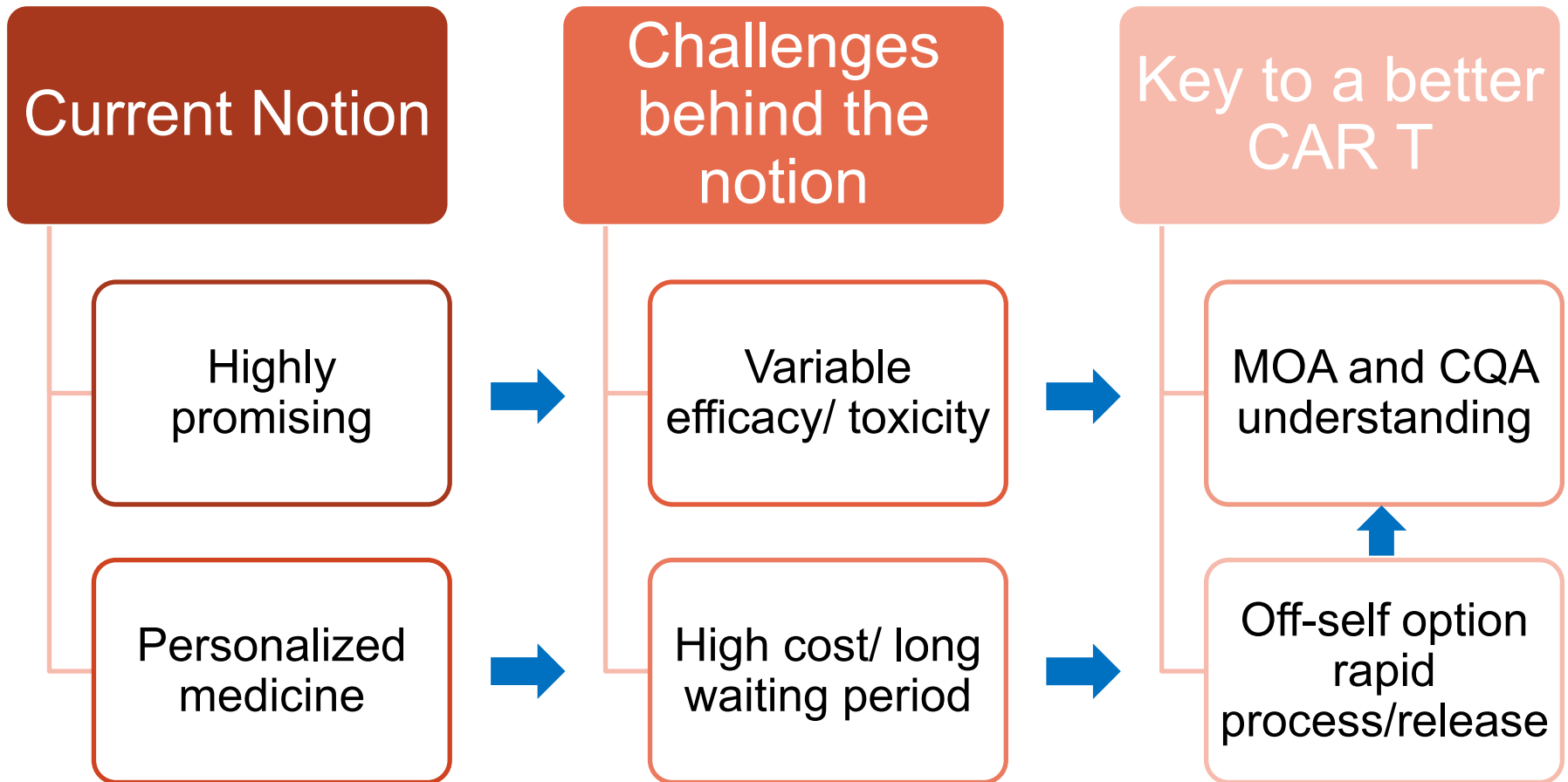




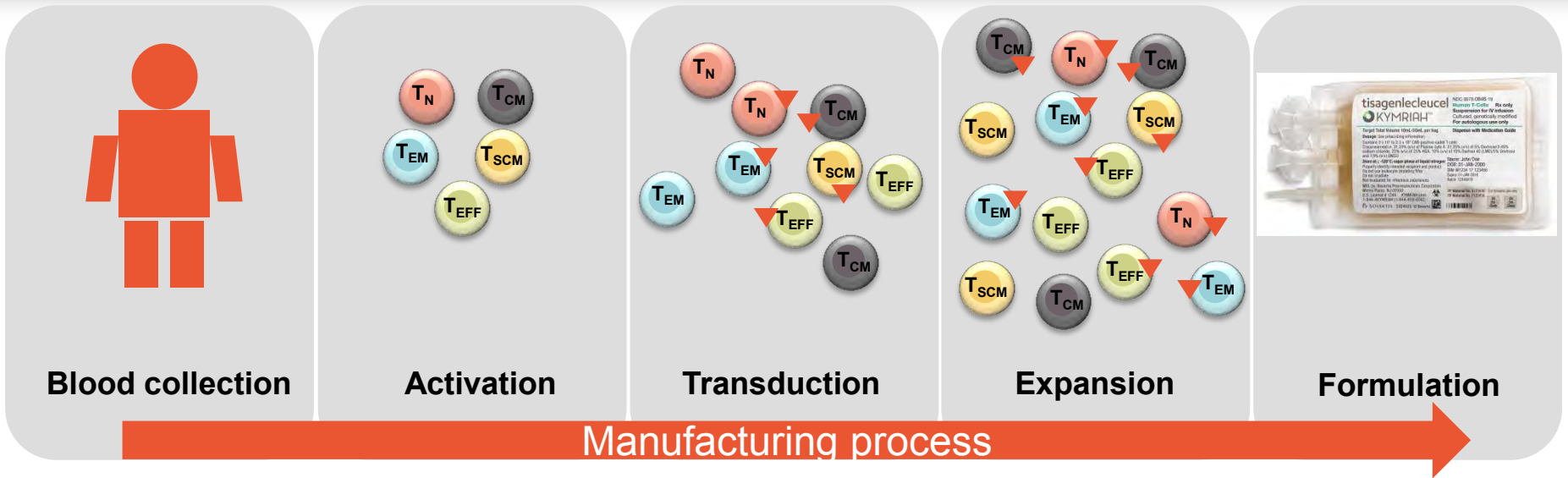
Advancing Product Analytics for CAR T Cell Therapies

Qiong (Chelsea) Xue
Cell Therapy
Takeda Pharmaceuticals

CAR T Cell Therapies

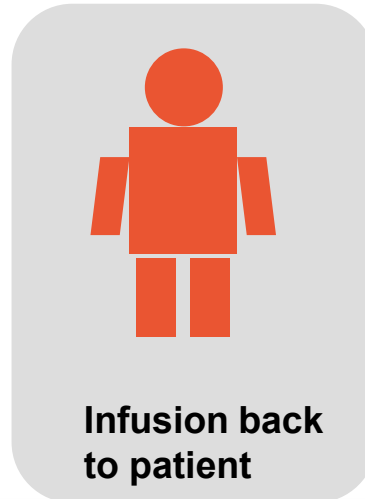


What's in the bag?



Complicated MOAs in patients:

- **Cytotoxic activity:** cytotoxines to promote cancer cell apoptosis
- **Proliferation:** interleukines to promote immune cell development and division
- **Immune cell activation:** inflammatory cytokines to promote inflammation and immune response; chemokines to recruit immune cells



Complex compositions in final products:

- T cells and non-T cells
- Transduced and untransduced
- CD4/CD8 ratio
- T_N , T_{CM} , T_{EM} , T_{SCM} , T_{EFF}
- Th1, Th2, Th9, Th17
- Multiple cytokine/chemokine
- Different degree of exhaustion
- Different degree of senescence

Chemistry, Manufacturing, and Control (CMC) Information for Human Gene Therapy Investigational New Drug Applications (INDs)

- FDA guideline, July 2018

“Your summary should also include a description of **critical quality attributes (CQAs) that are relevant to the safety and biological activity** of the product as they are understood at the time of submission. ... we recommend that **you evaluate a number of product characteristics during early clinical development to help you identify and understand the CQAs of your product.** This will also help ensure your ability to assess manufacturing process controls, manufacturing consistency, and product stability as product development advances.”



