

# Precision Medicine 2030

*FDA-MCERSI Workshop on Application of Artificial Intelligence and Machine Learning for Precision Medicine*

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**This presentation reflects the views of the speaker and should not be construed to represent FDA's policies**

# Agenda

***Where can AI methodologies be applied to support targeted drug development and precision patient care?***

- Landscape
- Opportunities
- Challenges



# A Brief History of Medicines



This collage illustrates the history of medicine through various stages and sources:

- Natural Sources:** Images of medicinal plants like opium poppy and willow bark, and a vial of Vaccinia Virus.
- Historical Medicines:** Bottles of Laudanum (opium tincture), Chloral Hydrate (sedative), and Ether (anesthetic).
- Chemical Structures:** A diagram of a cyclic peptide chain with labels for amino group (NH<sub>2</sub>), carboxyl group (COOH), alpha chain, and beta chain.
- Pharmaceuticals:** A bottle of Thorazine (75 mg) and a box of Penicillin mould, credited to Professor Alexander Fleming (1928).
- Modern Medicine:** A collection of colorful pills, vials of antibiotics, and a syringe.
- Biological Processes:** A 3D model of a protein-DNA complex and a diagram of a cell membrane with various receptors.
- Other:** A bottle of Aspirin and a box of Penicillin mould.

# “Precision Medicine”

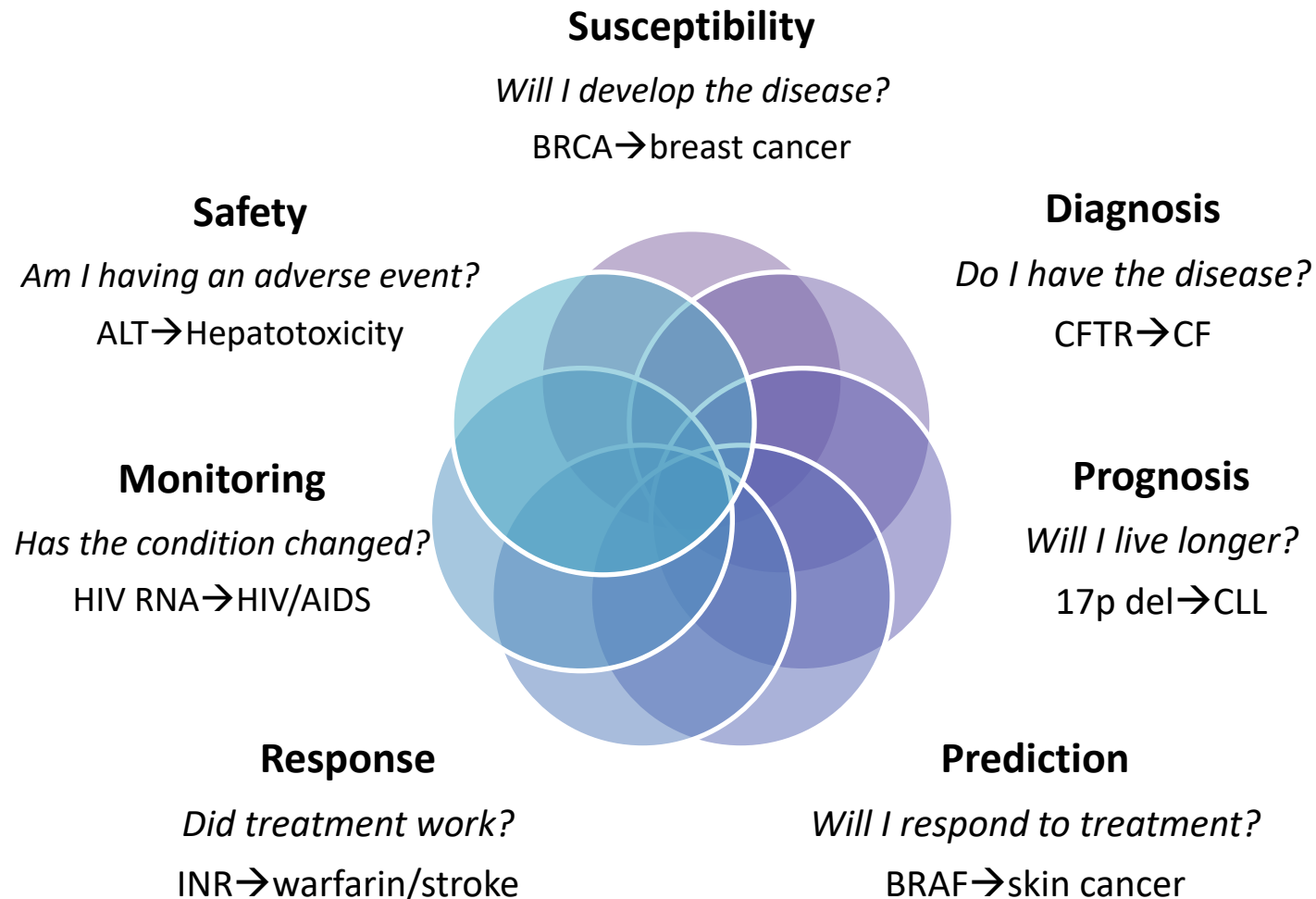
***An innovative approach to tailoring disease prevention and treatment that takes into account differences in people's genes, environments, and lifestyles<sup>1</sup>***

- Targeted therapies: Drugs or biologics intended for use with a genomic, proteomic, or other biomarker/tool that
  - Identifies patients who are eligible for treatment
  - Aids in determining the appropriate dosage
  - Allows for monitoring of response to individualize therapy<sup>2</sup>

