

Building Cognitive Computing for Healthcare

ARCC Conference 2017

May 26th 2017

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Clinical Lead and Architect on Watson for Genomics



Agenda

- Building Cognitive Computing
 - Defining the Importance of Natural Language Processing (NLP)
- Applying Cognitive to a Complex Problem
- Surfacing Valuable Insights in Big Data

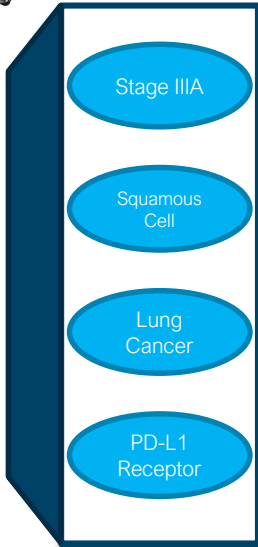
Moving Beyond Jeopardy: Bringing Cognitive Computing to Healthcare

Applying Cognitive Computing to Complex Problems

Define Attributes

Attribute
Extraction

Compare to Prior
Knowledge



?

What is the best treatment for a given patient?

Chief Complaint
The patient is a 67-year-old male who has been diagnosed with non-small cell squamous cell lung carcinoma (clinical stage T3, N2, M0 stage IIIA) with metastatic disease. The following history is provided for the patient's medical history.

History of Present Illness
The patient's medical history provided, the patient is a 67-year-old male with a 37 pack year smoking history. In 2018, he presented to his primary care physician with complaints of dry cough, chest pain, and a 10 lb weight loss over the last 6 months. He was initially treated with antibiotics and steroids, but symptoms did not improve. He was then referred to a pulmonologist where a CT scan of the chest was performed which showed a 4.5 cm x 3.5 cm peripheral enhancing nodule in the right lung middle lobe. He was treated with a course of antibiotics. His symptoms improved and he was treated with a second course of antibiotics. He was then referred to a pulmonologist where a PET scan was performed which showed a 4.5 cm x 3.5 cm enhancing nodule in the right lung middle lobe. The patient was then referred to a pulmonologist where a biopsy was performed which showed a 4.5 cm x 3.5 cm enhancing nodule in the right lung middle lobe. The patient was then referred to a pulmonologist where a biopsy was performed which showed a 4.5 cm x 3.5 cm enhancing nodule in the right lung middle lobe.

Data Set

Applying Cognitive Computing to Complex Problems

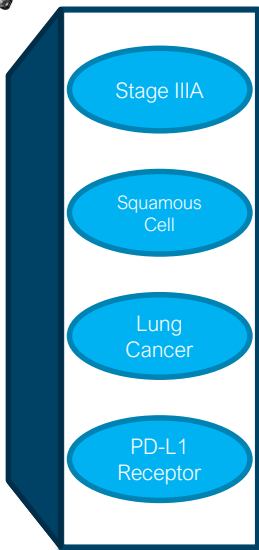
Define Attributes

Attribute Extraction

Compare to Prior Knowledge



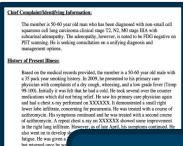
Clinical Expert



Natural Language Processing



What is the best treatment for a given patient?



Data Set

Chief Complaint/Identifying Information:

The member is 50-60 year old man who has been diagnosed with non-small cell squamous carcinoma clinical stage IIIA, adenopathy, histology consistent with squamous cell carcinoma on consultation on 11/11/11.

History of Present Illness:

Based on the medical records provided, the member is a 50-60 year old male with a 35 pack year smoking history. In 2010, he presented to his primary care physician with complaints of a dry cough, weight loss and grade fever (Temp 101.0 F). He felt that he had been ill for several over the counter cough syrup providing relief. He saw his primary care physician again in 2011 and was diagnosed with pneumonia. He was treated with a course of azithromycin. A repeat chest x-ray on XXXXXX. It demonstrated a small right lower lobe infiltrate concerning pneumonia. He was treated with a course of azithromycin. His symptoms continued and he was treated with a second course of azithromycin. A repeat chest x-ray on XXXXXX showed some improvement in the right lung infiltrate. However, as of late April, his symptoms continued. He also went on to develop shortness of breath, dyspnea on exertion, chest pain, and fatigue. He was given a 2 week course of a prednisone. His symptoms improved but returned once he was tapered off of the prednisone.

Applying Cognitive Computing to Complex Problems

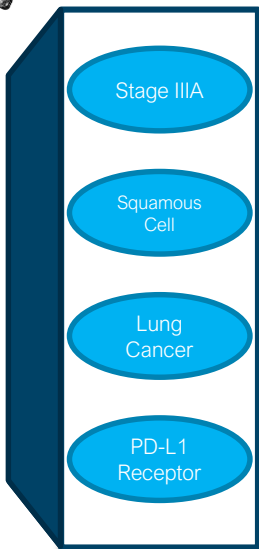
Define Attributes

Attribute Extraction

Compare to Prior Knowledge



Clinical Expert



Natural Language Processing

Chief Complaint/Identifying Information:

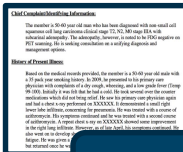
The member is 50-60 year old man who has been diagnosed with non-small cell squamous carcinoma clinical stage IIIA, with lymph node metastasis, adenopathy, histologically confirmed squamous cell carcinoma on consultation on 11/11/2014.

History of Present Illness:

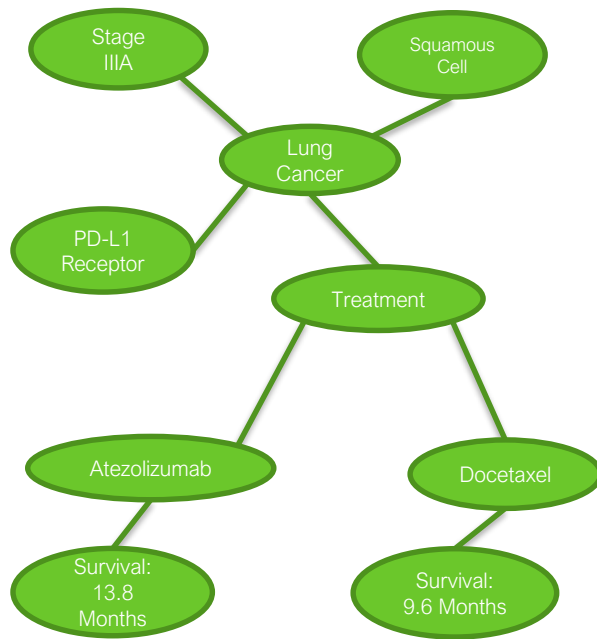
Based on the medical records provided, the member is a 50-60 year old male with a 35 pack year smoking history. In 2014, he presented to his primary care physician with complaints of a dry cough, chest pain, and low grade fever (Temp 100.0-100.4 F). He was treated with over the counter cough medicine which did not bring relief. He saw his primary care physician again in 2014 and was diagnosed with XXXXXX. It demonstrated a small right lower lobe consolidation concerning pneumonia. He was treated with a course of azithromycin. His symptoms continued and he was treated with a second course of azithromycin. A repeat chest x-ray on XXXXXX showed some improvement in the right lung infiltrate. However, as of late April, his symptoms continued. He also went on to develop shortness of breath, dyspnea on exertion, chest pain, and fatigue. He was given a 2 week course of a prednisone. His symptoms improved but returned once he was tapered off of the prednisone.



What is the best treatment for a given patient?



Data Set



Applying Cognitive Computing to Complex Problems

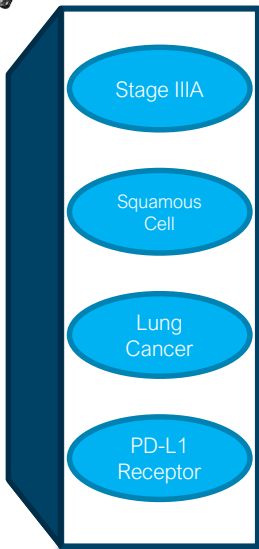
Define Attributes

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Clinical Expert



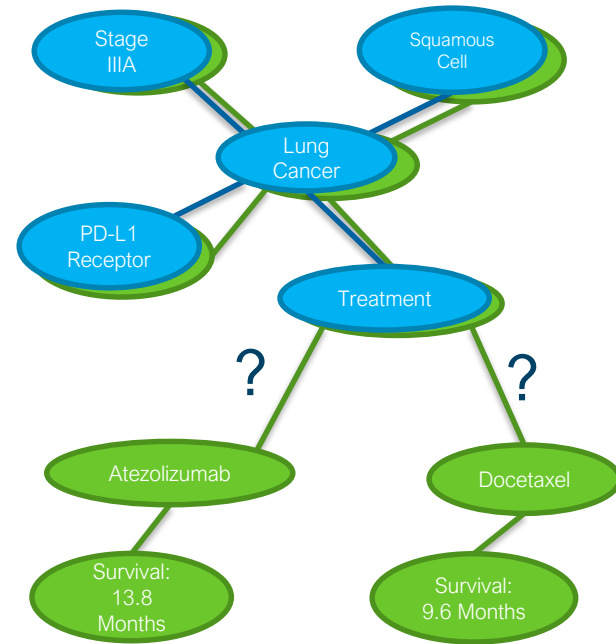
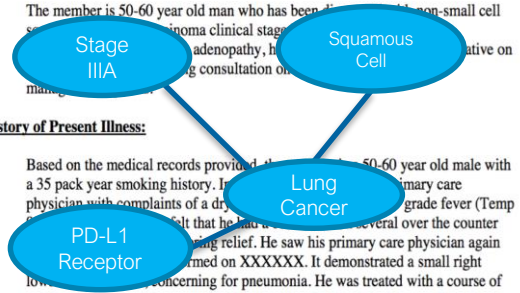
Natural Language Processing

Chief Complaint/Identifying Information:

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History of Present Illness:

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History of Present Illness:

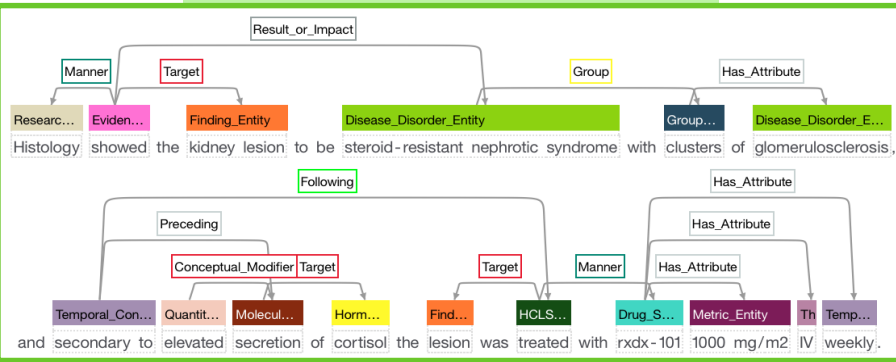
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Data Set

Differences in attribute extraction between Machine Learning and Rules-Based NLP

Machine Learning NLP

History showed the kidney lesion to be steroid-resistant nephrotic syndrome with clusters of glomerulosclerosis, and secondary to elevated secretion of cortisol the lesion was treated with rxdx-101 1000 mg/m2 IV weekly.



Disease	steroid resistant nephrotic syndrome
	<subtype> glomerulosclerosis
Therapy	rxdx-101

Rules-Based NLP

History showed the kidney lesion to be steroid-resistant nephrotic syndrome with clusters of glomerulosclerosis, and secondary to elevated secretion of cortisol the lesion was treated with rxdx-101 1000 mg/m2 IV weekly.

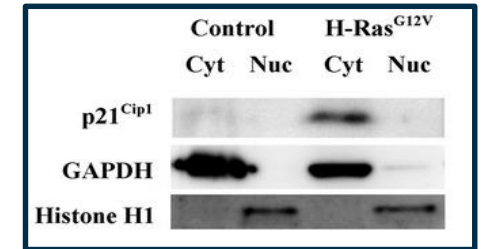
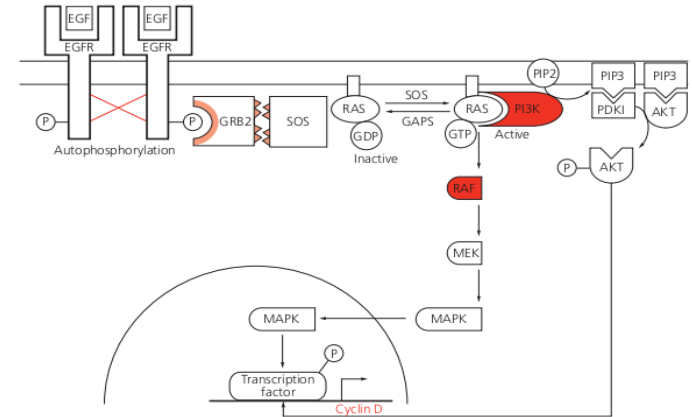
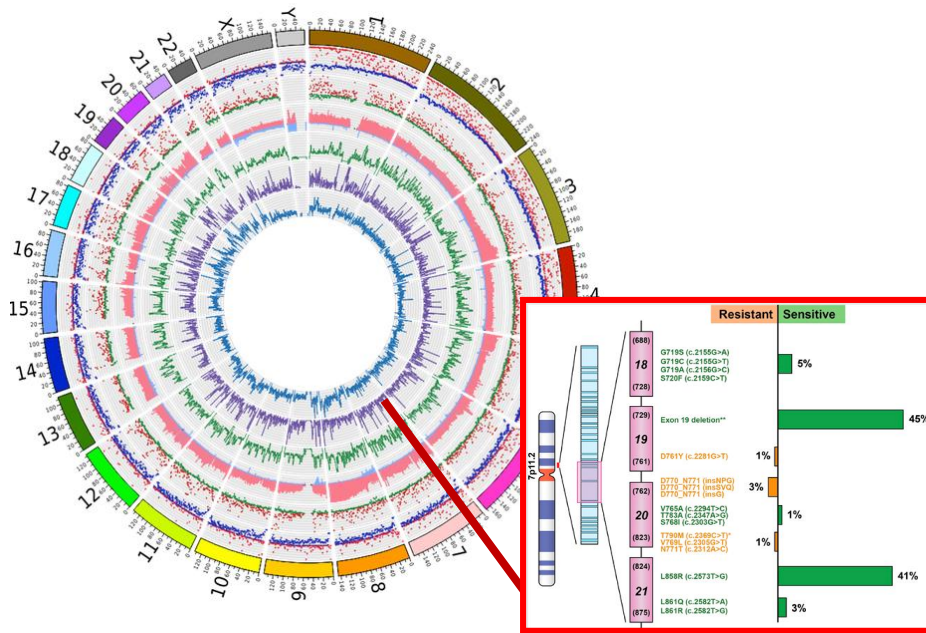
Rule: connect disease to therapy if within 10 words of each other



Disease	<conflict>	nephrotic syndrome
	<conflict>	glomerulosclerosis
Therapy	<incorrect>	cortisol

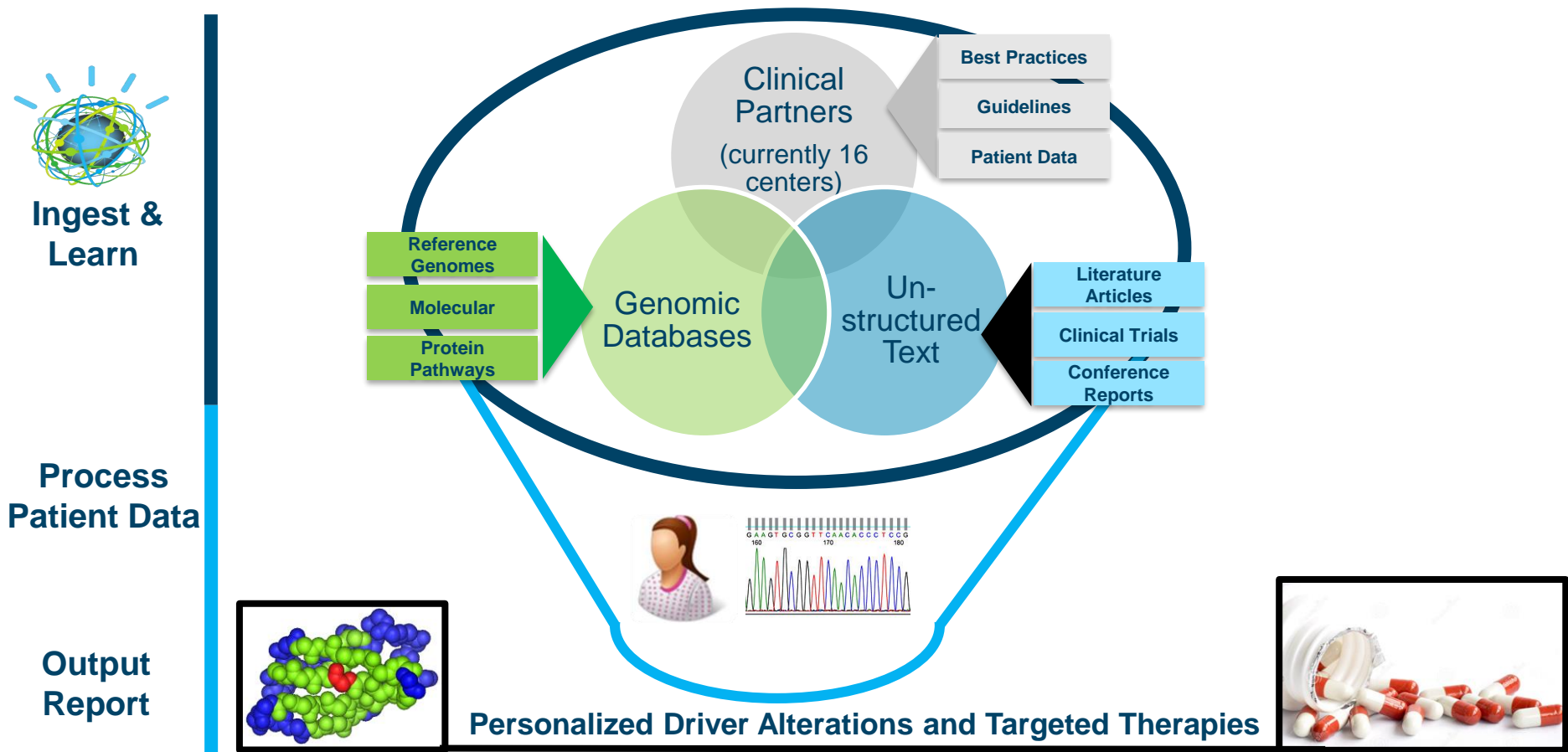
Applying Cognitive to Genomics

Impact on Medical Genomics



What's the best treatment given a tumor sequence?

