Model driven research in Brain Diseases Hans Lehrach

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Berlin Institute of Health
Charité
Alacris Theranostics



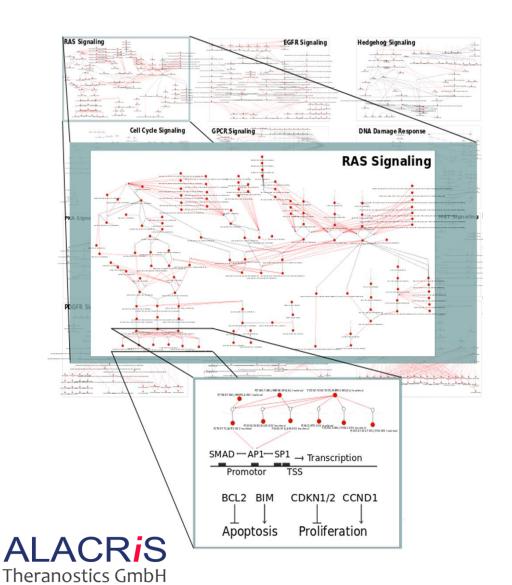
The case for model-driven research

 current research paradigm - identifies biological mechanisms with simple causality

 evolution generates a ,dark mass' of mechanisms not accessible to current research tools.

 Mechanistic models - compare experimental results and prediction to falsify hypothesis too complex to analyse by currently available strategies

ModCell™ - a generic mechanistic model of cellular signalling transduction



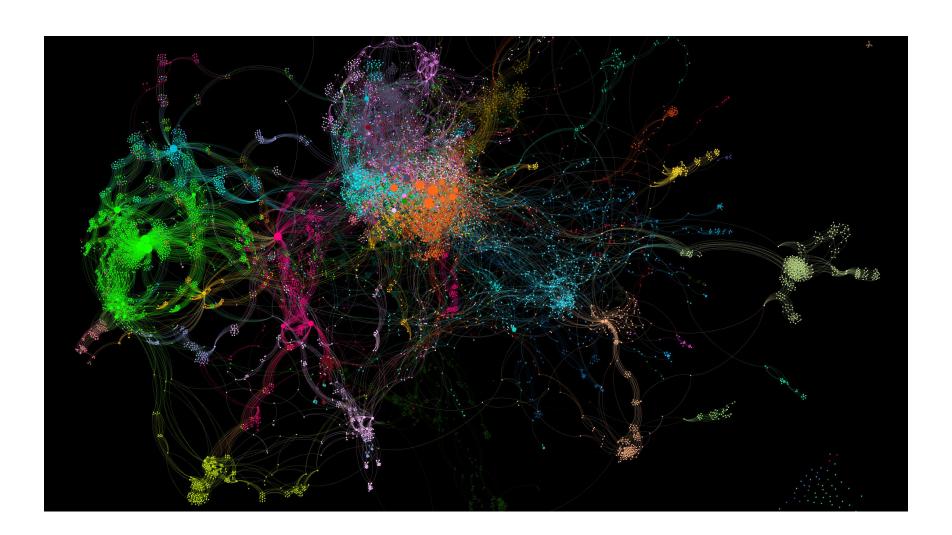
- 50 signalling pathways
- 20,000 reactions
- 800 genes
- 300 targeted drugs

Cancer pathways include:

RTK signaling
HEDGEHOG
NOTCH
WNT
TGFB signaling
Cytokine signaling
Death receptor signaling
GCPR/Hormone signaling
Ephrin signaling

Wierling et al., 2015 PMID:26464088

Human signalling pathways





We need samples for molecular analyses

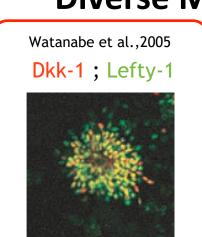
Fresh human samples

Deep brain stimulation-associated brain tissue imprints: a new *in vivo* approach to biological research in human Parkinsons disease

Affif Zaccaria, Ali Bouamrani, Stephan Chabardès, Michèle El Atifi, Eric Seigneuret, Johannes A. Lobrinus, Michel Dubois-Dauphin and Pierre R. Burkhard

Surrogate brain tissue:

Diverse Methods to Generate Brain Organoids

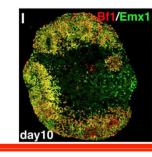


Elkabetz et al., 2008 Nog



Edri et al., 2015 Nog, LDN, SB431542

Gaspard et al.,2008 Intrinsic + Cyclop.



