



key point

- ★ A novel therapy to tackle tumor heterogeneity
- ★ A combination of antibody library technology and chimeric antigen receptor technology
- ★ Putting clonal selection theory into practice
- ★ A vivid interpretation of Whack-a-Mole game in the life science

Abstract

When we play **Whack-a-Mole**, we keep hitting them with our little hammers, but they always pop up on the other side of the machine, like neoantigens that are constantly appearing as tumors grow and evolve. To address this problem, we conceived and preliminarily tested the **CAR-NK92 library therapy** based on antibody library technology.

We have a conjecture that mRNAs encoding antibodies extracted from the peripheral blood of healthy humans, can construct the antibody library after negative selection, and the CAR-NK92 library consisting of NK-92 cells loading the antibody library can target all tumor antigens.

Background

A new strategy:
Expressing more CARs in one cell
AND
Constructing a CAR-immune cell library by expressing different CARs in different cells

Reason:

1. The barrier of tumor heterogeneity
2. Side effects like GVHD (graft versus host disease) and CRS (cytokine release syndrome)

Modeling

1. Limitations on the number and types of CARs that can be expressed in a single cell

We derived the relationship between the number of types of CARs expressed by individual immune cells and the rate of antigen evasion and over-misfiring of normal cells.

Method: Bioinformatics analysis

Result: Taking colon cancer cells as an example, the number of CAR types should be controlled within 4-8 types. And related research has shown that transfection of multiple CARs on a single cell is challenging and expression of multiple CARs on a single cell is prone to ligand non-dependent tonic signaling.

Decision: To express multiple CARs on a large number of immune cells!

2. Expressing multiple CARs on a large number of immune cells

2.1 CAR-NK92 suicide gene pathway feasibility model

Method: Improved CAR activation model, based on the original Makalkithan kinetic proofreading model

$$\frac{C_N}{C_T + R_T} = \frac{M}{M + K_D} \frac{1 - \frac{\alpha M}{\beta K_D}}{\frac{\alpha M}{\beta K_D}}$$

Result: The biological properties of CAR-NK92 cells in terms of sensitivity, specificity and time efficiency in the recognition of exogenous molecules were reflected more successfully and realistically. It has implications for predicting the activity of CAR-NK92 cell gene circuits in animal and clinical experiments.

2.2 AP1903 pro-apoptosis model

Method: Optimizes the linear model equation and obtains a relatively complete effect change equation to describe the relationship between efficacy, time and concentration of AP1903.

$$E = k_1 \times C_p + E_0$$

the linear model equation

$$E = E_0 \pm \frac{E_{\max} C_p}{C_{V_{\max}} + C_p}$$

a relatively complete effect change equation

Result: When the initial concentration was set at 0.05, 0.1, 0.5 and 1.0mg/kg and AP1903 was injected at an interval of 12h, the relationship of plasma pharmacodynamics with time was observed.

When the initial concentration was set at 0.05, 0.1, 0.5 and 1.0mg/kg and AP1903 was injected at an interval of 24h, the relationship of plasma pharmacodynamics with time was observed.

3. Cellular automata visualization simulation

Visual dynamic simulation of the CAR-NK92 library competing with tumour cells

HP

Pre-project Phase

General survey questionnaire and interview

Implementation Phase

Interview the doctor

Invite an expert

Publicize the project and Synthetic Biology

Follow up with previous interviewees

Cooperate with ZJUWnti on modeling

Educate high school students

Result

Verification of specific killing effect of CAR-NK92 cells

Verification of the ability of AP1903 to cause apoptosis when the engineered NK cells are not stimulated

Verification of the growth dynamics of logic-gated NK-92 cells

The gene circuit was effective in the cell-based CAR screening method and adding AP1903 was essential for the enrichment of CAR-NK-92 cells.

In these experiments, we determined the kinetics of the genetic circuit in which NFAT was coupled with KRAB activation in engineered NK-92 cells after CAR stimulation!

Design

Design thought

How can we deal with the large number of tumor antigens?

Engineered CAR-immune cells

Design plan

Selection of chassis cells

Kill switch circuit

NOTE: Engineering Optimization

opposite construct

Promoters are at opposite ends and at a long distance

Induced suicide switch

ICASP9

AP1903

dimerization

Inducing cell apoptosis

Future Prospect

Despite the long way to go from experiment to clinic, we make a preliminary outlook on the future of this project!

1. Draw 200 copies of peripheral blood from healthy volunteers.
2. Extract mRNAs encoding antibodies in B cells and use RT-PCR and SOE-PCR to form a scFv cDNA library.
3. Select leukocytes from volunteers on a large scale and obtain their HLA antigens as material for negative selection.
4. After negative selection, reserve the negative ones to form the final library that can be put into use.
5. Before the patient is treated, confirm that the scFv library has no affinity to the patient's HLA antigens.
6. Construct CAR-NK92 cells carrying the scFv library and infuse them into the patient for treatment.