2 (Alectinib/Brigatinib/Ceritinib)



eter MacCallum Cancer Foundation University of

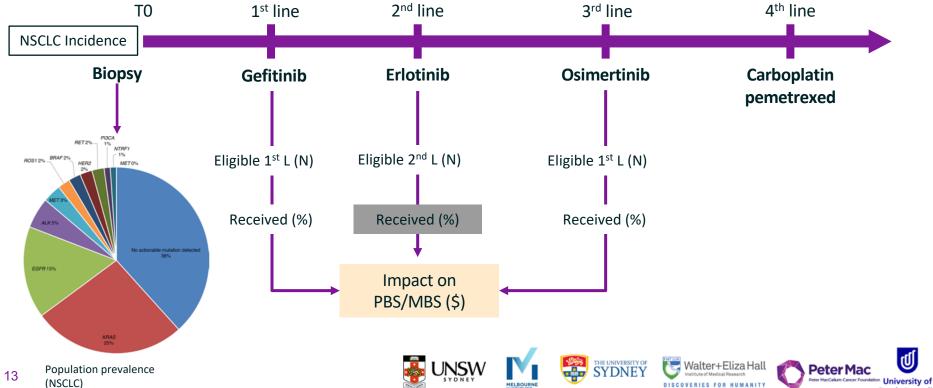
South Australia



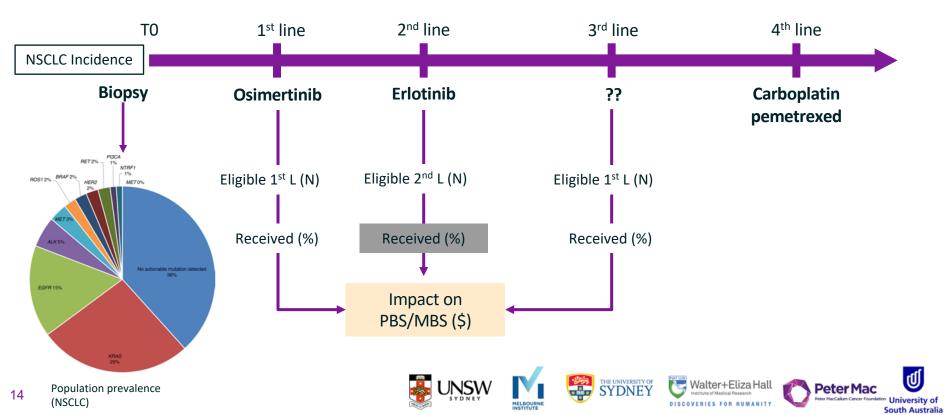
(NSCLC)



South Australia



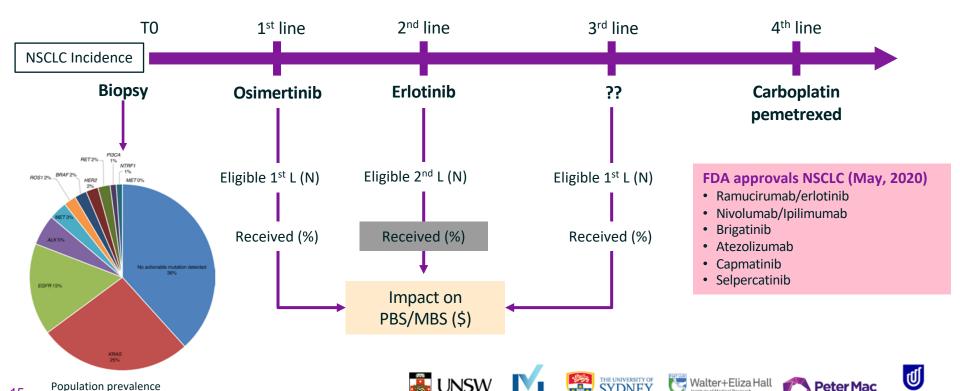






on University of

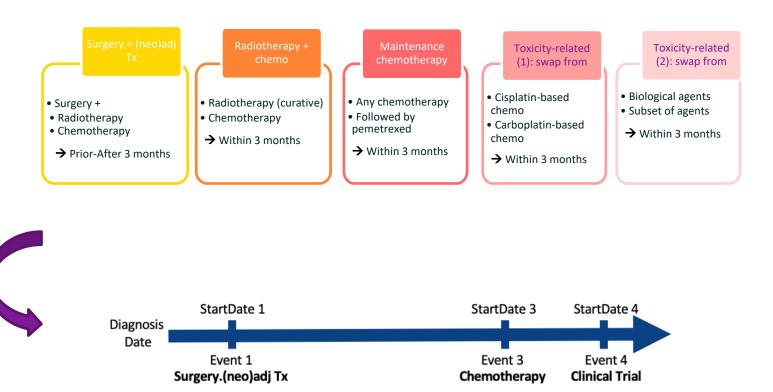
South Australia



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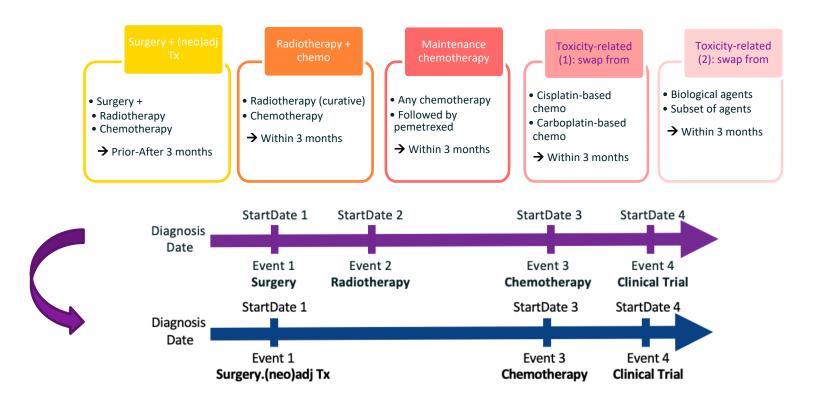
Method – Defining treatment lines