Eiraku et al., 2008

Dkk-1; Lefty-1

Shi et al., 2012 DM/Nog; SB431542 FGF2

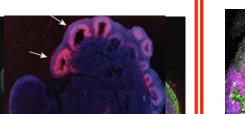


Day 35

PascQiettall., 2001137 LDM, 5643435452,2FGF-i



Llancasteretall,,200173 Intrimsrign(SicHIR)

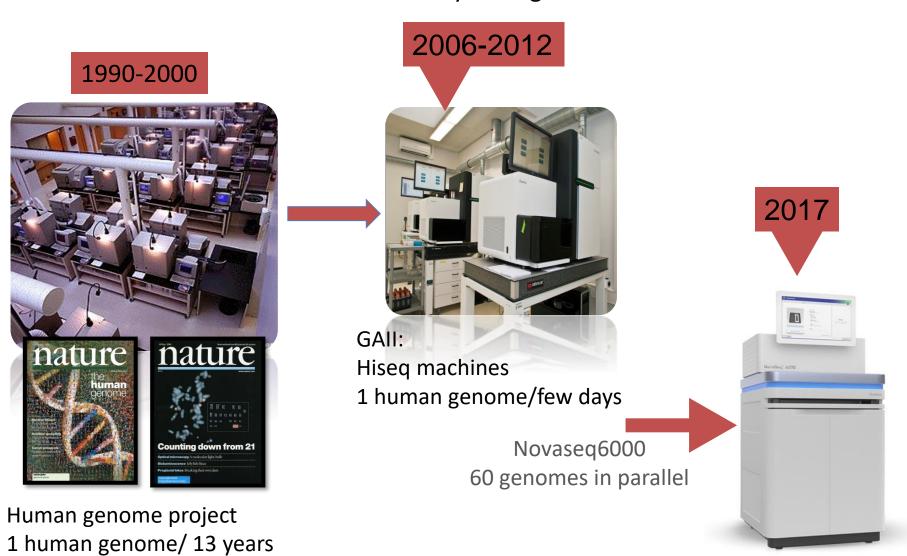


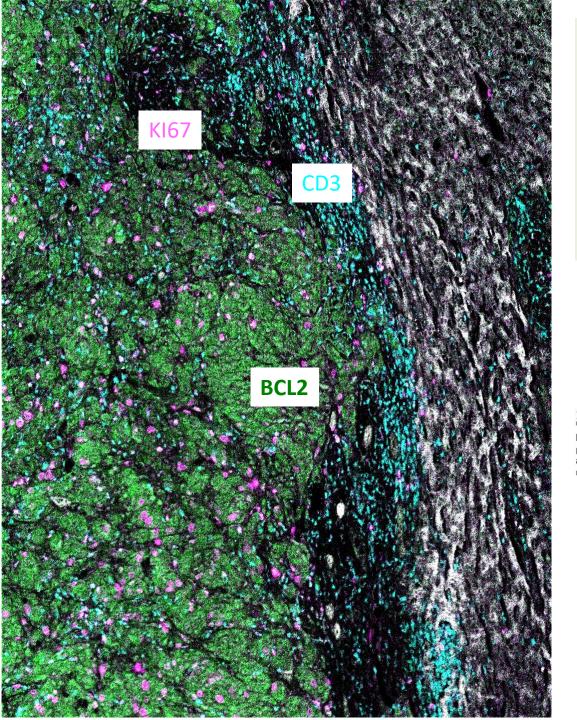
Keedststeigna, 200173 IWR: 1542

Maroof et al., 2013 XAV; LDN; SB431542; Purmorphamine

Genomics dominates "omics"

One NGS run generates as much information as one million runs on the sequencers used 10 years ago





Resolving RNAseq data spatially at the protein level by **Imaging Mass Cytometry** (IMC) 40 Ab in one experiment









Design

panels using IHC-validated antibodies conjugated to metal tags.

Stain

tissues (FFPE and frozen) or fixed cells with metalconjugated antibodies.