<sup>1</sup> [Precision Medicine | FDA](https://www.fda.gov/medical-devices/in-vitro-diagnostics/precision-medicine): <https://www.fda.gov/medical-devices/in-vitro-diagnostics/precision-medicine>

<sup>2</sup> Unofficial definition

# What Can Biomarkers Tell Us?



# Evolution of Precision Medicines

## Pharmacogenetics

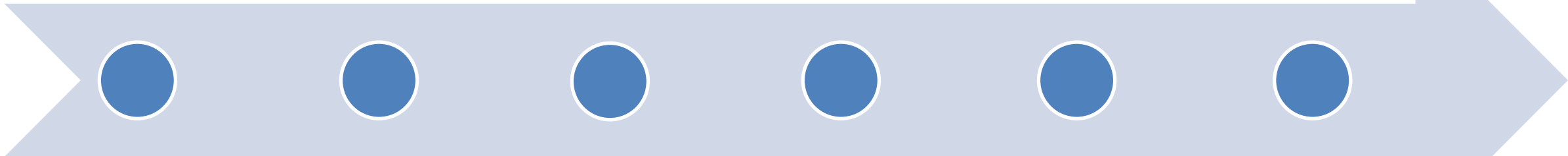
- Retrofit biomarkers of drug metabolism and response to guide dosing, patient selection, monitoring
  - *abacavir/HLA-B\*57:01*
  - *clopidogrel/CYP2C19*
  - *warfarin/VKORC1-CYP2C9*

## Reverse Translation

- Genetically informed target selection for new drug development
  - *alirocumab/PCSK9*
  - *evolocumab/PCSK9*
  - *setmelanotide/MC4R*

## Precision Medicines v2

- Tissue agnostic and other omic approaches for patient selection
  - *pembrolizumab/TMB*
  - *larotrectinib/NTRK*
  - *olaparib/HRR*



## Precision Medicines v1

- Prospectively codeveloped drugs and biomarkers for patient selection
  - *trastuzumab/HER2*
  - *afatinib/EGFR*
  - *ivacaftor/CFTR*

## Genetically Targeted Therapies

- RNA interference and other modalities that target the genome
  - *nusinersen/SMN2*
  - *patisiran/TTR*
  - *lumasiran/HAO-1*

## Individualized Therapies

- Production of drug designed to treat a very small number of patients (1 or 2)
  - *milasen/MFSD8*

# Pathways to Integrate Technology into Drug Development and Practice



Note: These pathways do not exist in isolation and many times parallel efforts are underway within or between pathways. All share common core concepts, are data-driven, and involve regulatory assessment and outcomes based on the available data.

# Medical Products and Biomarkers



## Selected CDER Drug Approvals in 2022 (of 37 new molecular entities)

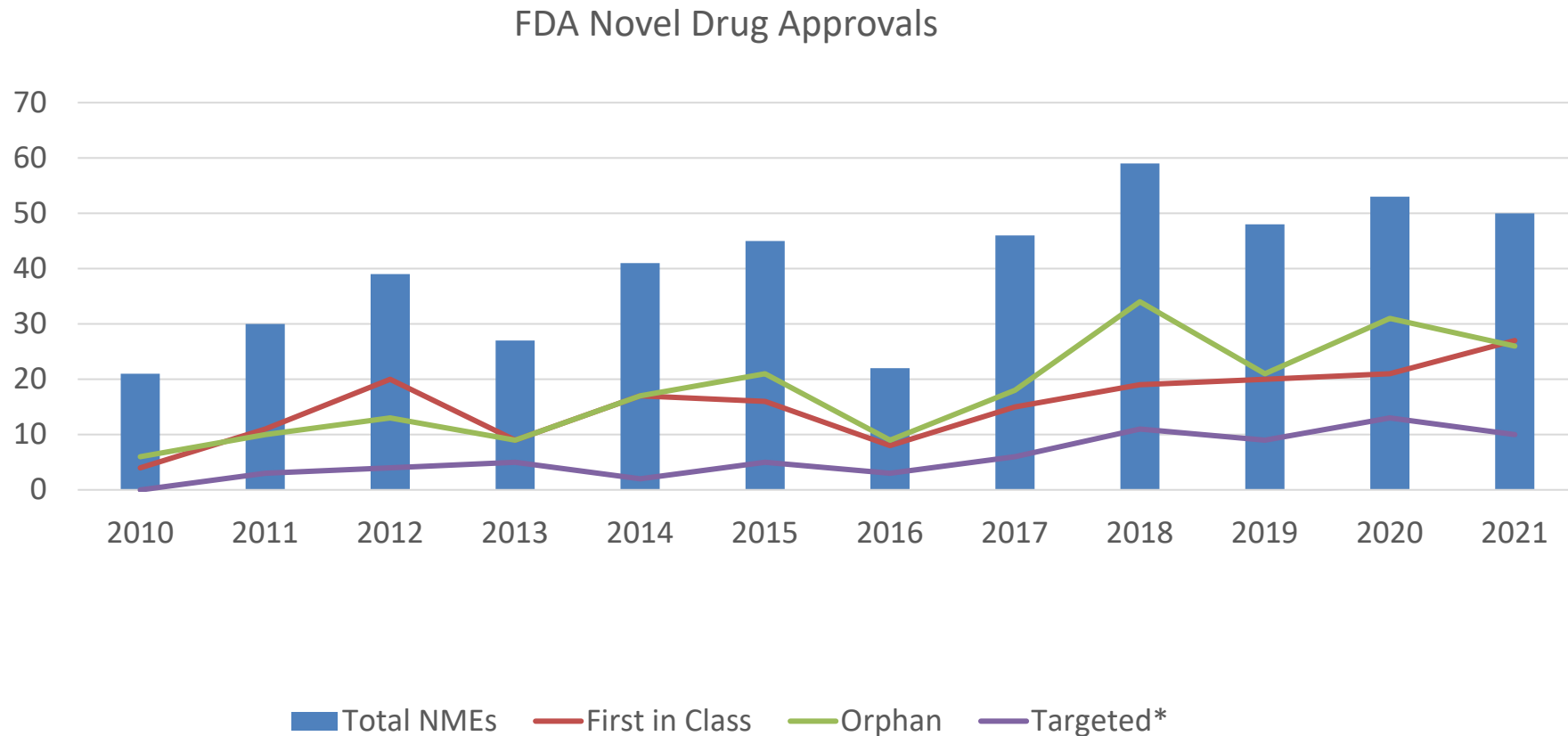
Drug	Disease or Condition	Biomarker	Use
Tebentafusp-tebn	Uveal melanoma	<i>HLA-A*02:01</i> †	Selection (subset)
Lutetium Lu 177 vipivotide tetraxetan	Prostate cancer	Prostate-specific membrane antigen	Selection (subset)
Futibatinib	Cholangiocarcinoma	<i>FGFR2</i> gene fusions or other rearrangements	Selection (subset)
Mirvetuximab soravtansine-gynx	Ovarian, fallopian tube, primary peritoneal cancer	Folate receptor alpha†	Selection (subset)
Olutasidenib	Acute myeloid leukemia	<i>IDH1</i> mutations†	Selection (subset)
Adagrasib	Non-small cell lung cancer	<i>KRAS</i> G12C mutation†	Selection (subset)
Abrocitinib	Atopic dermatitis	<i>CYP2C19</i> variants	Dosage
Mitapivat	Pyruvate kinase deficiency	<i>PKLR</i> variants	Selection (diagnosis)
Vutrisiran	Transthyretin amyloidosis polyneuropathy	<i>TTR</i> variants	Selection (diagnosis)
Olipudase alfa-rpcp	Acid sphingomyelinase deficiency	<i>SMPD1</i> variants	Selection (diagnosis)