EUROPEAN MEDICINES AGENCY
SCIENCE MEDICINES HEALTH

26 July 2018
EMA/CAT/GTWP/671639/2008 Rev. 1
Committee for Advanced Therapies (CAT)

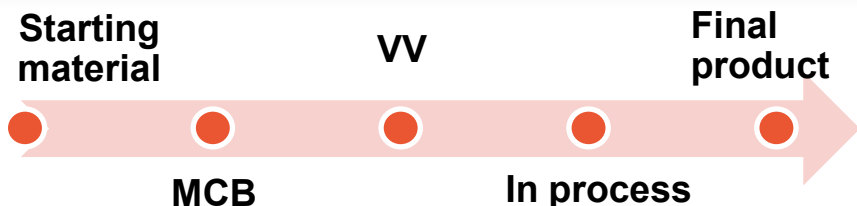
Guideline on quality, non-clinical and clinical aspects of medicinal products containing genetically modified cells

“Rigorous characterization of the genetically modified cell medical products (either alone or in combination with a medical device) is essential.”

The characteristics including:

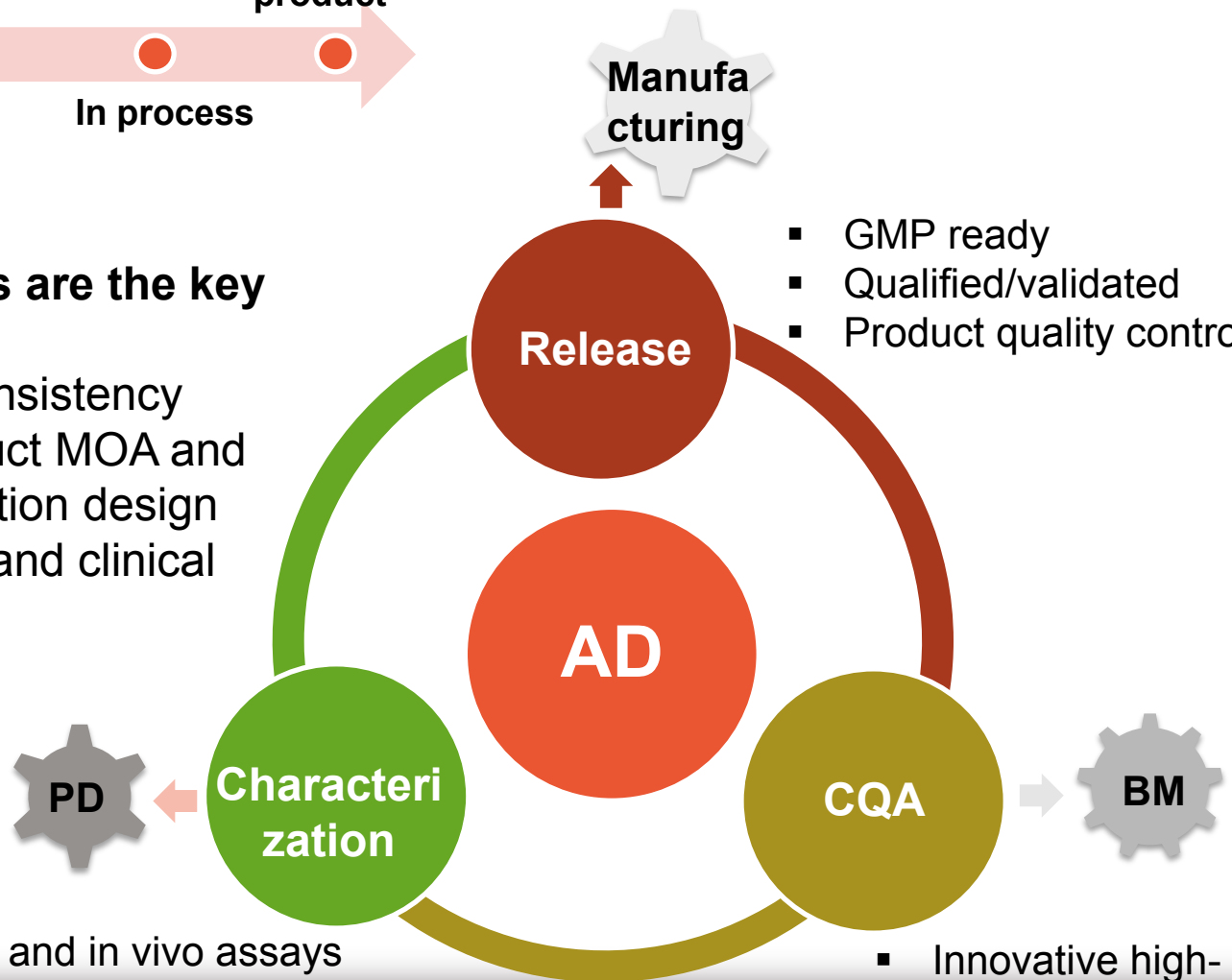
- Heterogeneity of the cell population (e.g. percentage of sub-populations)
- Proliferation and/or differentiation capacity of the genetically modified cells
- Cell functionality (other than proliferation/differentiation)
- Potency testing for products containing genetically modified T-cells against tumor cells (e.g. CAR-T cells) is preferably based on the cytotoxic potential of the T-cells)