Image

biomarkers using precise capture and detection with CyTOF technology.

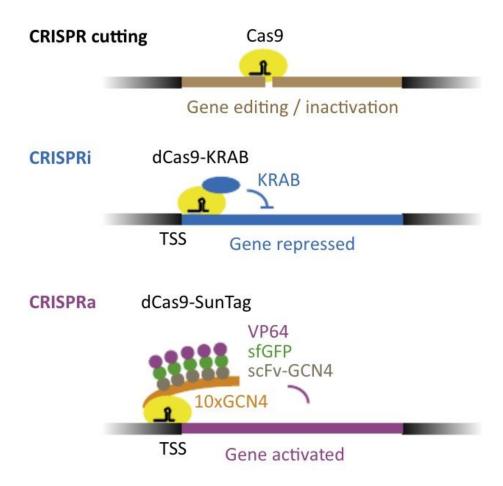
Analyze

using post-analytical imaging and secondary analysis software tools.



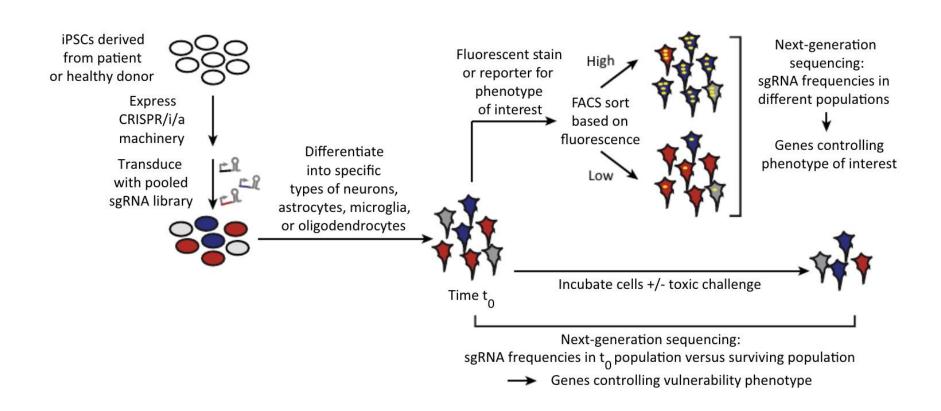
M.-L. Yaspo, Unpublished

CRISPR variants



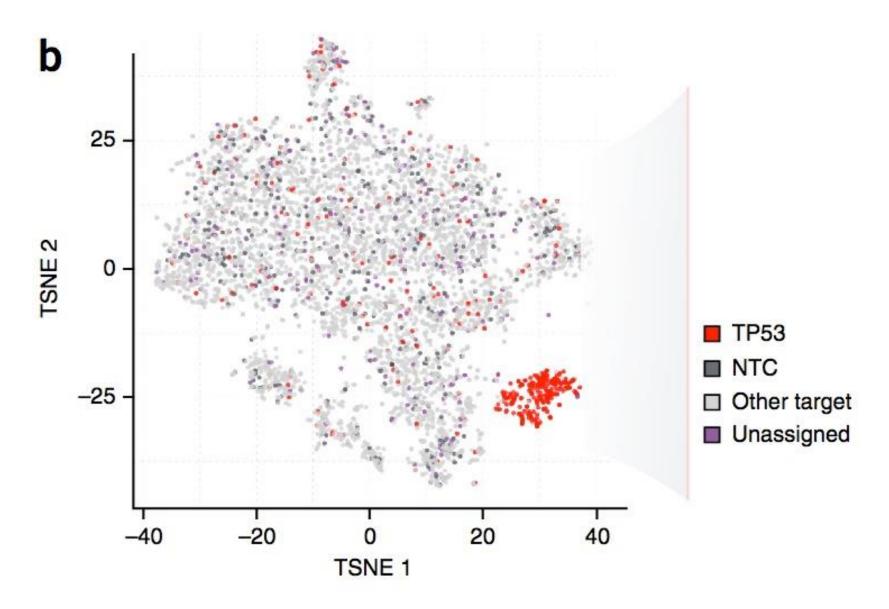
Martin Kampmann, Trends in Molecular Medicine, June 2017, Vol. 23, No. 6, 483-485

