

Präzision-Onkologie: Erfahrung für die ersten 1000 fortgeschrittenen Krebspatienten und was wir gelernt haben

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UNIVERSITÄT
BERN



Weill Cornell Medicine
Caryl and Israel Englander
Institute for Precision Medicine



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Weill Cornell Medicine, New York, NY, U.S.A.

 @MarkARubin1

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Precision Medicine Enters into the National Health Care Dialogue

Obama to Request Research Funding for Treatments Tailored to Patients' DNA

By ROBERT PEAR

WASHINGTON — President Obama will seek hundreds of millions of dollars for a new initiative to develop medical treatments tailored to individual patients, administration officials say.

The proposal, which President Obama described in greater detail in his budget in the coming weeks. The effort is likely to be one of the most significant of his administration.

"This is an incredible area of promise," said Senator Bill Cassidy, Republican of Louisiana and a gastroenterologist. "There will be bipartisan support."

Mr. Obama called it precision medicine, but the terms "personalized medicine" and "individualized medicine" are also widely used to describe the evolving field in which, for example, a doctor prescribes a medication that targets a specific mutation in a patient's genes.

The money would support biomedical research at the National Institutes of Health and the regulation of diagnostic tests by the Food and Drug Administration, officials at the two agencies said.

The tests analyze the DNA in normal or diseased tissue. Doctors use that information to identify patients with cancer or other diseases who are most like-

ly to benefit from a particular treatment — and those who would be harmed or not respond at all.

"In some patients with cystic fibrosis, a mutation in a gene called CFTR was discovered by a team that included Dr. Francis S. Collins, who is now director of the National Institutes of Health. The F.D.A. has approved a drug for patients with a genetic mutation responsible for some cases of the disease, which clogs the lungs with thick, sticky mucus.

A patient taking that drug, William Elder Jr., a 27-year-old medical student in Ohio, was a guest of Michelle Obama at the State of the Union speech.

Representative Fred Upton, Republican of Michigan and chairman of the Energy and Commerce Committee, and Representative Diana DeGette, a Colorado Democrat who is on the committee, welcomed Mr. Obama's proposal. After holding hearings and round-table discussions last year, they said they were drafting a bill to encourage biomedical innovations, including personalized medicine.

As a senator in 2006 and 2007, Mr. Obama offered a bill to do just that — the Genomics and Personalized Medicine Act. Senator Richard M. Burr, Republican of North Carolina, introduced a similar bill in 2011.

"Personalized medicine represents a revolutionary and exciting change in the way we think about and deliver health care," Mr. Obama said in a speech last year. He cited the drug Herceptin, for the treatment of a particularly aggressive form of breast cancer.

Scientists said they now viewed breast cancer not as a single disease, but rather as a group of several subtypes, each with a distinct molecular signature. This, they said, helps explain why some tumors respond better than others to specific cancer-fighting drugs.

"Most medical treatments have been designed for the average patient," said Jo Handelsman, associate director of the White House Office of Science and Technology Policy. "In too many cases, this one-size-fits-all approach is not effective."

Dr. Ralph Snyderman, a former chancellor for health affairs at Duke University, often described as the father of personalized medicine, said he was excited by the president's initiative.

"Personalized medicine has the po-

tential to transform our health care system, which consumes almost \$3 trillion a year, 80 percent of it for preventable diseases," Dr. Snyderman said.

Although the personalized medicine initiative is still in the early stages, officials said, personalized medicine can save money while producing better results. "It focuses resources on the right patients at the right time," Dr. Snyderman said.

Support for hundreds of millions of dollars for 'personalized medicine.'

will work," he said. "You can avoid wasting money on people who won't respond or will have an adverse reaction."

The new techniques can also help prevent disease by predicting patients' susceptibility, Dr. Snyderman said. "If an individual has a much greater likelihood of developing colon cancer, a genetically based disease, you can begin screening a much younger age, 30 rather than 50, for example."

Dr. Margaret A. Hamburg, the F.D.A.

commissioner, has reorganized her agency to speed the review of drugs and diagnostic tests used in personalized medicine.

Senator Cassidy said he was still in the early stages of his bill, but he said it would require "a much more nimble federal bureaucracy."

On Jan. 25, Mr. Obama released a proposal under which Medicare would cover genetic tests of tumors in some cases of cancer. The tests could help identify who are beneficiaries who would respond favorably to particular cancer drugs.

"This is a watershed event," said Dr. Bruce Quinn, a health-policy specialist at the law firm Foley Hoag. "It means that policy makers now believe these tests are worth paying for."

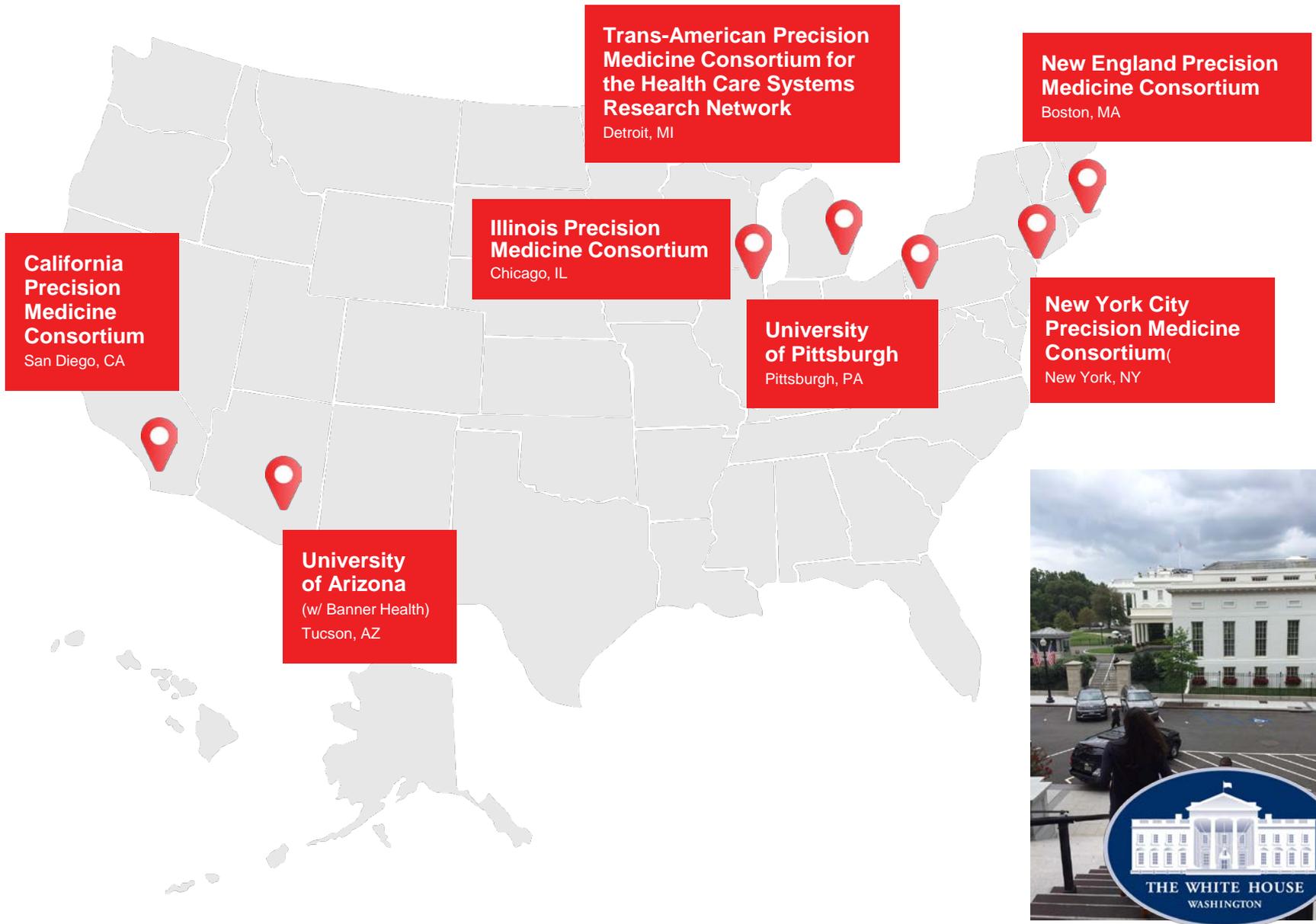
Mr. Obama's budget will also propose increased federal spending to combat antibiotic-resistant bacteria. The plan would nearly double spending from its current level of \$450 million a year.

White House officials described antibiotic resistance as a threat to public health and national security. They said at least 3,000 people in the United States die each year as a result of infections caused by such drug-resistant germs.

President Obama will seek hundreds of millions of dollars for a new initiative to develop medical treatments tailored to genetic and other characteristics of individual patients.

NYT Jan 25 2015

HPOs: Regional Medical Centers (RMCs)



PRÄZISIONSMEDIZIN



Präzisionsmedizin ist ein personalisierter Ansatz in der modernen Medizin, welcher individuelle Merkmale wie genetische Prädisposition, Umweltfaktoren oder Lebensstil der Patienten in die Behandlung miteinbezieht.

Die Initiative der UniBern / Inselspital in der Präzisionsmedizin strebt an, dass die richtige Behandlung zur richtigen Zeit zum richtigen Patienten gelangt.

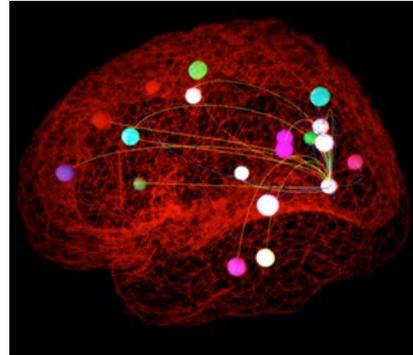
Kreismodell Precision Medicine



Big data transformiert die Wissenschaft...



High energy physics -
Large Hadron Collider



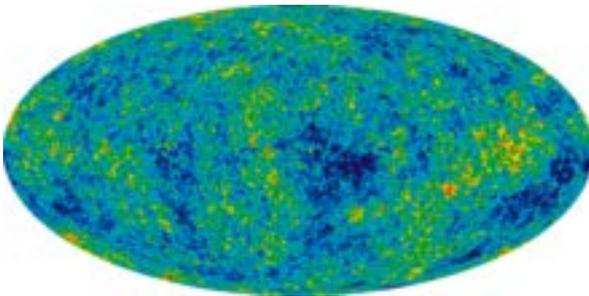
Neuroscience -
The Human Connectome Project