Analytical development for cell therapies



Integrated analytics are the key

- Product quality
- Manufacturing consistency
- Understand product MOA and drive next-generation design
- Patient selection and clinical prediction





- GMP ready
- Qualified/validated
- Product quality control

- In vitro and in vivo assays
- Assess process changes
- Ready to be included in IND

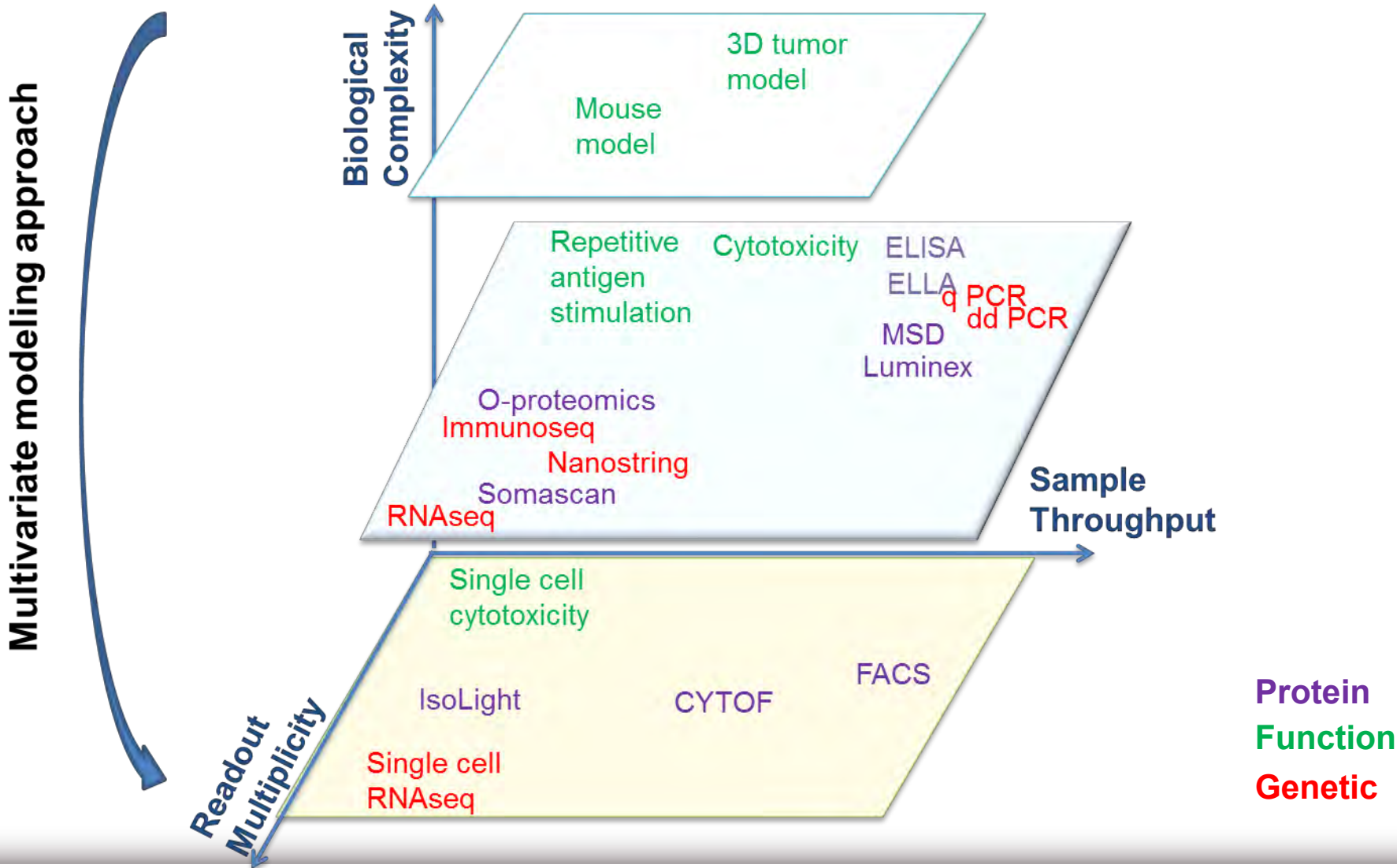
- Innovative high-dimensional platforms
- Enable correlative studies