Gene	Alteration	Alteration Type	Functional Significance	Annotation	Actionable Gene	Actionable Variant
PDGFRA	Y288C	SNV				
PTEN	K163*	Truncation				
FGFR2	K292M	SNV				

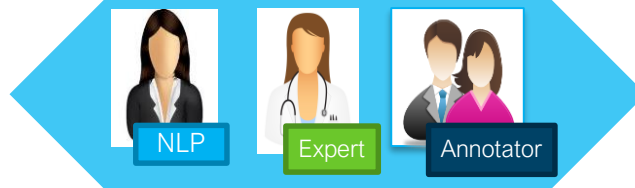




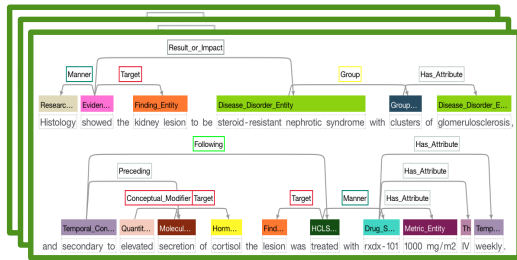
FAST Annotators working with Boston Children's Hospital physicians in March 2016

Our Approach to NLP

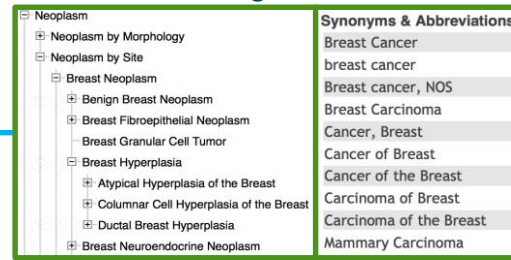
Focused Annotation Specialist Team (FAST)



Label Documents



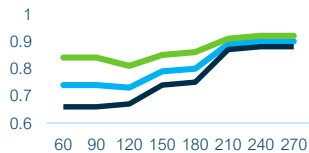
Standardize Ontologies and Dictionaries



Train Machine Learning



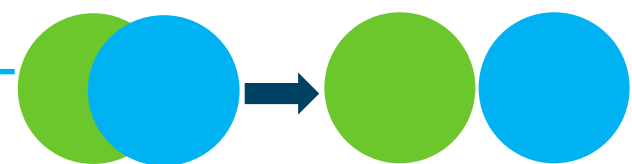
Measure Performance



Enable Integration & Reuse



Reduce Bias



Annotating mentions



ZOOM IN

REPLACE

CONCORDANCE

ATTRIBUTE VIEW

IN_PROGRESS

SAVE

CLOSE

Alpha...

14pt

1



Mention

24353160.txt

Epidermal growth factor receptor (EGFR) gene mutations (G719X, exon 19 deletions/insertions, L858R, and L861Q) predict favorable responses to EGFR tyrosine kinase inhibitors (TKIs) in advanced non-small cell lung cancer (NSCLC).

However, EGFR exon 20 insertion mutations (~10% of all EGFR mutations) are generally associated with insensitivity to available TKIs (gefitinib, erlotinib, and afatinib).

The basis of this primary resistance is poorly understood.

We studied a broad subset of exon 20 insertion mutations, comparing in vitro TKI sensitivity with responses to gefitinib and erlotinib in NSCLC patients, and found that most are resistant to EGFR TKIs.

The crystal structure of a representative TKI-insensitive mutant (D770_N771insNPG) reveals an unaltered adenosine triphosphate-binding pocket, and the inserted residues form a wedge at the end of the C helix that promotes the active kinase conformation.

Unlike EGFR-L858R, D770_N771insNPG activates EGFR without increasing its affinity for EGFR TKIs.

Unexpectedly, we find that EGFR-A763_Y764insFQEA is highly sensitive to EGFR TKIs in vitro, and patients whose NSCLCs harbor this mutation respond to erlotinib.

Analysis of the A763_Y764insFQEA mutant indicates that the inserted residues shift the register of the C helix in the N-terminal direction, altering the structure in the region that is also affected by the TKI-sensitive EGFR-L858R.

Entity

Mention

Type

SubType

Role

c	Cancer_Type
g	Gene/Protein
q	Location
r	Response
s	Substitution
t	Therapy
v	Variant_Type

Annotating relationships



ATTRIBUTE VIEW

IN_PROGRESS

SAVE

CLOSE

Alpha...

14pt

1



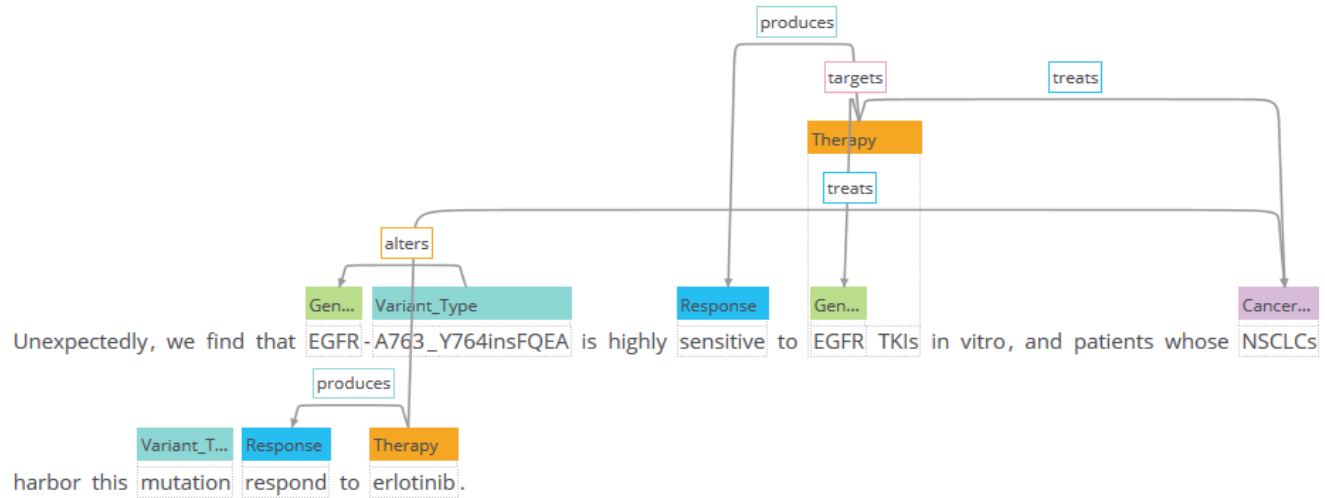
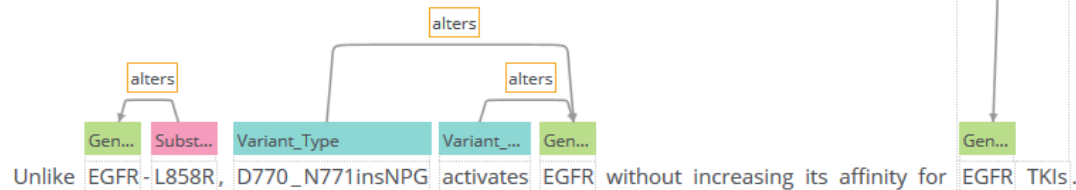
Mention



Relation



Coref



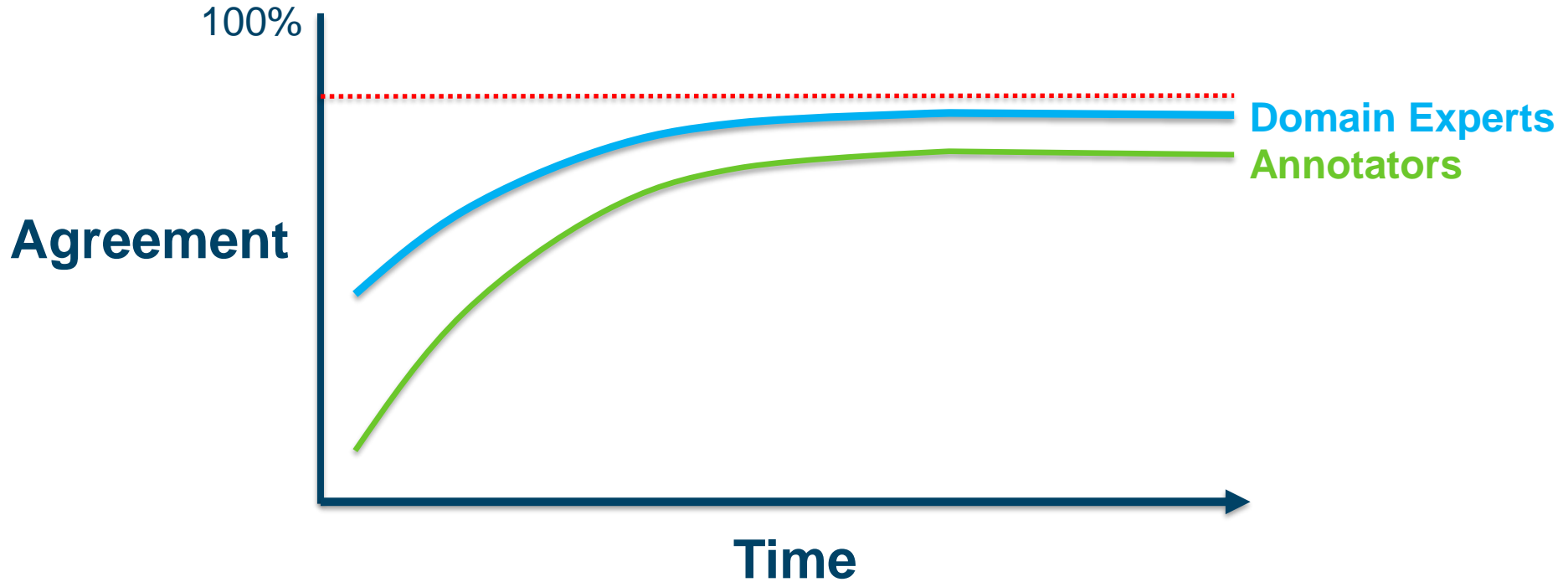
Relation Attributes

Type

- a alters
- d drives
- l located
- p produces
- h targets
- y treats

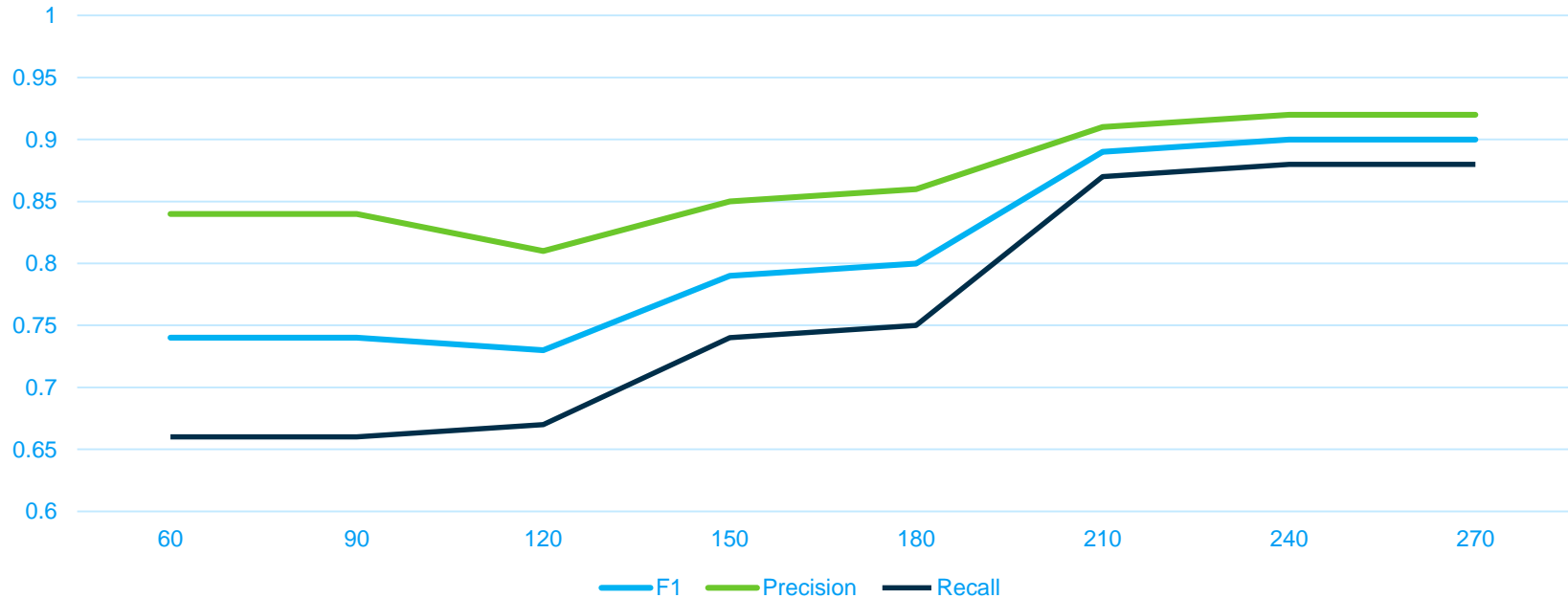
III

Ensuring Agreement: Measuring Kappa Scores



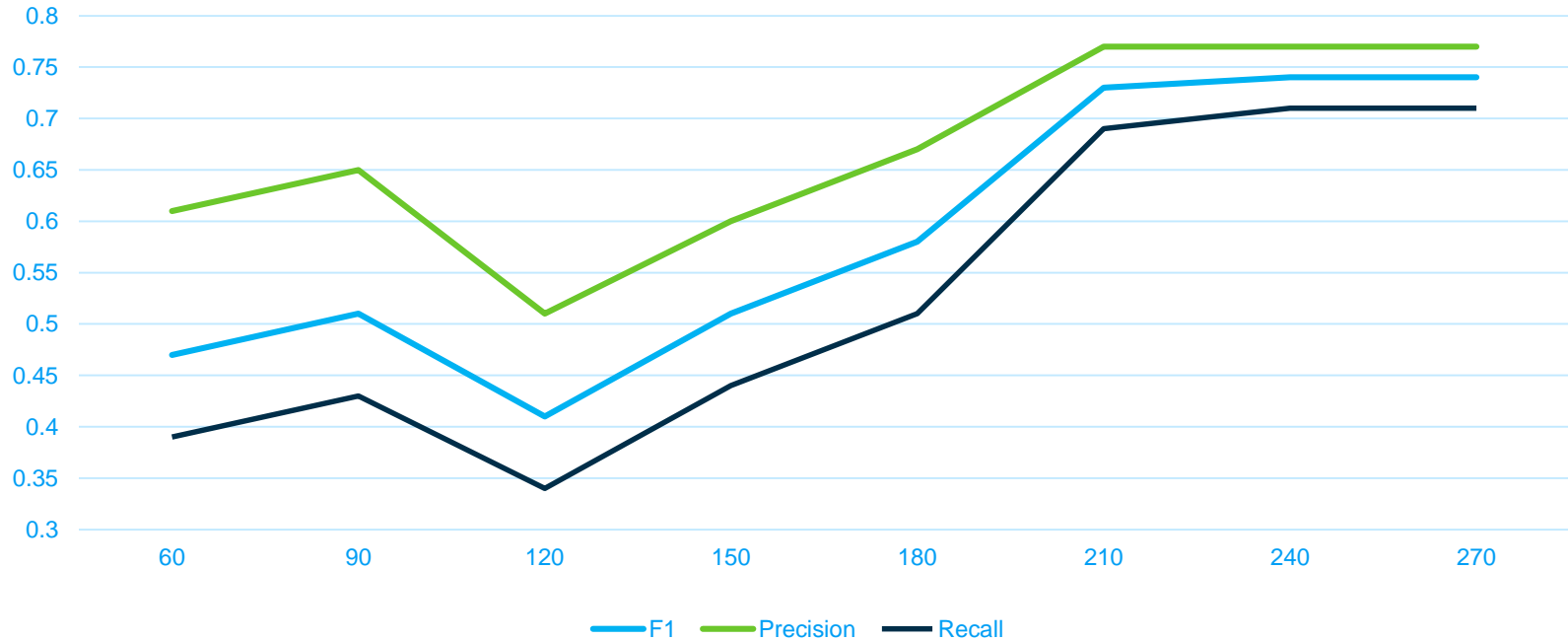
Concepts Detection: Compare to a standard and iterate

Model Performance for Labeling Concepts



Relationship Detection: Compare to a standard and iterate

Model Performance for Labeling Relationships



Maintaining Context Within and Across Documents

<text>Unexpectedly, we find that EGFR A763_Y764insFQEA is highly sensitive to EGFR TKIs in vitro, and patients whose NSCLCs harbor this mutation respond to erlotinib.</text> = "alters">

<relation rid="s6-r1" subtype="OTHER" type="alters">

<rel_entity_arg argnum="1" eid="E-39"/>

<rel_entity_arg argnum="2" eid="c1"/>

<relmentions>

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<rel_mention_arg argnum="1" mid="s6-m6">A763_Y764insFQEA</rel_mention_arg>