Using a curated registry, building a treatment pathway: rules based



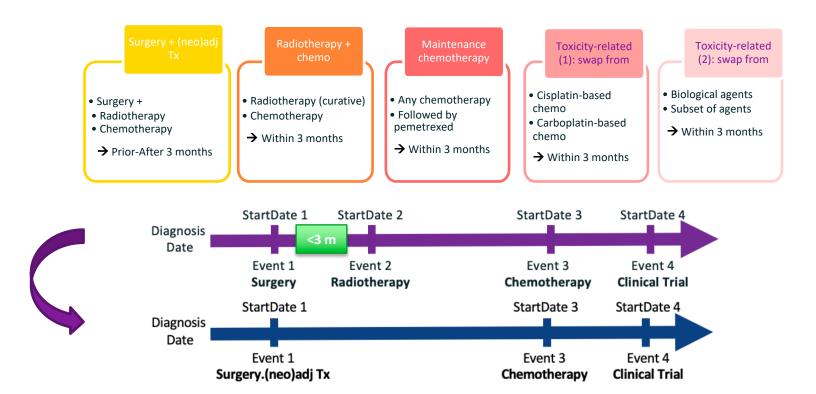
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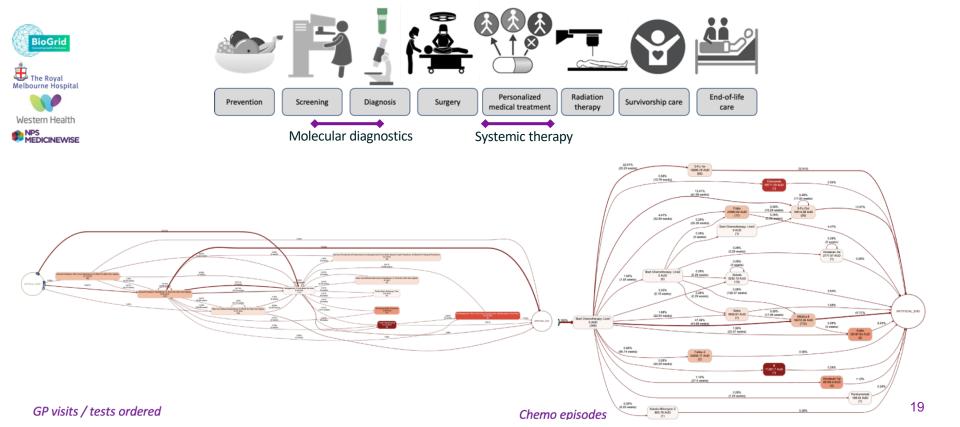


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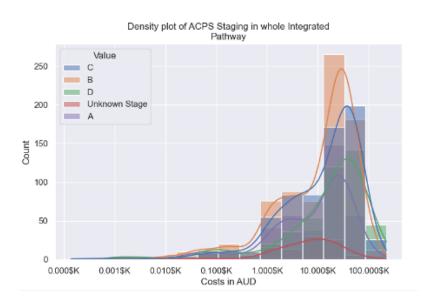
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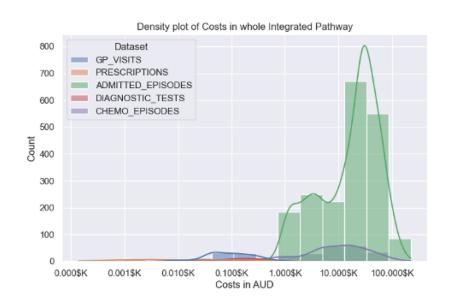


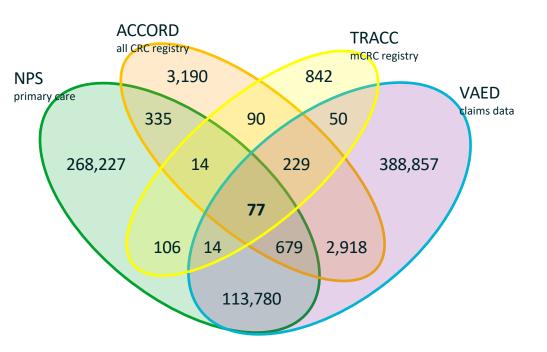
Analysis of the whole episodes of care: data driven