Product analytics for release testing



	 KYMRIAH [®] (tisagenlecleucel) <small>Suspension for IV infusion</small>	 YESCARTA [®] (axicabtagene ciloleucel) <small>Suspension for IV infusion</small>
Appearance	Visual inspection	Visual inspection
Identity	Presence of CAR transgene, by q-PCR	Presence of scFv heavy chain variable region, linker and CD28 sequences
Dose	Viable cell count; CAR expression by flow cytometry	Viable cell count; anti-CD19 CAR expression
Potency	Release of IFN γ in response to CD19-expressing target, by ELISA	Cell viability; anti-CD19 CAR expression
Purity/impurity	Percentage of viable T cells; percentage of viable CD19+ B cells; determination of residue beads; endotoxin	Gentamicin; endotoxin
Sterility	Pharmacopia (USP 71/EP/JP) growth method	Pharmacopia (USP 71/EP/JP) growth method
Mycoplasma	USP compendia method	USP compendia method; q -PCR
RCL	Determination of VSV-G DNA by q PCR; co-culture RCL	Co-culture RCR; presence of scFV heavy chain variable region, linker and CD28 sequences

Analytical platforms enabling comprehensive product characterization



Single cell analysis reveals CQA of CD19 CAR T for NHL

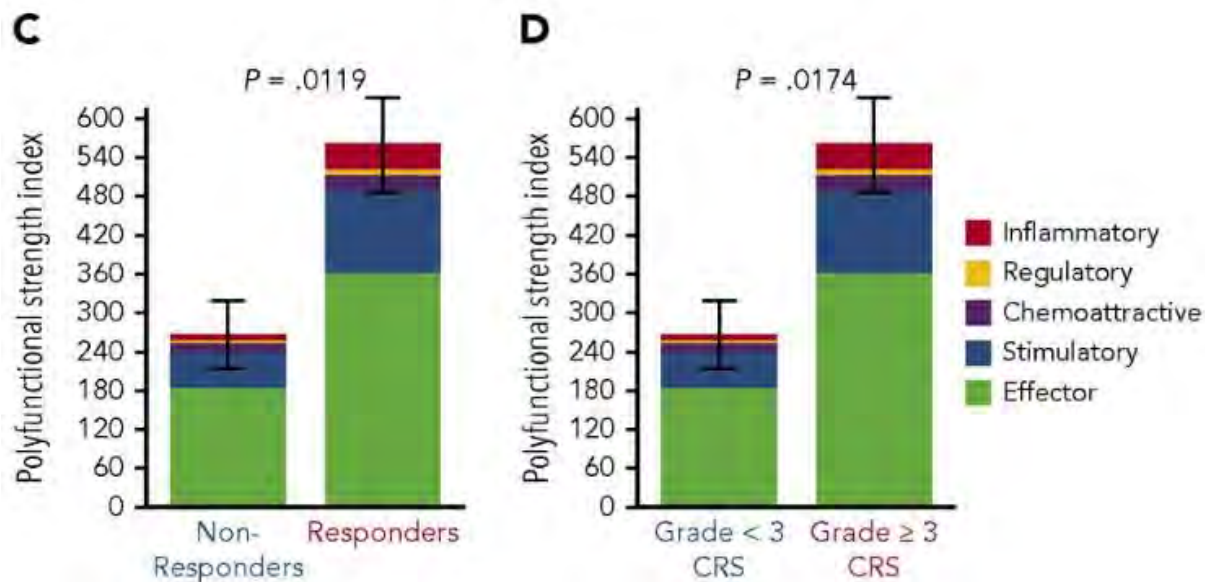


Prepublished online June 12, 2018;
doi:10.1182/blood-2018-01-828343



Preinfusion polyfunctional anti-CD19 chimeric antigen receptor T cells associate with clinical outcomes in NHL

John Rossi,^{1*} Patrick Paczkowski,^{2*} Yueh-Wei Shen,¹ Kevin Morse,² Brianna Flynn,² Alaina



Comprehensive analytics uncover CTL019 CAR T CQA for CLL



nature
medicine

LETTERS

<https://doi.org/10.1038/s41591-018-0010-1>



Determinants of response and resistance to CD19 chimeric antigen receptor (CAR) T cell therapy of chronic lymphocytic leukemia