The table includes 1) new molecular entities (NMEs), and selected supplements to previously approved drugs, for which the indication is restricted to a subset of patients with the disease/condition, and 2) NMEs for non-oncologic genetic diseases that mechanistically target the underlying pathophysiology

† Companion diagnostic



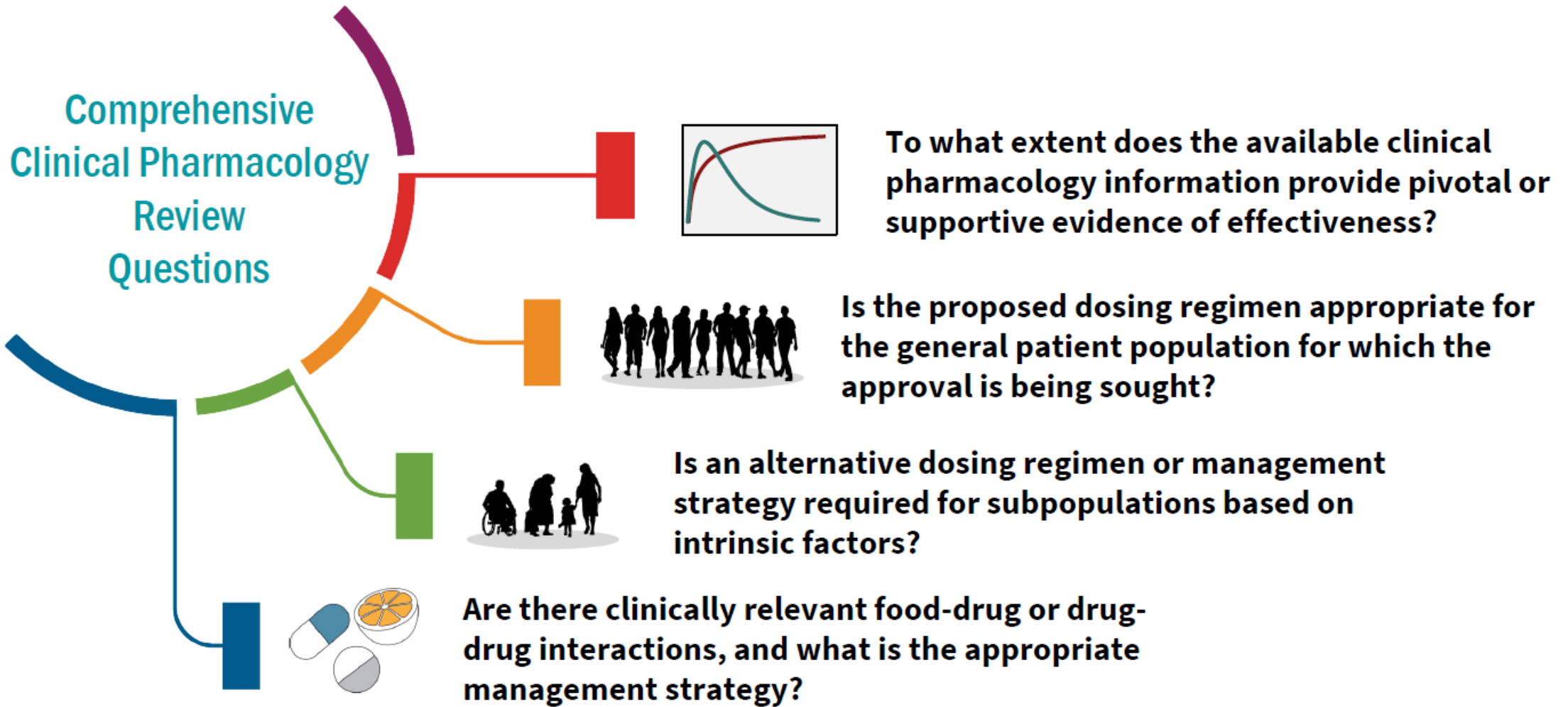
# Trends in Targeted Drug Development



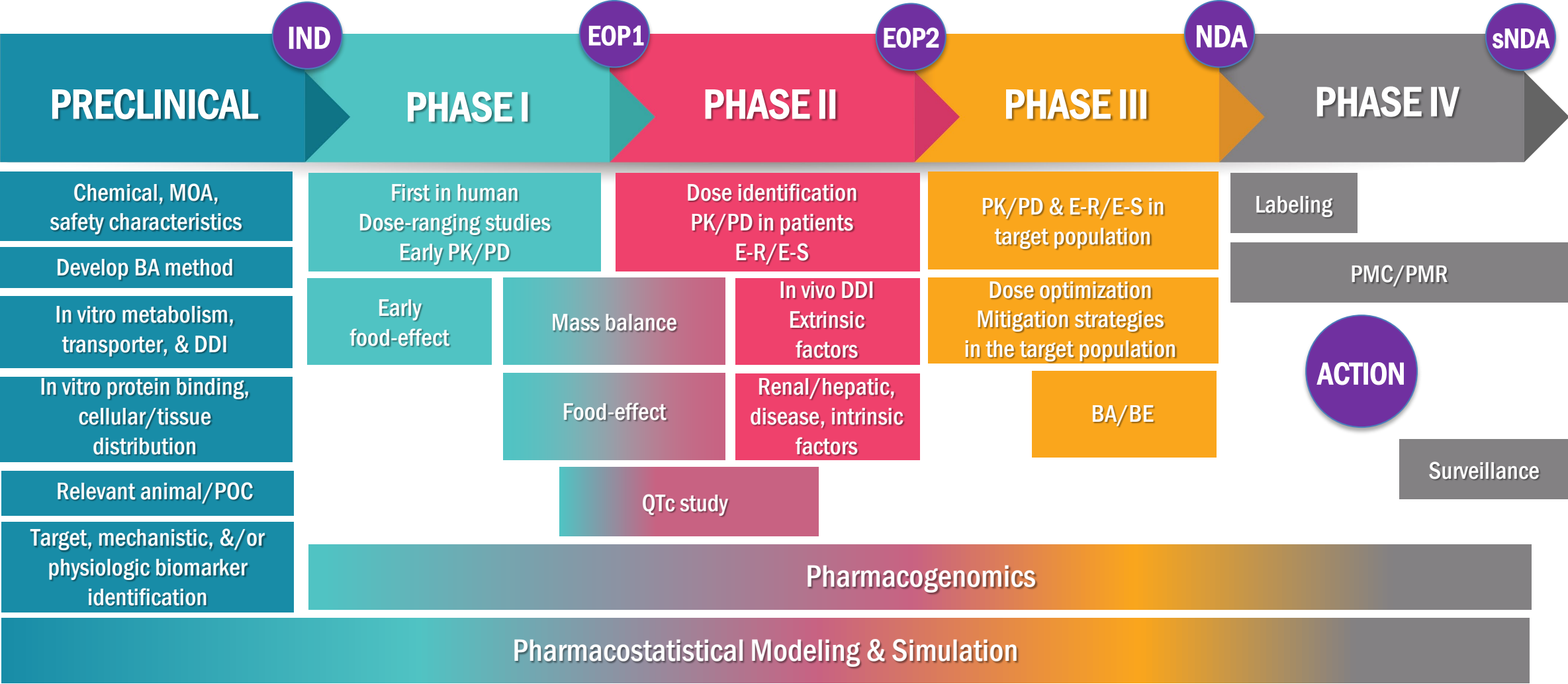