Ecology - Fluxnet



Genomics
DNA sequencer



Astronomy -
Sloan Digital Sky survey

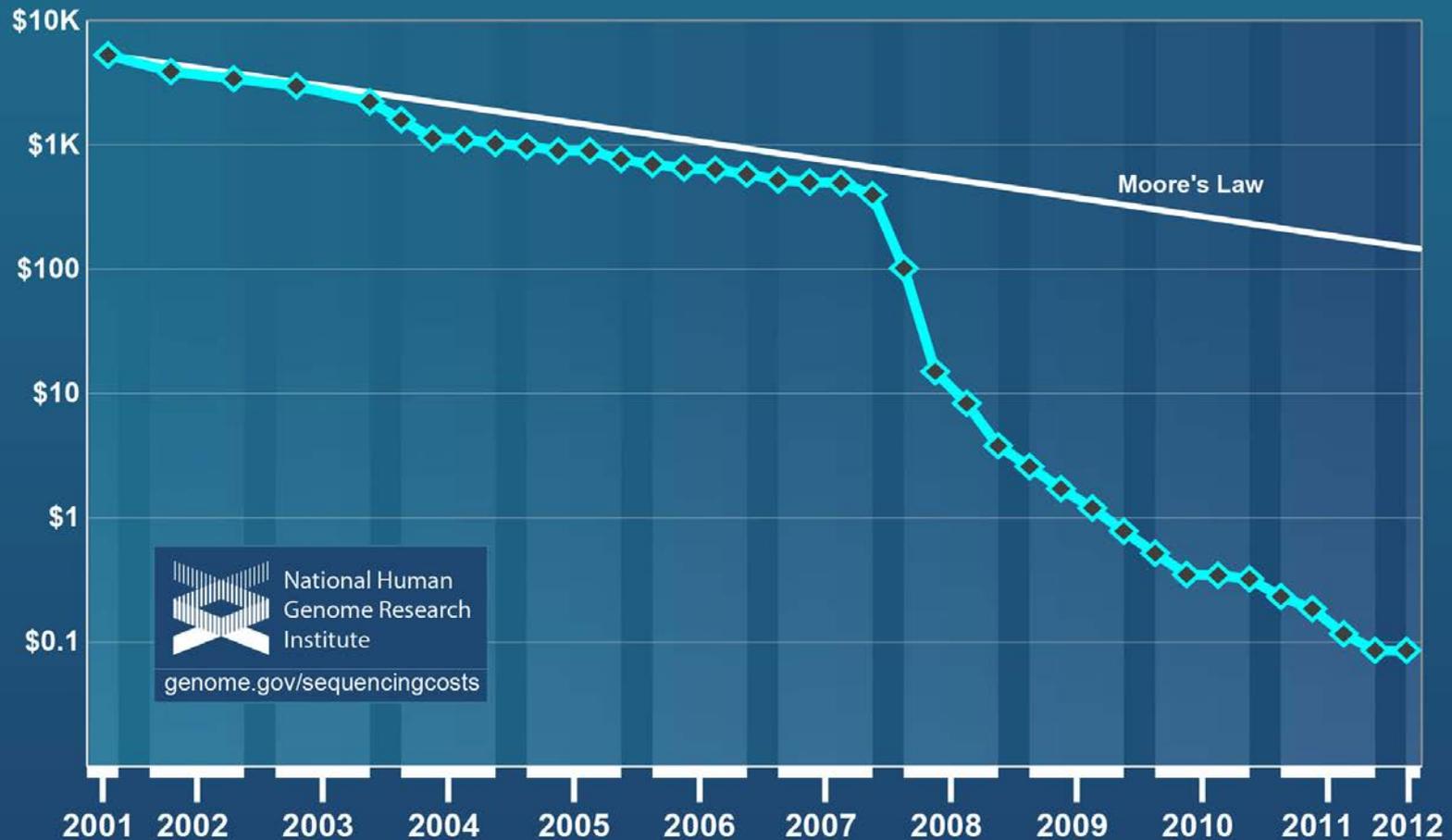


Knowledge of knowledge
Meta-data of scientific documents

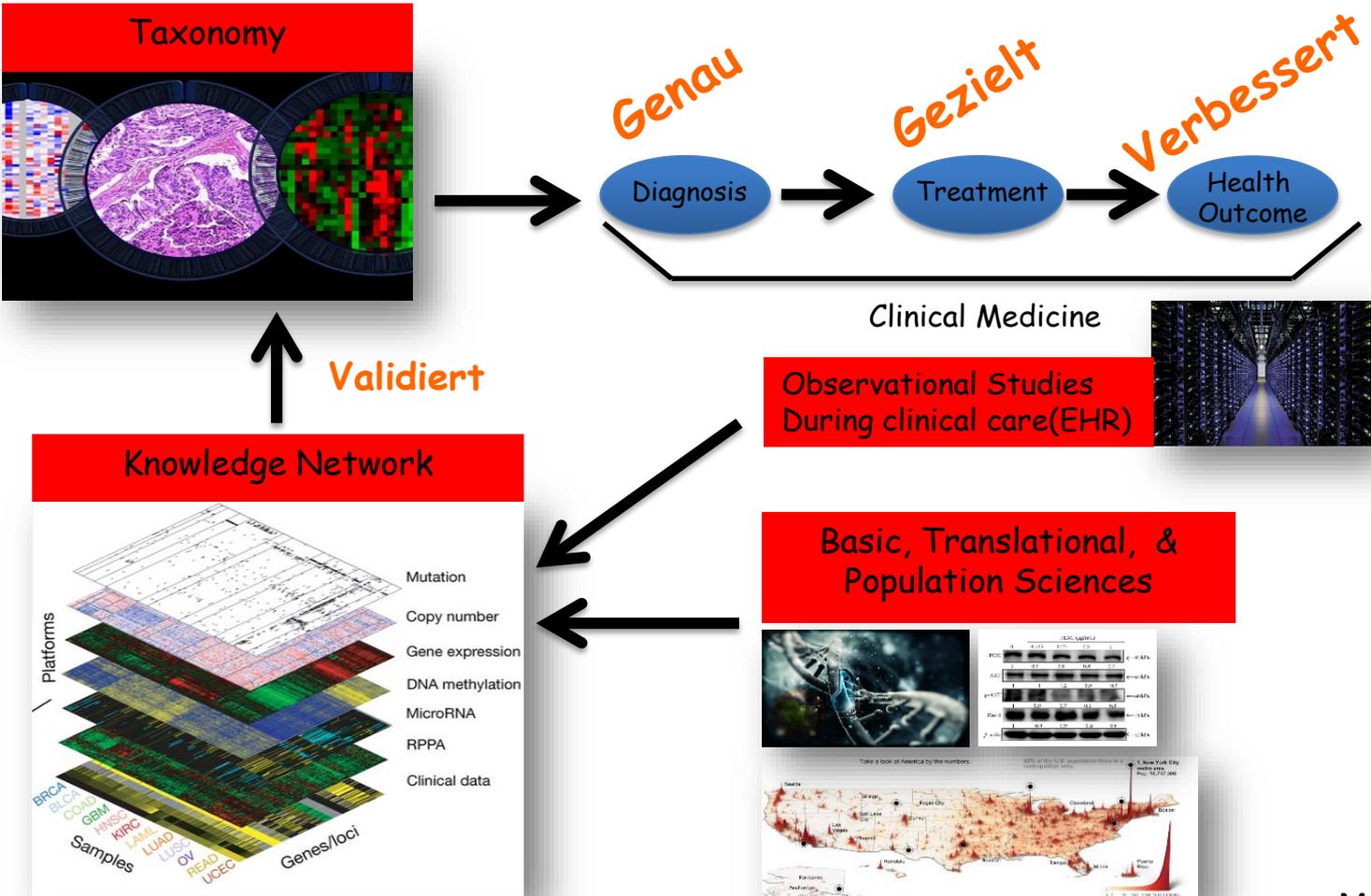
ISI Web of
KNOWLEDGE
Transforming Research

Datenexplosion in der Genforschung aufgrund sinkender Kosten

Cost per Raw Megabase of DNA Sequence



Das Versprechen und die Zukunft der Präzisionsmedizin



Modified
From National Academy
of Medicine

ZIEL

Eine **kontinuierliche Pflege**, in welcher wir den gesunden, kranken und wieder genesenen Zustand eines Patienten umfassend begleiten und erfassen - unabhängig der zugrundeliegenden Krankheit.

- Dies erlaubt uns, wertvolles wissenschaftliches Material zu sammeln, damit klinische Entscheide fundierter sind
- Um die Krankheit besser zu verstehen, brauchen wir Wissen über die Gesundheit
- Daten helfen uns nicht nur bei der Heilung von Krankheiten, sondern auch für die Erhaltung der Gesundheit



ZUSAMMENARBEIT

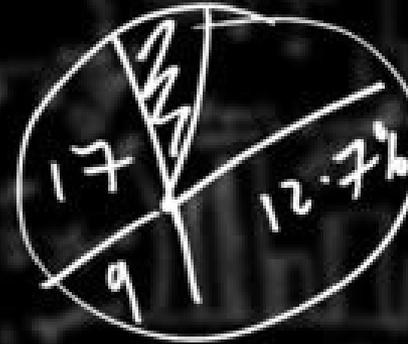
Präzisionsmedizin ist Teamwork – sie kombiniert die Kenntnisse und Fähigkeiten von ÄrztInnen, GenetikerInnen und DatenanalytInnen, um KlinikerInnen die besten Diagnose- und Behandlungsmöglichkeiten zu ermöglichen

- Teamwork heisst, wir arbeiten innerhalb von vielen unterschiedlichen Disziplinen, um eine möglichst grosse Expertise zu garantieren
- Aufgrund der riesigen Datenmengen aus genetischen Tests sind Bioinformatiker für unser Team essenziell
- Wir sammeln nicht nur Daten, wir nutzen sie und lernen aus ihnen

LERNSYSTEM FÜR DAS GESUNDHEITSWESEN

Wir schaffen ein Lernsystem für unser Gesundheitswesen, in dem die Behandlung aktiv überwacht und stetig verbessert wird

- Dies erlaubt uns, aktuelle Behandlungsstandards kontinuierlich zu verbessern.
- Die Verwendung von Genetik eröffnet völlig neue Wege zur Behandlung von Krankheiten
- Unser Ziel ist es, jeden Behandlungsplan eines Patienten so einzigartig zu gestalten, wie es auch das individuelle Genmaterial ist
- Wissenschaftliche Erkenntnisse werden direkt in die Ausbildung der ersten Generation von ÄrztInnen im Gebiet der Präzisionsmedizin einfließen



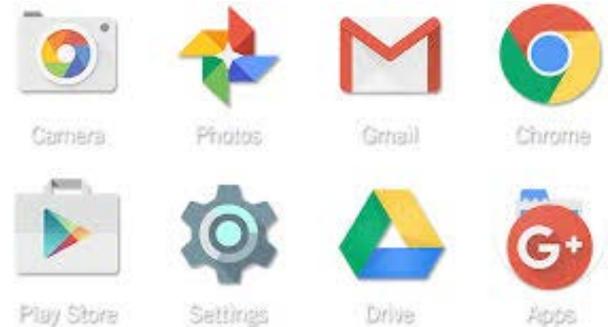
Kreismodell Precision Medicine



Small data too..



Wo bist Du?
Wo warst Du?
Was kaufst Du?
Wie oft schreibst Du Emails?
Wieviel Leute wohnen bei Dir?
Was suchst Du im Internet? Wann? Wie oft?



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Microsoft Surface Book mit Performance Base

N° 67

TEST FORCE

Small text at the bottom:
- Lieferung Schweiz
- In 15 Minuten
- In 15 Minuten
- Bestpreisgarantie

Wieviel Angst sollten wir haben?



Bedeutung von genomischem Datenschutz

- **Das allgemeine Dilemma der genomischen Privatsphäre**
 - Aus dem Informationsaustausch wird die Person möglicherweise in ihrer **Privatsphäre beeinträchtigt**, aber die **Gesellschaft profitiert** von der medizinischen Forschung
 - *Wie kann man Risiken und Belohnungen ausgleichen?*

An Introduction to the

All of Us Research Program



National Institutes
of Health

Updated July 7, 2017

@AllofUsResearch #JoinAllofUs

The Vision of President Obama



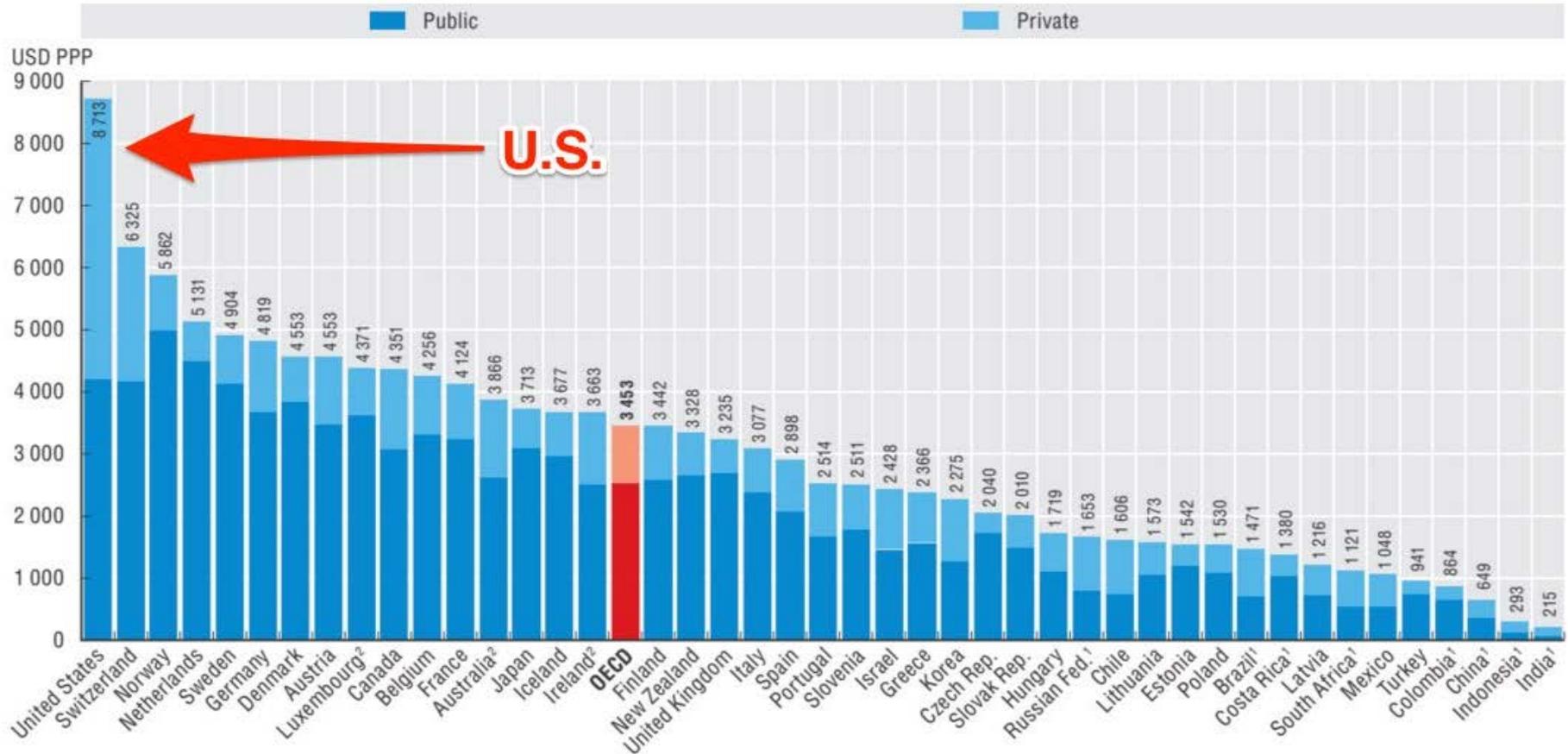
“My hope is that this becomes the foundation, the architecture, whereby in 10 years from now we can look back and say that we have revolutionized medicine.”

- President Barack Obama

US Spend Most on Health Care

Source: Organization for Economic Co-operation and Development

9.1. Health expenditure per capita, 2013 (or nearest year)



Infant Mortality by Country (2003)