CRISPR-phenotype screens



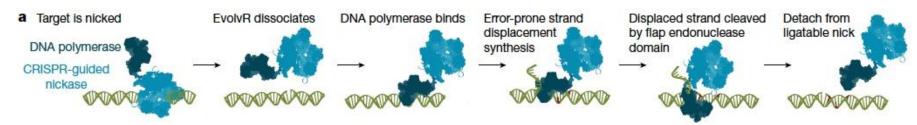
Martin Kampmann, Trends in Molecular Medicine, June 2017, Vol. 23, No. 6, 483-485

CrispR Droplet sequencing: CROPseq (Christoph Bock)



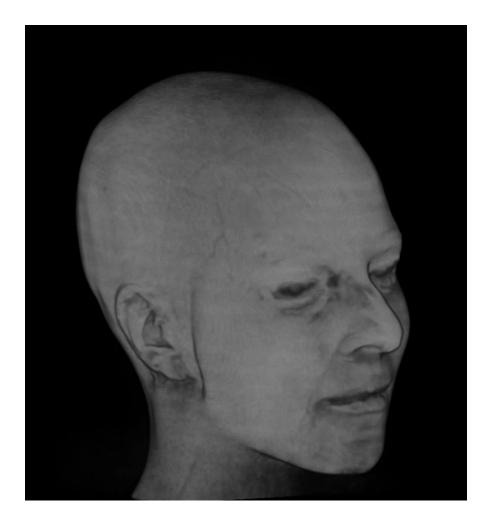
Hill et al., Nature Methods, Feb. 2018

Lineage tracing: EvolvR



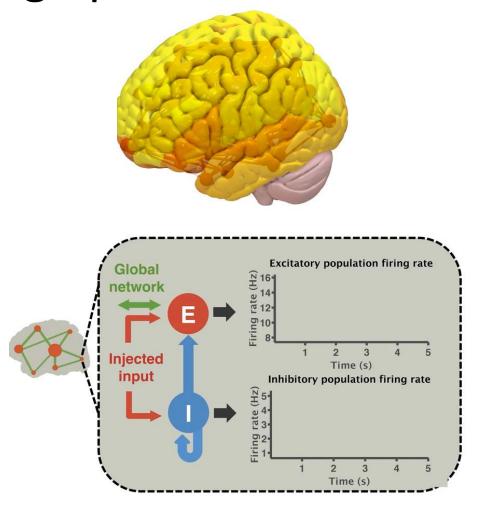
"Brain, New Approach to Brain Diseases" Workshop 21st November 2018

Brain network dynamics simulations



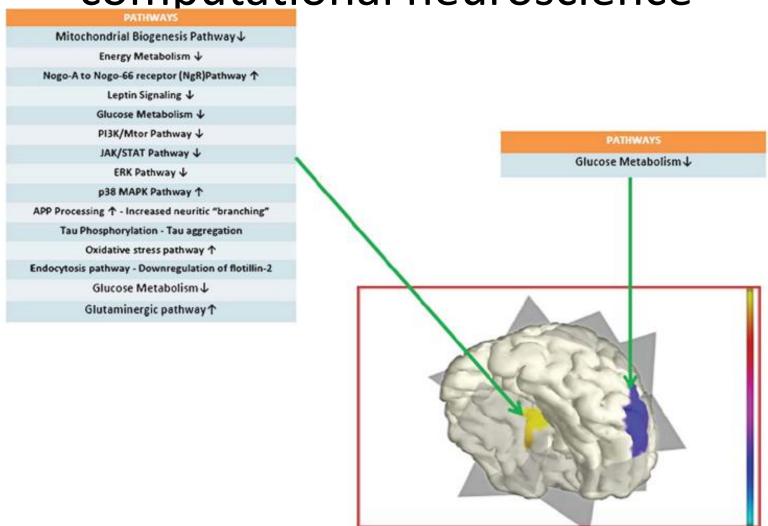
Schirner, Rothmeier, Jirsa, McIntosh, Ritter 2015 **Neuroimage** An automated pipeline for constructing personalized virtual brains from multimodal neuroimaging data