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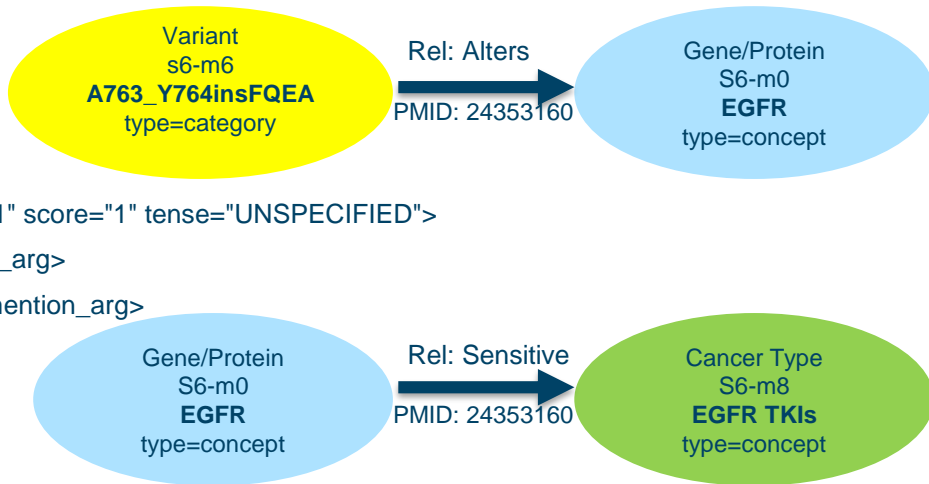
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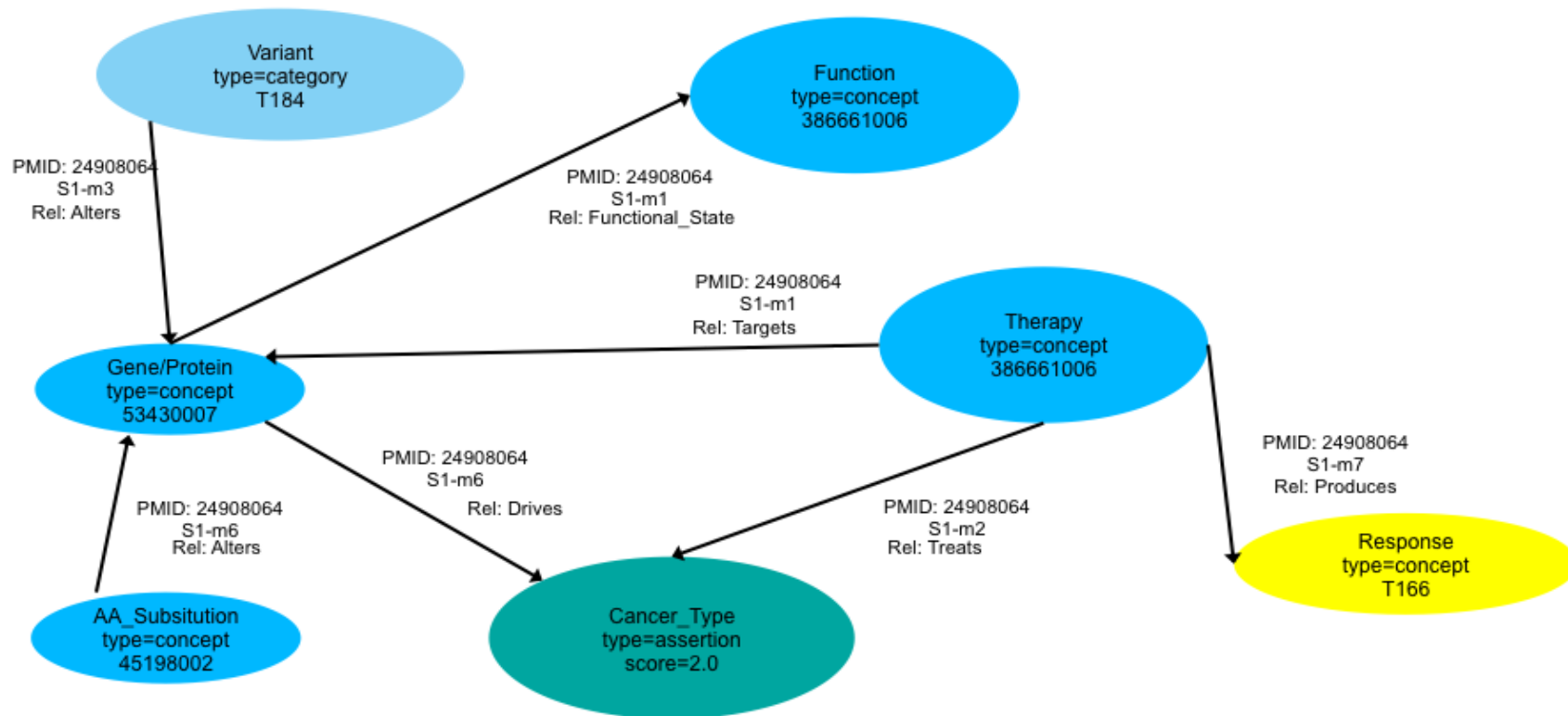
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<rel_mention_arg argnum="2" mid="s6-m8">EGFR TKIs</rel_mention_arg>



Clinical Knowledge: Example Graph



Clinical Knowledge: Extracted Data from Corpus of Documents

IBM Watson for Drug Discovery / Explore a Network

EGFR(G) x ERLOTINIB(Dr) x GEFITINIB(Dr) x AFATINIB(Dr) x CARCINOMA, NON SMALL C... x in Full Text Medical Journals Explore

Advanced Options Related Disease, drug, chemical 1904-2017 Biological Relationships: Non-PTM, Other, PTM

- Searched Entity
- Gene
- Disease
- Drug
- Chemical

Common Entities
No entities selected
5 to 641 documents

Up- & Downstream Entities
No direction specified

Gene Ontology
No ontology categories set

Reset all filters to default

Clinical Knowledge: Extracted Data from Corpus of Documents

IBM Watson for Drug Discovery / Explore a Network

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Common Entities
No entities selected
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Output Sample from Knowledge Graph Query

Gene	Variant	Variant_Type	Functional_Impact	Cancer_Type	NCI Code	Therapy	Response	PMID	Evidence_Level
KIT	exon 9 p.449-514	mutation	gain-of-function	Gastrointestinal Stromal T	C3868	sorafenib	sensitivity	22270258	Level 2B
KIT	exon 11 p.550-592	mutation	gain-of-function	Gastrointestinal Stromal T	C3868	sorafenib	response	22270258	Level 2B
KIT	exon 17 p.788-828	mutation	gain-of-function	Gastrointestinal Stromal T	C3868	nilotinib	sensitivity	2119758, 214560	Level 2B
ABL1	T315I	missense mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	AURK inhibitors + BCR-ABL1 inh	response	22772060	Level 2
ABL1	T315I	missense mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	axitinib	response	25686603	Level 2
ALK	F856S	missense mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	crizotinib	sensitivity	26032424	Level 3
ALK	A348D	missense mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	crizotinib	sensitivity	26032424	Level 3
BCR-JAK2	rearrangement	fusion gene	gain-of-function	Acute Lymphoblastic Leuk	C3167	ruxolitinib	sensitivity	22897847	Level 3
CRLF2	rearrangement	fusion gene	gain-of-function	Acute Lymphoblastic Leuk	C3167	BET inhibitors	sensitivity	22904298	Level 3
CRLF2	rearrangement	fusion gene	gain-of-function	Acute Lymphoblastic Leuk	C3167	mTOR inhibitors	sensitivity	22955920	Level 3
FBXW7	any	mutation	loss-of-function	Acute Lymphoblastic Leuk	C3167	steroids in early setting	response	20861909	Level 2A
IL7R	237-255_indel or S185C	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	ruxolitinib (in SH2B3 delete	sensitivity	2897847, 229559	Level 3
IL7R	237-255_indel or S185C	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	mTOR inhibitors (in SH2B3 del	sensitivity	22955920	Level 3
JAK1	S646F	missense mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	ruxolitinib	sensitivity	22955920	Level 3
JAK2	rearrangement	fusion gene	gain-of-function	Acute Lymphoblastic Leuk	C3167	ruxolitinib	sensitivity	2875628, 228994	Level 3
JAK2	R683	missense mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	ruxolitinib	sensitivity	2805579, 229559	Level 3
KRAS	codon 12	missense mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	MEK inhibitors	sensitivity	18701506	Level 3
NOTCH1	rearrangement	fusion gene	gain-of-function	Acute Lymphoblastic Leuk	C3167	Gamma secretase inhibitor	sensitivity	2688224, 230339	Level 3
NOTCH1	PEST domain p.2245-2536	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	Gamma secretase inhibitor	response	202006 (abstr 65	Level 2B
NOTCH1	PEST domain p.2245-2536	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	Gamma secretase inhibitor	sensitivity	29778842, 225108	Level 3
NOTCH1	PEST domain p.2245-2536	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	Gamma secretase inhibitors + mTOR	sensitivity	19246562	Level 3
NOTCH1	PEST domain p.2245-2536	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	Gamma secretase inhibitors + CDK4	sensitivity	19318552	Level 3
NOTCH1	PEST domain p.2245-2536	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	steroids early setting	response	209, 20861920, 20	Level 2A
NOTCH1	HD p.1529-1723	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	Gamma secretase inhibitor	response	202006 (abstr 65	Level 2B
NOTCH1	HD p.1529-1723	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	Gamma secretase inhibitor	sensitivity	29778842, 225108	Level 3
NOTCH1	HD p.1529-1723	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	Gamma secretase inhibitors + mTOR	sensitivity	19246562	Level 3
NOTCH1	HD p.1529-1723	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	Gamma secretase inhibitors + CDK4	sensitivity	19318552	Level 3
NOTCH1	HD p.1529-1723	mutation	gain-of-function	Acute Lymphoblastic Leuk	C3167	steroids early setting	response	209, 20861920, 20	Level 2A

Leveraging Literature Articles to Support Clinical Care

Gene	Alteration	Alteration Type	Functional Significance	Annotation	Actionable Gene	Actionable Variant
PDGFRA	Y288C	SNV				
PTEN	K163*	Truncation				
FGFR2	K292M	SNV				

Leveraging Literature Articles to Support Clinical Care

Gene	Alteration	Alteration Type	Functional Significance	Annotation	Actionable Gene	Actionable Variant
PDGFRA	Y288C	SNV	Activating	Activating Mutation: Exogenous expression of the Y288C mutant in BaF3 cells increased the number of viable cells measured an average of 5.76 fold more compared with cells expressing wildtype PDGFRA.	Yes	Yes-Functional Genomics
PTEN	K163*	Truncation	Unknown	Likely loss of function: This mutation likely leads to loss of function due to premature stop codon in the phosphatase domain. Missense mutations in K163 resulted in an ≈80-fold reduction in its membrane affinity (Das et al., 2003)	Yes	Yes – Inferred
FGFR2	K292M	SNV	Unknown	Potentially activating. The functional significance of this mutation is unknown. However, it is located within the Ig-like domain type 3, and other mutations in this domain can create an autocrine, feed-forward FGF signaling loop resulting in FGFR2 activation.	Yes	Potentially

Summary

Empowering Healthcare Professionals with Cognitive Computing

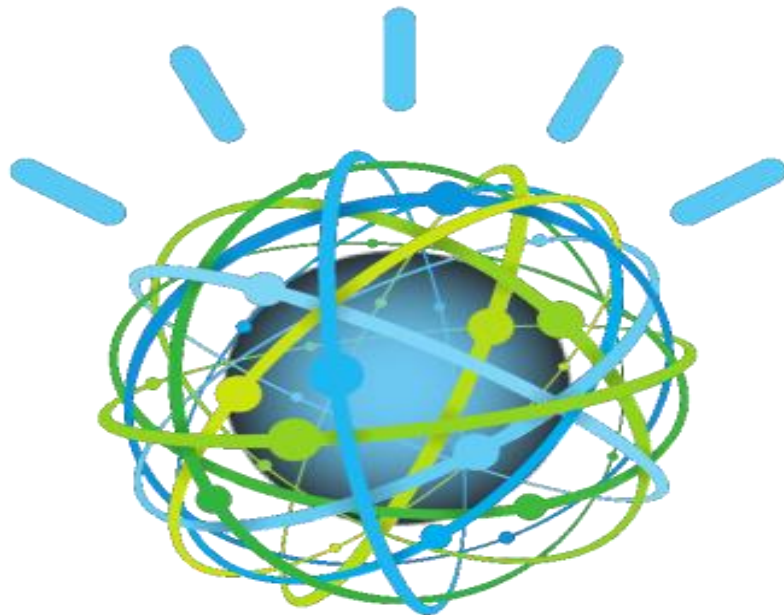
The Process of Building

- Work Closely with domain experts
- Leverage tools (like NLP) to extract unstructured concepts and their associations
- Combine unstructured and structured data to create knowledge
- Validate and iterate with domain experts

Value of Applying to Healthcare

- Increased efficiency
- Reduced bias
- Scaling expert knowledge
- Innovating across a complex domain

With Cognitive computing, the extraction of information from clinical data and published literature can empower healthcare professionals to stay up-to-date, focus on the complex problems, and spend more time with caring for patients.



Questions?
(remember, I'm not Watson..)

PRECISION ONCOLOGY FOR ALBERTA: INNOVATING HEALTH SYSTEM READINESS.

ARCC: Canadian Centre for Applied Research and
Cancer Control

Toronto: May 25-26, 2017

Tania Bubela

ALMDx/PACEOMICS Team

Alberta Health Services

- Don Juzwishin

Faculty of Medicine & Dentistry

University of Alberta

- Christopher McCabe
- Judith Hugh (+AHS)
- Michael Mengel (+AHS)
- Deborah James
- Stacey Hume (+AHS)

Research staff/students:

Michael Paulden, Mark Bieber, Westerly Luth, Katherine Fu, Kiah Van der Loos, Monica Wang, Yael Mansour, Negar Razavilar



Outline

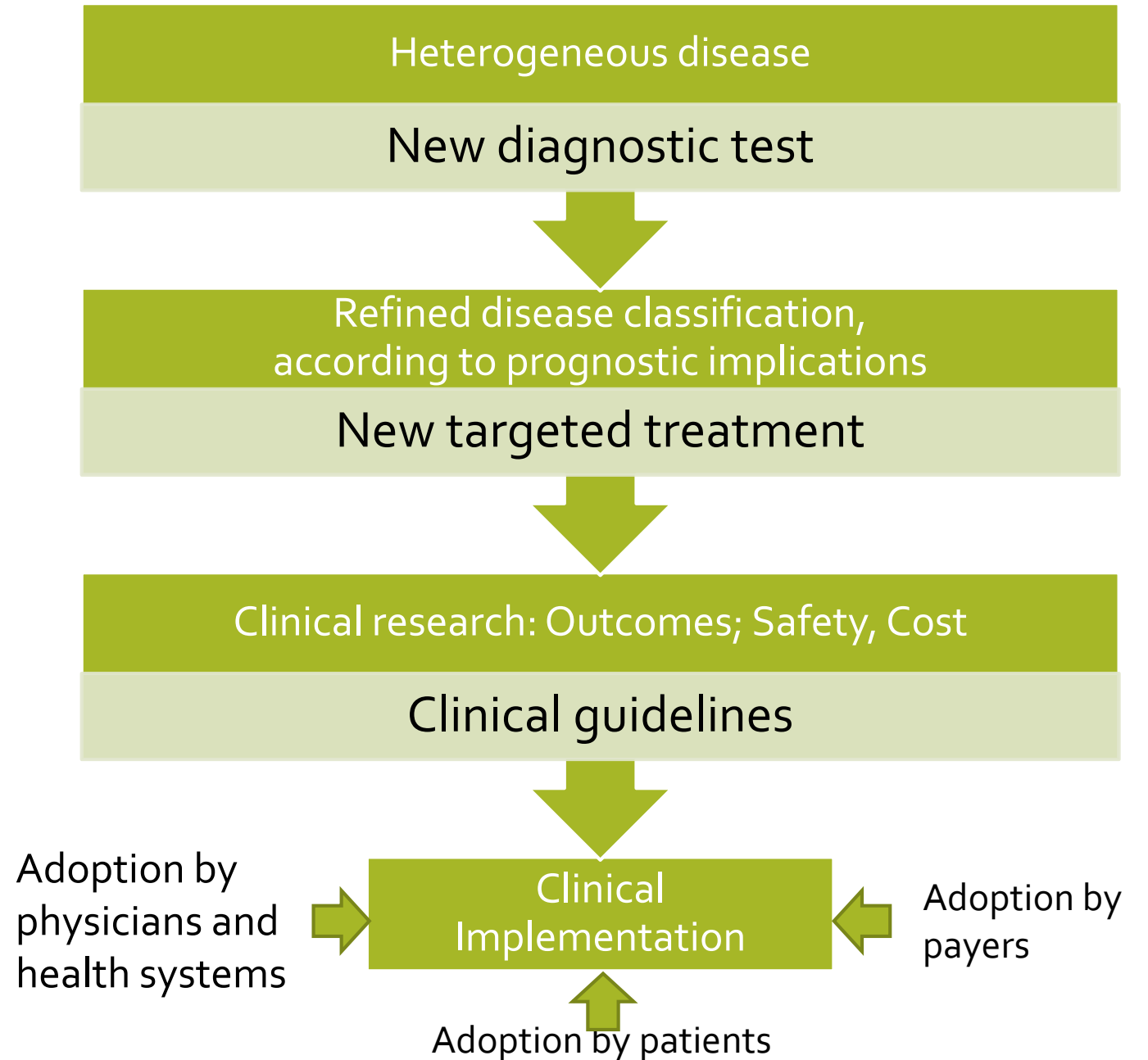
- What is precision medicine?
- Precision medicine landscape in Alberta
- Supporting precision medicine in two contexts:
 - Adoption of new technologies wherever developed
 - Local innovation
- Mike Paulden will present a case example from of Life-Cycle Technology Management in Oncology
- Conclusions and lessons learned on barriers and enablers

PRECISION MEDICINE FOR ALBERTA?

ALMDX/PACEOMICS TEAM

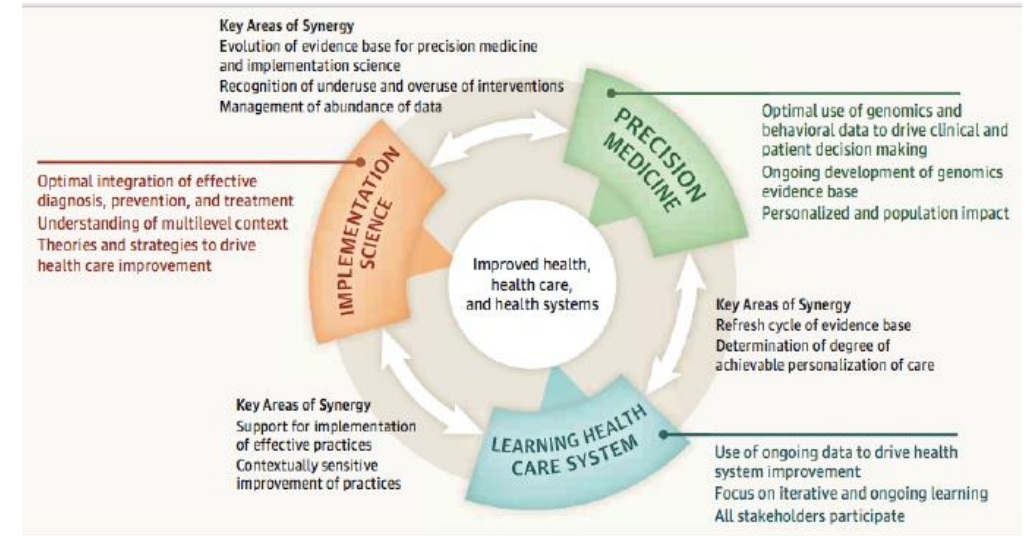
Precision Medicine: Personalized, Problematic & Promising