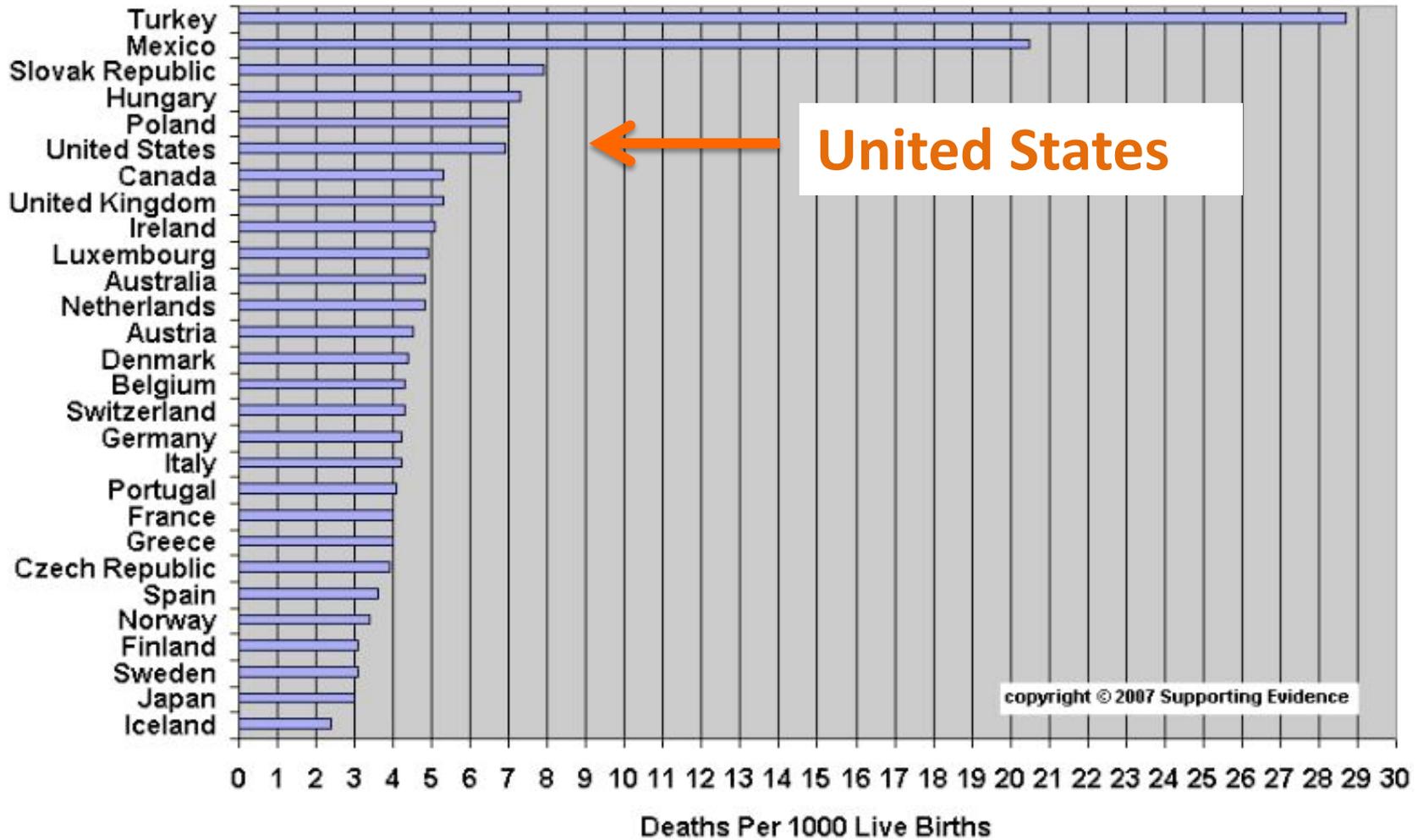
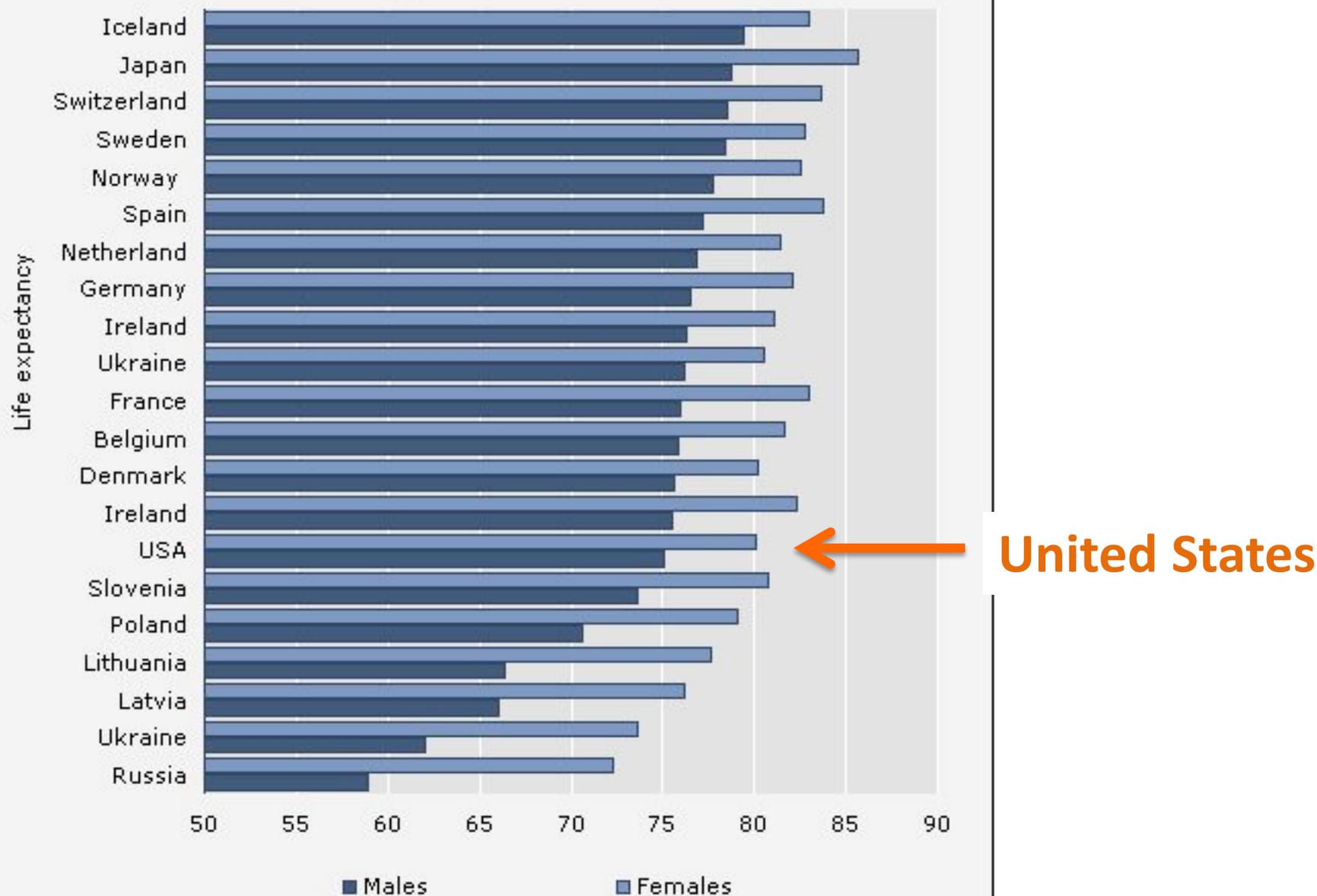


Figure 1. Life expectancy in selected countries of the world, 2005-2006 (Iceland), 2004 or 2005 in other countries





Etablierte Programminfrastruktur

DATA AND RESEARCH CENTER (DRC)

Big data capture, cleaning, curation, & sharing in secure environment

Vanderbilt, Verily, Broad Institute

BIOBANK

Repository for processing, storing, & sharing biosamples (35+M vials)

Mayo Clinic

PARTICIPANT CENTER

Direct volunteer participant enrollment, digital engagement innovation, & consumer health technologies

Scripps Research Institute (with multiple partners)

PARTICIPANT TECHNOLOGY SYSTEMS CENTER

Website & mobile apps for participants

Vibrent Health

HEALTH CARE PROVIDER ORGS (HPOs)

Clinical & scientific expertise network, enrollment & retention of participants

20+ regional med centers, FQHCs, VA, future awards to grow network

COMMUNICATIONS & ENGAGEMENT

Comms, marketing, & design expertise; engagement coordination & community partners network

Wondros, HCM, future awards to grow network of community partners

All of Us Research Program



The All of Us Anthem

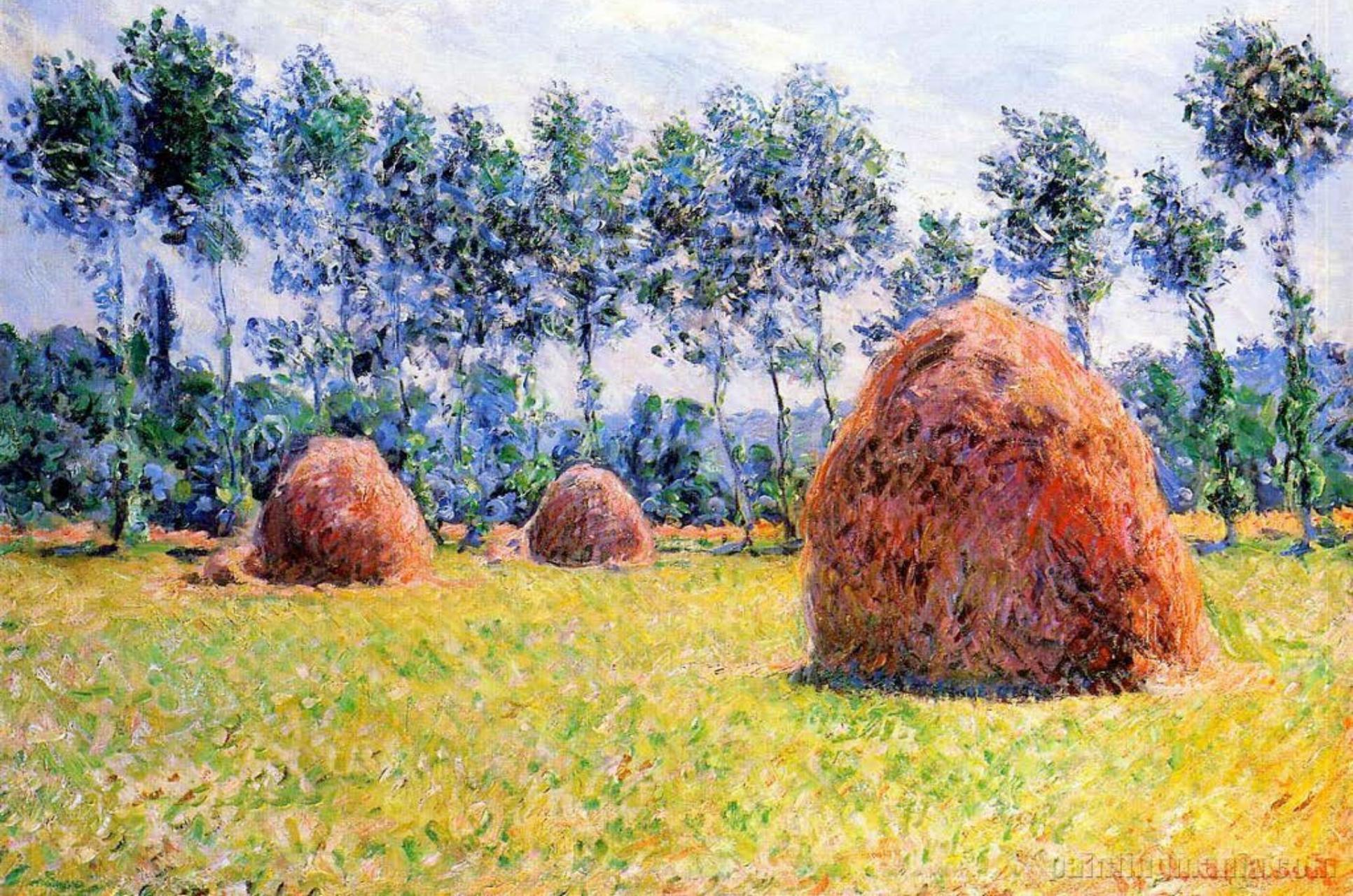


Der Nutzen der Teilnahme bei *All of Us*

- Eine **Gelegenheit** etwas über Ihre Gesundheitsindikatoren zu erfahren und Zugang zu Ihren eigenen Daten zu haben
- Eine Gelegenheit, **Krankheiten zu bekämpfen** und die Gesundheit zukünftiger Generationen zu fördern
- Eine Gelegenheit, **Teil einer Bewegung zu sein**, für eine präzisere, persönlichere und effektivere Gesundheitsvorsorge



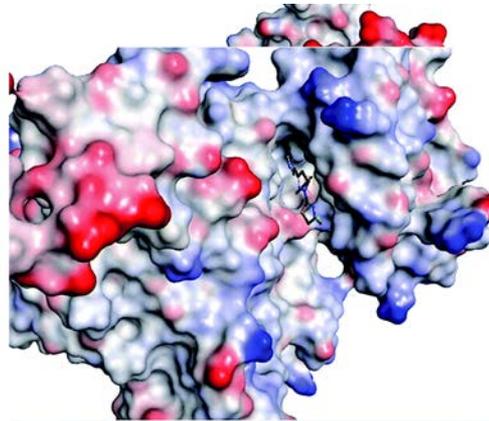
Präzisionsmedizin: Lehren aus dem Feld



Präzisionsmedizin: Lehren aus dem Feld



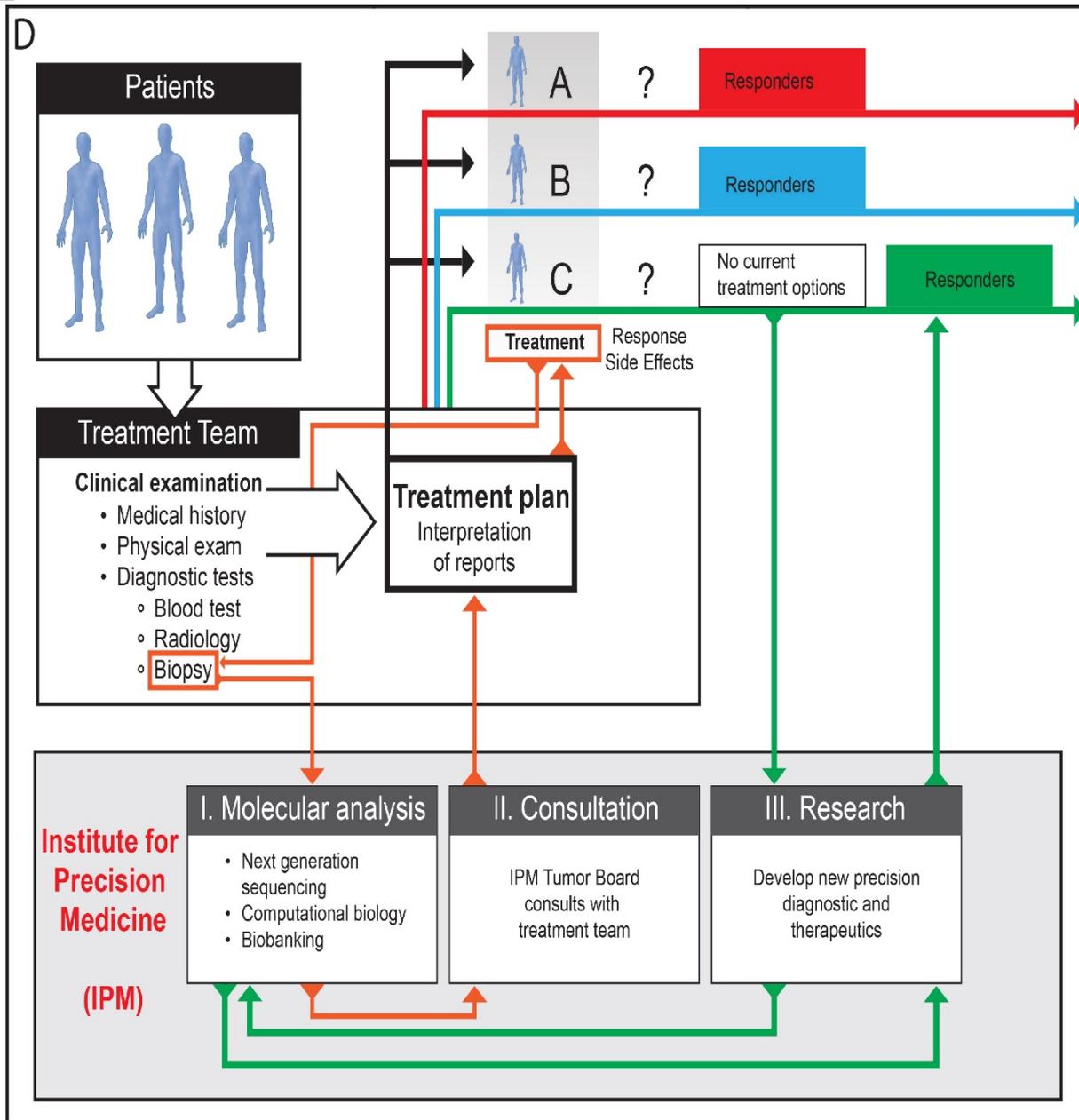
Drug



Mutation



Dramatic Response



Klinische Versuche

Hypothese:
Präzisionsmedizin
kann helfen, um
klinische Studien
zu verbessern

Weill Cornell Medicine PM Experience: Precision Cancer Care in Real Time

Research

Original Investigation

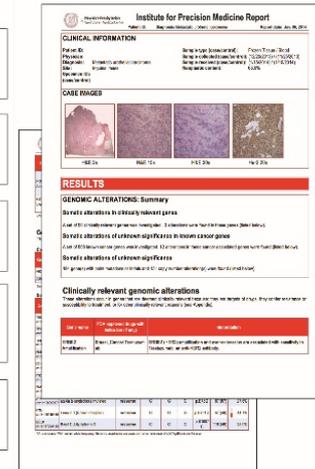
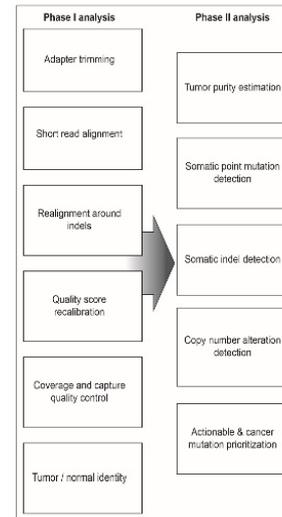
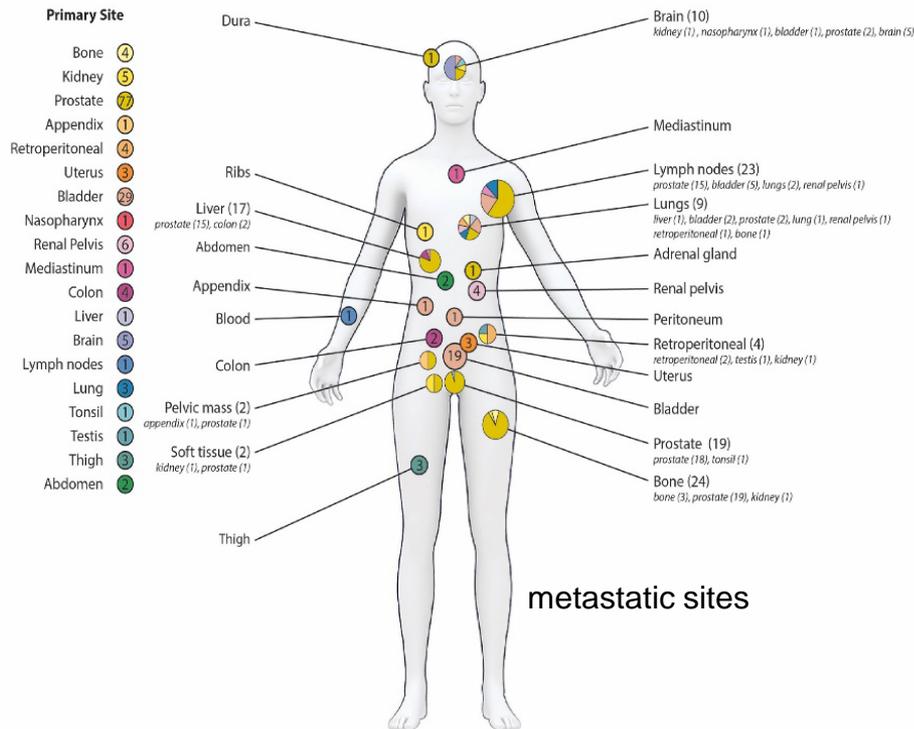
Whole-Exome Sequencing of Metastatic Cancer and Biomarkers of Treatment Response