Joseph A. Fraietta^{1,2,3}, Simon F. Lacey^{1,2,3,9}, Elena J. Orlando^{4,9}, Iulian Pruteanu-Malinici⁴, Mercy Gohil²,

Transcriptional profiling - RNAseq

- CR: memory-related genes, IL-6/STAT3 signatures
- NR: effector differentiation, glycolysis, exhaustion and apoptosis

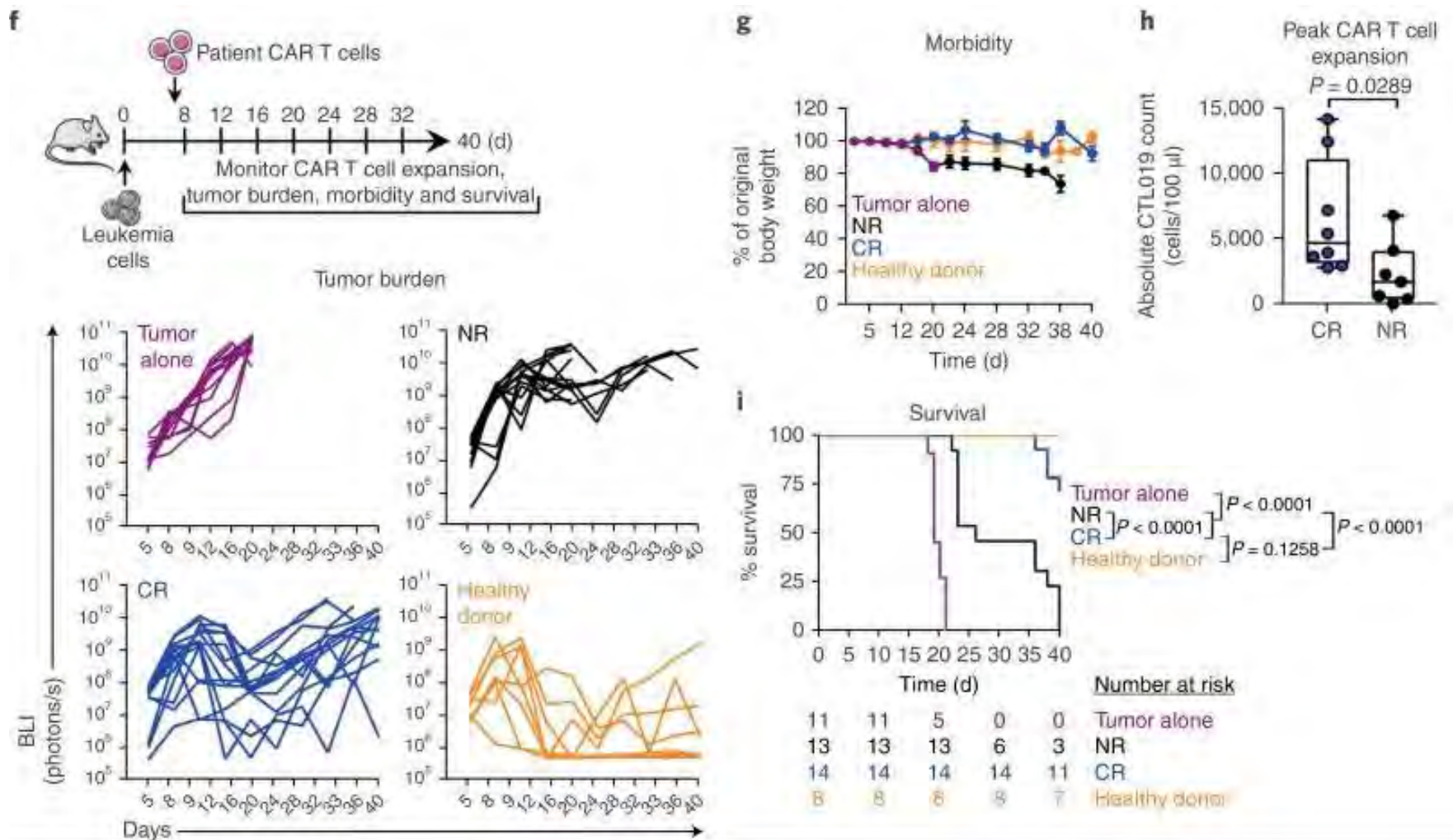
Phenotypic - FACS

- CR: %CD8+CD45RO-CD27+
- NR: %CD8+PD-1+; %CD8+PD-1+TIM-3+; %CD8+PD-1+LAG3+

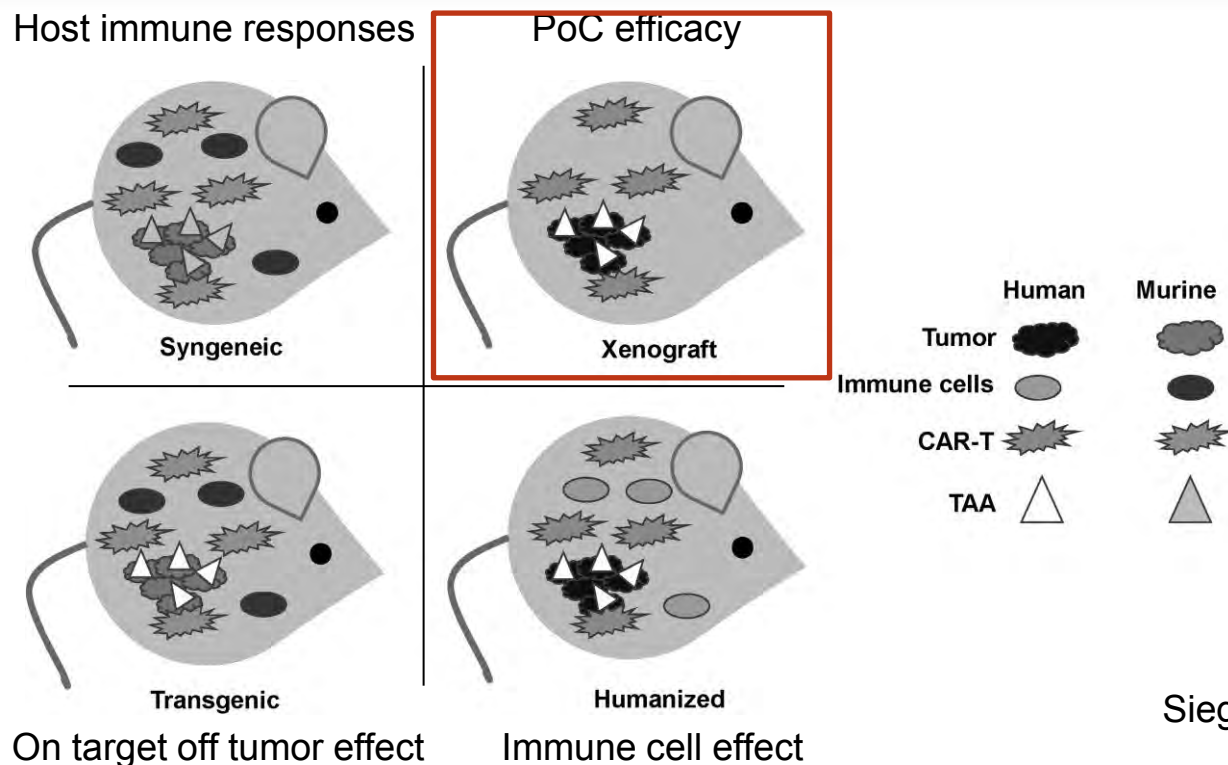
Functional - cytokine/ chemokine by Luminex

- CR: highly functional CAR T cells produced STAT3-related cytokines IL-6, IL-17, IL-22, IL31, CCL20

Capability of mouse models in differentiating an efficacious vs. non-efficacious products



Mouse model used for product characterization



Measurements:

- 1) Efficacy comparison between products generated from different processes
- 2) In vivo CK – serial blood sample for flow cytometry/qPCR
- 3) Cytokine/chemokine production
- 4) Tumor/tissue pathology

Moving beyond Kymriah and Yescarta



Future CAR Ts

Hematopoietic

Solid tumor

- Armored CAR T  Memorial Sloan Kettering Cancer Center
- Smart CAR T 

Autologous

Allogeneic

- Healthy donor derived



- iPSC derived



Memorial Sloan Kettering Cancer Center

- $\gamma\delta$ T cell derived



Additional analytical considerations for future CAR T therapies



New design of CAR T constructs:

- Expected effect need to be confirmed at the cellular level
- The regulation of transgene expression or the intended elimination of the genetically modified cells should be evaluated for proper function
- For secreted gene products the distribution and persistence of the transgene product should be included in the analysis

Genome editing:

- Induced off-target changes should be identified using appropriate bioinformatics tools for in silico screening as well as deep sequencing techniques. This is expected to include appropriate screening of genome-edited cells for off-target effects that may be missed by in silico prediction.
- The persistence of genome editing tools in the cells should be evaluated.