\*Targeted drugs for the purpose of this figure are new molecular entities (NMEs) for which the initially approved indication is restricted to a subset of patients who are identified through molecular testing (and excludes genotype-based dosing and infectious disease subsets).

# **OPPORTUNITIES AND CHALLENGES**

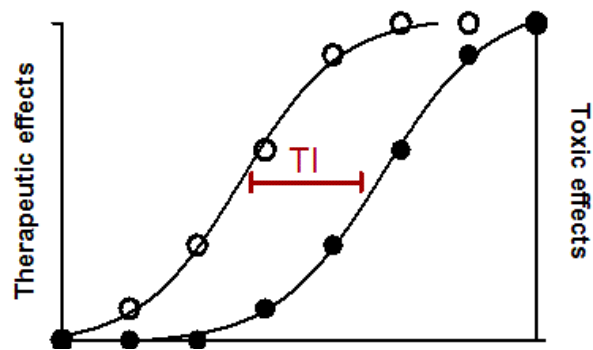
# Clinical Pharmacology Fundamentals



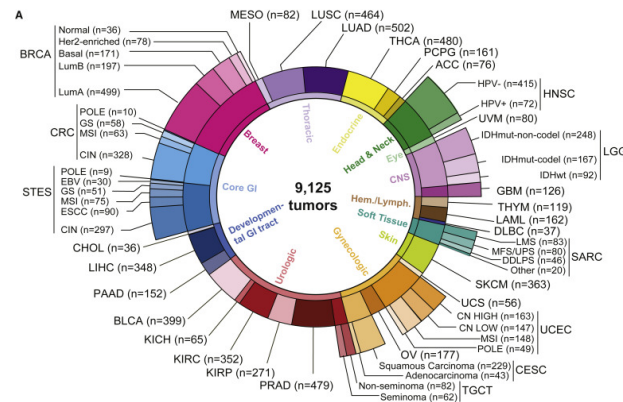
# Drug Development and Regulation: Progressive Reduction of Uncertainty