Himisha Beltran, MD; Kenneth Eng, MS; Juan Miguel Mosquera, MD; Alexandros Sigaras, MS; Alessandro Romanel, PhD; Hanna Rennert, MD; Myriam Kossai, MD; Chantal Pauli, MD; Bishoy Faltas, MD; Jacqueline Fontugne, MD; Kyung Park, MD; Jason Banfelder, MCh.E; Davide Prandi, PhD; Neel Madhukar, BS; Tuo Zhang, PhD; Jessica Padilla, MA; Noah Greco, MBA; Terra J. McNary, BA; Erick Herrscher, BA; David Wilkes, PhD; Theresa Y. MacDonald, BS; Hui Xue, PhD; Vladimir Vacic, PhD; Anne-Katrin Emde, PhD; Dayna Oschwald, PhD; Adrian Y. Tan, MD; Zhengming Chen, PhD, MPH; Colin Collins, PhD; Martin E. Gleave, MD; Yuzhuo Wang, PhD; Dimple Chakravarty, PhD; Marc Schiffman, MD; Robert Kim, BS; Fabien Campagne, PhD; Brian D. Robinson, MD; David M. Nanus, MD; Scott T. Tagawa, MD; Jenny Z. Xiang, MD; Agata Smogorzewska, MD, PhD; Francesca Demichelis, PhD; David S. Rickman, PhD; Andrea Sboner, PhD; Olivier Elemento, PhD; Mark A. Rubin, MD

Beltran et al., JAMA Oncology May 28 2015
Rennert et al., npj Genomic Medicine 2016

Seit Oktober 2015 WES Testing bei WCM

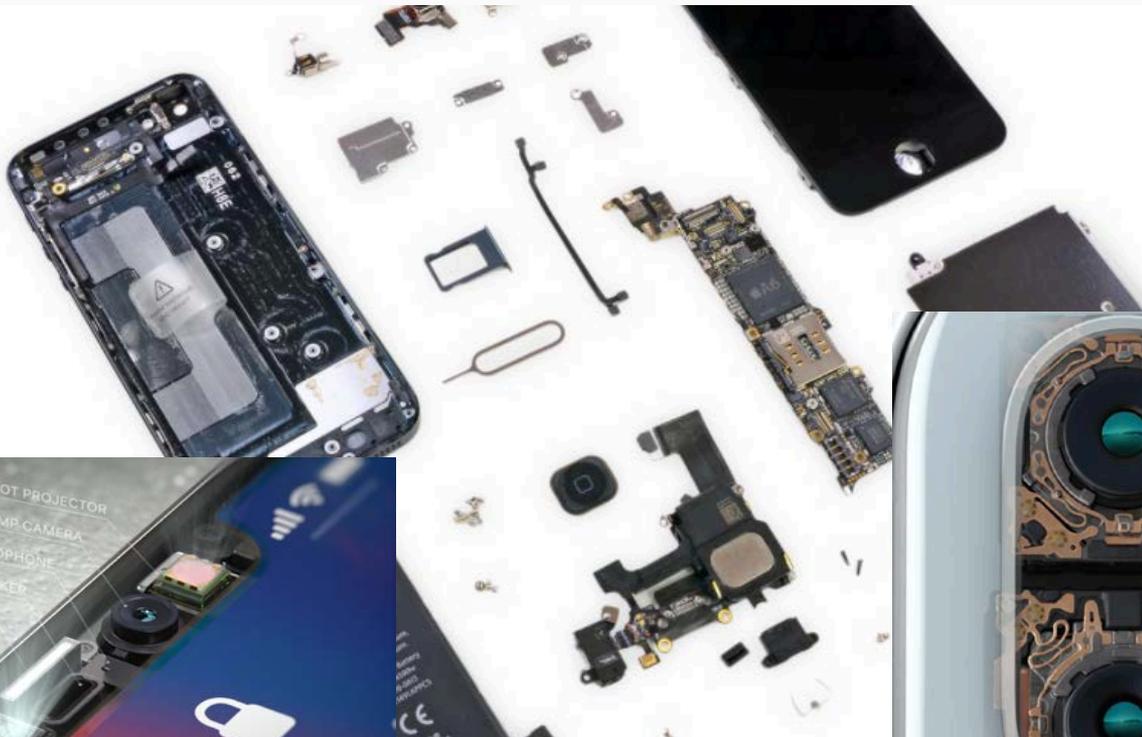
EXaCT-1 Genom-Tests für fortgeschrittenen Krebs Über 1000 getestete Patienten



WES+Analysis pipeline

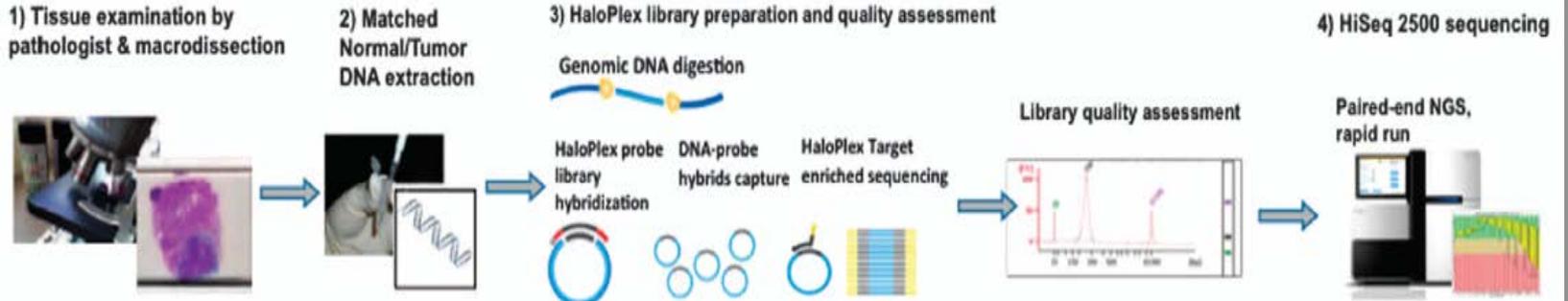


patient reports

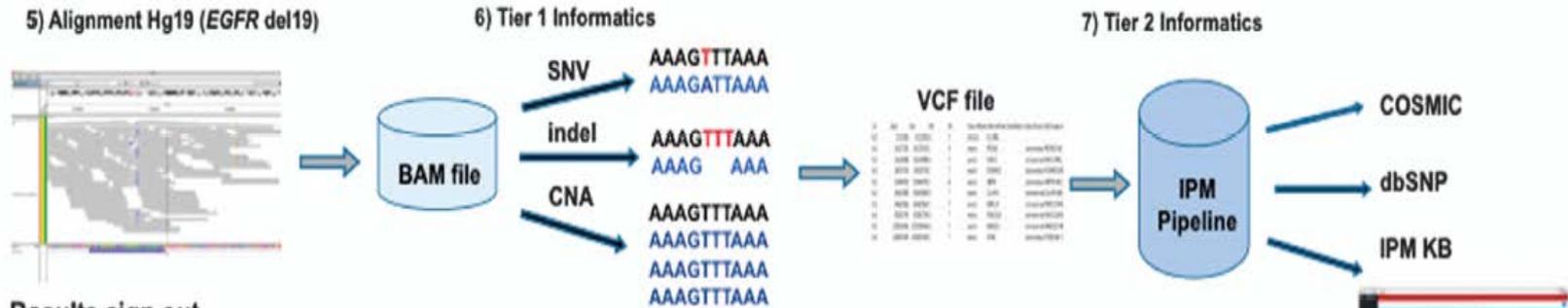


EXaCT-1 Workflow

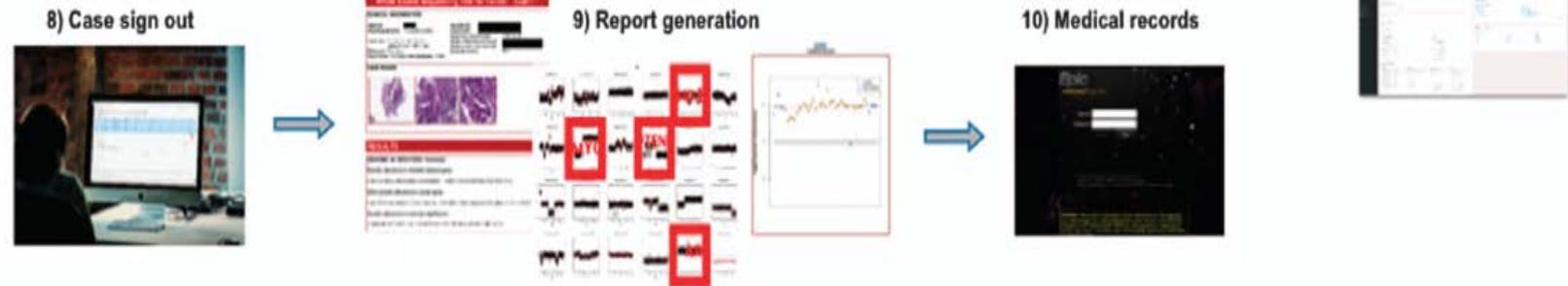
Sample preparation and sequencing



Data analysis and variant tiering



Results sign out



Integrierter Bericht mit detaillierter Interpretation



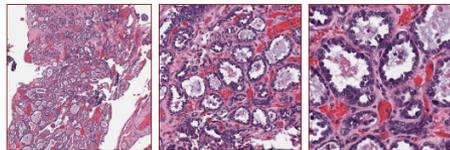
Institute for Precision Medicine Report - Preliminary

Patient ID: PM266 Diagnosis: Clear cell carcinoma Report date: Dec. 05, 2014

CLINICAL INFORMATION

Patient ID: PM266
 Physician: Himisha Beltran M.D.
 Diagnosis: Clear cell carcinoma
 Site: Pelvic mass
 Specimen IDs (case/control): PM266_ZA2_1_Case_HALO / PM266_ZC2_1_Ctrl_HALO
 Sample type (case/control): FFPE / FFPE
 Sample collected (case/control): (3/18/2014) / (3/18/2014)
 Sample received (case/control): (11/14/2014) / (11/14/2014)
 Neoplastic content: 56.6%

CASE IMAGES



RESULTS

GENOMIC ALTERATIONS: Summary

Somatic alterations in clinically relevant genes

A set of 49 clinically relevant genes was investigated. 2 alterations were found in these genes (listed below).

Somatic alterations of unknown significance in known cancer genes

A set of 509 known cancer genes was investigated. 8 alterations in these cancer associated genes were found (listed below).

Somatic alterations of unknown significance

13 gene(s) with point mutations or indels and 41 copy number alteration(s) were found (listed below).

Clinically relevant genomic alterations

These alterations occur in genes that are deemed clinically relevant because: they are targets of drugs, they confer resistance or susceptibility to treatment, or for other clinically relevant reasons (see Appendix).



Institute for Precision Medicine Report - Preliminary

Patient ID: PM266 Diagnosis: Clear cell carcinoma Report date: Dec. 05, 2014

Gene name	FDA approved drugs with indication (if any)	Interpretation
FGFR3 focal amplification	none	FGFR3 amplification may be associated with response to the multitargeted tyrosine kinase inhibitor pazopanib (Liao et al, 2013, Cancer Res).

VAF: variant allele frequency

Notes

The status of alterations in gene(s) KRAS is indeterminate because the coverage was below the optimal levels of this method (<10 reads). Hence, analysis of the alteration(s) with an independent methodology will be performed.

Genomic alterations of unknown significance in cancer genes

These alterations occur in genes that are cancer associated, but their impact on the disease is unknown (see Appendix).

Copy number alterations

Gene name	Description	Classification of alteration	Altered region
FH	fumarate hydratase	LARGE SCALE AMPLIFICATION	chr1:223,533,597-249,212,519
H3F3A	H3 histone, family 3A	LARGE SCALE AMPLIFICATION	chr1:223,533,597-249,212,519
BCL7A	B-cell CLL/lymphoma 7A	FOCAL AMPLIFICATION	chr12:122,468,644-123,419,896
STAT3	signal transducer and activator of transcription 3 (acute-phase response factor)	FOCAL AMPLIFICATION	chr17:40,039,428-40,673,093
YWHAE	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide (14-3-3 epsilon)	FOCAL AMPLIFICATION	chr17:649,687-1,968,405
AKT2	v-akt murine thymoma viral oncogene homolog 2	FOCAL AMPLIFICATION	chr19:39,759,400-40,947,690
WHSC1	Wolf-Hirschhorn syndrome candidate 1(MMSET)	FOCAL AMPLIFICATION	chr4:1,316,228-2,160,908

Genomic coordinates are based on human reference GRC37/hg19. Large scale alterations involve at least 50 genes.

Somatic mutations and indels

Gene name	Gene description	Classification	Reference Allele	Tumor Allele 1	Tumor Allele 2	AA change	Tumor (Normal) read depth	Tumor VAF
ARID1A chr1:27094361	AT rich interactive domain 1A (SWI-like)	nonsense	G	G	A	p.W1024*	53 (55)	41.5%

AA: amino-acid; VAF: variant allele frequency; Genomic coordinates are based on human reference GRC37/hg19 and are 1-based.