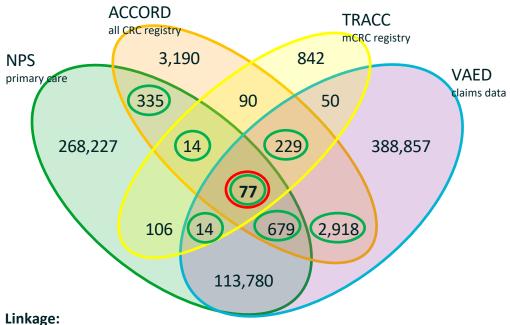


Cost analysis over entire care episodes



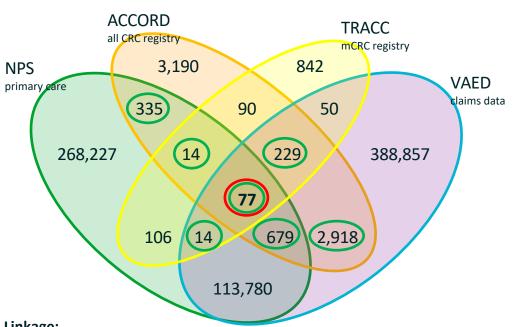






- 1. VAED-ACCORD-NPS-TRACC (N=77)
- 2. ACCORD-NPS (N=1,105)
- 3. VAED-ACCORD (n=3,903)
- 4. VAED-ACCORD-NPS (n=4,223)



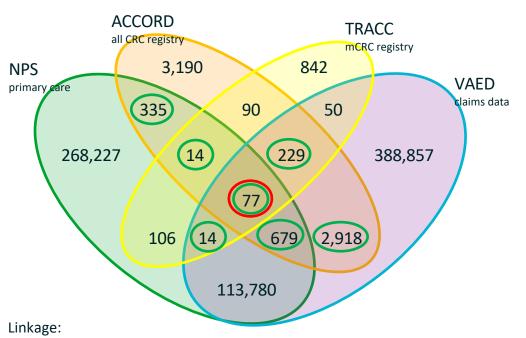


	Melbourne Hospital	Western Health
ACCORD Clinical registry	Royal Melbourne n=1,058	Western Health n=1,044
Coverage	June 2006- June 2017	June 2011- June 2018
% surgery <1 year	90% (n=954)	83% (n=865)
% chemo offered and accepted	30% (n=323) 90% (n=292)	28% (n=295) 90% (n=265)
% chemo received	27.5% (n=292)	25.4% (n=265)
Alive	64% (n=673)	75% (n=782)

Linkage:

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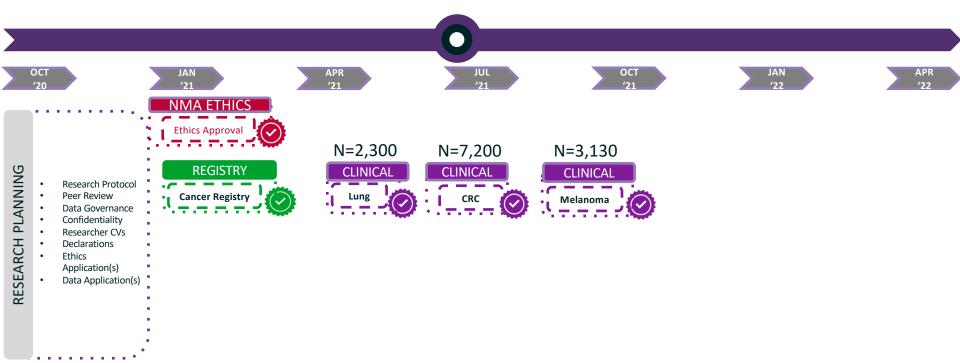
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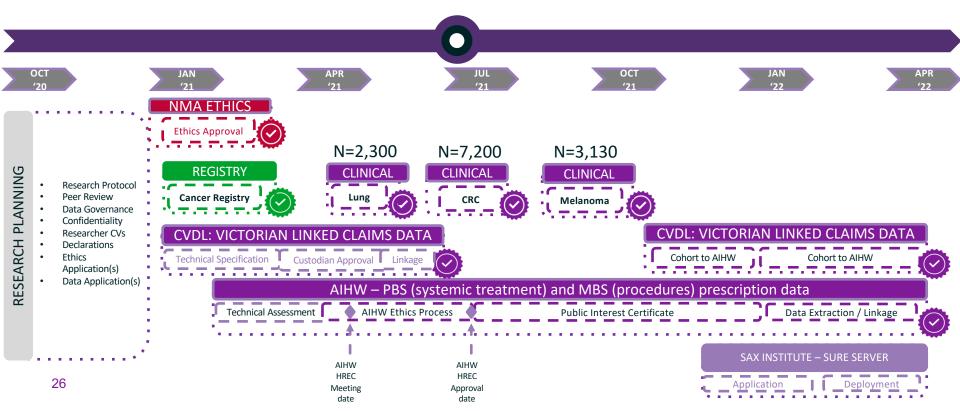
An (enduring) linked population level dataset





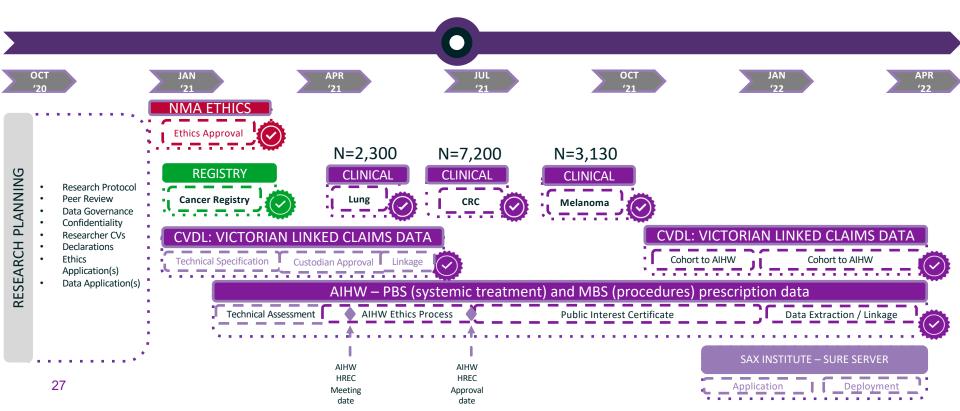
An (enduring) linked population level dataset