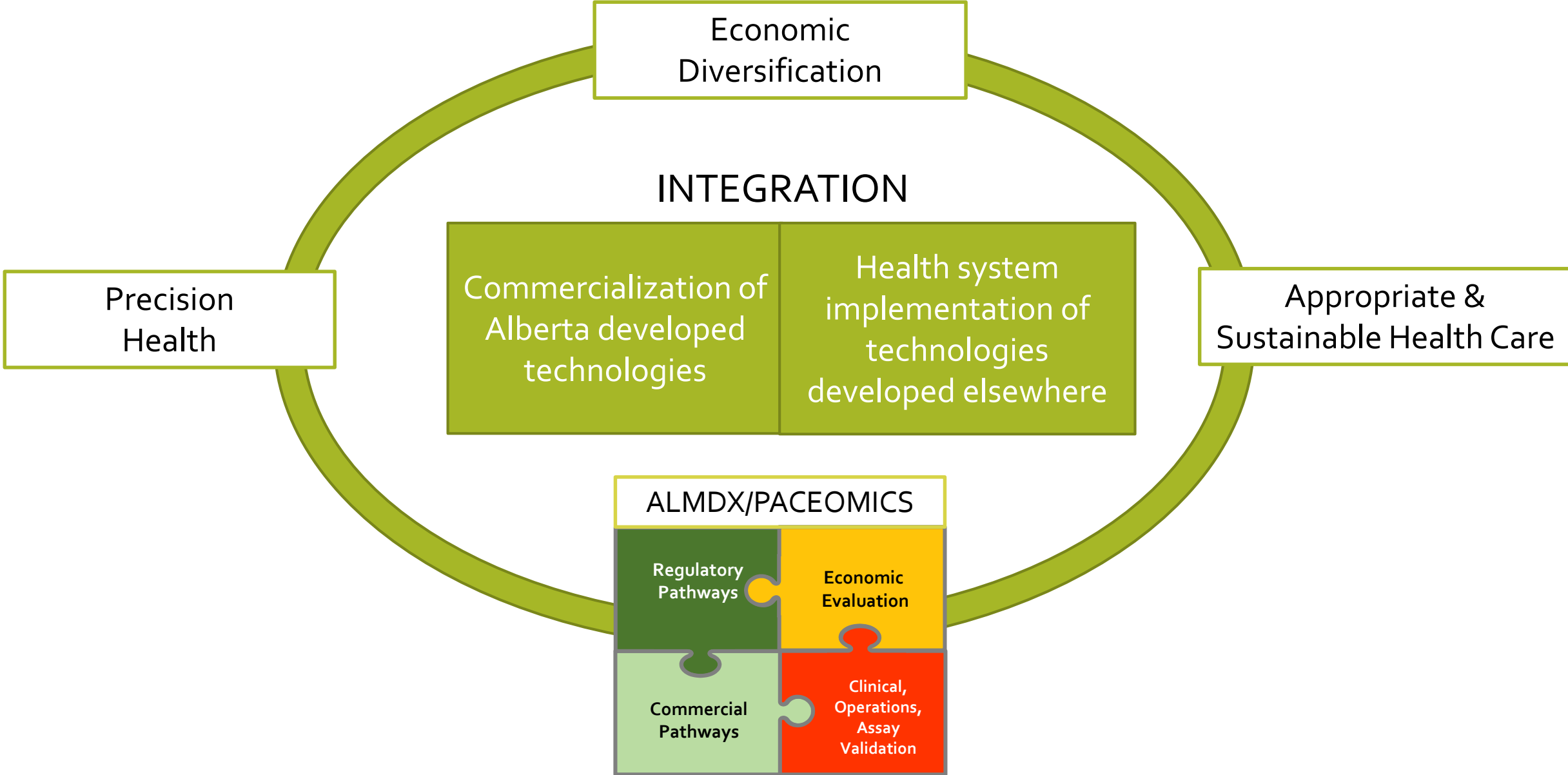
(Jameson & Longo *NEJM* 2015)



Current Challenges in Alberta

- Consensus / Joint Strategy and Vision
- Managing implementation of PM technologies
 - Ascribing 'value' to new technologies and approaches
 - Defining/implementing new funding strategies
 - 'On-ramps', decision criteria, procurement mechanisms and 'off-ramps'
 - Off-ramps for established but low value technologies – Choosing Wisely, Appropriateness of Care, Utilization management
- Changing legislation, policies, and 'behavior'
- Connecting the ecosystem
 - Re-allocating internal budgets to reward nimble portfolios





Key Areas of Synergy

Evolution of evidence base for precision medicine and implementation science
Recognition of underuse and overuse of interventions
Management of abundance of data

Optimal integration of effective diagnosis, prevention, and treatment
Understanding of multilevel context
Theories and strategies to drive health care improvement



Improved health, health care, and health systems

Key Areas of Synergy

Support for implementation of effective practices
Contextually sensitive improvement of practices



Key Areas of Synergy

Refresh cycle of evidence base
Determination of degree of achievable personalization of care

Use of ongoing data to drive health system improvement
Focus on iterative and ongoing learning
All stakeholders participate

PRECISION MEDICINE

Optimal use of genomics and behavioral data to drive clinical and patient decision making
Ongoing development of genomics evidence base
Personalized and population impact

Precision Medicine in Oncology

Courtesy of Dr. Jeffrey S. Ross, Albany Medical College
Medical Director, Foundation Medicine Inc.

Evolution of Targeted Therapies

1970: ER testing and hormonal therapy for breast cancer

1990: cytogenetics/FISH testing and therapy for Heme malignancies

1998: *HER2* testing and Trastuzumab for breast cancer

2001: *BCR-ABL* testing and Imatinib for chronic myelogenous leukemia

2003: *EGFR* mutation testing and Erlotinib for non-small cell lung cancer (NSCLC)

2007: *KRAS* mutation testing and Cetuximab/Panitumumab for colorectal cancer

2010: *EML4-ALK* testing and Crizotinib in NSCLC

2011: *BRAF* mutation testing and Vemurafenib in melanoma

2013: *HER2* mutation identification and anti-Her2 targeted therapy in
NSCLC, breast cancer and micropapillary urothelial cancer

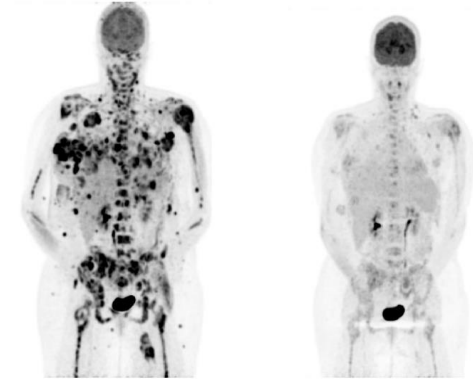
2013 *NTRK1* fusion testing and Crizotinib in NSCLC

2014: *PD1* and *PDL1* and immunotherapies

As of 2017: 207 Pharma pipelines for CDx – 158 in cancer

Next Generation Pathology to Precision Oncology

Patient with *BRAF* V600E mutated malignant melanoma treated
with selective inhibitor of BRAF (Vemurafenib)



Pre-therapy

15 days Post-therapy

Slide courtesy of Dr. Grant McArthur

Ion Torrent PGM
sequencer

Next-Gen
Sequencing

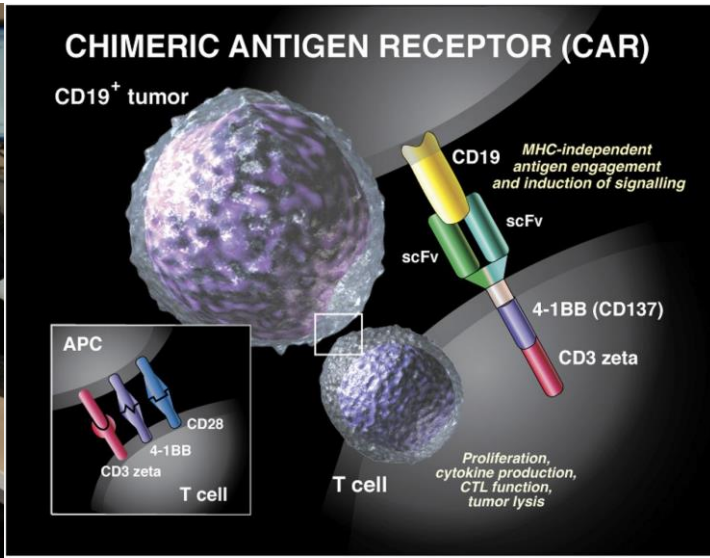


Complete Regression in Pediatric Glioblastoma

Molecularly matched targeted therapy against *BRAF* V600E



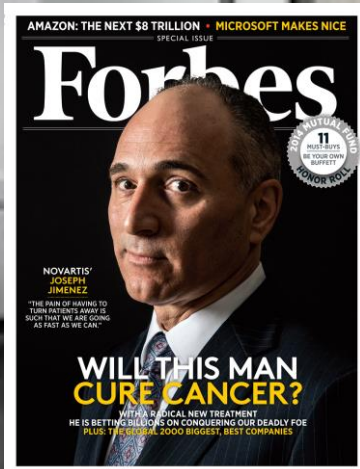
Cancer Cellular Immunotherapy



Pharma & Healthcare / #Biotech
AUG 31, 2016 @ 08:32 AM 18,697 VIEWS

Novartis Dissolves CAR-T Unit, Cutting 120 Positions

Matthew Herper, FORBES STAFF
I cover science and medicine, and believe this is biology's century. FULL BIO

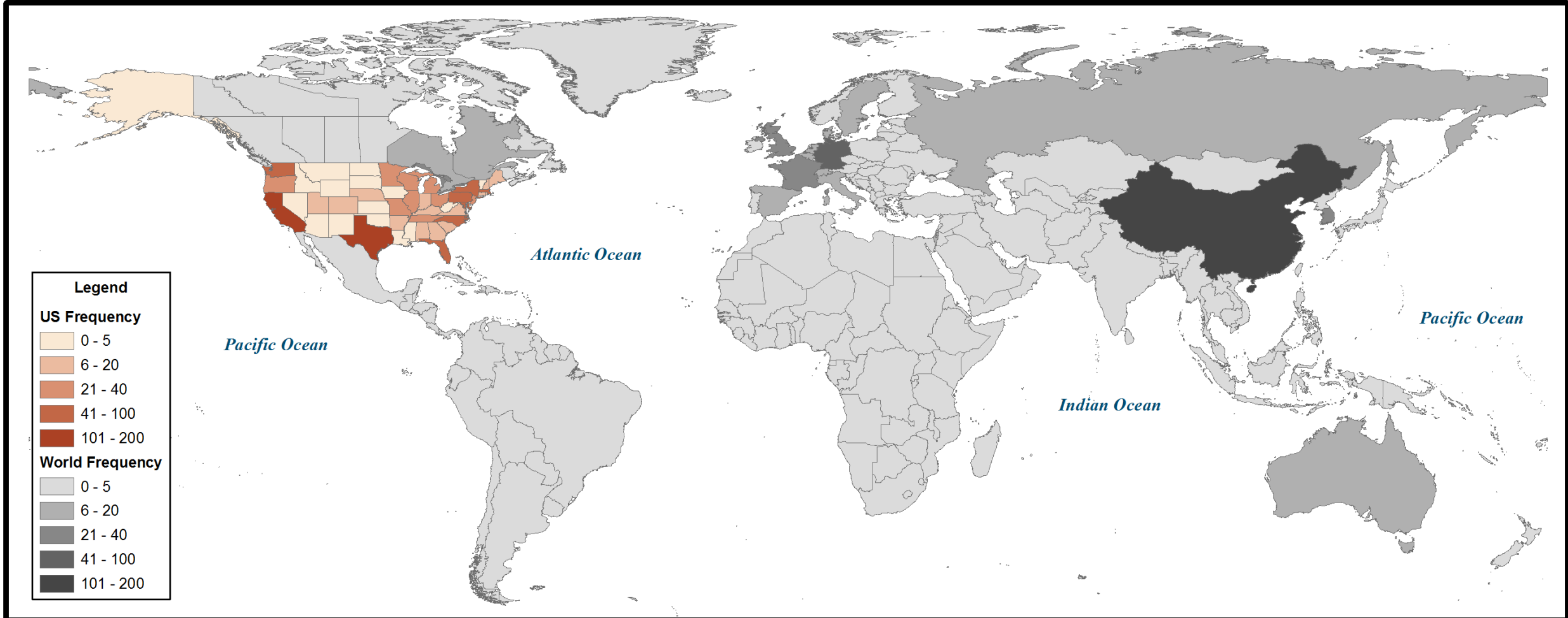


Signage is displayed in front of the Novartis AG Institutes for BioMedical Research building in Cambridge, Massachusetts, on Aug. 5, 2016. Photographer: Scott Eisen/Bloomberg

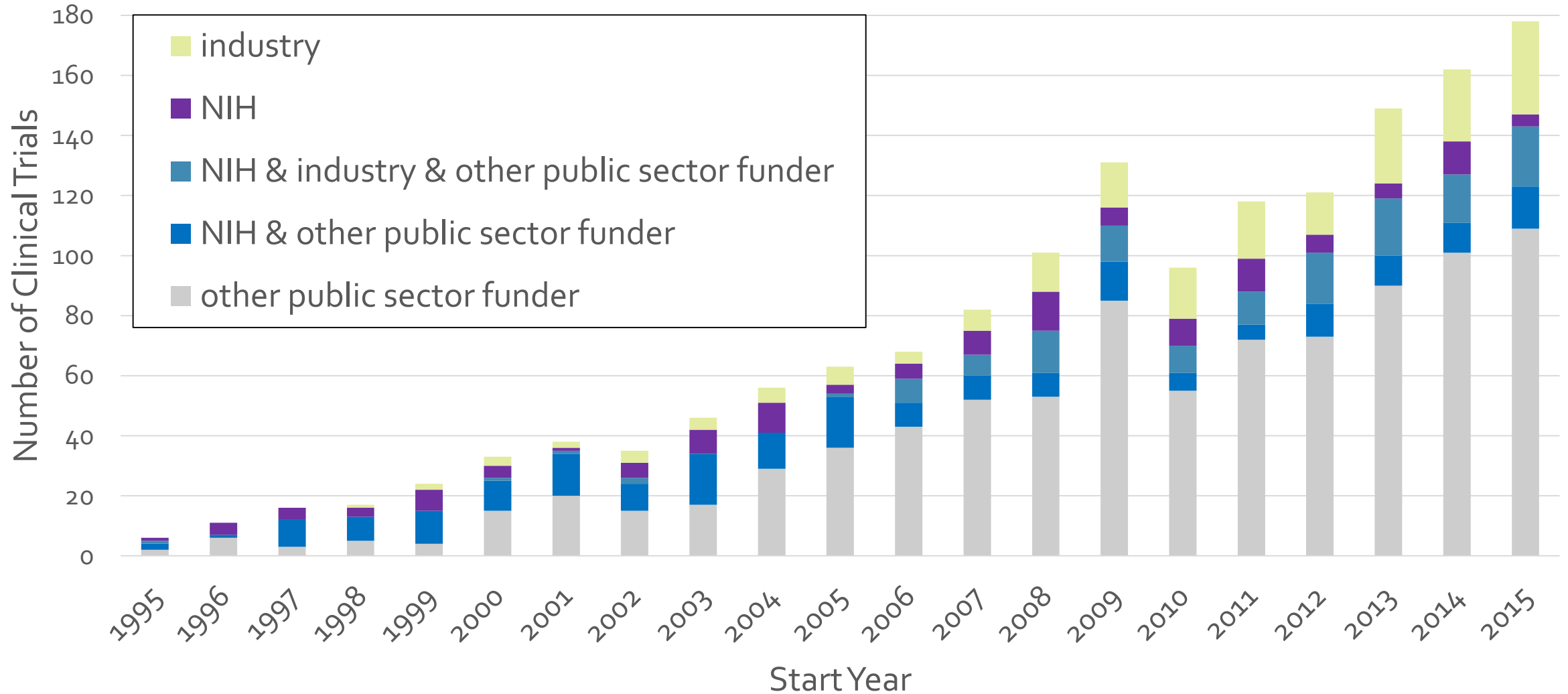
Novartis-Penn Center for Advanced Cellular Therapeutics

Novartis is shutting down the business unit it created to develop white blood cells that can attack certain types of cancer, but