- Guideline on quality, non-clinical and clinical aspects of medicinal products containing genetically modified cells, EMA, July 2018

Additional analytical considerations for future CAR T therapies

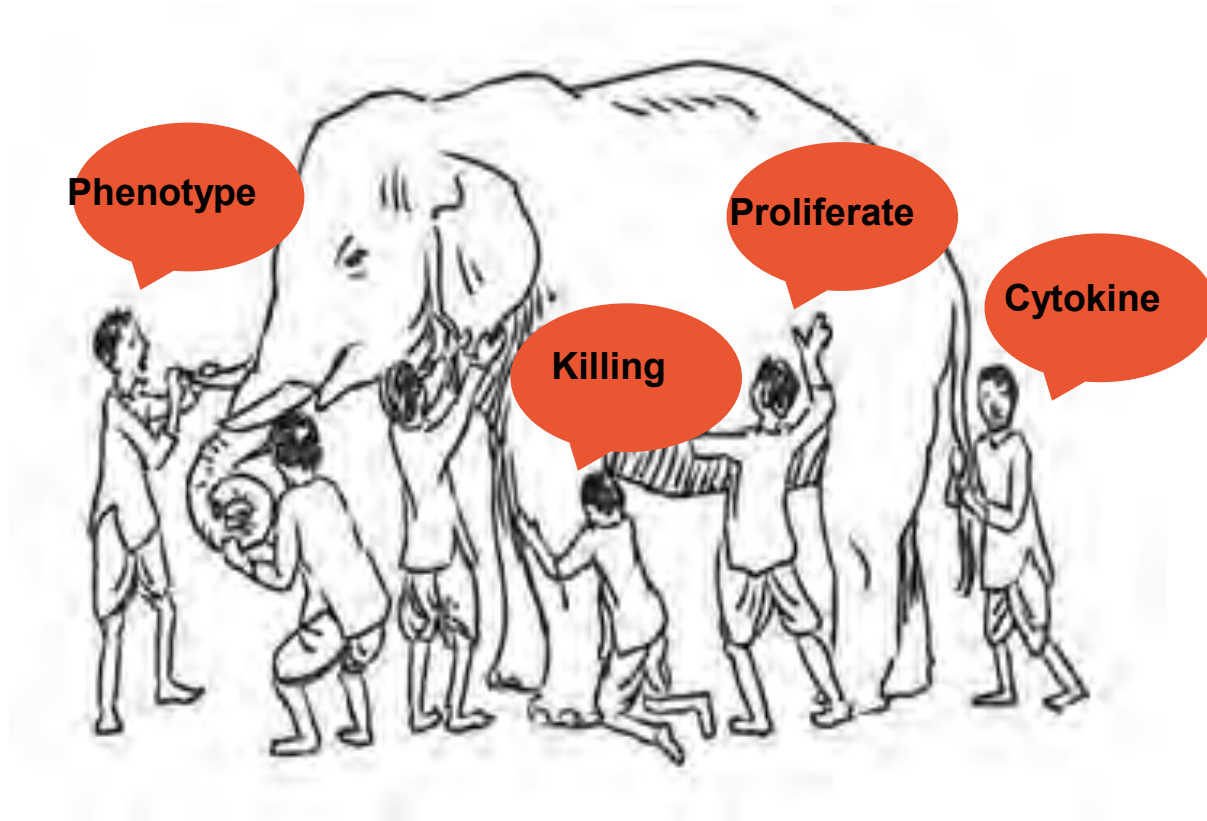


iPSC:

- iPS cells may carry an inherent risk of tumourigenicity
- The nonclinical qualification of the level of undifferentiated iPS cell impurities need to be addressed
- Reprogramming may induce epigenetic changes in the cells with consequences that are not yet fully understood. A variety of high-throughput methods are available for evaluation of the genetic and epigenetic profiles of the iPS cell lines and their derivatives.

- Guideline on quality, non-clinical and clinical aspects of medicinal products containing genetically modified cells, EMA, July 2018

Comprehensive product characterization is the key to true understanding of CAR T therapies



Modified from www.williepietersen.com

Thank you