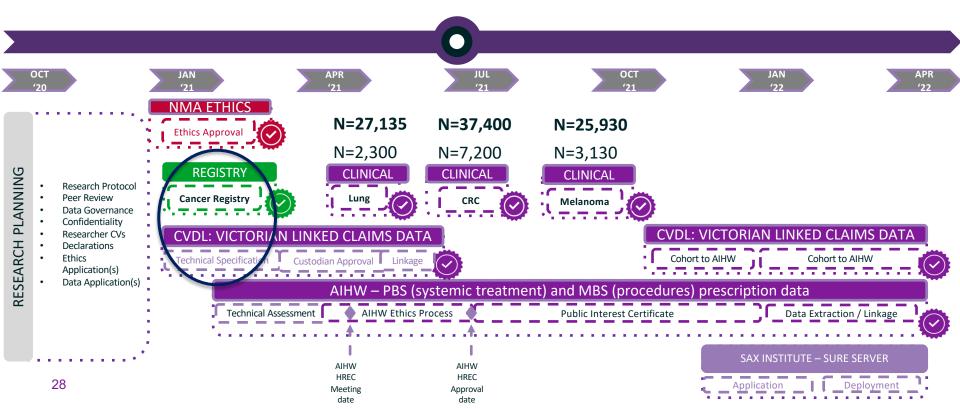
Challenges with data linkage II (data gaps)





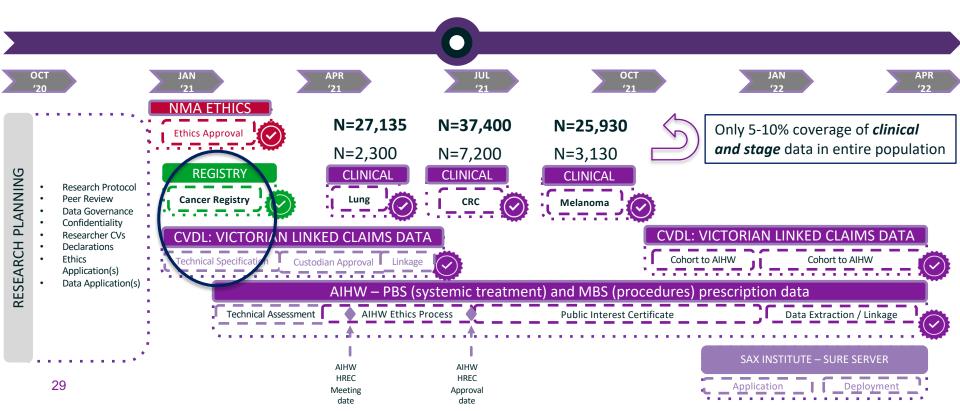
Challenges with data linkage II (data gaps)





Challenges with data linkage II (data gaps)





Machine learning for extracting and linking data

Mapping treatment episodes in relatively "clean" clinical datasets

Purpose: The analysis of specific treatments (sequences) and correlation with survival

Issue: Rule-based using clinical expertise is labor intensive

Use ML: Process mining is used to identify unique sequences of events

Combining various (clinical, claims, administrative) datasets through linkage

Purpose: The analysis of whole episodes of care, to investigate disparities in access to

care and outcomes or health service utilisation across different settings

Issue: "Data gaps", such as missing clinical information (stage, comorbidities) at

population level

Use ML: ML methods can be used to infer clinical information from other data (e.g.

pharmacy prescription data to infer toxicities)

How reliable is population level research if we only have complete data for a fraction of the patients? E.g. can we reliably estimate cost of care delivery from a subset of patients using process mining or ML methods?



If my dataset only has only 5% patients with complete clinical information (stage, comorbidities, ECOG, PROMs), there is insufficient statistical power to do any meaningful regression analyses

- Yes
- No



Inferring stage information or estimating immunotherapy related toxicities from drug dispensing data using machine learning will never be accepted by clinicians

- Yes
- No



We should not be using machine learning, instead we should solve data privacy and security issues to enable faster and better use of all existing data

- Yes
- No



Machine Learning Extraction and RWD Generation at Scale



Corey M. Benedum, PhD, MPH Flatiron Health

@DrCoreyBenedum

coreybenedum in





Disclaimers

Corey Benedum is an employee of Flatiron Health, an independent subsidiary of Roche Group. He holds stock ownership in Roche.