Inferring unobservable mechanisms through personalized simulation



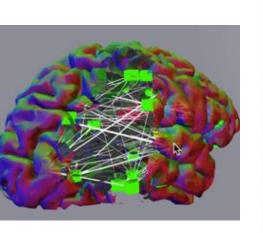
Schirner, McIntosh, Jirsa, Deco, Ritter 2018 **eLife**Inferring multi-scale mechanisms using brain network modelling

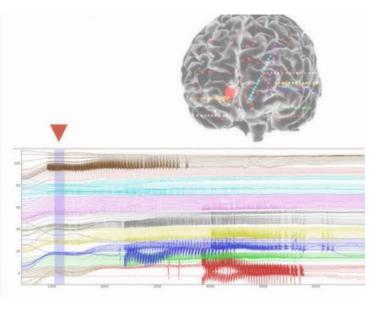
Liniking systems biology and computational neuroscience

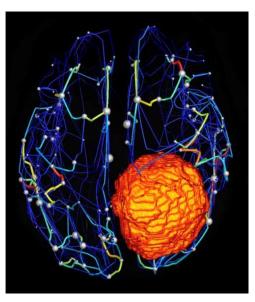


Iyappan et al. 2016 J Alzheimer's Disease. Towards a Pathway Inventory of the Human Brain for Modeling Disease Mechanisms Underlying Neurodegeneration

Applications in Stroke, Epilepsy, Tumors, Dementia, Schizophrenia...

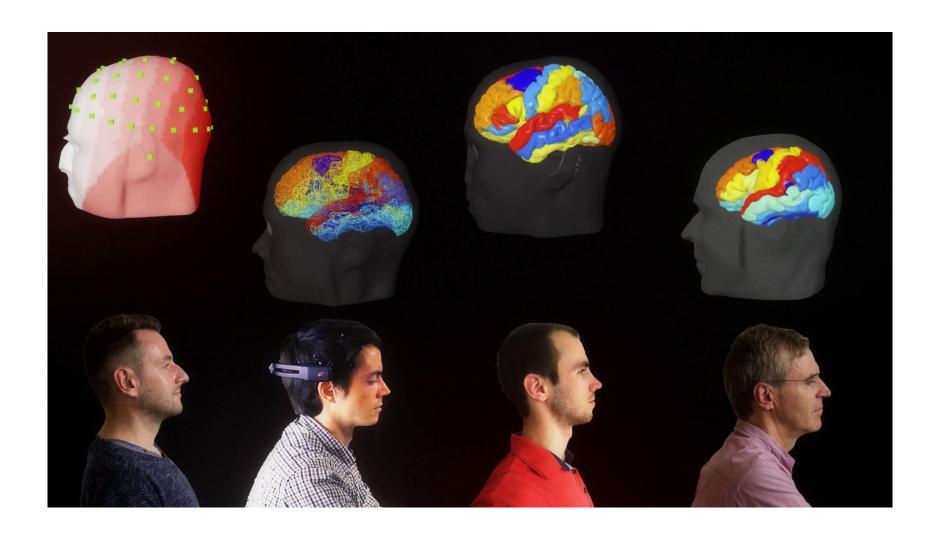






E.g. Proix et al. 2017 Brain; Jirsa et al. 2014 Brain; Zimmermann et al. 2018 Neuroimage Clinical; Aerts et al. eNeuro 2018

100 000 virtual brains from uk biobank



Examples

- Parkinson's disease: models based on –omics analysis of cells recovered during therapeutic deep brain stimulation
- MS: deep immune status, modelling the immune system, analysis of monocygotic twins with/without MS
- Migraine: GWAS modelling, brain simulation (Petra Ritter, Charité)
- •

FET-Flagship project application www.digitwins.org



Without DigiTwins (www.digitwins.org), we will, over the next framework program

- Spend €10 trillion on healthcare
- Spend ~€2 trillion on drugs, to which the individual patients do not respond, and their consequences
- Have the equivalent of the population of Milan die of adverse drug reaction
- Come 7 years closer to the implosion of the European healthcare system
- Make little progress on biological mechanism with complex causality
- Save €1 bn over 10 years, the equivalent of the amount we spend in healthcare every 5 hours (and probably waste every 10 hours).

If you think this could be relevant to Europe and the European Citizens, please mail (lehrach@molgen.mpg.de) or call (+491723839194) me (and tell your colleagues).