HORIZON SCANNING: GLOBAL IMMUNOTHERAPY CLINICAL TRIALS

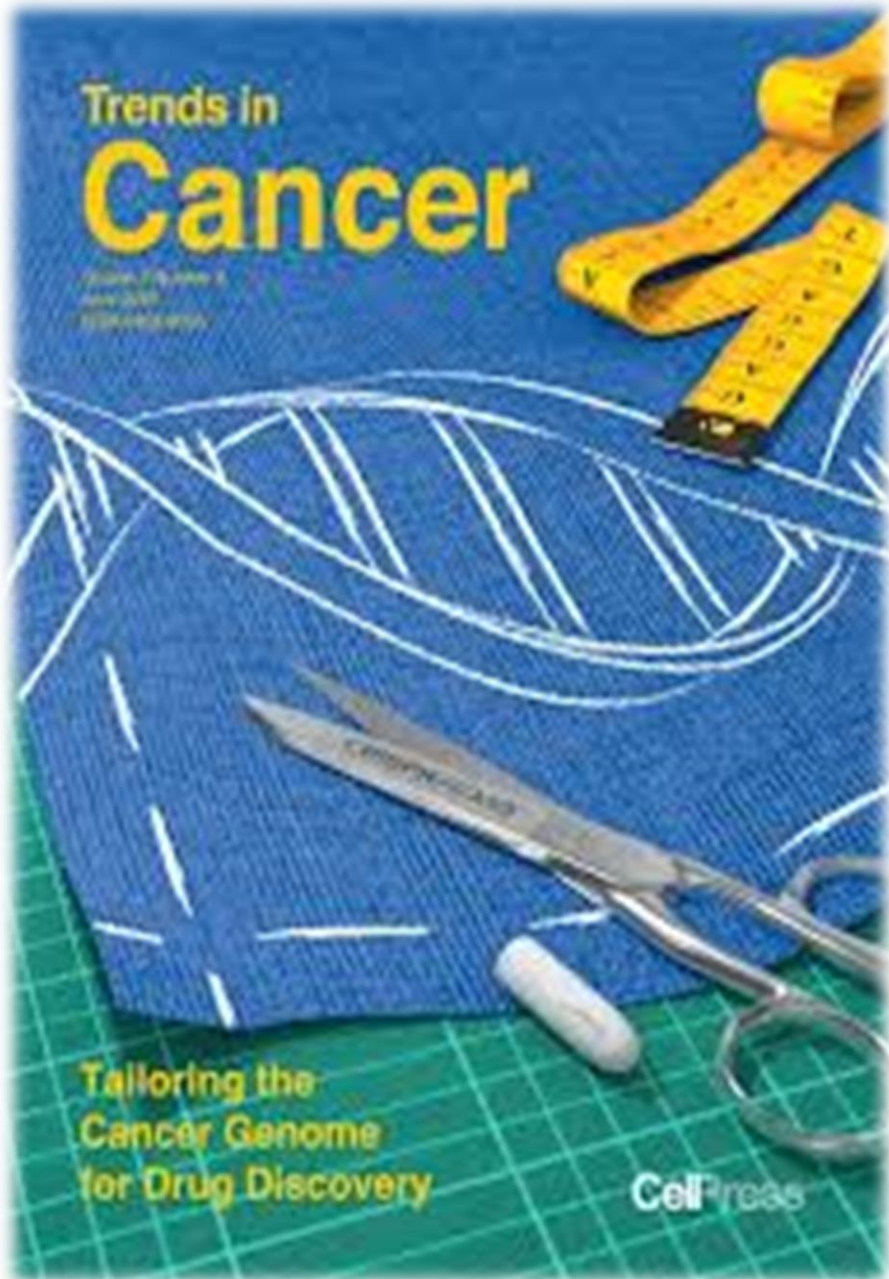


SPONSORS OF 1554 CI CLINICAL TRIALS

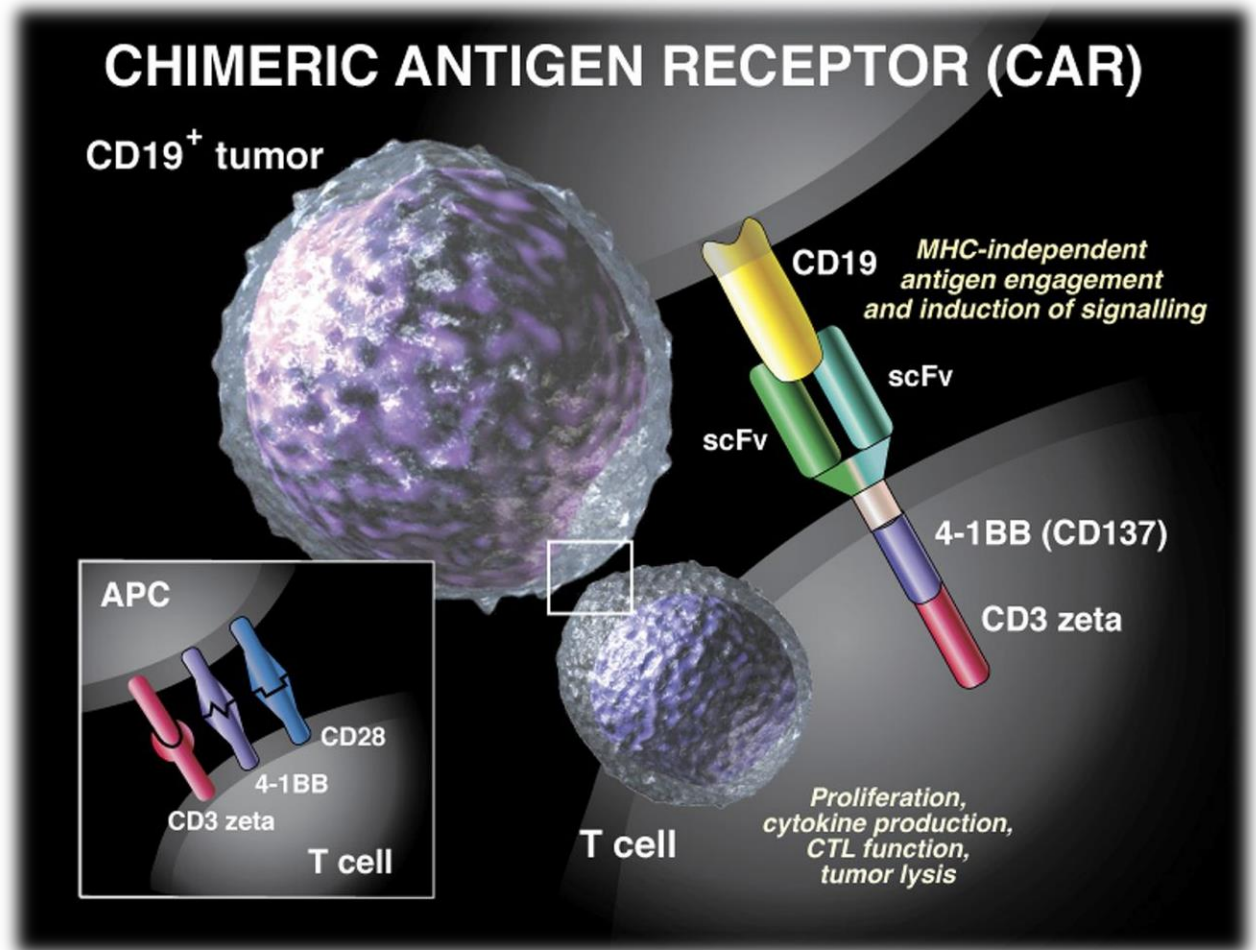




FIRST-EVER CRISPR TRIAL POINTS TO LOOMING PROBLEMS



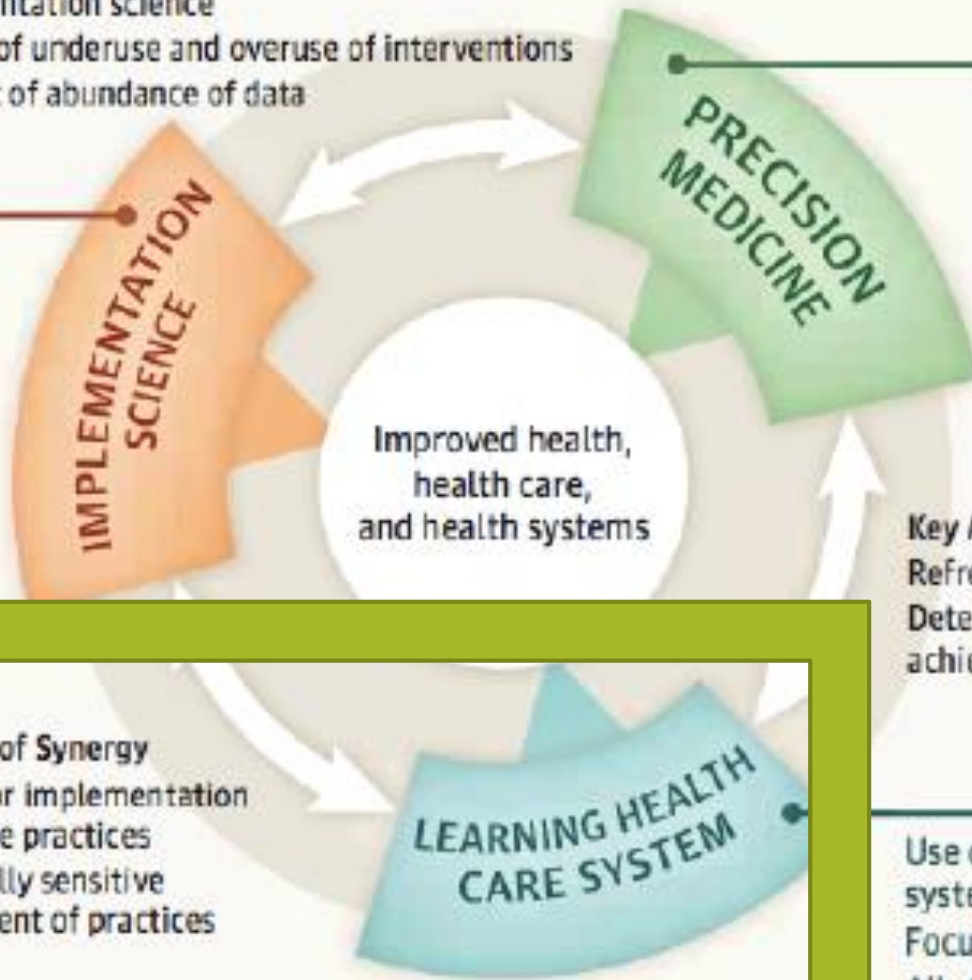
CRISPR and CAR-T



Key Areas of Synergy

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Theories and strategies to drive health care improvement



Optimal use of genomics and behavioral data to drive clinical and patient decision making
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Refresh cycle of evidence base
Determination of degree of achievable personalization of care

Key Areas of Synergy

Support for implementation of effective practices
Contextually sensitive improvement of practices

Use of ongoing data to drive health system improvement
Focus on iterative and ongoing learning
All stakeholders participate

International Environment



ExAC

the gnomAD team assembled
2.9 petabytes
of raw data...
that's equal to **331 years**
of movie streaming

254.2 million
rare and common genomic variants,
159.8 million
of which were novel
were discovered using gnomAD

gnomAD contains
sequences from over
140,000
people

gnomAD contains data from
people hailing from
each of the
6 inhabited
continents

THE PRECISION MEDICINE INITIATIVE®



LONGER-TERM GOALS

Create a research cohort of **> 1 million American volunteers** who will share genetic data, biological samples, and diet/lifestyle information, all linked to their electronic health records if they choose.



NEAR-TERM GOALS

Intensify efforts to apply precision medicine to **cancer**.

Innovative **clinical trials**
of targeted drugs for
adult, pediatric cancers

Use of
combination
therapies

Knowledge to
overcome **drug**
resistance



Capital Investment: Beijing Genomics Institute



Complex Legislative Framework

- **Health Information**

- *Personal Information Protection and Electronic Documents Act* R.S. 2000 c. 5 [PIPEDA]
- *Health Information Act*, R.S.A. 2000, c. H-5. [HIA]

- **Personal Information Held by Private Sector**

- *Personal Information Protection and Electronic Documents Act*, R.S. 2000, c. 5 [PIPEDA]
- *Personal Information Protection Act*, S.A. 2003, c. P-6.5 [PIPA]

- **Personal Information Held by Federal Government**

- *Privacy Act*, R.S. 2000, c. P-21.

- **Personal Information Held by the Alberta Government**

- *Freedom of Information and Protection of Privacy Act*, R.S.A. 2000, c. F-25. [FOIP]

Custodians and the Controlled Arena

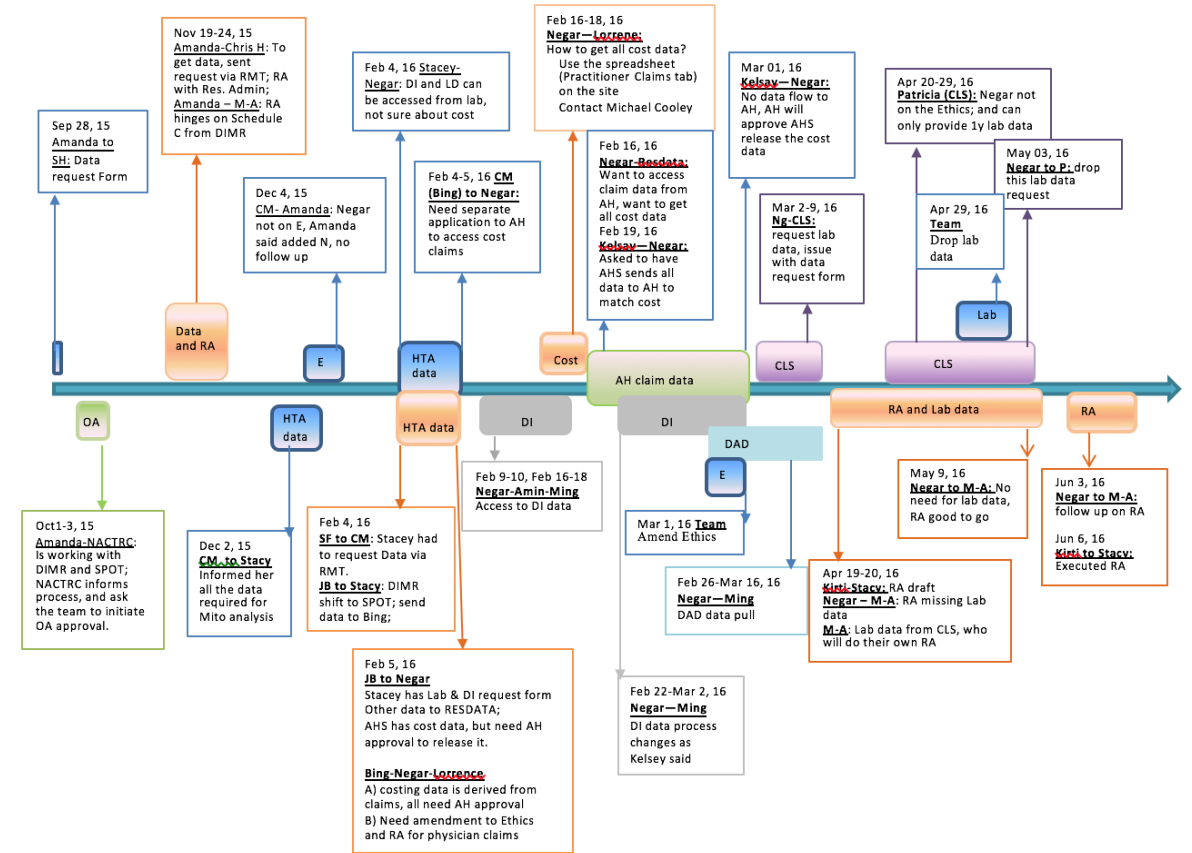
- There is a “controlled arena” around the custodians.
- Individually identifying health information can move from one custodian to another for the purposes authorized

Movement of identifying health information outside of the controlled arena is more restricted

- **A researcher under certain conditions**
 - Research agreements
 - Ethics

Barriers to data access

- **Lack of process clarity**
 - Who is responsible for triggering data requests?
 - Who are the appropriate contact points?
 - What information does each database contain?
- **Role of data custodian**
- **Miscommunication**



Key Areas of Synergy

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IMPLEMENTATION SCIENCE

Improved health, health care, and health systems

PRECISION MEDICINE

Optimal use of genomics and behavioral data to drive clinical and patient decision making
Ongoing development of genomics evidence base
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Key Areas of Synergy

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LEARNING HEALTH CARE SYSTEM

Use of ongoing data to drive health system improvement
Focus on iterative and ongoing learning
All stakeholders participate

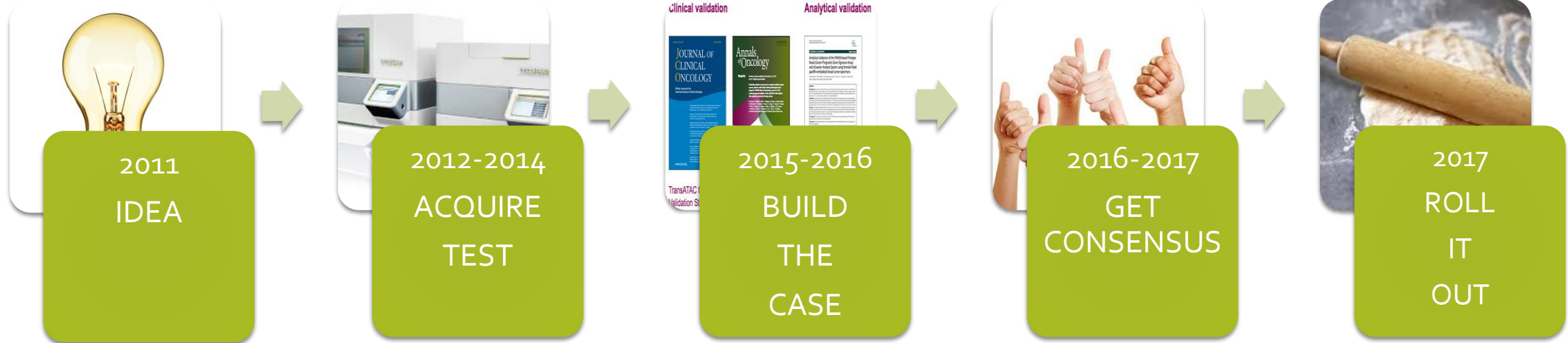
Key Areas of Synergy

Support for implementation of effective practices
Contextually sensitive improvement of practices

Challenges:

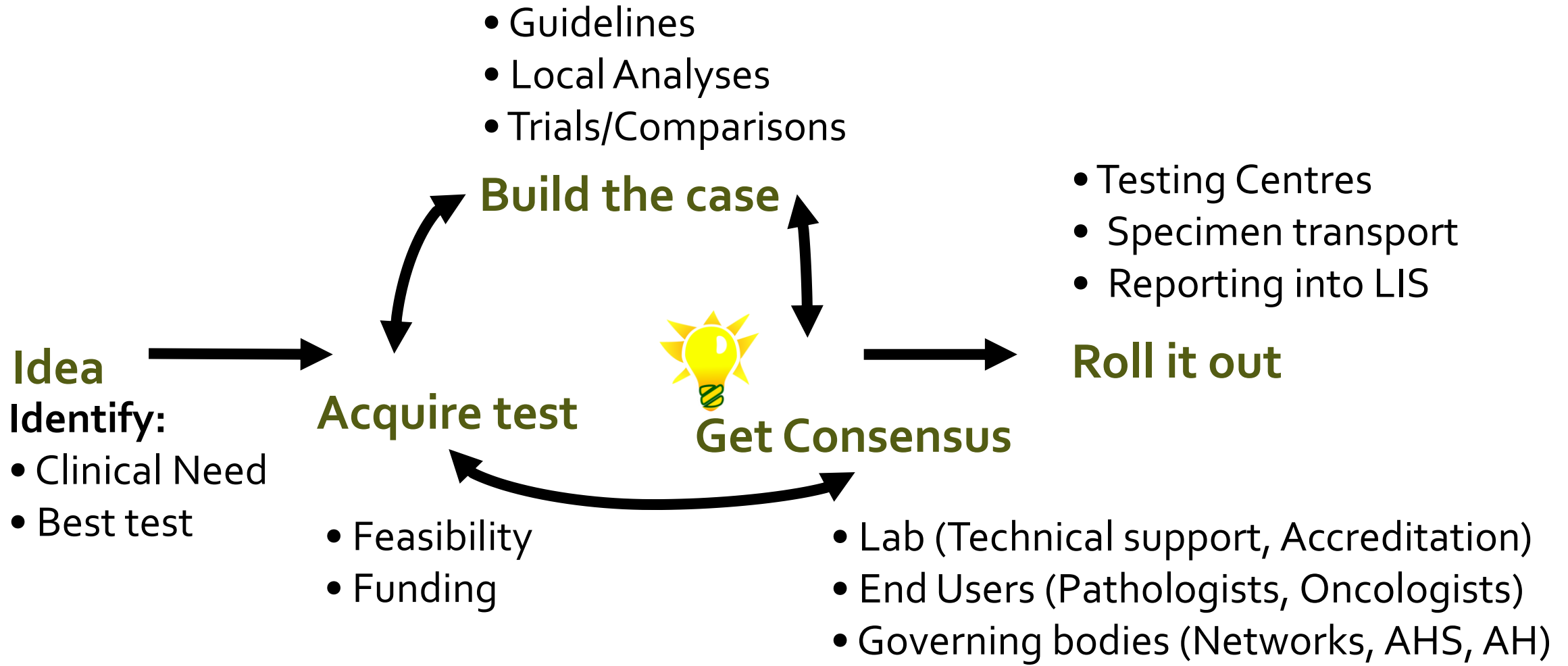
Moving from Oncotype Dx™ to Prosigna™

- CFI purchase of test platform
- ALMDx & Cancer SCN local test



Guidelines
Local Analyses
Trials/
Comparisons

What is the Best way to Replace an Existing Test?



CONCLUSIONS ON PRECISION MEDICINE FOR ALBERTA

ALMDx/PACEOMICS Team

Identified challenges

- **Business models & regulatory pathways**
 - Laboratory Developed Test
 - Licensed Software
 - Distributed Kit
- **Capacity limitations**
 - Sequencing infrastructure
 - Bioinformatics capacity
- **AHS Organizational structure, lack of:**
 - Competitive billing
 - Outreach/advertising
 - Innovation incentives
 - Transparent re-investments between operational silos
- **Organizational culture**
 - Not-for-profit
 - Clinician opinion and behaviour
- **Laws and policies**
 - Procurement
 - Privacy/Health Information
- **Data access (essential for PM)**
 - Data aggregation across databases for analytics
 - Complex operational approval structures
 - Lack of procedural clarity and transparency

Opportunities

- **Life-cycle evaluation of new technologies**
 - 'Fend off' low value new diagnostics
 - Adopt through *dynamic* health-economic and operational evidence generation and implementation process
 - Evidence-based utilization management: 'right test at the right time in the right patient' enable adoption and decommissioning of tests/technologies
- **Support local innovators**
 - Support SME as an effective beta-site and real world incubator
 - Commercialization and regulatory affairs
 - Provincial procurement policies
- **Impact on the Alberta Economy**
 - Jobs in Alberta through repatriation of high-complexity diagnostics
 - Build expertise in implementation sciences i.e. knowledge-based jobs
 - Become competitive in the global diagnostic testing market: research & innovation
 - Attractive partner to industry





THANK YOU

Innovation in Cancer Care: The Promise and the Peril

Mike Paulden, Assistant Professor,
School of Public Health, University of Alberta

@mikepaulden paulden@ualberta.ca mikepaulden.com

Overview

1. How might **economists** define “**innovation**” in cancer care?
2. What are some of the “**promises**” associated with this approach to defining “innovation”?
3. What are some of the “**perils**”?
4. Case study: **Oncotype DX** vs **Prosigna** for guiding adjuvant chemotherapy decisions in early breast cancer

How might economists define “innovation”?