Genomic alterations of unknown significance

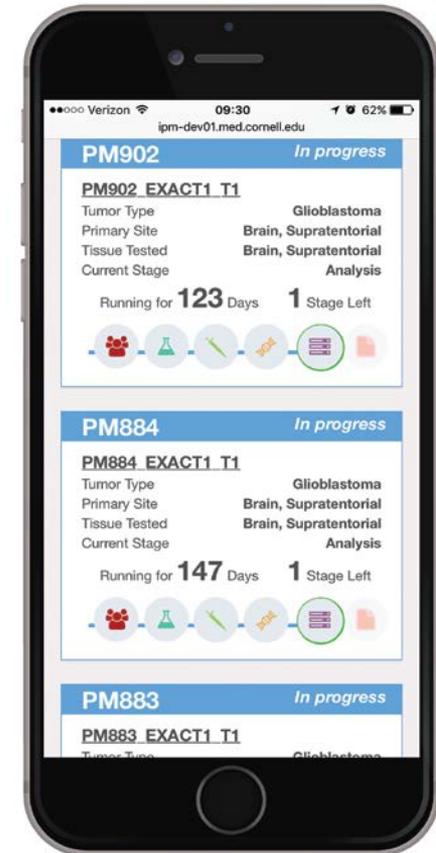
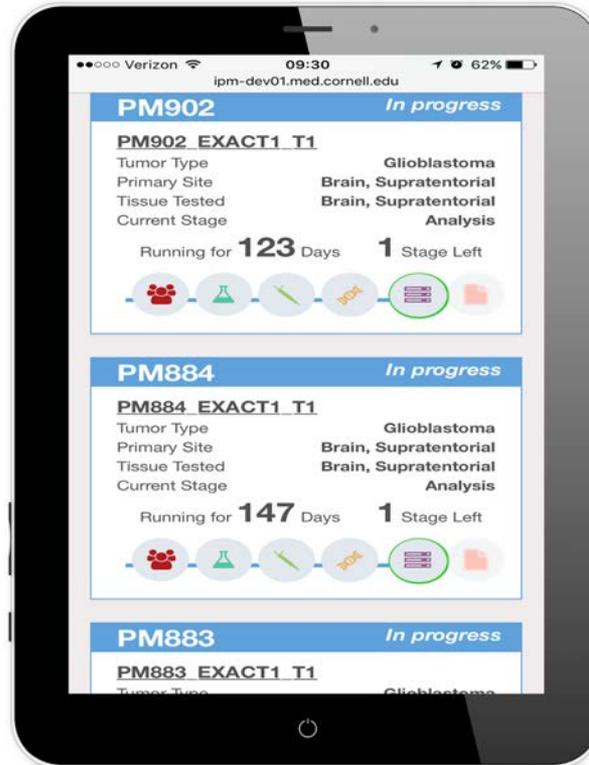
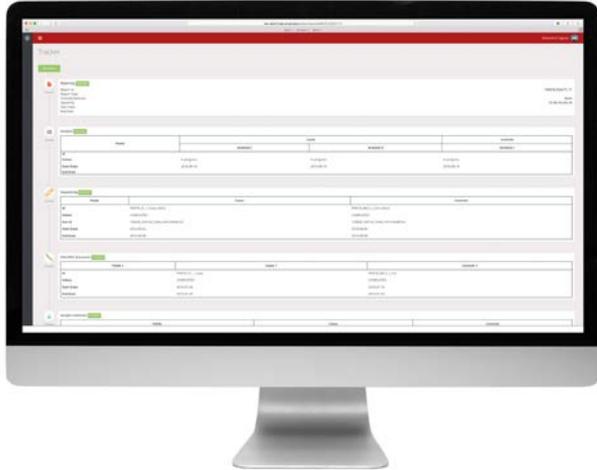
These alterations are not known to have any effect on the disease, but are here reported in the event that in the future progress in scientific knowledge could determine their role (see Appendix).

Alle Patienten- und Genomdaten wurden gespeichert. Nicht nur Teildaten.

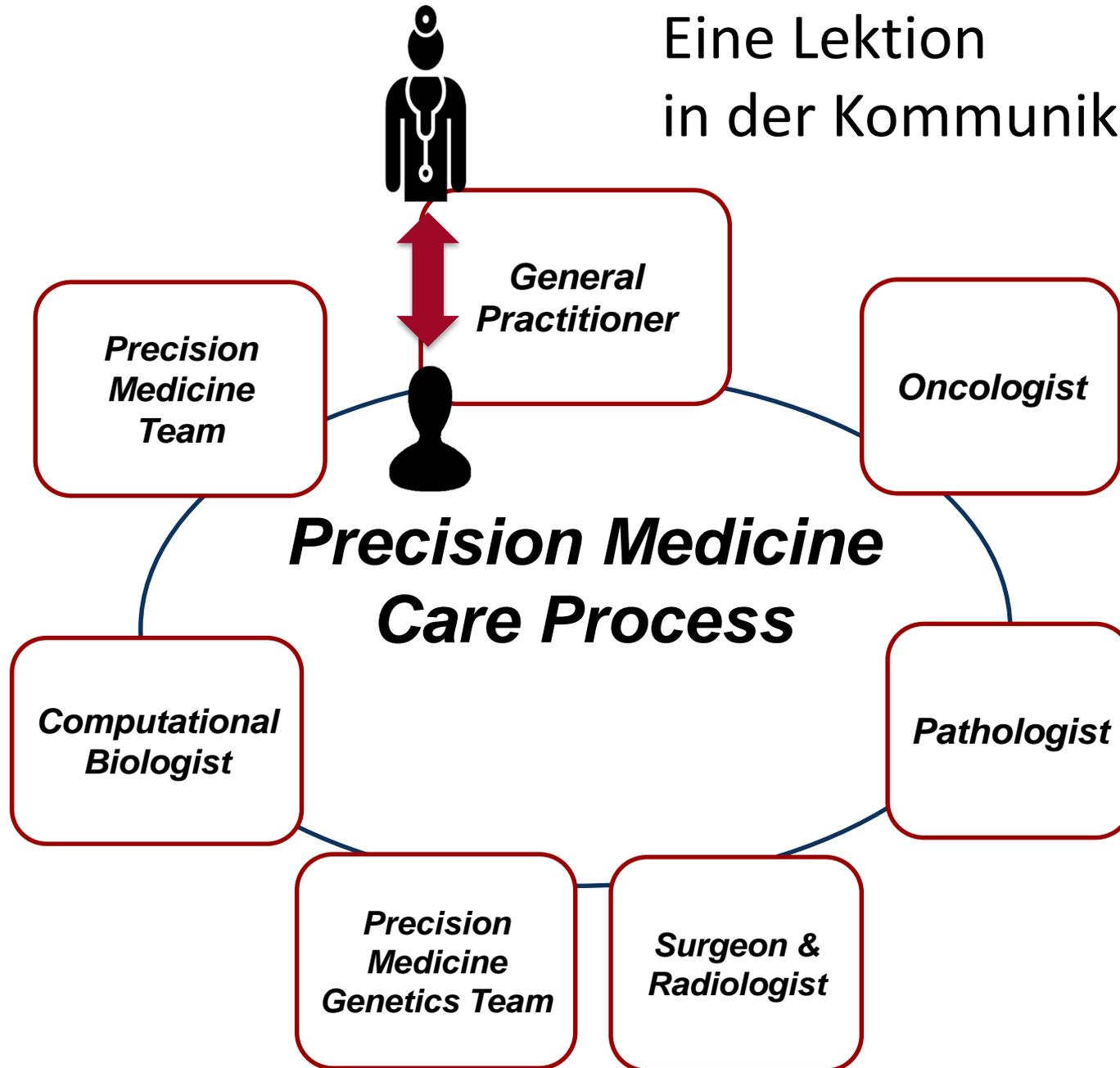
Patient, doctor, admin specific dashboards



Tracker – Mobile



Eine Lektion in der Kommunikation

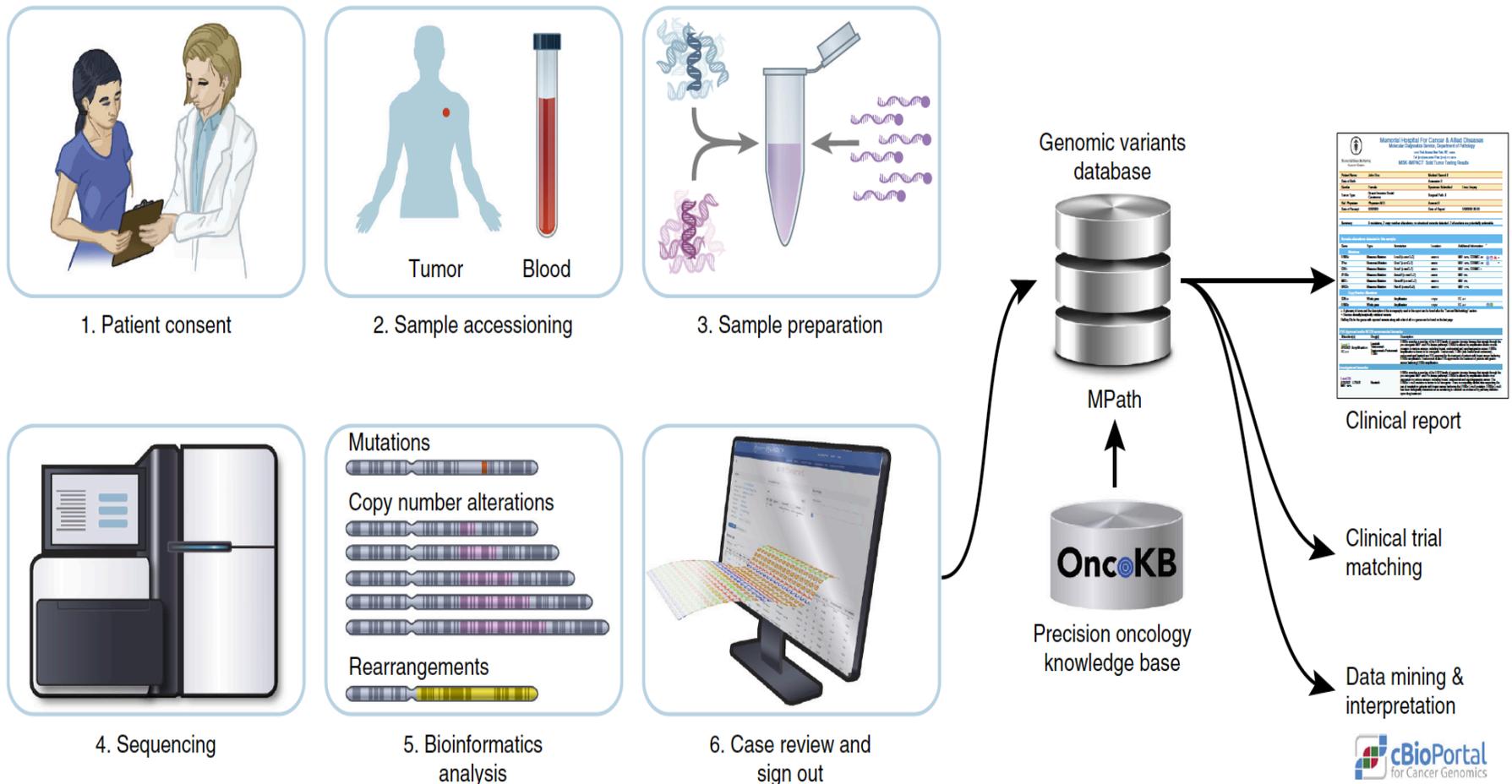


The image features a central brown oval containing white text. This oval is surrounded by twelve stylized, grey, human-like figures arranged in a circle. Each figure has a different colored oval on its chest, representing a diverse group of participants. The colors include yellow, red, black, teal, and brown. The figures are positioned as if they are sitting around a table, engaged in a discussion.

Diskussion mit
Pflegepersonen und
PM-Teammitgliedern
zur Entwicklung eines
Konsultationsberichts

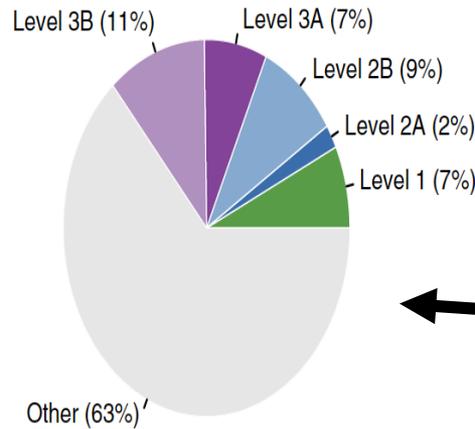
Präzisionsmedizin-Tumorboard

Wie sieht die genomische Analyse von 10.000 Patienten aus?



Wie viele Patienten können auf Basis **der Daten** eine neue Behandlung erhalten?

Level 1	FDA-recognized biomarker for an FDA-approved drug in the same indication
Level 2A	Standard of care biomarker for an FDA-approved drug in the same indication
Level 2B	Standard of care biomarker for an FDA-approved drug in another indication
Level 3A	Compelling clinical evidence supporting the biomarker as being predictive of drug response in the same indication
Level 3B	Compelling clinical evidence supporting the biomarker as being predictive of drug response in another indication



Nur 7% Level 1



Wie helfen wir Patienten, die keine Optionen mehr haben?

Number of biomarkers: PIK3C, EGFR, BRAF, ERBB, AK1, IDH, KR, PTE, ME, FG, NR, GNA, HRA, KRA, TP, ERBB, GNA, PTC, TSC, Other, ERBB, CDK, MDM, CCND, FG, FG, ME, PIK3C, Other, PTE, SMARCB, Other, AL, ROS, R, NTRK, FG, NTRK, Other

TIME

What if your immune system could be taught to kill cancer?

Inside the brutally selective, hugely expensive, lifesaving trials of immunotherapy.

By Alice Park

Science

Breakthrough of the Year

Cancer Immunotherapy

T cells on the attack

2015 TOP CANCER DOCTORS

Newsweek

07.31.2015-08.07.2015

SPECIAL HEALTH ISSUE

EXPLORING THE YOUNG GENOME

SUPER RESPONDERS TO THE RESCUE

CURING CANCER

MAKING MEDICAL BRAND NAMES

THE FRAGILE LACK OF WEDS FOR RUGS

THE DRUG IS TOO DAMN EXPENSIVE!

AMAZON: THE NEXT \$8 TRILLION • MICROSOFT MAKES NICE

SPECIAL ISSUE

Forbes

NOVARTIS' JOSEPH JIMENEZ

"THE PAIN OF HAVING TO TURN PATIENTS AWAY IS SUCH THAT WE ARE GOING AS FAST AS WE CAN."

WILL THIS MAN CURE CANCER?