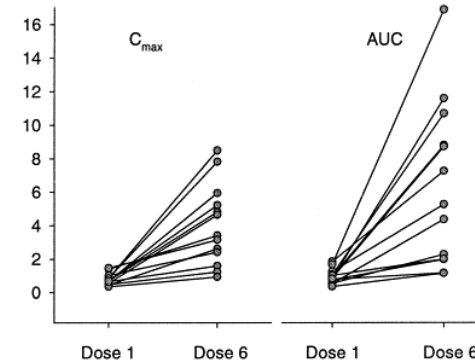
# Resolving Variability in Drug Response



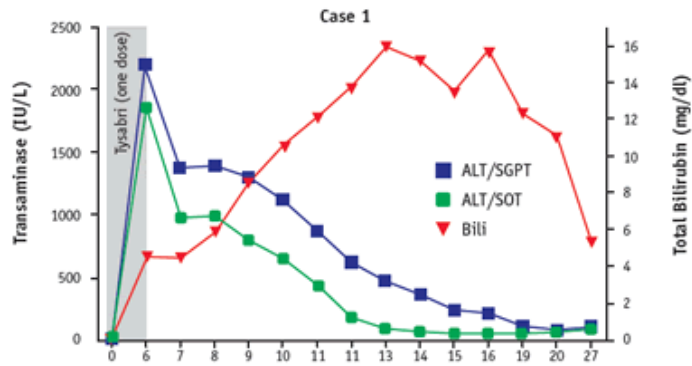
Therapeutic Index



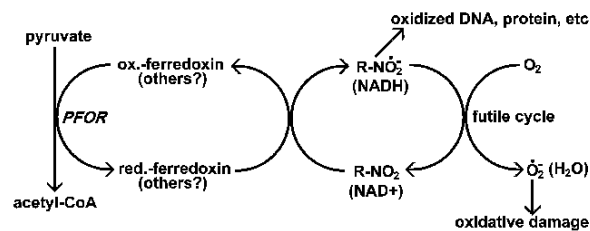
Pathobiology



Variability



Safety



Mechanism of Action

Subgroup	n Patients Meeting Primary Endpoint / N of Patients (%)		Mean percentage difference (95% CI)
	Mavacantan	Placebo	
<b>Age, years</b>			
<49	10/27 (37%)	6/25 (24%)	13% (-11.7 to 37.8)
50-64	21/51 (41%)	13/63 (21%)	21% (3.7 to 37.3)
≥65	14/45 (31%)	3/40 (8%)	24% (7.8 to 39.4)
<b>Sex</b>			
Female	19/57 (33%)	5/45 (11%)	22% (6.9 to 37.5)
Male	26/66 (39%)	17/63 (21%)	19% (4.3 to 33.6)
<b>Body-mass index, kg/m<sup>2</sup></b>			
<30	35/77 (46%)	16/77 (21%)	25% (10.3 to 39.0)
≥30	10/48 (22%)	6/51 (12%)	10% (-4.9 to 24.8)
<b>LEUF at baseline</b>			
<75%	25/69 (36%)	11/70 (16%)	21% (6.3 to 34.7)
≥75%	20/54 (37%)	11/58 (19%)	18% (1.7 to 34.4)
<b>NHYA class at baseline</b>			
II	29/88 (33%)	16/95 (17%)	16% (3.7 to 28.5)
III	16/35 (46%)	6/33 (18%)	28% (6.4 to 48.6)
<b>β blocker usage at baseline</b>			
Yes	28/94 (30%)	20/95 (21%)	9% (-3.6 to 21.1)
No	17/29 (59%)	2/33 (6%)	53% (32.9 to 72.2)
<b>Type of exercise testing</b>			
Bicycle	15/55 (27%)	11/58 (19%)	8% (-7.2 to 23.8)
Treadmill	30/68 (44%)	11/70 (16%)	26% (13.8 to 43.0)
<b>NT-proBNP at baseline, ng/L</b>			
≤ median of 710 ng/L	18/55 (33%)	13/68 (19%)	14% (-1.9 to 29.1)
> median of 710 ng/L	24/65 (37%)	9/58 (16%)	21% (6.4 to 36.4)

Heterogeneity

# Advancing Precision Medicine Through AI



## Data sources

- Clinical trials
- Observational studies
- Health records
- Claims

## Sampling

- Fluids
- Cells
- Tissues
- Sounds
- Air
- Images
- Video
- Audio
- Pressures
- Electrical signals
- Outcomes

## Measures

- Anthropometrics
- Protein expression
- Gene variants
- Movements
- Colors
- Voltages
- Symptoms

## Impact

- Novel drug targets
- Risk assessment
- Diagnostic criteria
- Prognosis
- Dose selection
- Response prediction
- Monitoring
- Clinical decision support