Defining “innovation” requires a clear objective

Typically, economists make **two key assumptions**:

Health system has a **constrained budget** - we can't afford everything

Given this, resources are allocated to **satisfy some objective**

Conventional objective is to “**maximize population health**”

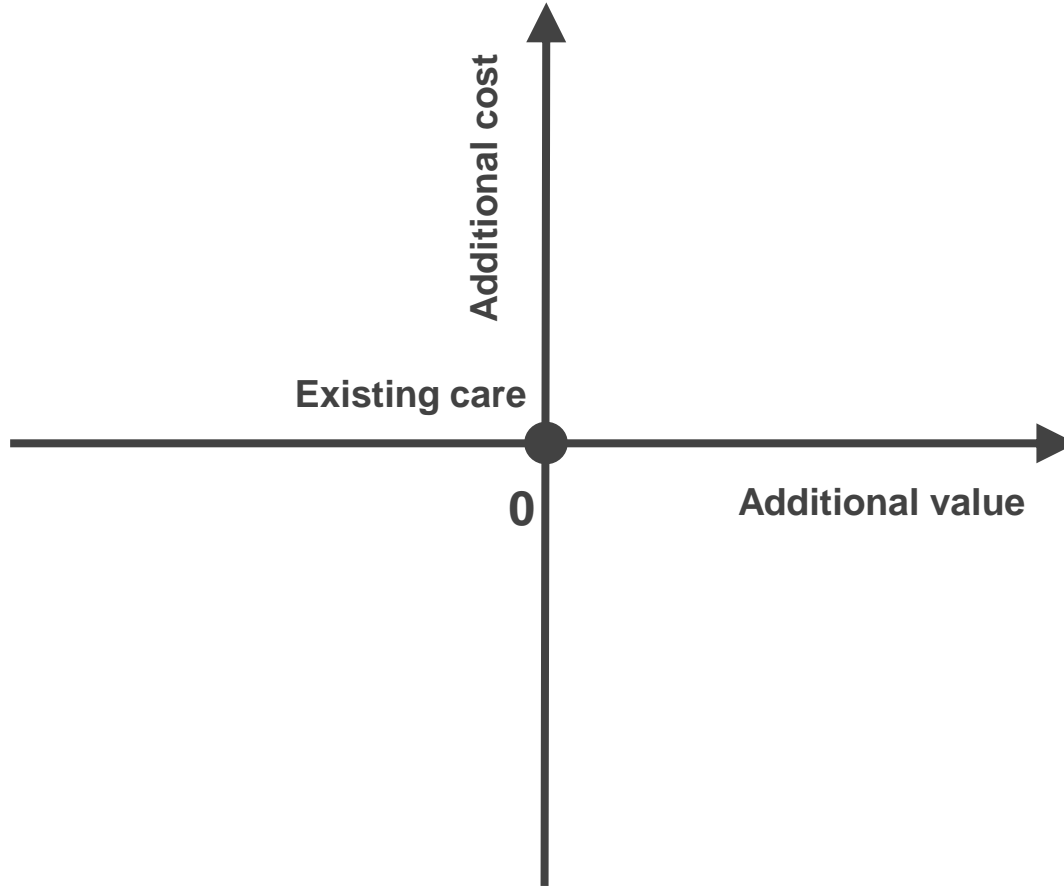
Often measured using **quality-adjusted life years (QALYs)**

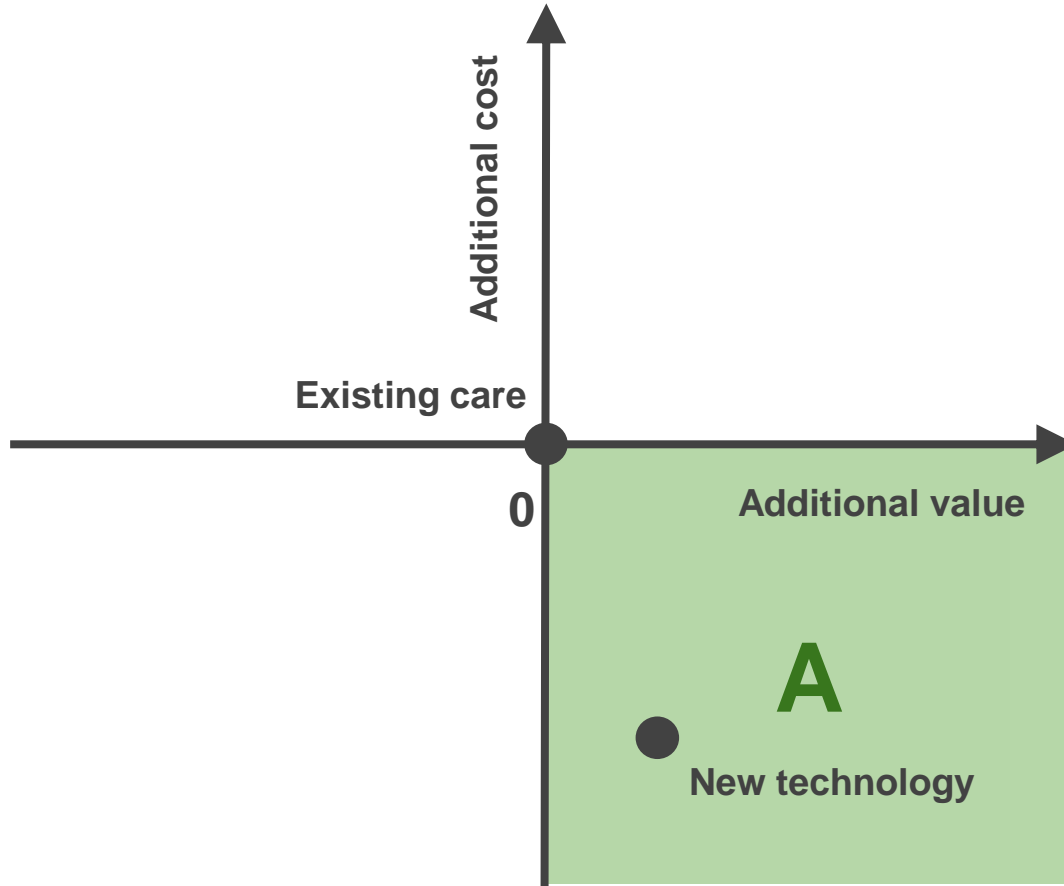
But **other measures** of health can be used

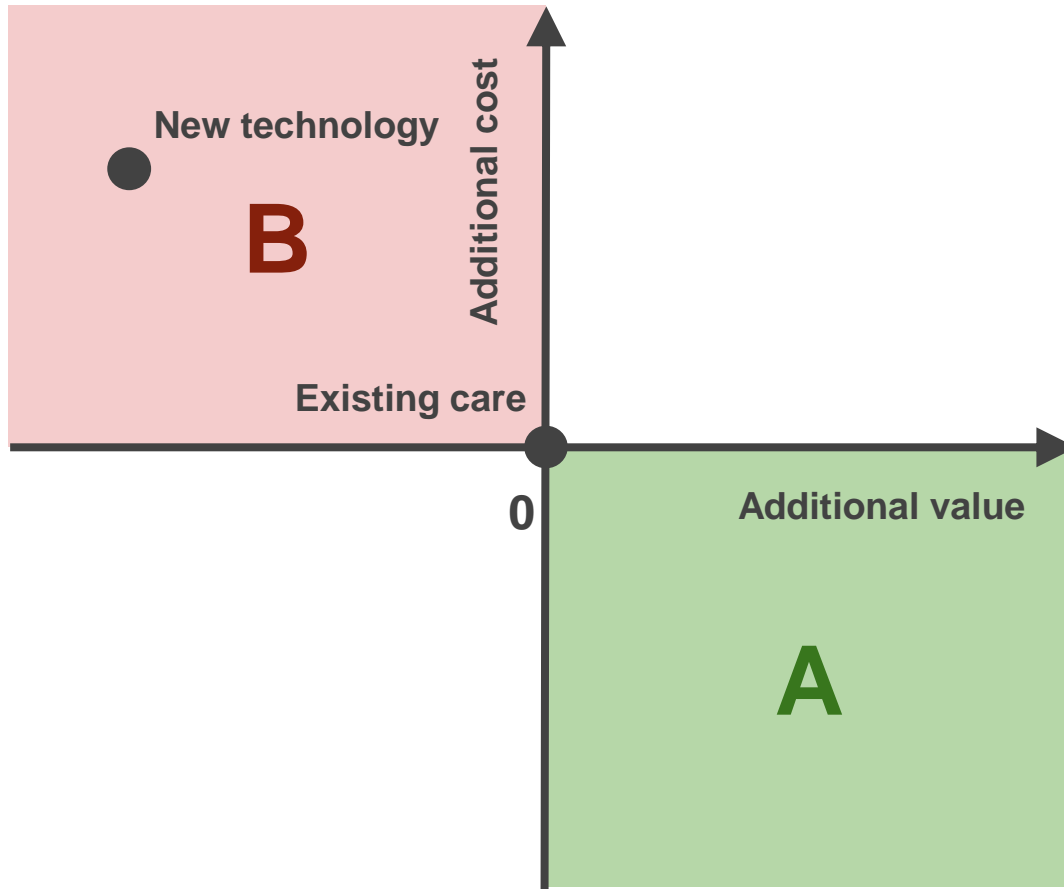
Alternative objectives may include **non-health outcomes**

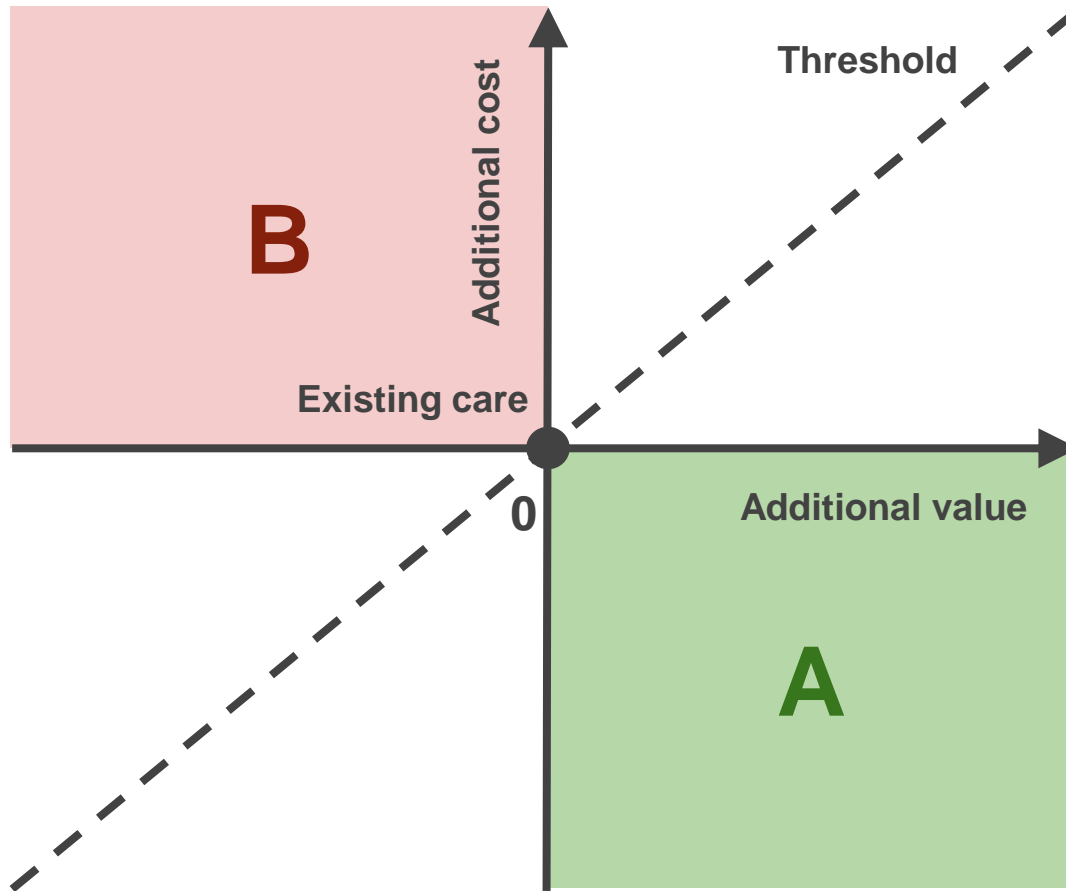
Mike Paulden, University of Alberta @mikepaulden paulden@ualberta.ca mikepaulden.com Slide 4 of 33

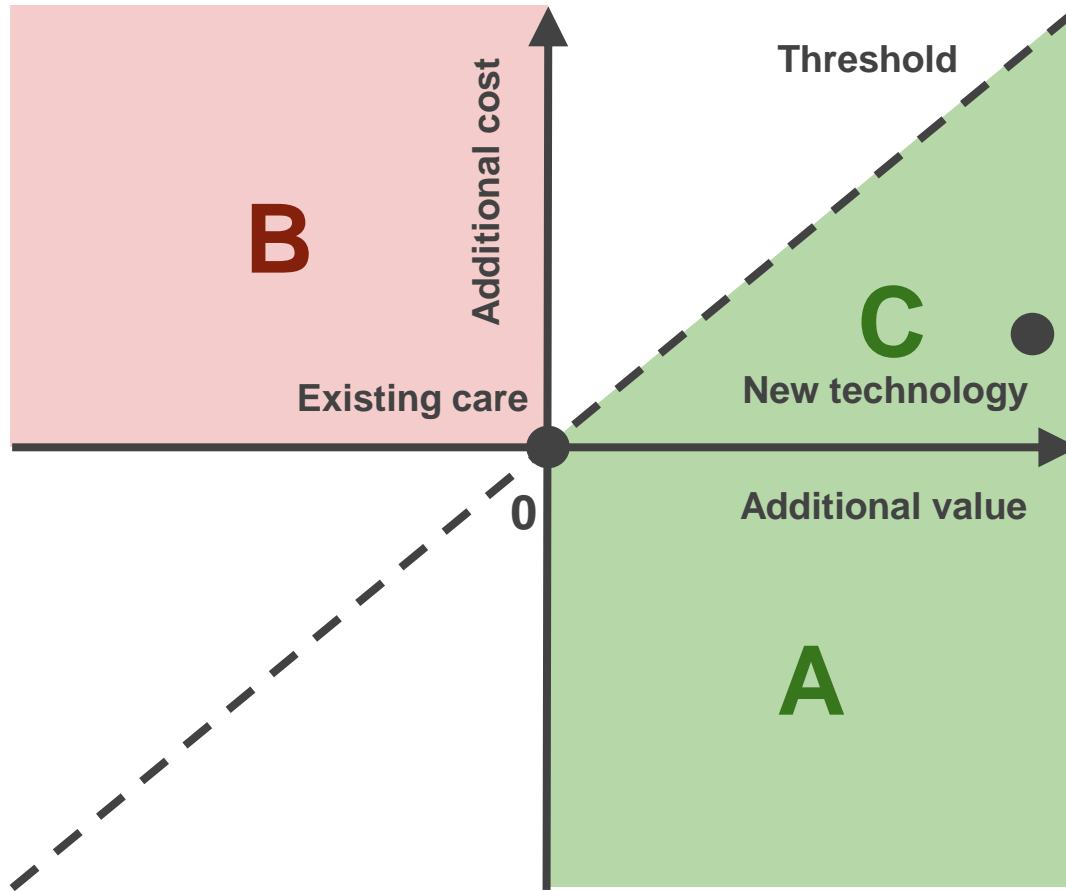
Can consider generic objective of “**maximizing value**”