WITH A RADICAL NEW TREATMENT HE IS BETTING BILLIONS ON CONQUERING OUR DEADLY FOE

PLUS: THE GLOBAL 2000 BIGGEST, BEST COMPANIES

2015 MUTUAL FUND HONOR ROLL

11 MUST-BUYS BE YOUR OWN BUFFETT!

OUTLOOK

nature

THE INTERNATIONAL WEEKLY JOURNAL OF SCIENCE

Antitumour immunity enhanced by inhibiting PD-1/PD-L1 and identifying mutant neo-antigens

PAGES 498, 508, 563, 568, 572 & 577

IMMUNE-CHECKPOINT BLOCKADE IN CANCER

PEER REVIEW

ACCEPT YOUR OWN PAPER

13 new immune cancers are duping the system

PAGE 498

MICROSCOPY

THE CASE FOR AIMING HIGHER

Atomic resolution at there for the taking

PAGE 497

ENERGY

'NIGHT-TIME' COOLING BY DAY

New materials enable radiative cooling in sunlight

PAGE 548

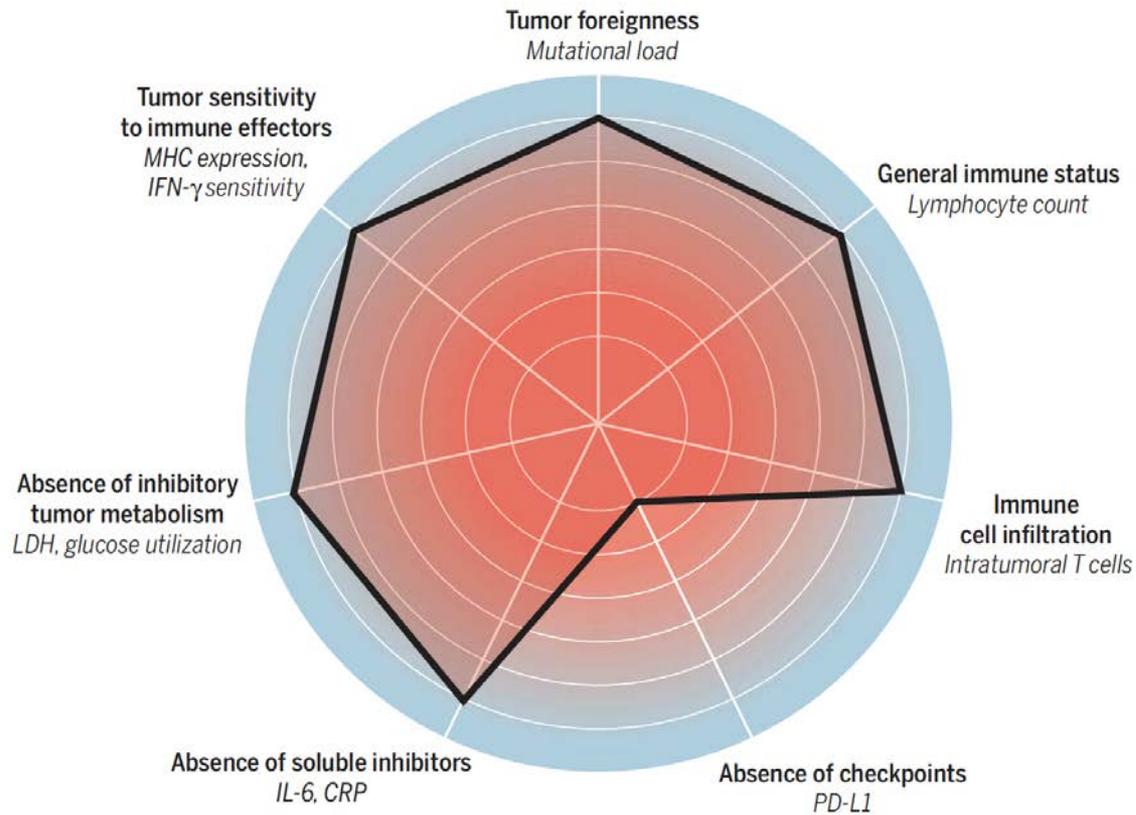
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UK £12.50

US \$24.00

BENEFITS OF IMMUNOTHERAPY



CANCER IMMUNOLOGY

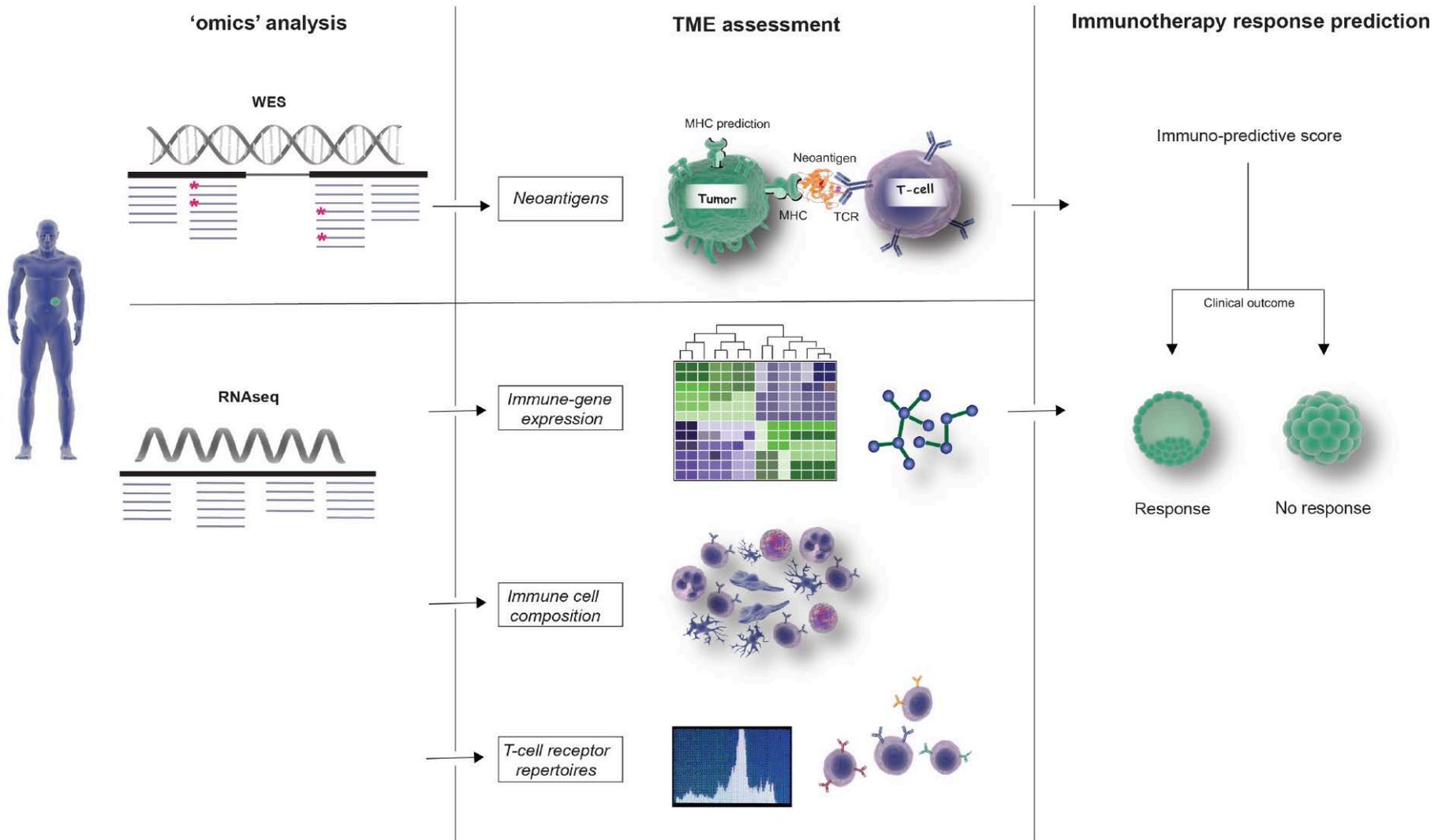
The “cancer immunogram”

Visualizing the state of cancer-immune system interactions may spur personalized therapy

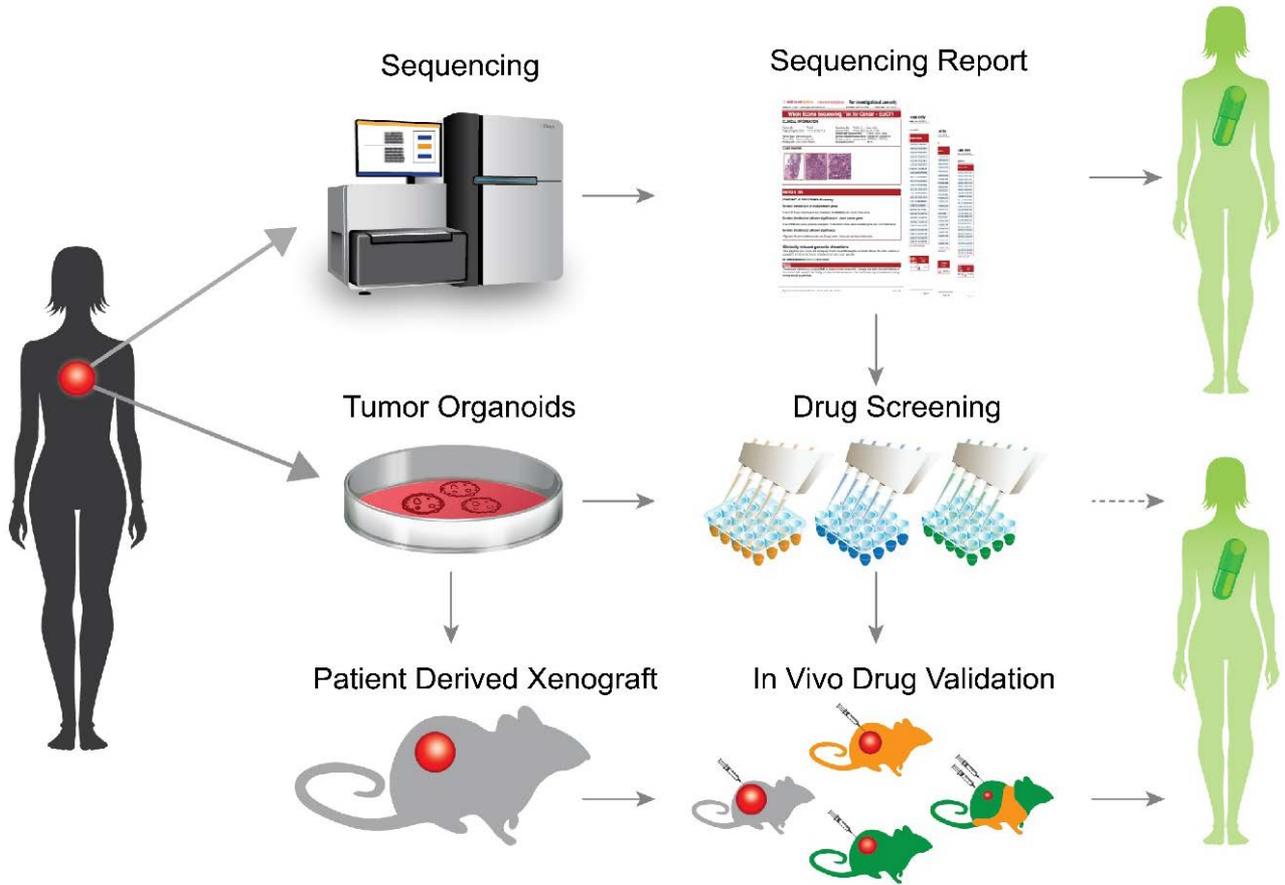
By Christian U. Blank,^{1,2} John B. Haanen,^{1,2} Antoni Ribas,³ Ton N. Schumacher²

Science May 2016

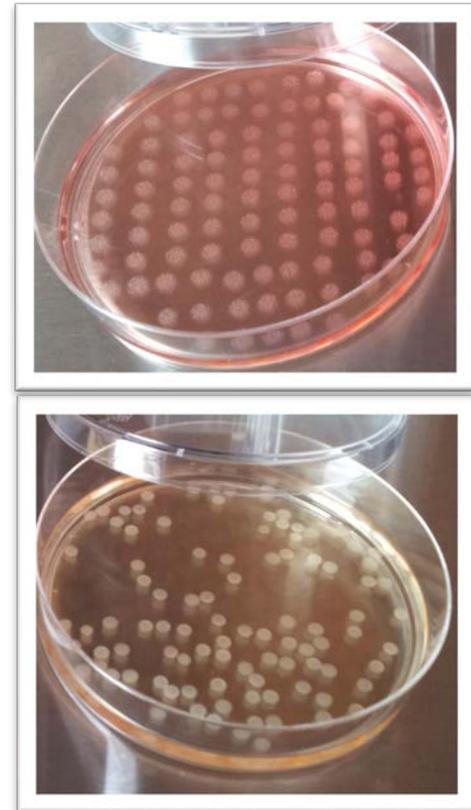
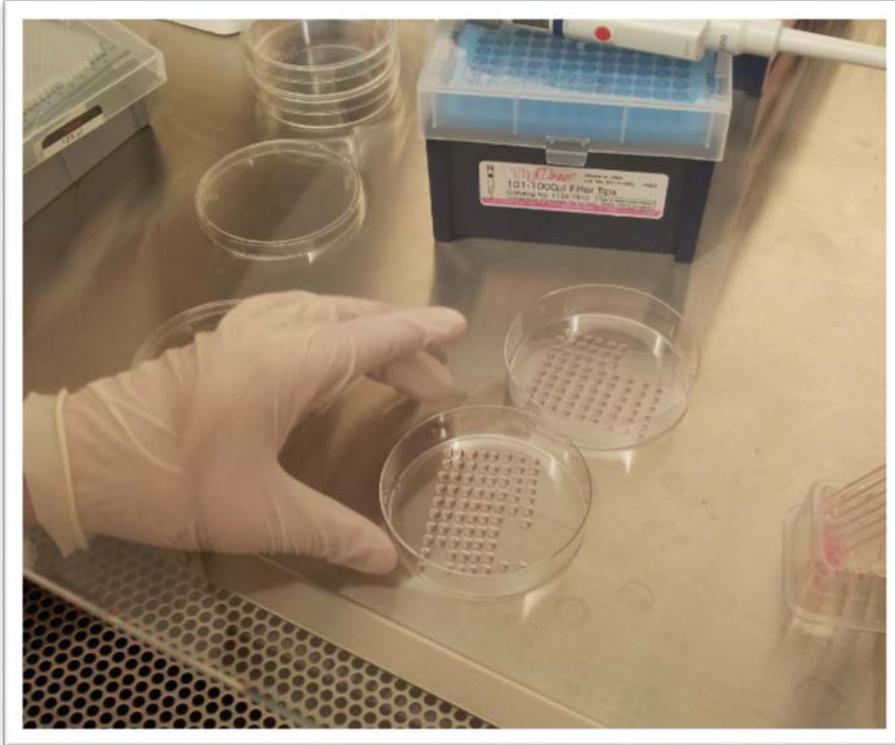
Wir brauchen **viele Daten**, um zu verstehen, welche Patienten von dieser teuren Behandlung profitieren.