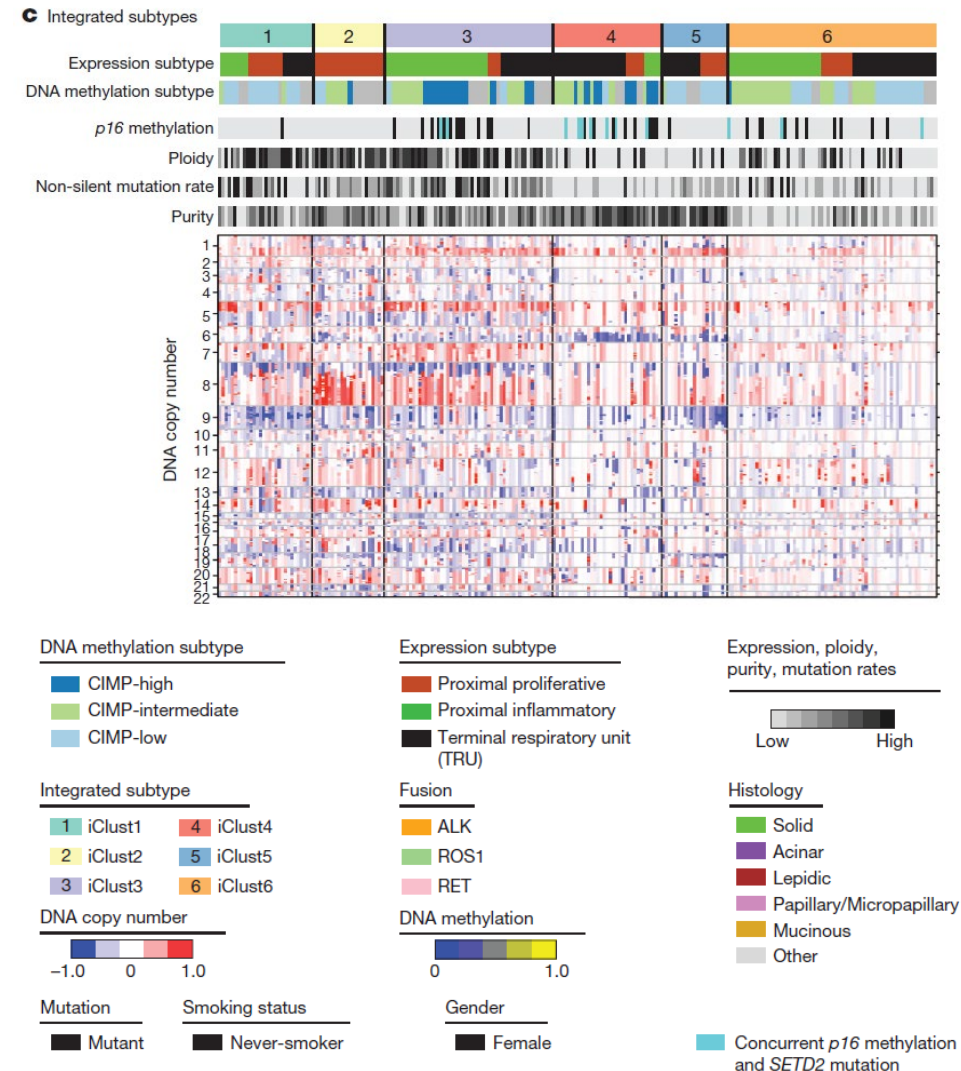
# Multidimensional Profiling

## Lung Adenocarcinoma



Drug	NSCLC Molecular Subgroup
Mobocertinib	EGFR exon 20 insertion mutations
Sotorasib	KRAS G12C mutation
Amivantamab-vmjw	EGFR exon 20 insertion mutations
Lorlatinib	ALK expression
Cemiplimab-rwlc	PD-L1 high (no EGFR/ALK/ROS1)
Tepotinib	MET exon 14 skipping alterations
Osimertinib	EGFR exon 19 deletions, L858R, T790M
Pralsetinib	RET fusions
Ramucirumab	EGFR exon 19 deletions, L858R
Erlotinib	EGFR exon 19 deletions, L858R
Nivolumab	PD-L1 high (no EGFR/ALK)
Ipilimumab	PD-L1 high (no EGFR/ALK)
Brigatinib	ALK rearrangements
Atezolizumab	PD-L1 high
Capmatinib	MET exon 14 skipping mutations
Selpercatinib	RET fusions

(selected approvals)



TCGA Research Network. Nature 2014.

# Multidimensional Profiling

## Anticoagulant Dose Prediction



### DOSAGE AND ADMINISTRATION

The appropriate initial dosing of COUMADIN varies widely for different patients. Not all factors responsible for warfarin dose variability are known, and the initial dose is influenced by:

- Clinical factors including age, race, body weight, sex, concomitant medications, and comorbidities
- Genetic factors (CYP2C9 and VKORC1 genotypes) [see [CLINICAL PHARMACOLOGY \(12.5\)](#)]

Select the initial dose based on the expected maintenance dose, taking into account the above factors. Modify this dose based on consideration of patient-specific clinical factors. Consider lower initial and maintenance doses for elderly and/or debilitated patients and in Asian patients [see [USE IN SPECIFIC POPULATIONS \(8.5\)](#) and [CLINICAL PHARMACOLOGY \(12.3\)](#)].