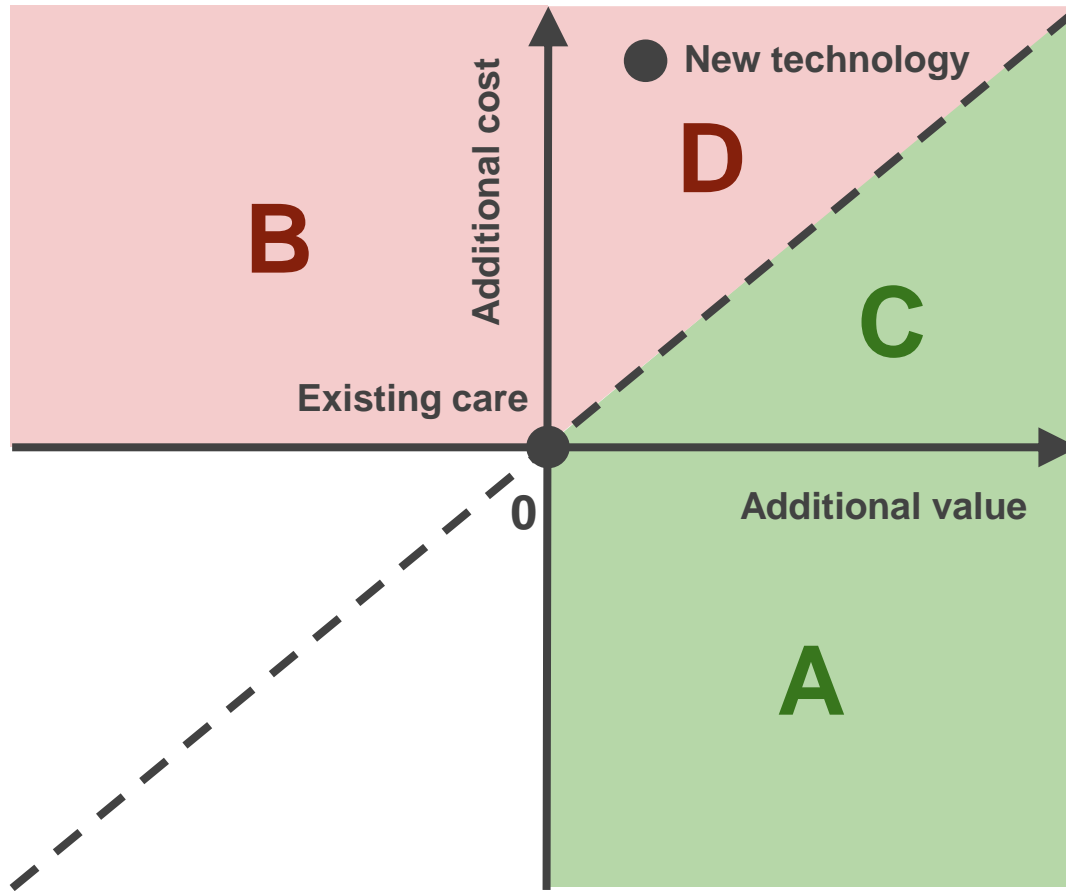


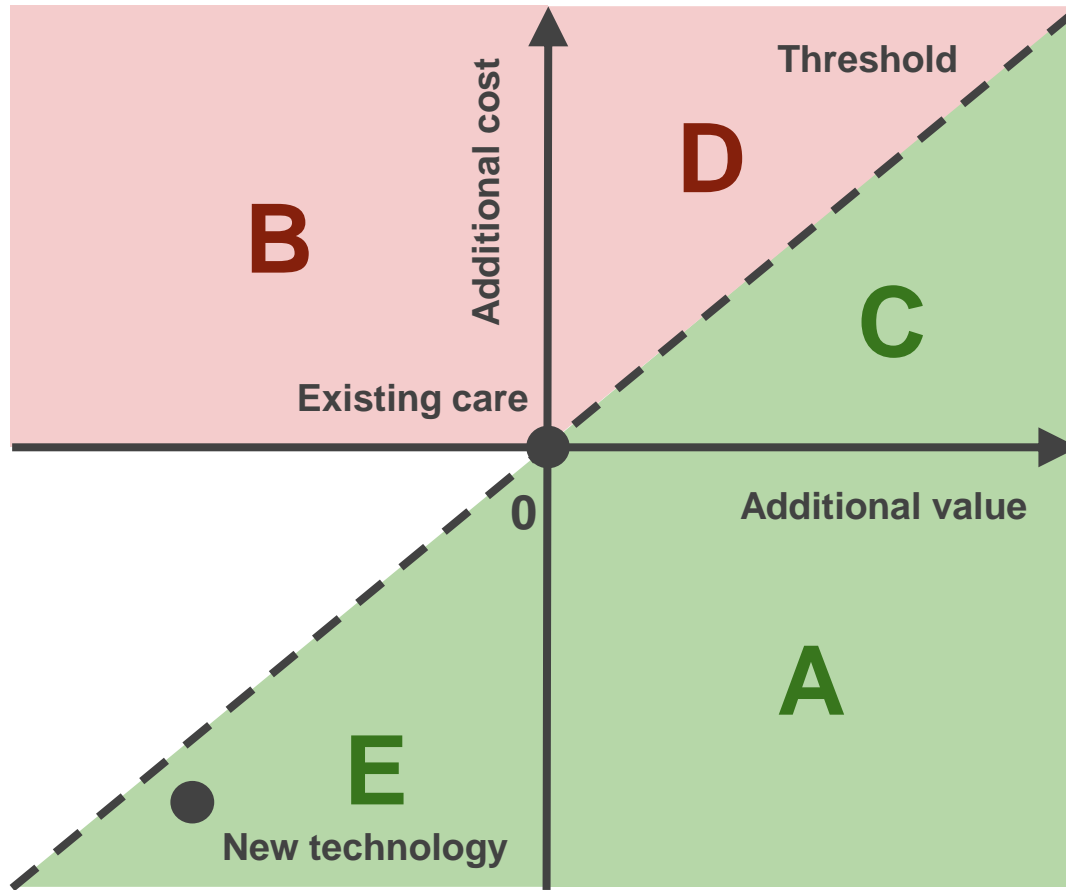


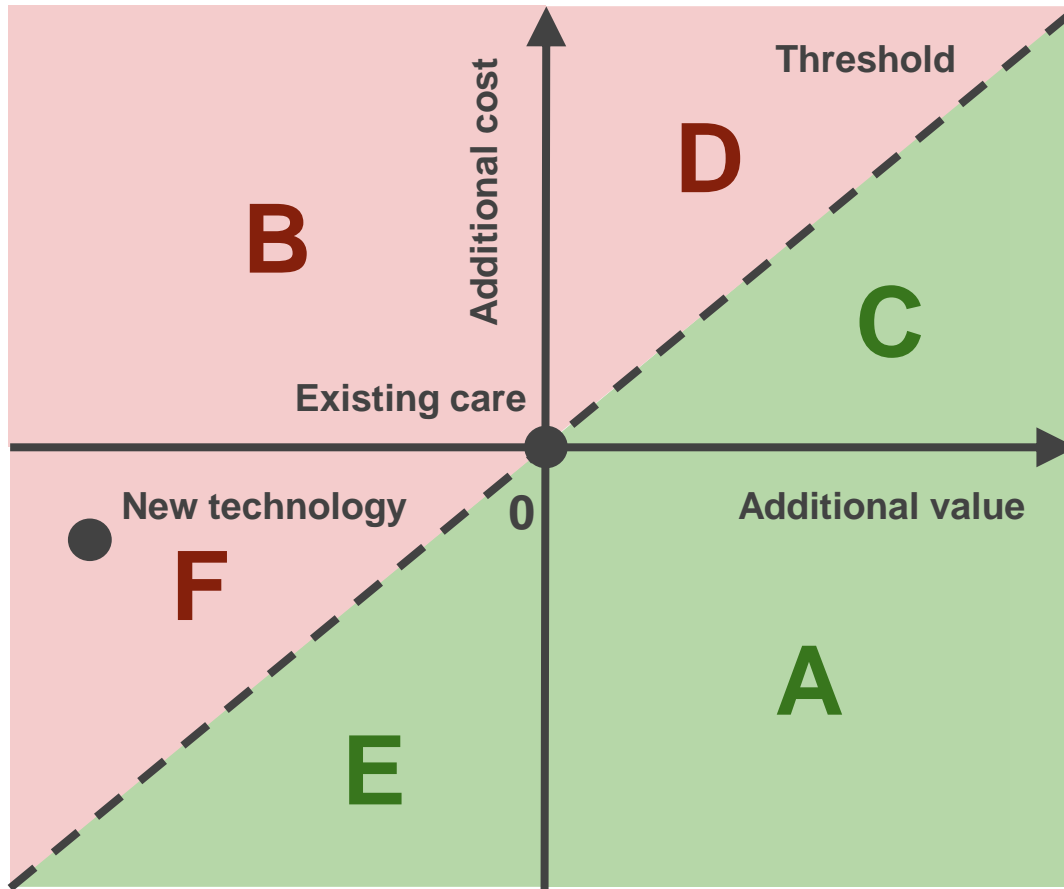




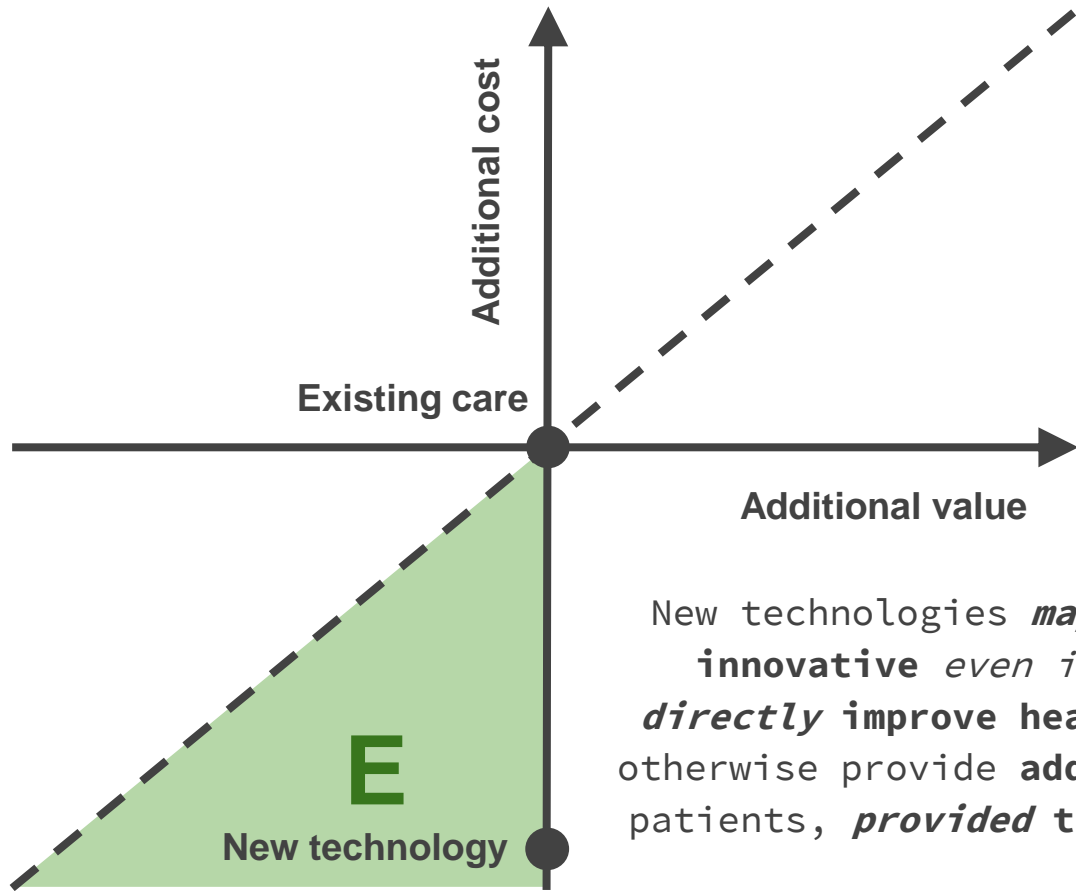




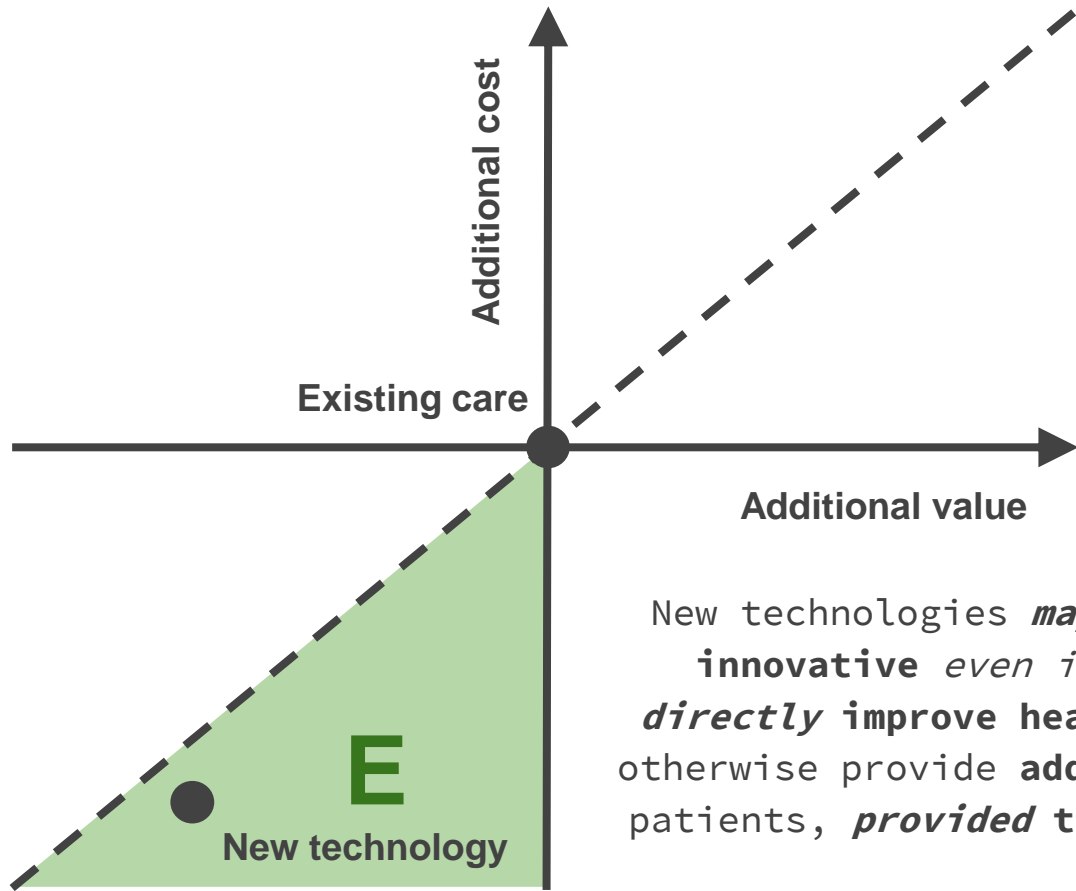




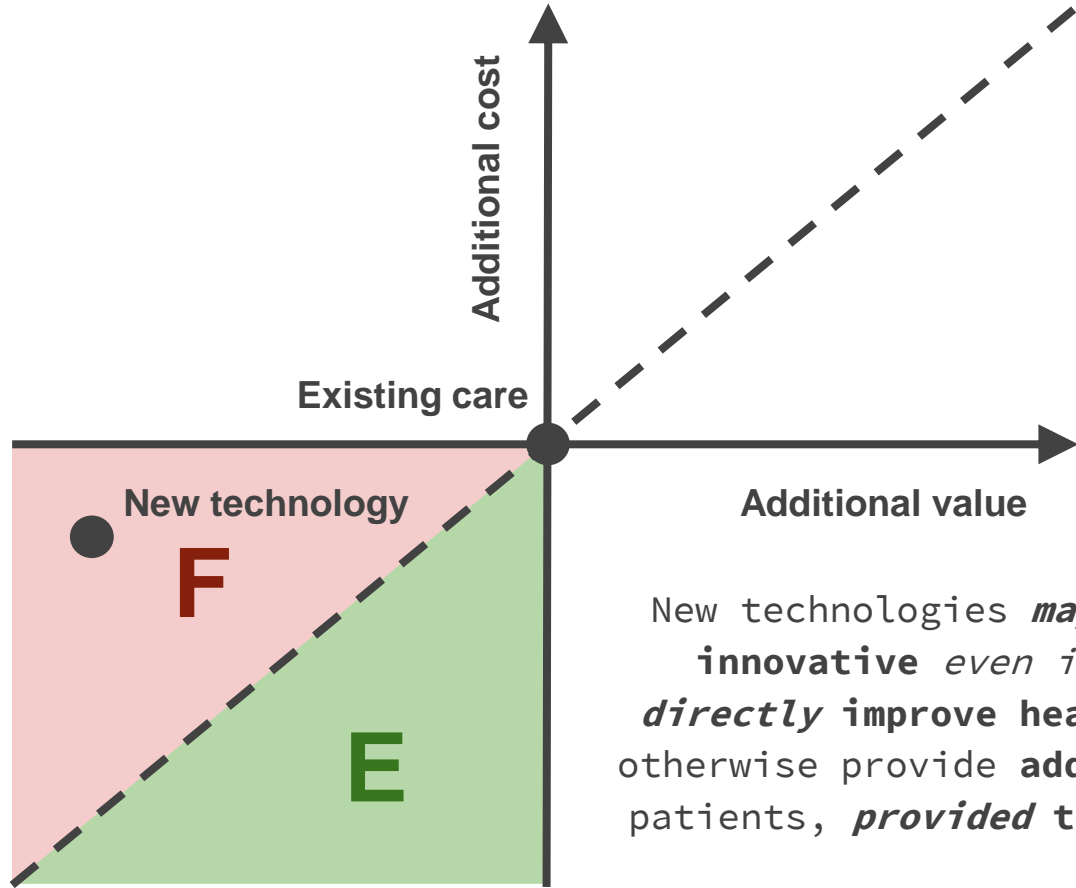
What are some of the “promises”?



New technologies *may* be considered **innovative** *even if* they do not **directly** improve health outcomes or otherwise provide **additional value** to patients, *provided* they reduce costs

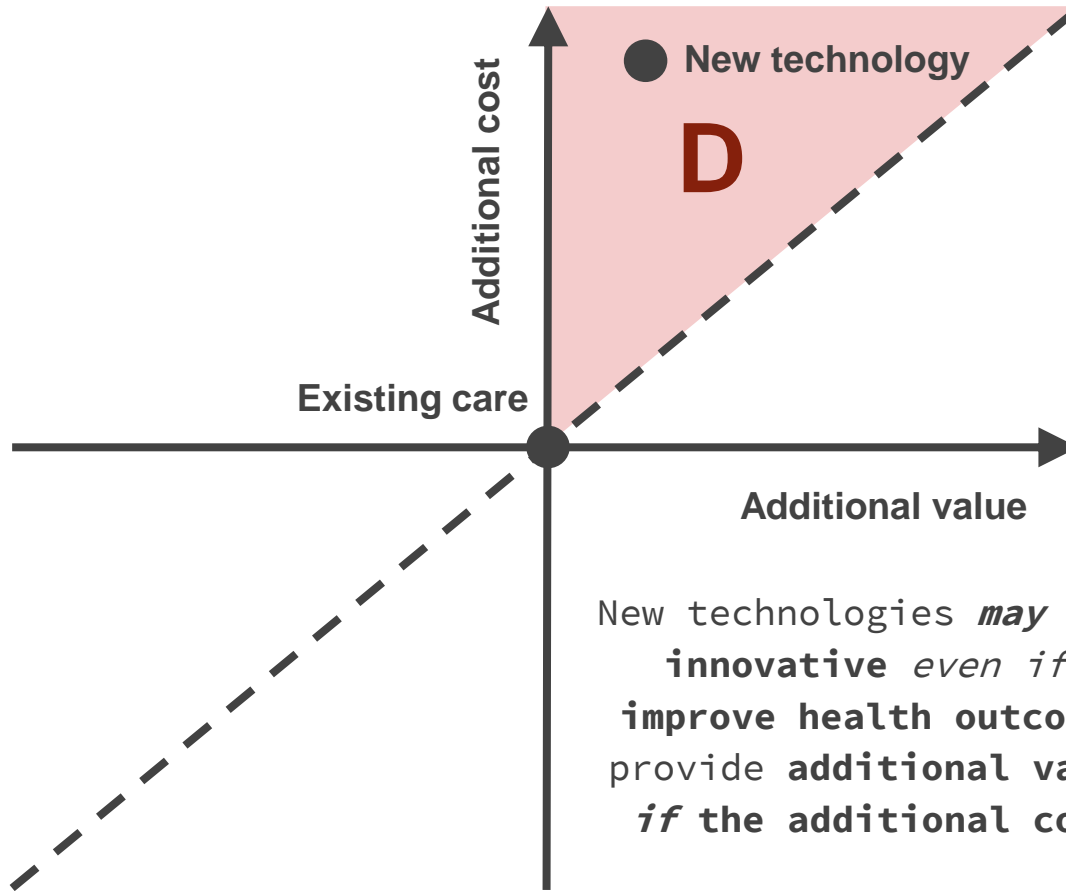


New technologies *may* be considered **innovative** *even if* they do not *directly* improve health outcomes or otherwise provide **additional value** to patients, *provided* they reduce costs

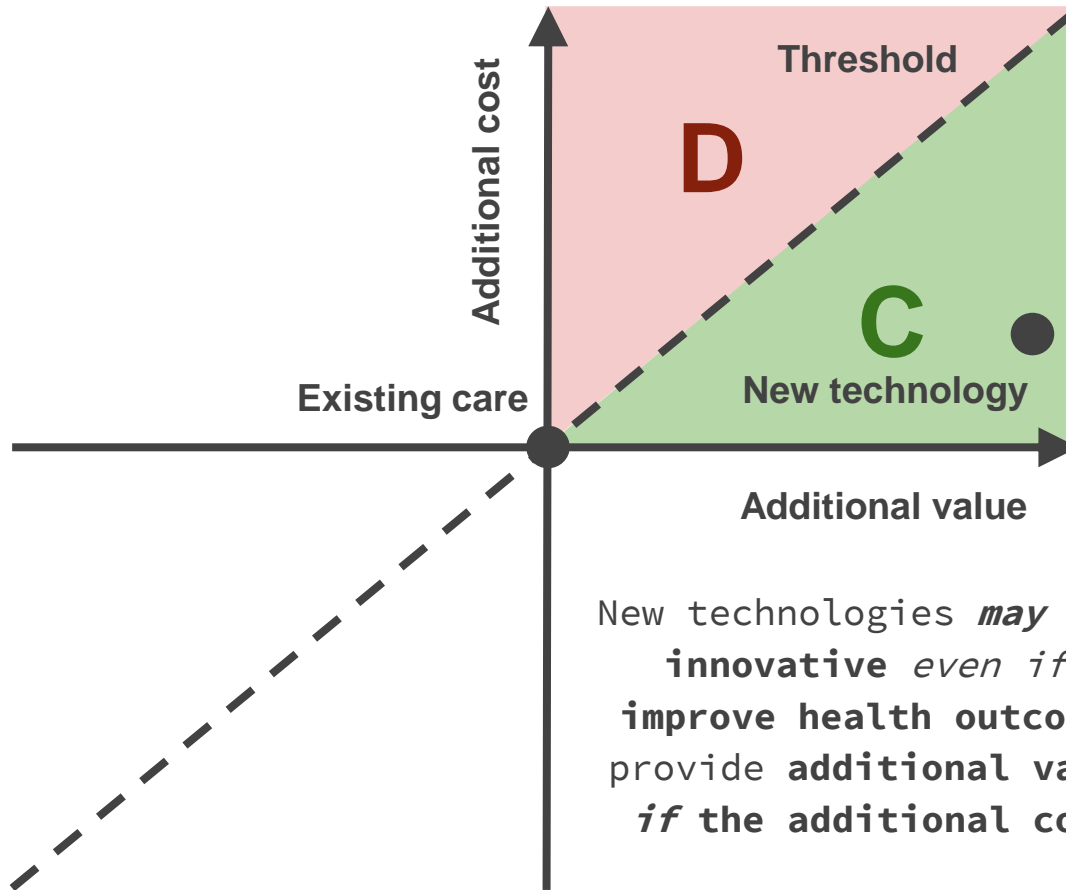


New technologies *may* be considered **innovative** *even if* they do not **directly** improve health outcomes or otherwise provide **additional value** to patients, *provided* they reduce costs

What are some of the “perils”?