Personalized Cancer Models to Guide Precision Medicine



Organoids in drug testing: In vitro – Mini Sarcoma Production



Pauli et al, March 2017



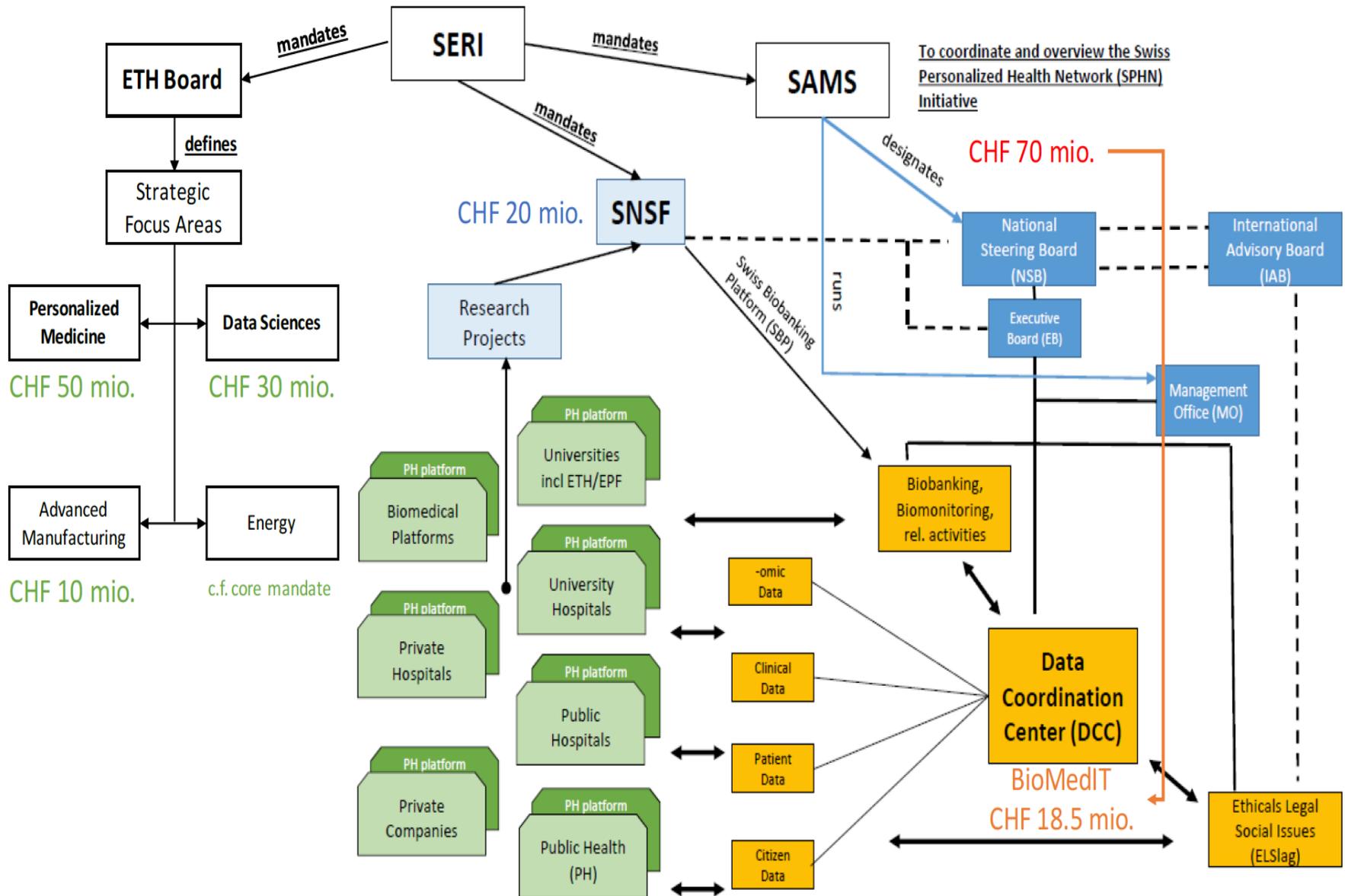
TOBLERONE

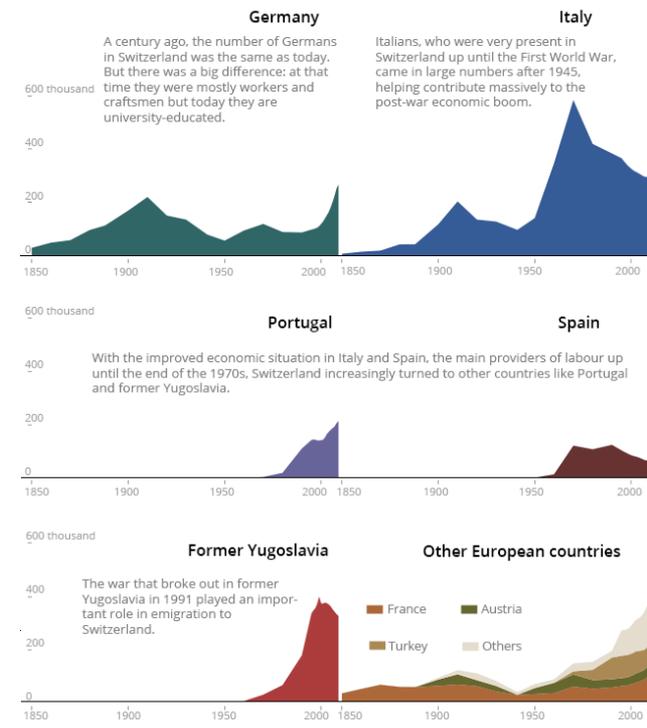
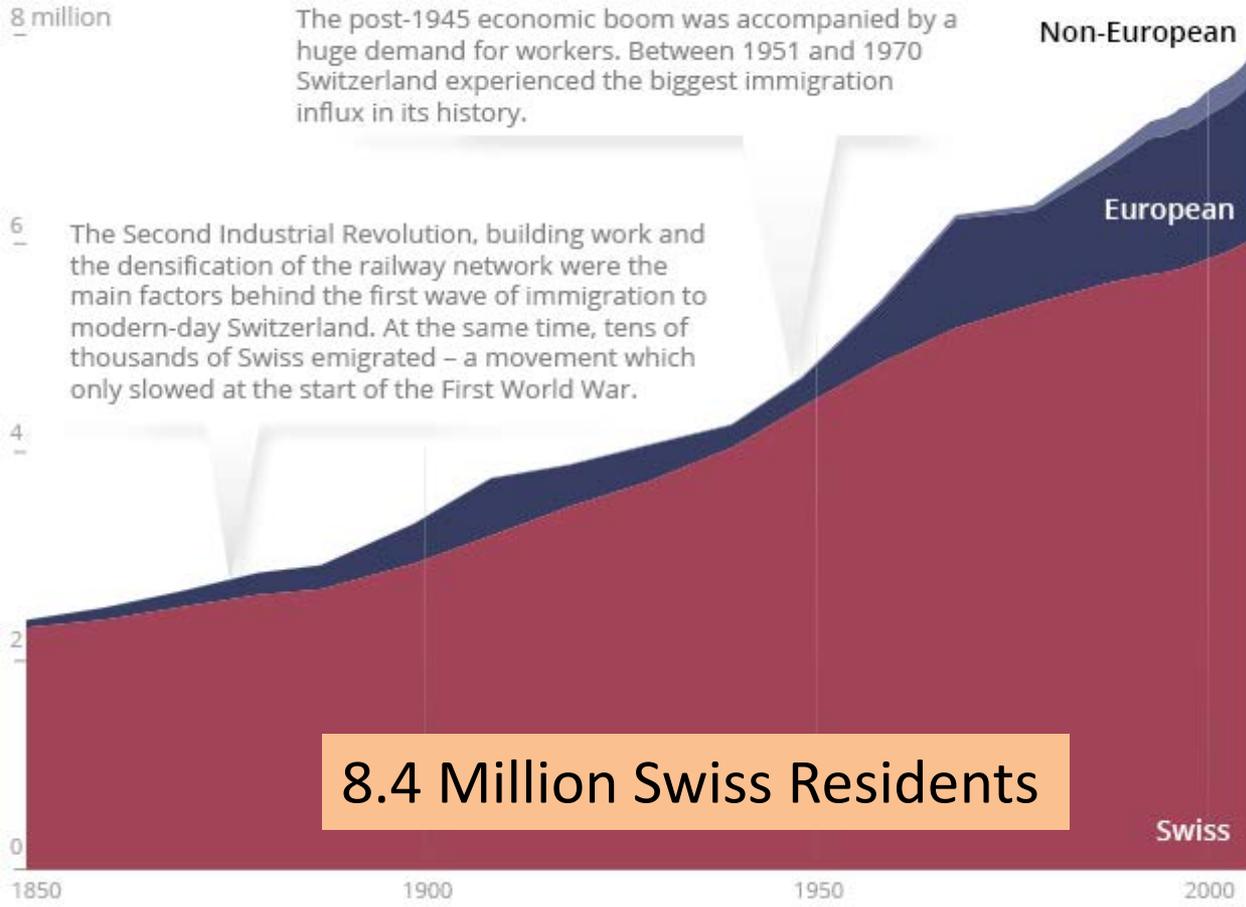
© Hättingh Künzli+Frey AG, CH-3122 Schönholzi/Bern

Die Schweiz: Rund 8,5 Millionen Einwohner

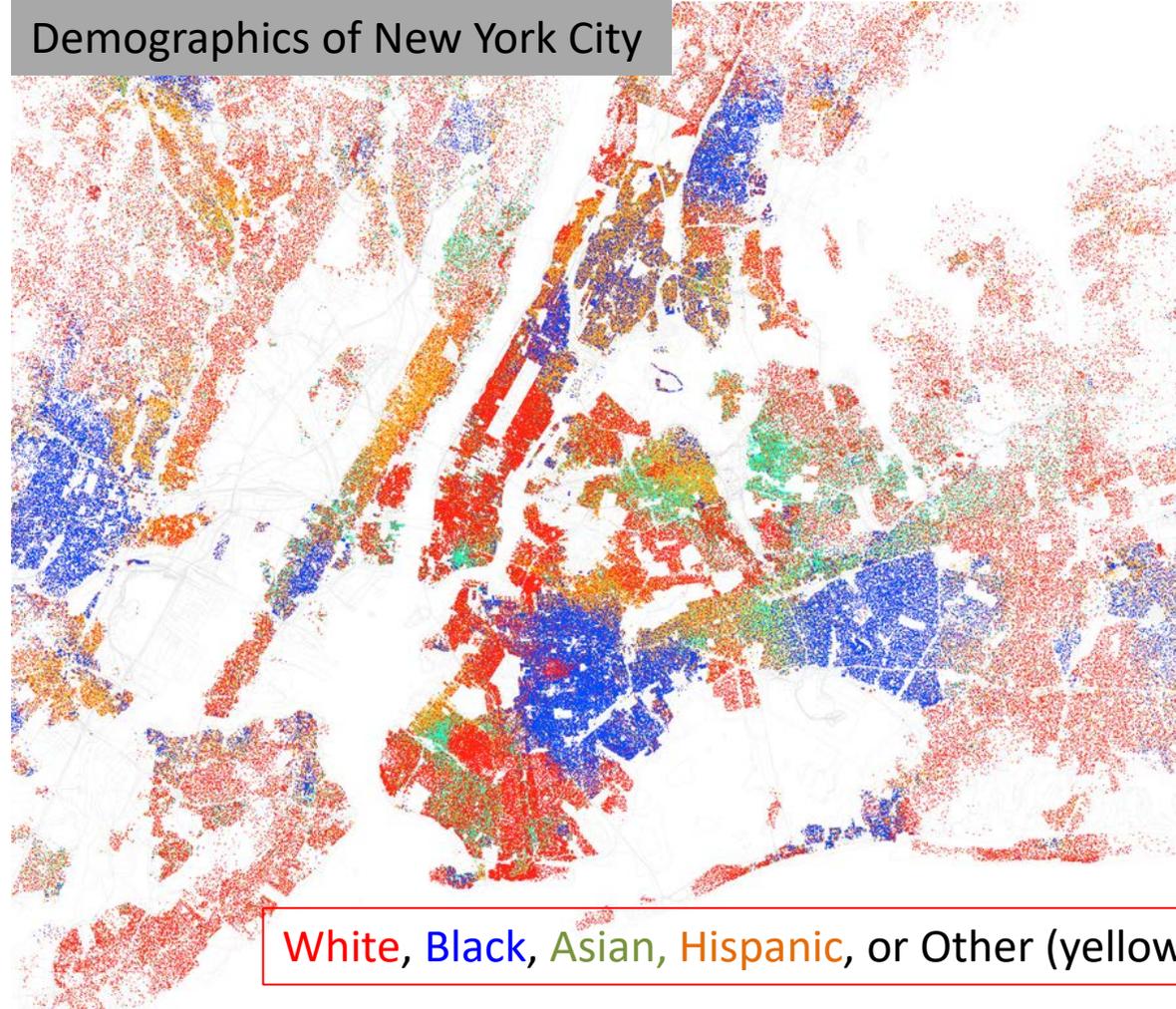


Wie wird die Präzisionsmedizin in der Schweiz aussehen?





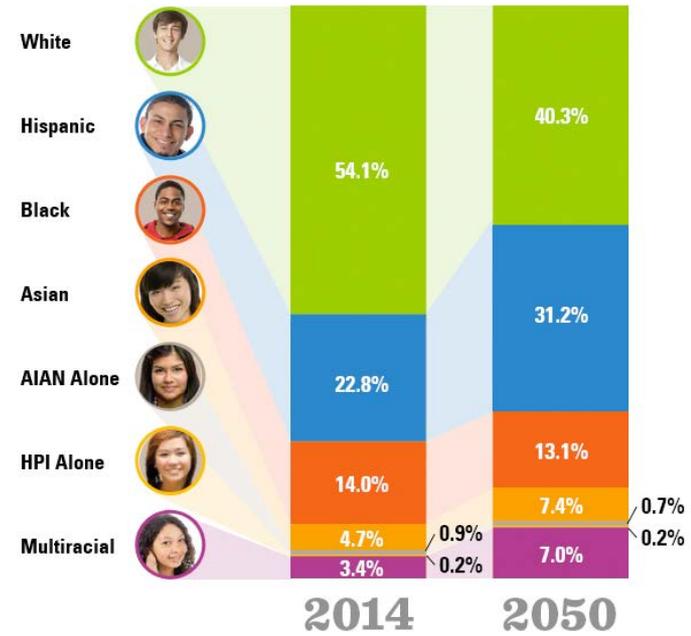
Demographics of New York City



By Eric Fischer - Race and ethnicity 2010: New York City (USA)