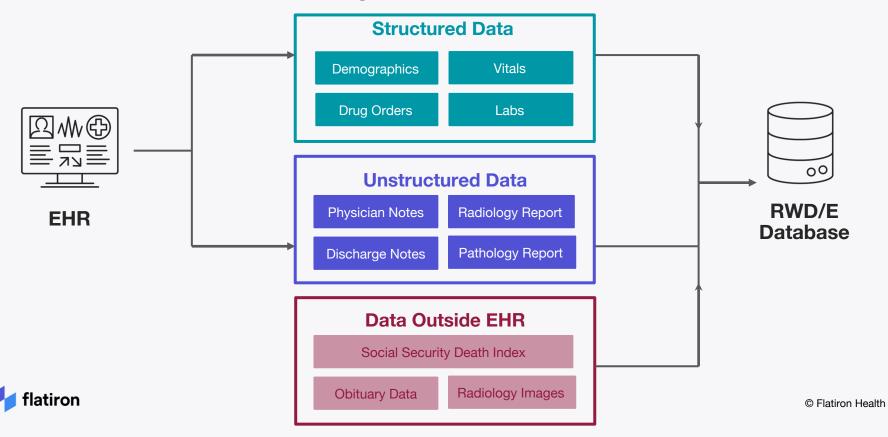
The value of realworld evidence

- Compare Treatment Effectiveness
- Understand Treatment Effectiveness
- HTA decision making
- Measure safety and effectiveness of off-label treatments
- Identify disparity in care

And much more...



Real-world evidence generation

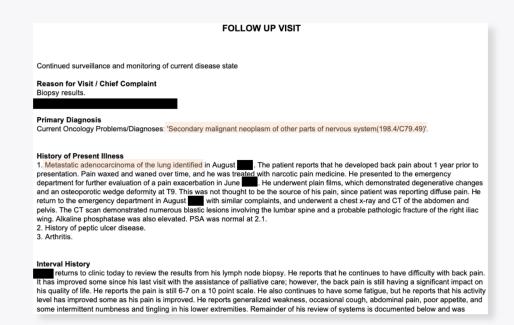


Challenge:

Critical data elements come from unstructured data

Several data elements critical for outcomes research are stored in unstructured data sources.

Abstracting this information is a **costly** and **resource intensive** task.





The promise of clinical ML for RWE

Real-world data and analytics organizations are looking to machine learning (ML) to efficiently extract data found in unstructured data at scale.













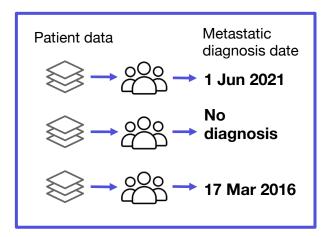
Keys to accurate information extraction with ML

- High quality labels that are designed with clinical expertise and are consistently / accurately collected.
- 2. Large volume of labels obtained from trained clinical experts performing chart review (abstraction)



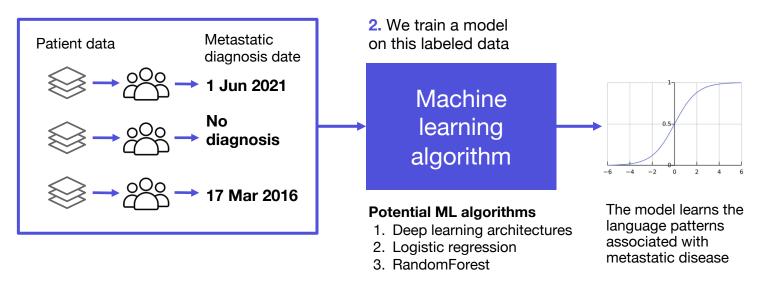


1. Abstractors label some of the patients



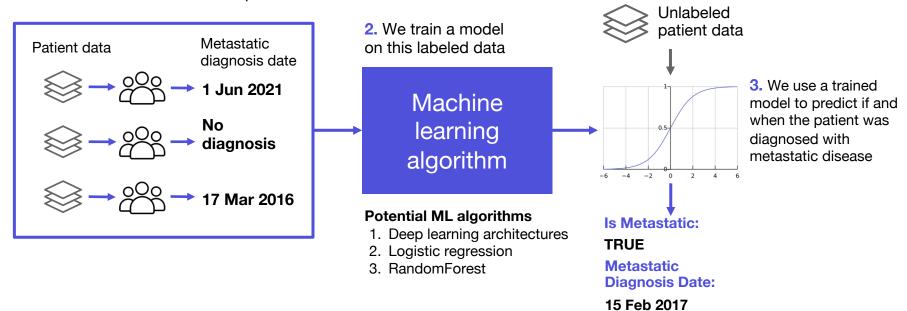


1. Abstractors label some of the patients





1. Abstractors label some of the patients





DETAILED LOOK INSIDE 2 1. Abstractors label some of the patients Unlabeled 2. We train a model patient data Metastatic Patient data on this labeled data diagnosis date 3. We use a trained Jun 2021 Machine model to predict if and when the patient was learning diagnosed with diagnosis algorithm metastatic disease **Potential ML algorithms** Is Metastatic: Deep learning architectures **TRUE** 2. Logistic regression Metastatic 3. RandomForest **Diagnosis Date:** 15 Feb 2017



Model Building Process:

- 1. construct list of relevant search terms
- 2. filter to sentences with informative terms.
- 3. assign dates to sentences
- 4. create model input from sentence-date pairs











- Metastatic
- Mets
- Recurrent
- Stage
- relapsed

...



Model Building Process:

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- Metastatic
- Mets
- Recurrent
- Stage
- relapsed

...

Clinic note: 20 Mar 2018

Name: John Doe DOB: 6/15/1952

History of Present Illness:

65 year old male w h/o Stage 4 lung adeno ca

(EGFR neg, ALK neg, ROS1 neg, BRAF neg, PDL1 high expression (60%) > cisplatin/alimta/pembro > 05/22/2017

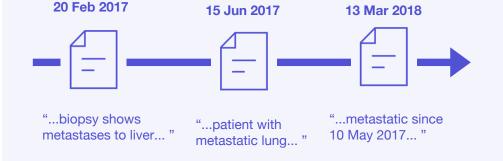
Completed XRT L iliac bone, 01/26/2018 Completed XRT Anterior Subcutaneous Chest Nodule

...