New technologies *may not be considered innovative* even if they *directly improve health outcomes* or otherwise provide **additional value** to patients, *if the additional cost is too great*



New technologies *may not be considered innovative* even if they *directly improve health outcomes* or otherwise provide **additional value** to patients, *if the additional cost is too great*

Political and ethical issues

If policy makers wish to **incentivise development of innovative technologies** and **improve population health**, they must be **willing to reject technologies in region D** – those which provide **added value**, but which **cost so much** that they **displace *even more value*** elsewhere

Politically and ethically **difficult**, since **beneficiaries** of new technologies are generally **known**, but the **patients who bear the cost of their adoption** are generally **unknown**

Dynamic issues

For technologies to be considered **innovative**, they must provide **sufficient additional value**, at a **sufficiently low additional cost**, compared to all other **comparators**

But the set of comparators can **change over time**

New technologies may result in technologies *previously* considered innovative **no longer** considered innovative

Over time, policy makers need to **reassess** whether technologies are **still innovative** and be **prepared to**

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Case study: Oncotype DX vs Prosigna



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Cost-Effectiveness of the 21-Gene Assay for Guiding Adjuvant Chemotherapy Decisions in Early Breast Cancer

Mike Paulden, MSc^{1,*}, Jacob Franek, MHS², Ba³ Pham, MSc¹, Philippe L. Bedard, MD^{3,4}, Maureen Trudeau, MD^{5,6}, Murray Krahn, MD⁷

¹Health Economics and Technology Assessment (THETA) Collaborative, University of Toronto, Toronto, ON, Canada; ²Medical Advisory Secretariat, Ontario Ministry of Health and Long-Term Care, Toronto, ON, Canada (previous affiliation); Health Quality Ontario, Toronto, ON, Canada (current affiliation); ³Ontario Ministry of Health and Long-Term Care, Toronto, ON, Canada (previous affiliation); ⁴University Health Network - Princess Margaret Cancer Centre, Toronto, ON, Canada; ⁵Department of Medicine, Division of Medical Oncology and Hematology, University Health Network - Princess Margaret Cancer Centre, Toronto, ON, Canada; ⁶Department of Medicine, University of Toronto, Toronto, ON, Canada; ⁷Division of Medical Oncology and Hematology, Sunnybrook Health Sciences Centre - Odette Cancer Centre, Toronto, ON, Canada; ⁸Department of Medicine, University of Toronto, Toronto, ON, Canada

ABSTRACT

Objectives: Adjuvant chemotherapy decisions in early breast cancer are complex. The 21-gene assay can potentially aid such decisions, but costs US \$4175 per patient. Adjuvant Online is a freely available decision aid. We evaluate the cost-effectiveness of using the 21-gene assay in conjunction with Adjuvant Online, and of providing adjuvant chemotherapy conditional upon risk classification. **Methods:** Markov decision model simulated risk classification, probabilistic Markov decision model simulated risk classification, and the natural history of breast cancer in a hypothetical cohort of 50-year-old women with lymph node-negative, estrogen receptor- and/or progesterone receptor-positive, human epidermal growth factor receptor 2/HER2-negative early breast cancer. Cost-effectiveness was considered from an Ontario public-payer perspective by deriving the lifetime incremental cost (2012 Canadian dollars) by the quality-adjusted life-year (QALY) for each strategy, and the per quality-adjusted life-year cost-effective, assuming a willingness-to-pay of \$60,000 per QALY. **Results:** The 21-gene assay has an incremental cost per QALY in patients at low, intermediate, or high

Adjuvant Online! risk of \$22,440 (probability cost-effective 78.40%), \$2,526 (99.40%), or \$1,111 (99.82%), respectively in patients at low (high) 21-gene assay risk. Adjuvant chemotherapy increases (decreases) high 21-gene assay risk health outcomes. For patients at costs and worsens (improves) health outcomes. For patients at intermediate 21-gene assay risk and low, intermediate, or high intermediate 21-gene assay risk, chemotherapy has an incremental cost of \$44,088 (0.59%), \$1,776 (17.65%), or \$1,778 (82.93%), respectively. **Conclusions:** The 21-gene assay appears cost-effective, regardless of Adjuvant Online risk. Adjuvant chemotherapy appears cost-effective for patients at intermediate or high 21-gene assay risk, although this finding is uncertain in patients at intermediate 21-gene assay and low Adjuvant Online risk.

Keywords: breast cancer, chemotherapy, cost-effectiveness analysis, decision making, pharmacogenetics.

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Introduction

Adjuvant chemotherapy decisions for women with early stage breast cancer are complex. At present, decisions are informed by clinical judgment, often supplemented through the use of Adjuvant Online [1]. Adjuvant Online is a free online diagnostic tool that estimates a woman's risk of breast cancer-specific mortality (BCSM) or relapse on the basis of information entered by the physician, including the woman's age, comorbidities, tumor size, estrogen receptor status, number of involved lymph nodes, and progesterone receptor status. Results can be categorized as proposed course of treatment [1]. Results can be categorized as "low" (BCSM < 5%), "intermediate" (5% < BCSM < 17%), or "high" (BCSM > 17%) risk [1]. A validation study of Adjuvant Online has found a high degree of correlation between predicted and observed survival [4].

An alternative predictive tool has recently become available. The 21-gene assay (Endocept Dx; Genomic Health, Redwood City, CA) quantifies the expression of 21 genes in breast cancer tissue by

performing reverse transcription polymerase chain reaction on formalin-fixed paraffin-embedded tumor blocks that are obtained during initial surgery. Results are summarized by a "Recurrence Score" (RS) between 0 and 100, with scores categorized as "low" (RS < 10), "intermediate" (10 < RS < 30), or "high" (RS > 30) risk [5]. RS < 10, "intermediate" (10 < RS < 30), or "high" (RS > 30) risk [5]. It has been validated both in women with estrogen receptor-positive early stage breast cancer that is lymph node-negative, and in women with estrogen receptor-positive breast cancer that is lymph node-positive, as a means to predict the risk of distant recurrence and magnitude of chemotherapy benefit when added to endocrine therapy [6–9].

As of April 2012, the 21-gene assay cost US \$4175 per patient [10]. Its cost-effectiveness is therefore a matter of considerable policy interest. There is also uncertainty as to the clinical-policy interest of providing adjuvant chemotherapy, and cost-effectiveness of providing adjuvant chemotherapy, and cost-effectiveness analysis that comprehensively addresses both of these issues. An earlier version of our analysis formed part of a

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<http://dx.doi.org/10.1016/j.jval.2013.03.1625>

Paulden M, Franek J, Pham B, Bedard PL, Trudeau M, Krahn M. Cost-effectiveness of the 21-gene assay for guiding adjuvant chemotherapy decisions in early breast cancer. Value Health. 2013;16: 729–739. doi:10.1016/j.jval.2013.03.1625

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Economic evaluation of Prosigna and Oncotype DX for
guiding adjuvant chemotherapy treatment in Alberta

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Not to be circulated without authors' permission.

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Tel: 780 492 4202

Razavilar N, Hollman C, Paulden M, Hugh J, McCabe C. Economic evaluation of Prosigna and Oncotype DX for guiding adjuvant chemotherapy treatment in Alberta. Report produced for Alberta Health Services by McCabe Research Group, Faculty of Medicine and Dentistry, University of Alberta. May 2016.

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Oncotype DX appeared cost-effective vs usual care...

Ontario-based economic analysis published in 2013

Compared to **usual care** (adjuvant chemotherapy guided using Adjuvant! Online or clinical judgement) **Oncotype DX** was found to **increase costs** (\$3,505 per patient) and **improve health outcomes** (0.22 QALYs per patient), resulting in an incremental cost-effectiveness ratio (ICER) of **\$15,932 per QALY**

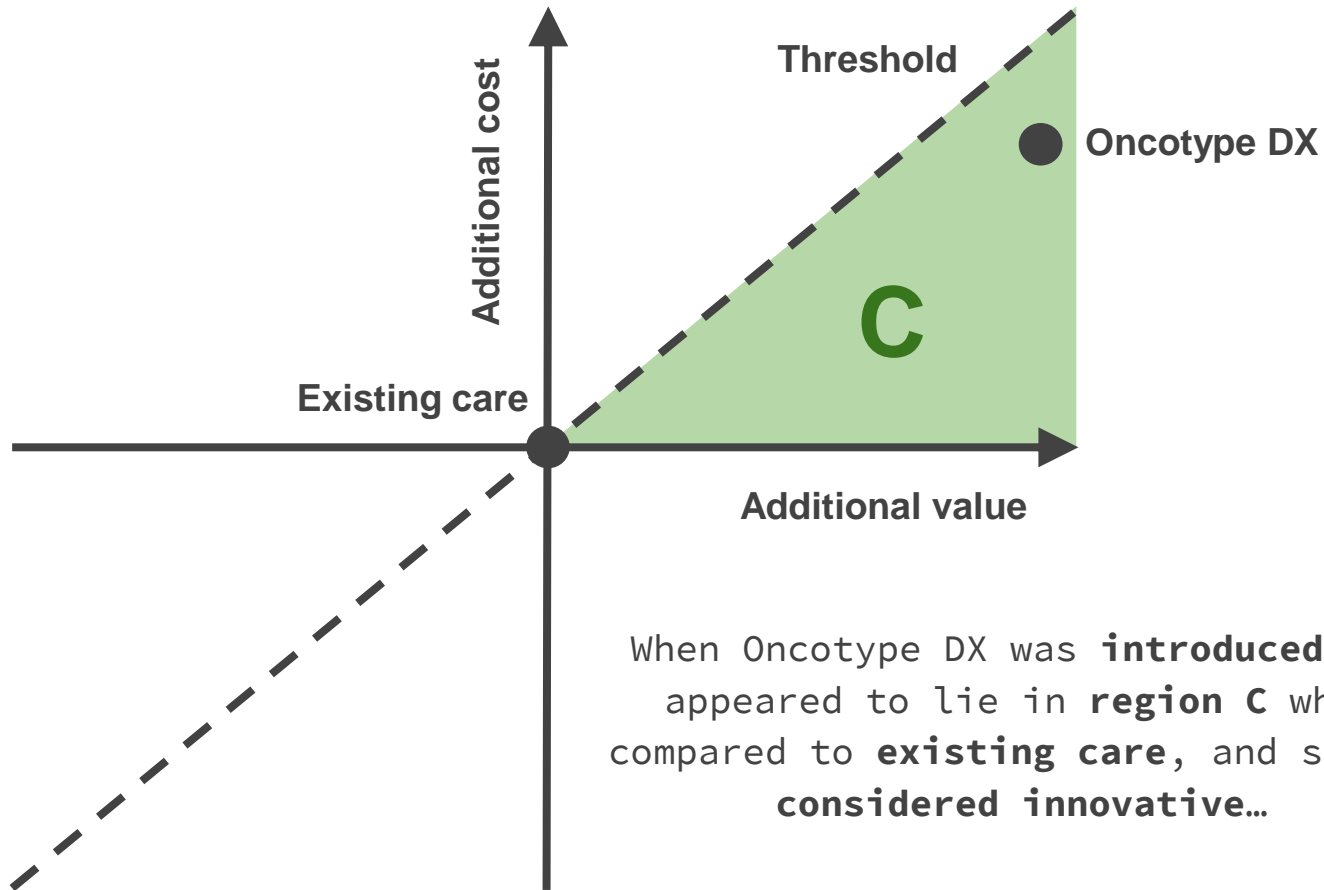
Compared to a conventional cost-effectiveness **threshold** of \$50,000 per QALY, **Oncotype DX appeared cost-effective**

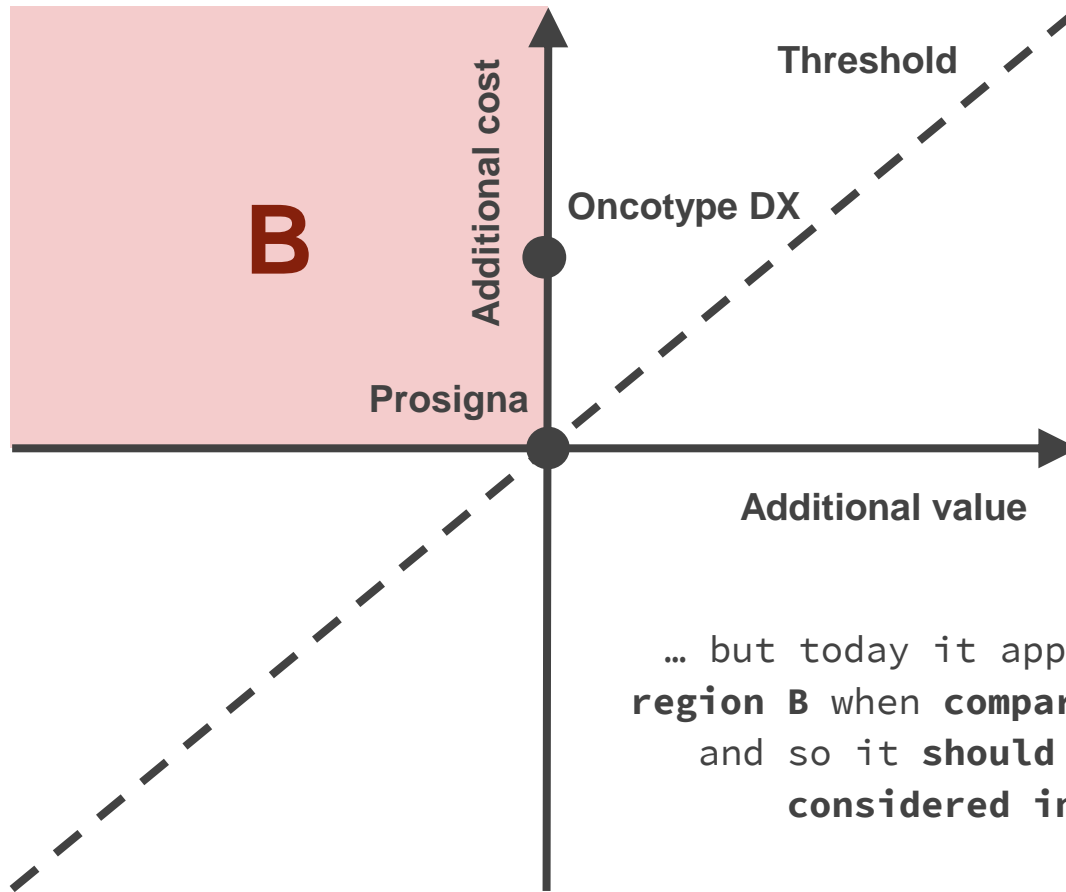
... but Oncotype DX did not appear cost-effective vs Prosigna

Alberta-based economic analysis reported in 2016

Compared to **Prosigna**, **Oncotype DX** was found to **increase costs** (\$1,240 per patient) and **worsen health outcomes** (0.003629 QALYs per patient), resulting in Oncotype DX appearing **dominated** (higher costs and worse outcomes)

It follows that **Oncotype DX is not cost-effective** compared to Prosigna





... but today it appears to lie in **region B** when compared to Prosigna, and so it **should no longer be considered innovative**

Policy response

Prosigna has **further benefits** compared to Oncotype DX:

Fewer intermediate (**indeterminate**) results

Superior prediction of **long-term treatment benefit** (TransATAC study)

Repatriation of **out-of-province testing** (in-house vs shipped to US)

Yet **barriers exist** to its implementation in practice

Lack of **life-cycle HTA processes** (Oncotype DX not re-assessed)

Appropriate utilization and **physician compliance** (reluctance to disinvest in Oncotype DX and adopt Prosigna in its place)

Conclusion

Conclusions

Health interventions or technologies might be considered **“innovative”** if they **improve population health outcomes**

If any **additional costs** are borne by the health system, this results in **forgone health care** for other patients

It follows that an **expensive** new technology *might not be considered innovative* - *even if it adds value* - since it *may displace more population health than it provides*

A technology that **reduces health expenditures** *might be*

considered innovative, even if it has no additional value

Conclusions

It also follows that technologies *previously considered innovative* may **no longer appear innovative** once cheaper and/or more valuable **comparators** become available

Continuing to use older technologies when **more innovative alternatives** are available may **diminish population health**

Policy makers should adopt a **life-cycle approach to HTA** and be **willing to disinvest in existing technologies**

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Any questions?