**Table 1: Three Ranges of Expected Maintenance COUMADIN Daily Doses Based on CYP2C9 and VKORC1 Genotypes<sup>†</sup>**

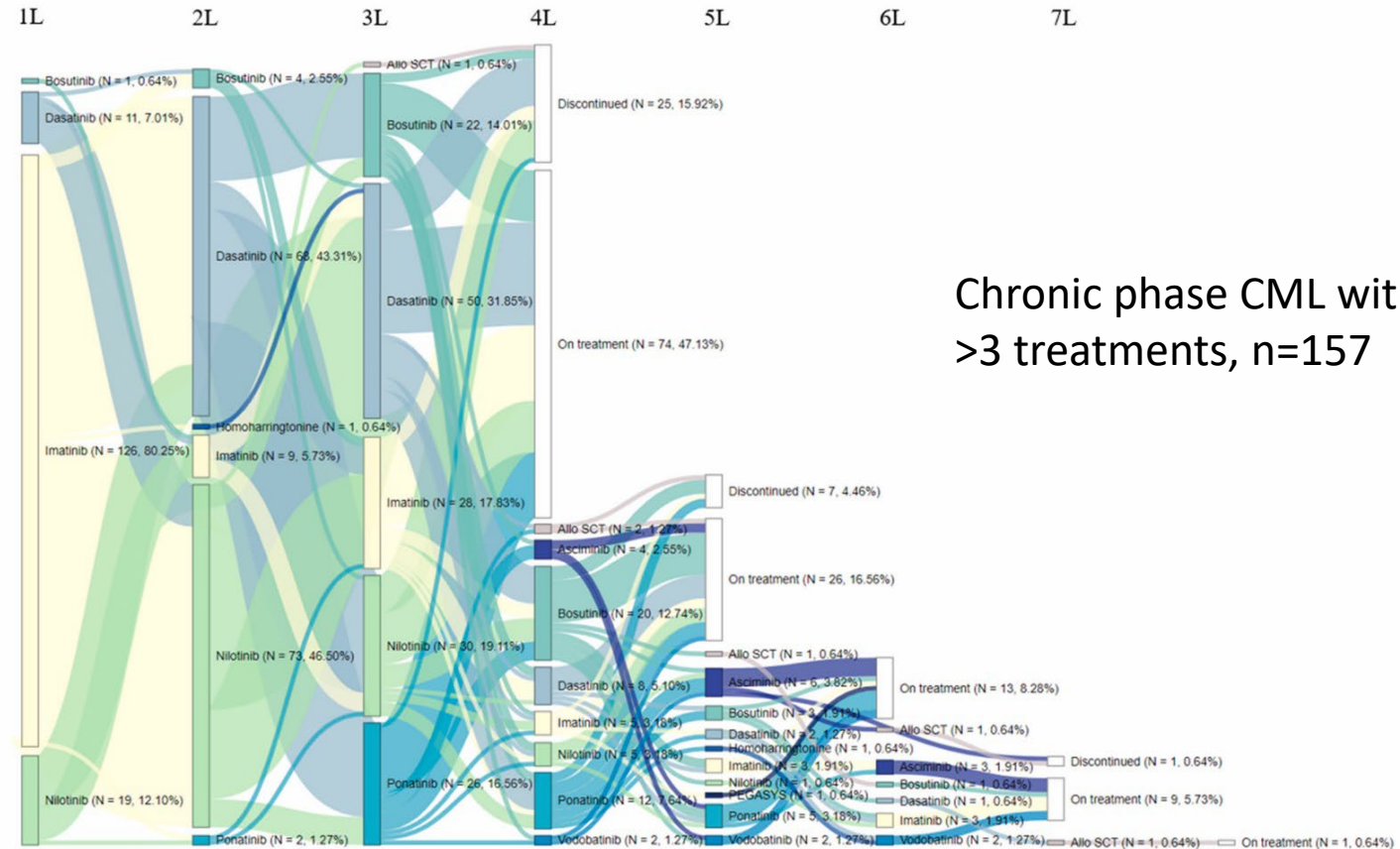
VKORC1	CYP2C9					
	*1/*1	*1/*2	*1/*3	*2/*2	*2/*3	*3/*3
GG	5-7 mg	5-7 mg	3-4 mg	3-4 mg	3-4 mg	0.5-2 mg
AG	5-7 mg	3-4 mg	3-4 mg	3-4 mg	0.5-2 mg	0.5-2 mg
AA	3-4 mg	3-4 mg	0.5-2 mg	0.5-2 mg	0.5-2 mg	0.5-2 mg

<sup>†</sup>Ranges are derived from multiple published clinical studies. VKORC1 -1639G>A (rs9923231) variant is used in this table. Other co-inherited VKORC1 variants may also be important determinants of warfarin dose.

The screenshot shows the 'WARFARINDOSING' website interface. The main heading is 'WARFARINDOSING' with the URL 'www.WarfarinDosing.org'. On the left is a navigation menu with links: Warfarin Dosing, Clinical Trial, Outcomes, Hemorrhage Risk, Patient Education, Contact Us, References, Glossary, and About Us. The main content area is divided into two sections: 'Required Patient Information' and 'Genetic Information'. The 'Required Patient Information' section includes fields for Age, Sex, Ethnicity, Race, Weight (lbs or kgs), Height (feet and inches or cms), Smokes, Liver Disease, Indication, Baseline INR, Target INR, Amiodarone/Cordarone® Dose, Statin/HMG CoA Reductase Inhibitor, Any azole (eg. Fluconazole), and Sulfamethoxazole/Septtra/Bactrim/Cotrim/Sulfatrim. The 'Genetic Information' section includes dropdown menus for VKORC1-1639/3673, CYP4F2 V433M, GGXC rs11676382, CYP2C9\*2, CYP2C9\*3, CYP2C9\*5, and CYP2C9\*6. At the bottom, there is a checkbox for 'Accept Terms of Use' and a red button labeled '> ESTIMATE WARFARIN DOSE'. A footer note indicates 'User: Patient: Version 3.0 Build : May 14, 2016'.