Model Building Process:

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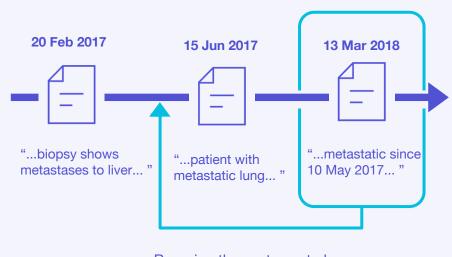




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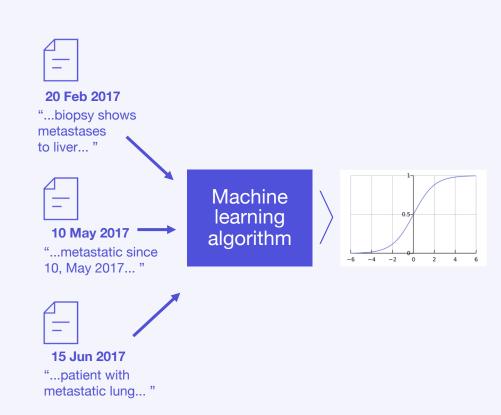


Reassign the sentence to have a timestamp matching the date referred to in the sentence

Model Building Process:

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Models must be generalizable to the target population

Potential Negative Outcome

Model is not generalizable leading to low performance and bias

Solutions

- Clearly define the target population
- Understand how training data are derived from this population to ensure representativeness





Models must be fair

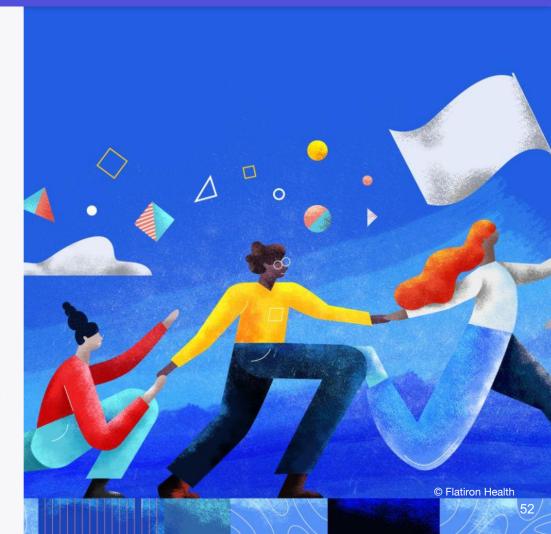
Potential Negative Outcome

Model performs poorly among certain subpopulations resulting in inadvertent exclusion of historically marginalized populations

Solutions

- Training data should balance diversity and representativeness of target population
- Model training and testing data should include enough examples from select subgroups





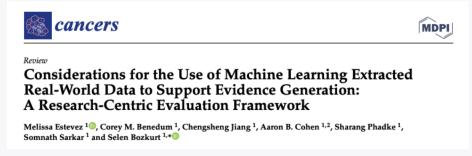
Models must be holistically and transparently evaluated

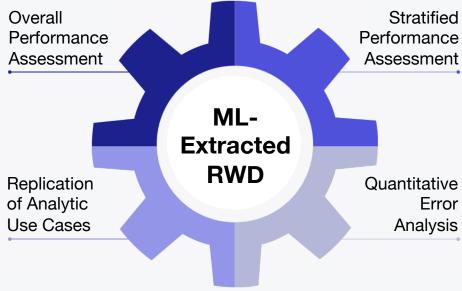
Potential Negative Outcome

Model errors may lead to biased study results and incorrect decisions / analytic conclusions.

Solutions

- Evaluate ML models and ML generated RWD
- Quantitative bias analyses and other bias correction methods





Presentation of Flatiron Health's replication of analytic use cases:

Nov 8, 15:00: Sondhi et al. Can ML-Extracted Variables Reproduce Real World Comparative Effectiveness Results From Expert-Abstracted Data? A Case Study in Metastatic Non-Small Cell Lung Cancer Treatment (**Poster RWD112**)

Nov 9, 10:00: Benedum et al. Machine Learning-Accelerated Outcomes Research: A Real-World Case Study of Biomarker-Associated Overall Survival in Oncology (Session 314: Applications of Machine Learning and Artificial Intelligence in Real-World Studies)

Poll Question

What would be the main barrier to adoption if you had access to ML-extracted data for HEOR?

