COST Action 17140 WG3 Meeting March 2022, Bellinzona

Ulf Dietrich Kahlert, PhD

Tissue-specific progenitor cells derived from gene-engineered human

induced pluripotent stem cells for early-stage drug development





UNIVERSITÄTSMEDIZIN MAGDEBURG





Lost in Translation



Therapeutic group		Phase 1 to Phase 2			Phase 2 to Phase 3			Phase 3 to approval			Overall	
		Total phase transitions	POS _{1,2} , %	(SE, %)	Total phase transitions	POS _{2,3} , %	SE, %	Total phase transitions	$POS_{3,APP}, \%$	(SE, %)	POS, %	(SE, %)
Oncology	No biomarker	9349	28.0	(0.5)	4773	17.4	(0.5)	1159	33.6	(1.4)	1.6	(0.2)
	With biomarker	1136	43.5	(1.5)	742	38.8	(1.8)	77	63.6	(5.5)	10.7	(1.9)
	All	10485	29.7	(0.4)	5515	20.3	(0.5)	1236	35.5	(1.4)	2.1	(0.2)





Seyhan et al. **TMC**. Dowden et al, **Nat. Rev. Drug Disc.**



"Inconsistency in study material is main contributor to irreproducibility." (46%)





Freedmann et al., Plos Biol.

Patient-derived cell line as fundamental study tool in early stage projects







Inconsistency in large pharmacogenomic studies

Benjamin Haibe-Kains^{1,2}, Nehme El-Hachem¹, Nicolai Juul Birkbak³, Andrew C. Jin⁴, Andrew H. Beck⁴*, Hugo J. W. L. Aerts^{5,6,7}* & John Quackenbush^{5,8}*

CCLE (Broad, USA) vs CGP (Wellcome, UK)







Haibe-Kains et al., Nature

Genetic and transcriptional evolution alters cancer cell line drug response

Uri Ben–David¹, Benjamin Siranosian¹, Gavin Ha^{1,2}, Helen Tang¹, Yaara Oren^{1,3}, Kunihiko Hinohara^{1,2}, Craig A. Strathdee¹, Joshua Dempster¹, Nicholas J. Lyons¹, Robert Burns², Anwesha Nag², Guillaume Kugener¹, Beth Cimini¹, Peter Tsvetkov¹, Yosef E. Maruvka¹, Ryan O'Rourke^{1,2}, Anthony Garrity¹, Andrew A. Tubelli¹, Pratiti Bandopadhayay^{1,2,3}, Aviad Tsherniak¹, Francisca Vazquez¹, Bang Wong¹, Chet Birger¹, Mahmoud Ghandi¹, Aaron R. Thorner², Joshua A. Bittker¹, Matthew Meyerson^{1,2,3}, Gad Getz^{1,4}, Rameen Beroukhim^{1,2,3,5,7}* & Todd R. Golub^{1,2,3,6,7}*

Lab-to lab validation

106 cell lines

- 27% CNV could not redetected
- 22% of genome altered by subclonal events

MCF-7-M (27 strains)

- Ten chromosome arms (25% of the genome) were differentially gained or lost
- 283 genes with copy number gains and 405 genes with copy number losses in at least one strain. Only a small minority of these changes (13% of gains and 21% of losses) were detected in all strains.



Ben-David et al., Nature











1100 PDX samples across 24 cancer types, compared to parental cell model (incl. primary model)

We found that large (>5-Mb) CNAs arose rapidly in PDXs: 60% of the PDX models acquired at least one large chromosomal aberration within a single *in vivo* passage, and 88% acquired at least one large aberration within four passages









>Alternative preclinical modeling strategies are warranted:

- 1. to overcome the issue of heterogeneity of results
- 2. aiming to increase frequency of clinical translationability





Patient-derived stem cell models of cancer as reliable source













Biotechnology

Systems & Synthetic Biology · Nanobiotech · Medicine Journal

RESEARCH ARTICLE | 🔂 Open Access

Progenitor cells derived from gene-engineered human induced pluripotent stem cells as synthetic cancer cell alternatives for in vitro pharmacology

Constanze Uhlmann, Ann-Christin Nickel, Daniel Picard, Andrea Rossi, Guanzhang Li, Barbara Hildebrandt , Gabriele Brockerhoff, Farina Bendt, Ulrike Hübenthal, Michael Hewera, Hans-Jakob Steiger, Dagmar Wieczorek, Aristoteles Perrakis, Wei Zhang, Marc Remke, Katharina Koch, Julia Tigges, Roland S. Croner, Ellen Fritsche, Ulf D. Kahlert 🗙 ... See fewer authors

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Lineage differentiation







Uhlmann et al., BTJ

In vivo tumorgenicity potential ?







Krause et al., in prep.

substance	NPC WT	NPC -EV	NPC -GLI1	NPC -c- MYC	NPC - TP53R175 H	NPC - EGFRvIII	NPC– EGFRvIII/ TP53R175H
Panobinostat	0.05164	0.2940	0.4568	0.1237	0.01986	1.367	0.2856
Vinblastine sulfate	0.01518	0.003848	0.4806	0.01955	0.02161	7.166	3.291
Lomustine	13.96	188.8	2410	217.10	191.30	1364	270
Duvelisib	14.04	75.29	53.4	48.68	732.90	1062	322.50
Paracetamol	Х	Х	х	Х	Х	х	x
Staurosporine	0.1118	0.5545	0.1986	0.2675	1.657	1.639	0.1599









Uhlmann et al**., BTJ**



500ng gRNA, 500ng Cas9





Rees & Liu, Nature Methods

• Single cell of origin

Reduction of heterogeneity?

• Genetic reduction





• Biomarker specific

Better molecular targeted /anti- cancer stem cell specific test system ?

 stem cell/ progenitor cell specific







isogenic controlled conditions amendable for multi-lineage differentiation comparisons

oncogenic potential of onco-proteins, early vs. late stage cancer progression models, Targeting cancer stem cell niche...





Preclinical testing matrix

Synthetic alternatives



- **Diversity (gender, immune, ethnic...)**
- Patient-specific iPS generation
- (episomal transformation, via electroporation, NEPA)
- > multi-factor and adjustable

Patient-derived systems





Off-target assessment

Pre-analytic issue: patient recruitment, consent, OR surveillance and reminder biomaterial logistics Cooperation pathologist





Translational MES biobank (since 01/22)

- · Cooperation with surgical partners Roland Croner and Aris Perrakis & surgical team of the clinic for surgery OVGU
- Inspired by German Institute for Normalization (D.I.N.) standards (Prof. Kahlert is a permanent member "Biobanks/ Bioresources" - NA 063-09-02-02 AK)
- Comprised high quality tissue and blood acquisition (minimal pre-analytical time window)
- · Association of clinical data at time point of operation and follow-up
- Development of functional models of healthy and tumor tissue to present comprehensive, patient-matched in vitro banking to conduct off-target assays of drug candidates in immunological and genetically matched background associated to tissue
- · Pancreas, Colon, Rectum, Liver, hepatic metastasis







Thank you







ulf.kahlert@med.ovgu.de www.mes.ulf-kahlert.com



COST WG Workshop 16/17 June Magdeburg