# Where Do AI Approaches Fit?



Notes: “Discontinued” indicates that patients discontinued their last line of treatment; “On treatment” indicates that patients were still on their last line of treatment at the time of data collection.

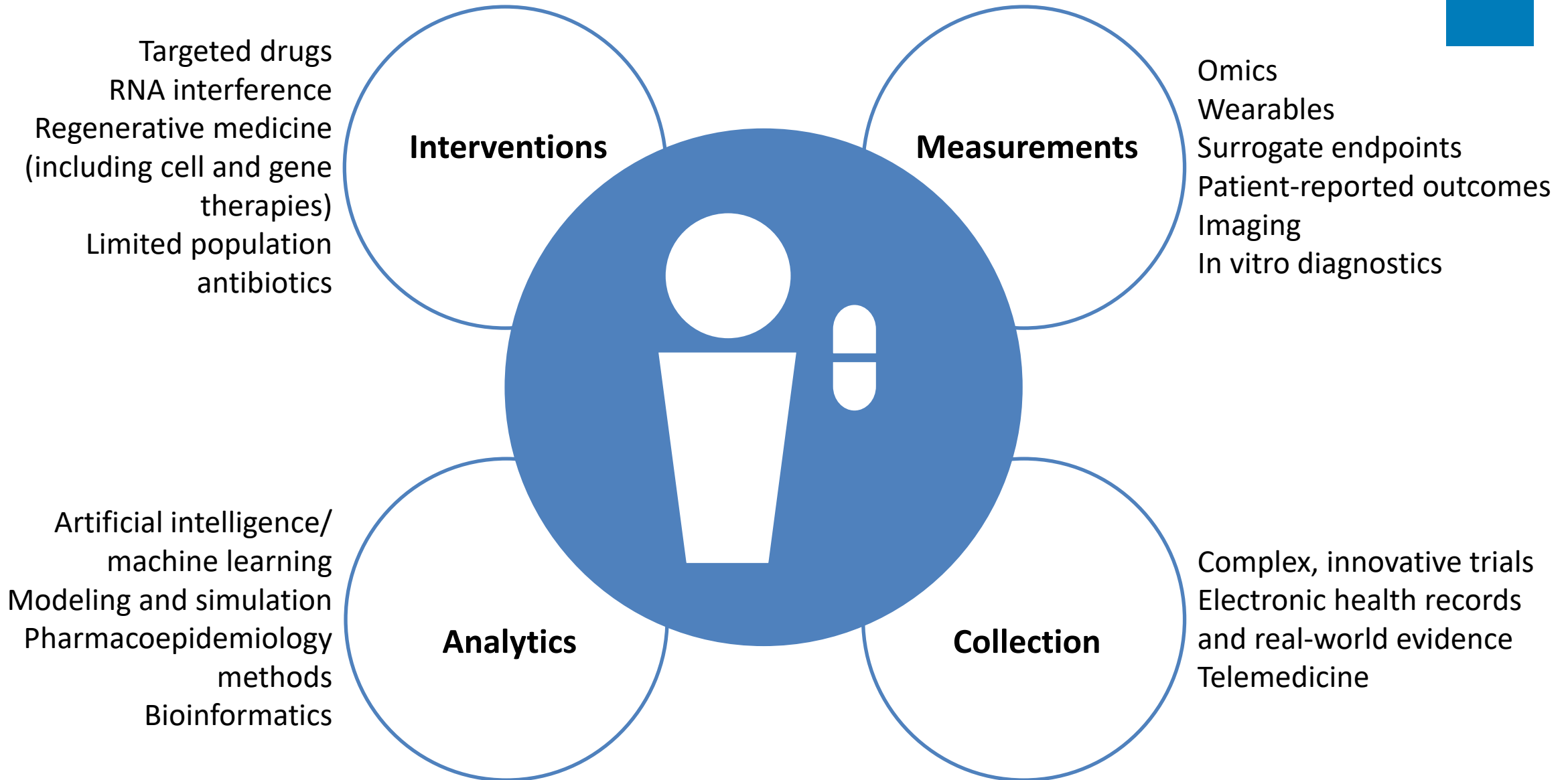
Abbreviations: 1L: first-line; 2L: second-line; 3L: third-line; 4L: fourth-line; 5L: fifth-line; 6L: sixth-line; 7L: seventh-line; Allo-SCT: allogeneic stem cell transplant.

# How Can We Advance Use of AI Approaches in Precision Medicine?\*



<b>Barriers</b>	Lack of standardization and interoperability
	Data privacy and security
	Limited data availability and quality
	Resistance to change
	Lack of regulatory guidance
<b>Solutions</b>	Data integration and standardization
	Improved algorithms and models
	Collaboration and partnerships
	Clinical validation and implementation
	Investment in AI research

\* According to ChatGPT  
OpenAI. "ChatGPT". *Welcome to ChatGPT*, [chat.openai.com/](https://chat.openai.com/).



# Summary

- Precision medicine approaches rely on technologies that improve mechanistic understanding of disease and drug response
- AI and complex data analytics can augment use of existing tools to further reduce biological complexity and optimize drug development
- AI tools in practice can help resolve multiple factors to support individualized therapeutic decision making