- 1. Concerns of data quality
- 2. Explainability / interpretability of models
- 3. Lack of formal regulatory guidance
- 4. Other barriers
- 5. I am already an ML-extracted data user

Thank you

Additional Collaborators: Blythe Adamson, Aaron B. Cohen, Melissa Estevez, Erin Fidyk, Sheila Nemeth

Corey Benedum

Quantitative Scientist

Machine Learning

Flatiron Health

@DrCoreyBenedum



coreybenedum in

IMPROVING MEDICAL DECISION MAKING: EVIDENCE AND UNCERTAINTY CONSIDERATIONS

Natalia Kunst, PhD

University of Oslo, Norway

Yale University Schools of Public Health and Medicine, USA

The Center for Healthcare Research in Pediatrics (CHeRP), Harvard Medical School, USA



Yale School of Public Health









Natalia.kunst@medisin.uio.no



Medical decision making



Medical decision making

Prognoses about future outcomes





Medical decision making

Prognoses about future outcomes







Medical decision making

Imperfect evidence



Prognoses about future outcomes





Trade-offs

Medical decision making

Imperfect evidence







Prognoses about future outcomes





Trade-offs

Medical decision making

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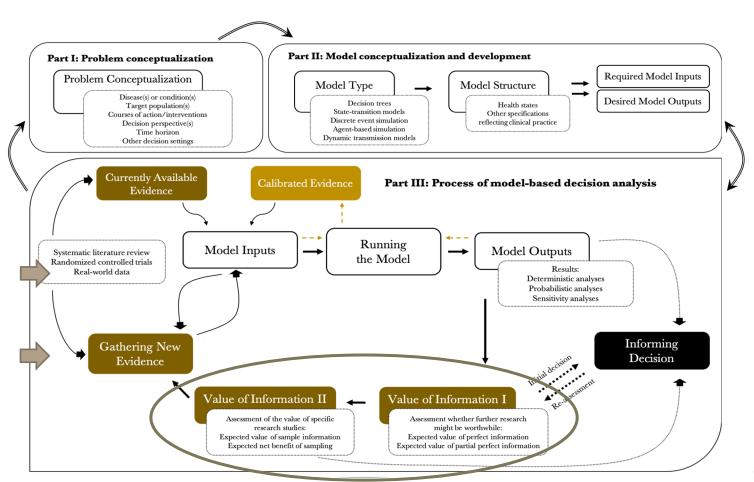








ITERATIVE DECISION-MAKING FRAMEWORK IN HEALTH AND MEDICINE



VALUE OF INFORMATION

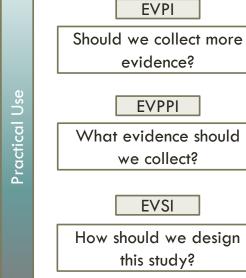
- Uncertainty in every decision
- There's a probability of making the wrong decision
- What are the consequences of making the wrong decision?
 - Costs?
 - Forgone benefits?
- Size of population being affected by the decision

VALUE OF INFORMATION

- How likely we are making the wrong decision and how bad it is to make the wrong decision
 - Uncertainty regarding our model parameters driven by the limited amount of information
- Cost of uncertainty (i.e., expected loss based on current information)
- There is an opportunity cost in the sense that we expect to have made a better decision had we had additional information/greater certainty
- Expected benefit of research
 - How valuable it is to collect additional evidence that enables us to reduce our uncertainty about the parameters

EVPI Expected Value of **Perfect Information EVPPI** Expected Value of Partial **Perfect Information EVSI** Expected Value of Sample Information

Computationa



Computational ease

EVPI Expected Value of **Perfect Information EVPPI** Expected Value of Partial **Perfect Information EVSI** Expected Value of Sample Information





Practical Use



Should we collect more evidence?

EVPPI

What evidence should we collect?

EVSI

How should we design this study?

Computational ease



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Computational ea





Practical Use

EVPI

Should we collect more evidence?

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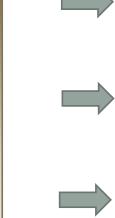
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Computational ease

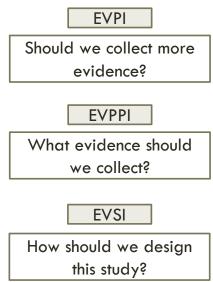
Practical Use

EVPI Expected Value of **Perfect Information EVPPI** Expected Value of Partial **Perfect Information EVSI** Expected Value of Sample Information





Practical Use



PARAMETER UNCERTAINTY AND POTENTIAL BIAS

- Populating the model inputs with appropriate and relevant evidence is necessary to ensure model credibility
- Sometimes the existing evidence may be insufficient to inform some of the relevant model inputs, thereby reducing or inhibiting the model's usefulness
- As indicated by the ISPOR VOI Task Force, when the risk of bias or appropriate technique for data analysis is unclear, the existing guidelines to aid characterization of uncertainty about methodological choice should be followed. These include:
 - Bilcke J. et al. Accounting for Methodological, Structural, and Parameter Uncertainty in Decision-Analytic Models: A Practical Guide. Med Decis Making. 2011; 31: 675-692
 - Jackson C.H. et al. Structural and parameter uncertainty in Bayesian cost-effectiveness models. J R Stat Soc Ser C Appl Stat. 2010; 59: 233-253

CANCER EPIDEMIOLOGY, BIOMARKERS & PREVENTION | RESEARCH ARTICLE

Estimating Population-Based Recurrence Rates of Colorectal Cancer over Time in the United States



Natalia Kunst 1,2,3,4 , Fernando Alarid-Escudero 5 , Eline Aas 1 , Veerle M.H. Coupé 3 , Deborah Schrag 6 , and Karen M. Kuntz 7

ABSTRACT

Background: Population-based metastatic recurrence rates for patients diagnosed with nonmetastatic colorectal cancer cannot be estimated directly from population-based cancer registries because recurrence information is not reported. We derived populationbased colorectal cancer recurrence rates using disease-specific survival data based on our understanding of the colorectal cancer recurrence-death process.