Changing USA Ethnicity

Changing race/ethnicity of America's adolescents

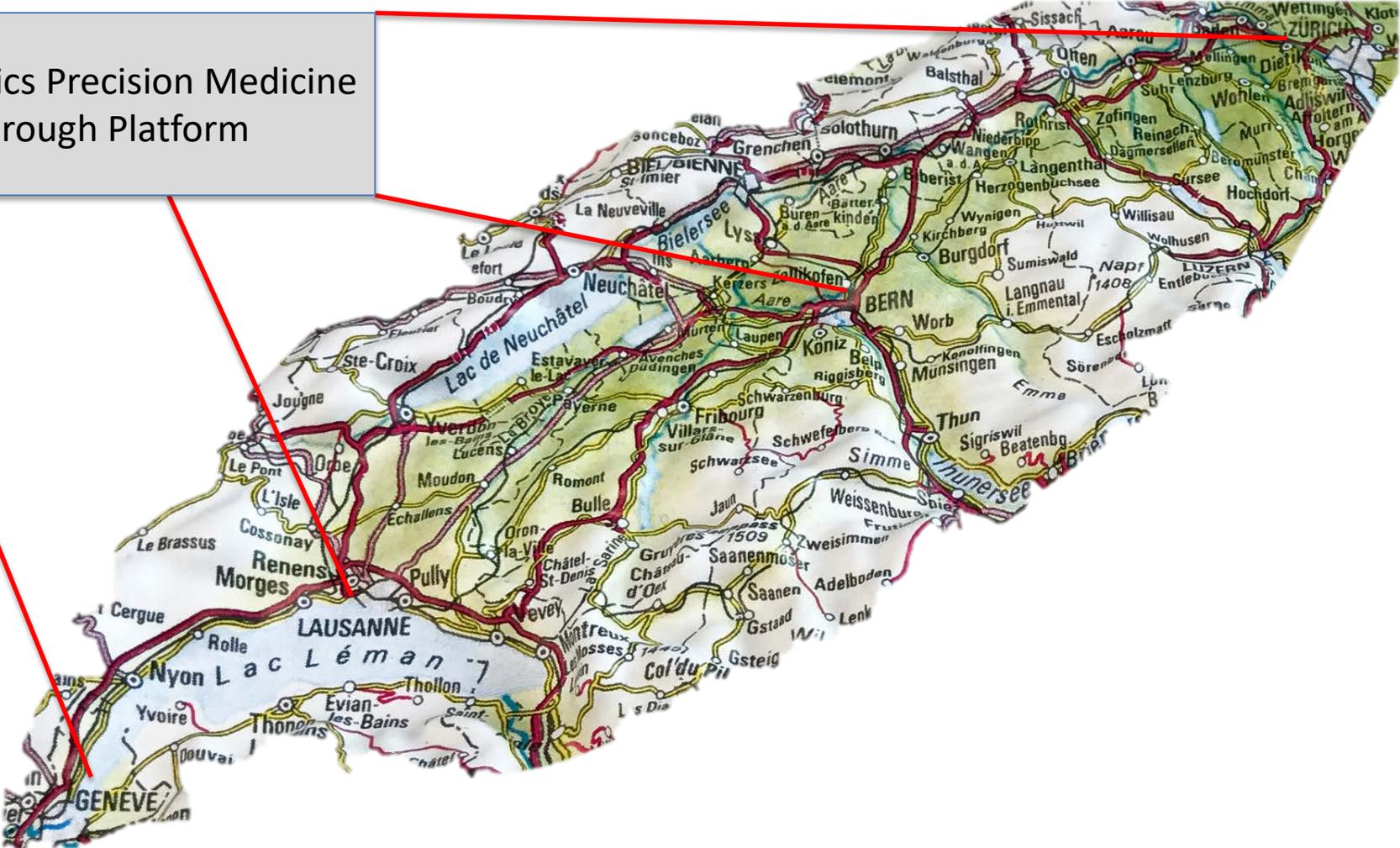


AIAN

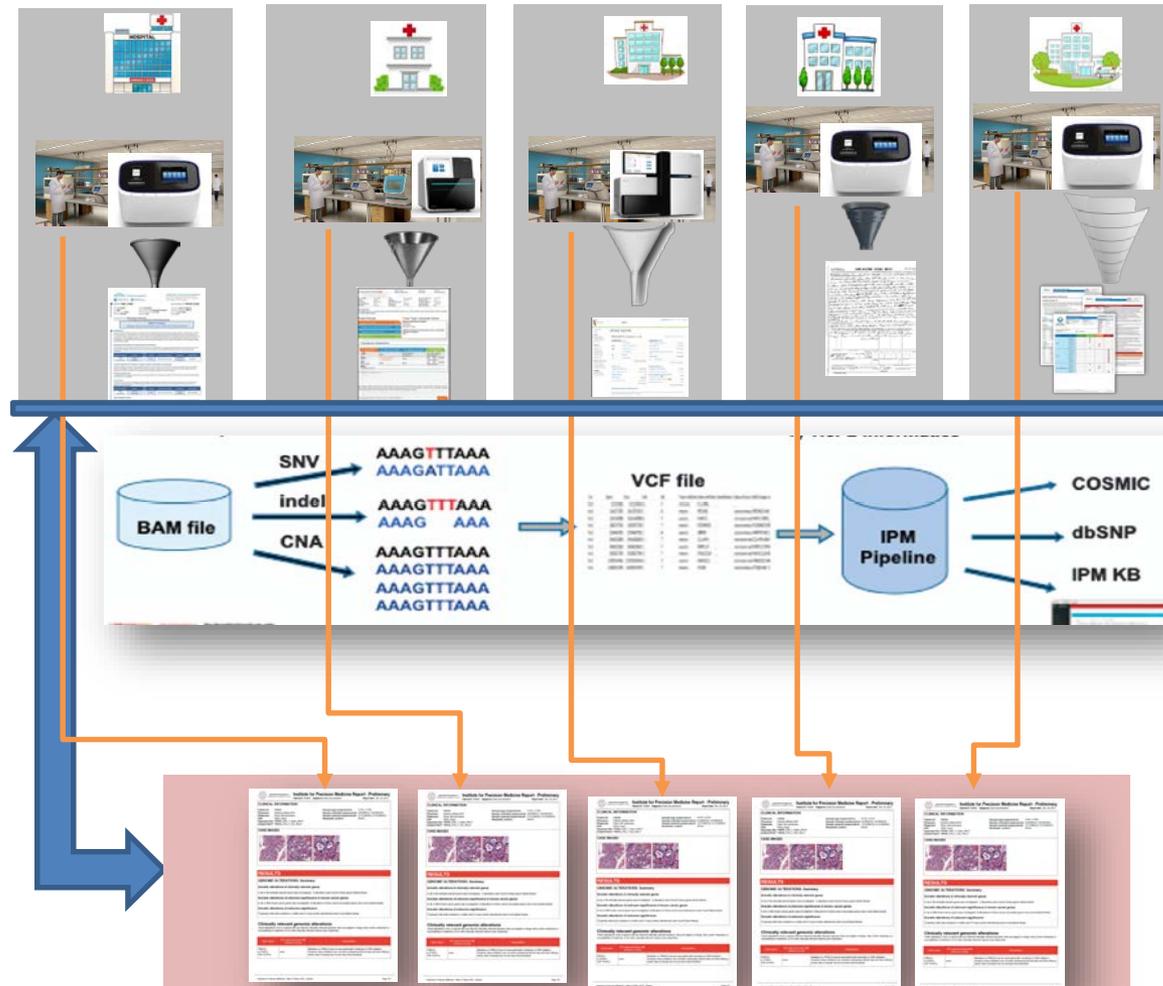
American Indian and Alaska Native

By Mugeek Vidalondon

Genomics Precision Medicine
Breakthrough Platform



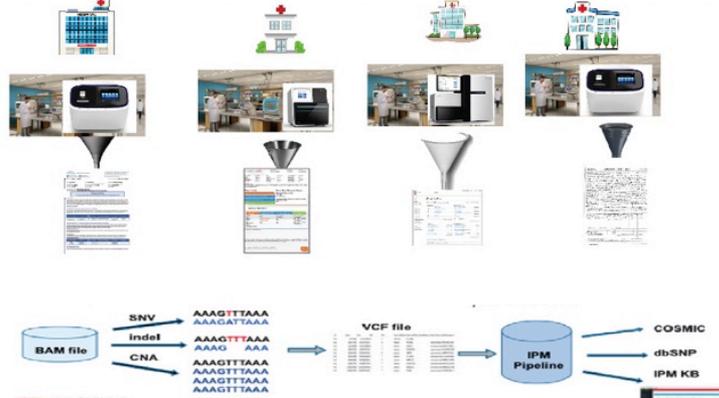
z.b.,
Wie sieht **genomische Analyse** heute in der Schweiz aus?



*Wie können wir den Wert der **genomischen Daten** in der Schweiz verbessern?*

SPHN/PHRT Proposal from Bern, Zurich, Lausanne, and Geneva

UNIVERSITY HOSPITAL UNIFORM REPORTING



WP1. Determine optimal configuration for uniform genomics reporting to enable reliable sharing of clinical and research data. (SIB/Others)



WP2. Harmonize Swiss uniform genomics reporting format(s) for clinical and research interoperability. (Patholink SNF & Onco SPHN)

DATA SHARING FOR RESEARCH & CLINICAL

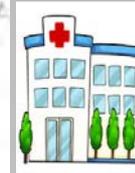
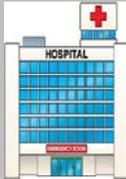
This complex block illustrates data sharing for research and clinical purposes. It features a screenshot of a data sharing interface with various filters and a bar chart. To the right, there is an illustration of a person wearing a magnifying glass over a DNA sequence, symbolizing research and clinical analysis. Below the illustration, a colorful DNA sequence is shown, representing genomic data.

WP3. Establish external SOCIBP sites and improve use/operator features for broader clinical and research connectivity (SIB/SAKK/Intern'l)

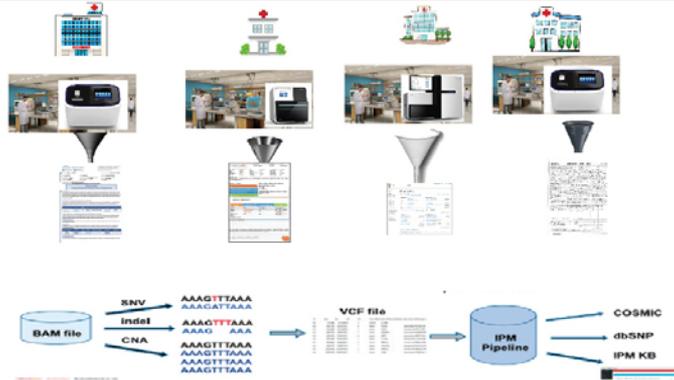




Swiss Genomics Today: Many Solutions and Lack of Uniform Reporting



UNIVERSITY HOSPITAL UNIFORM REPORTING

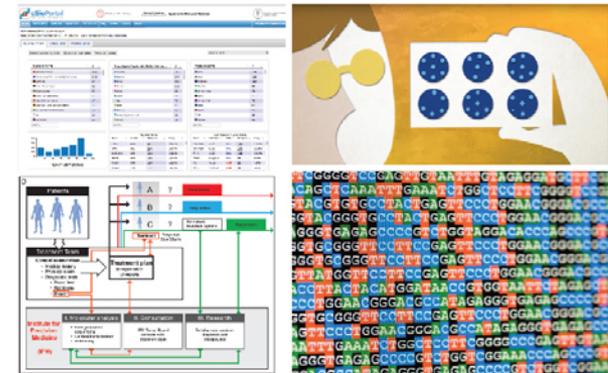


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DATA SHARING FOR RESEARCH & CLINICAL



WP3. Establish external SOCIBP sites and improve use/operator features for broader clinical and research connectivity (SIB/SAKK/Intern'l)



Swiss Molecular Pathology and Tumor Immunology Breakthrough Platform (SOCIBP)

SPHN Driver Project

Universität Bern / Inselspital

Mark A Rubin (Main Applicant)

Tobias Grob (WP2)

Rémy Bruggmann (WP1,3)

Universitätsklinik Zürich

Holger Moch (Co-Applicant, WP2)

PHRT Project

ETH-Zürich

Gunnar Rätsch (Co-Applicant, WP3,5,6)

EPFL-Ludwig Institute

George Coukos (Co-Applicant, WP4,6)

Co-Applicants (WP2)

CHUV Lausanne

Laurence de Leval

Hôpitaux Universitaires Genève

Thomas McKee

Proposed

Other Interactions

Swiss Institute of Bioinformatics

PATHOLINK (SNF)

Englander Institute of PM (Cornell)

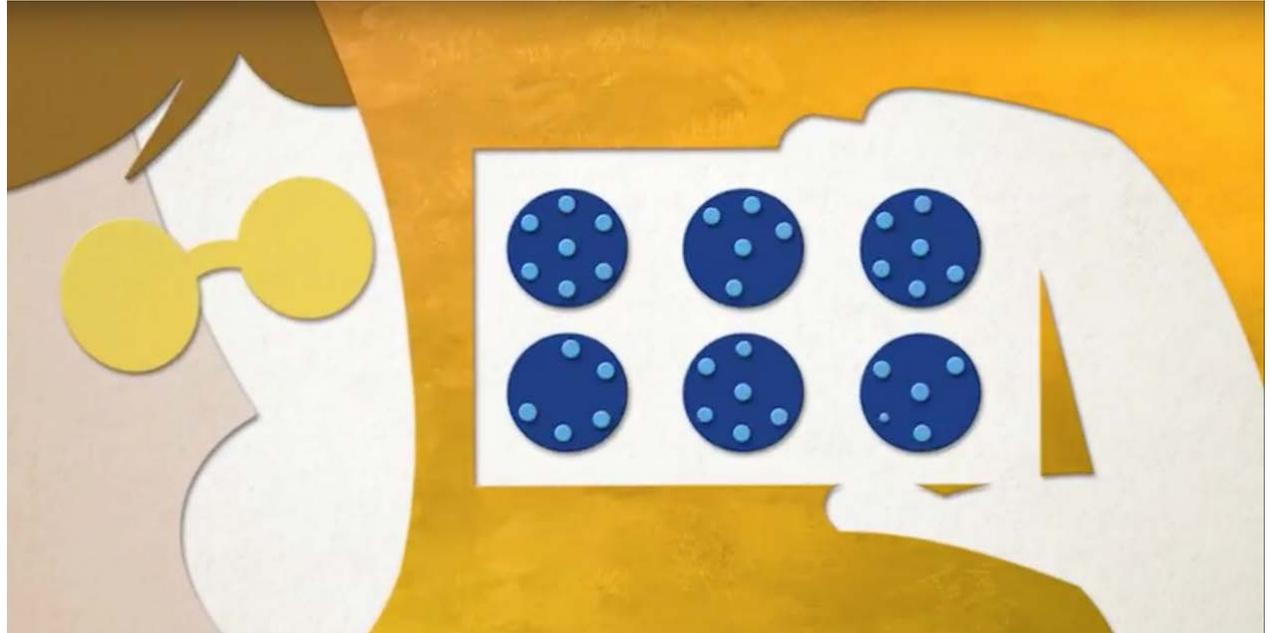
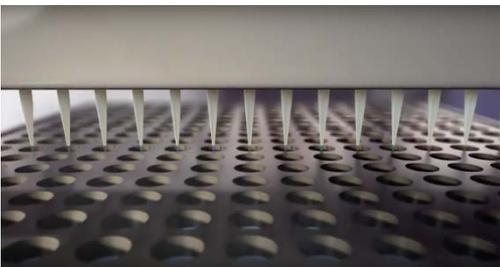
Other SPHN/PHRT Projects (see table)

Was sind die Kosten?

$F(\text{PM}) = \text{Social and economic PM Model}$

			None	\$100
			Weeks	\$1000
			Months	\$10,000
			Years	\$100,000
Condition	Diagnosis	Treatment	Benefit	Cost

Danke für Ihre Aufmerksamkeit



mark.rubin@dbmr.unibe.ch