Methods: We used a statistical continuous-time multistate survival model to derive population-based annual colorectal cancer recurrence rates from 6 months to 10 years after colorectal cancer diagnosis using relative survival data from the Surveillance, Epidemiology, and End Results Program. The model was based on the assumption that, after 6 months of diagnosis, all colorectal cancer-related deaths occur only in patients who experience a metastatic recurrence first, and that the annual colorectal cancer-specific death rate among patients with recurrence was the same as in those

diagnosed with de novo metastatic disease. We allowed recurrence rates to vary by post-diagnosis time, age, stage, and location for two diagnostic time periods.

Results: In patients diagnosed in 1975–1984, annual recurrence rates 6 months to 5 years after diagnosis ranged from 0.054 to 0.060 in stage II colon cancer, 0.094 to 0.105 in stage II rectal cancer, and 0.146 to 0.177 in stage III colorectal cancer, depending on age. We found a statistically significant decrease in colorectal cancer recurrence among patients diagnosed in 1994–2003 compared with those diagnosed in 1975–1984 for 6 months to 5 years after diagnosis (hazard ratios between 0.43 and 0.70).

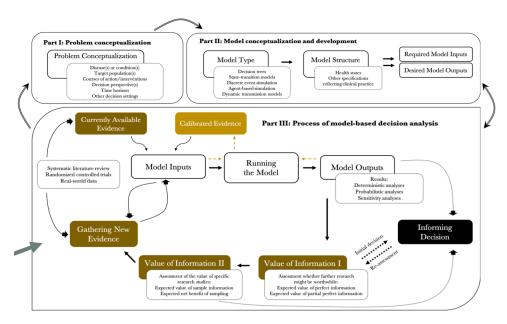
Conclusions: We derived population-based annual recurrence rates for up to 10 years after diagnosis using relative survival data.

Impact: Our estimates can be used in decision-analytic models to facilitate analyses of colorectal cancer interventions that are more generalizable.

Introduction

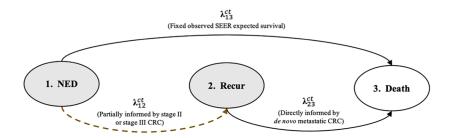
Improvements in colorectal cancer care have prolonged patient survival since 1975 (1, 2), but many patients still develop (metastatic) recurrence (3–5), from which patients can die from their disease. second cancer in that the former has the same type of cancer cells as the primary cancer, as opposed to the latter, which is unrelated to the primary cancer. The focus of our study was on distant recurrences.

Current evidence on recurrence rates comes from randomized controlled trials (RCT) in which disease-free survival is a common



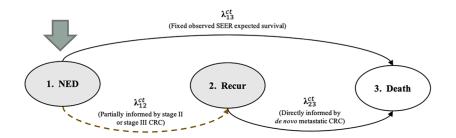
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Methods: Statistical multistate survival modeling techniques using data from the Surveillance, Epidemiology, and End Results (SEER) Program



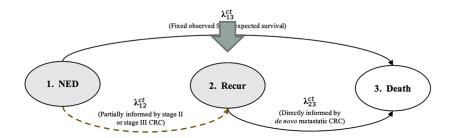
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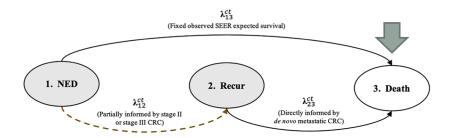


NED: No evidence of disease

Recur: Symptomatic distant recurrence

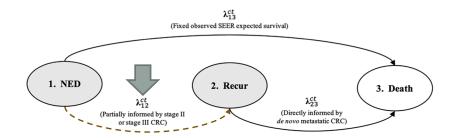
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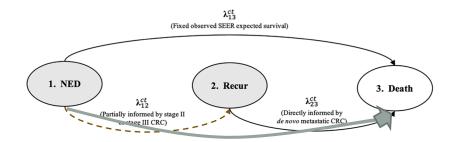
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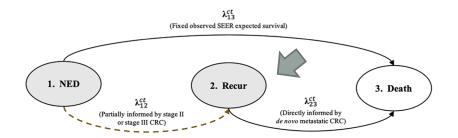
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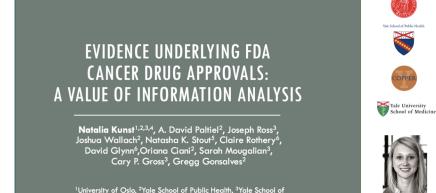
Results:

- The estimated **population-based** colorectal cancer recurrence rates were **higher** than the previously available trial-based estimates.
- The 10-year cumulative risk from **population-based** data vs. from trial-based estimates was:
 - Stage II colorectal cancer: 8.8-22.4% higher
 - Stage III colorectal cancer: 3.9-18.4% higher
- Potential bias in the effectiveness and cost-effectiveness evaluation

(FDA) REGULATORY DECISIONS

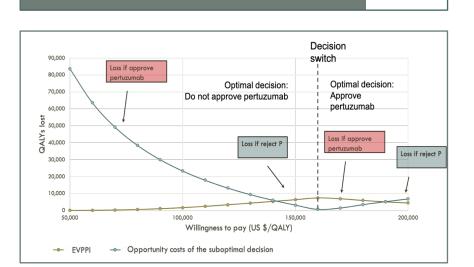
Accelerated approval decisions made with limited, preliminary data have:

- High uncertainty, and
- Significant downstream societal costs, if these decisions are made in error



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