

CANCER

Combination of metabolic intervention and T cell therapy enhances solid tumor immunotherapy

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Treatment of solid tumors with T cell therapy has yielded limited therapeutic benefits to date. Although T cell therapy in combination with proinflammatory cytokines or immune checkpoints inhibitors has demonstrated preclinical and clinical successes in a subset of solid tumors, unsatisfactory results and severe toxicities necessitate the development of effective and safe combinatorial strategies. Here, the liposomal avasimibe (a metabolism-modulating drug) was clicked onto the T cell surface by lipid insertion without disturbing the physiological functions of the T cell. Avasimibe could be restrained on the T cell surface during circulation and extravasation and locally released to increase the concentration of cholesterol in the T cell membrane, which induced rapid T cell receptor clustering and sustained T cell activation. Treatment with surface anchor-engineered T cells, including mouse T cell receptor transgenic CD8⁺ T cells or human chimeric antigen receptor T cells, resulted in superior antitumor efficacy in mouse models of melanoma and glioblastoma. Glioblastoma was completely eradicated in three of the five mice receiving surface anchor-engineered chimeric antigen receptor T cells, whereas mice in other treatment groups survived no more than 64 days. Moreover, the administration of engineered T cells showed no obvious systemic side effects. These cell-surface anchor-engineered T cells hold translational potential because of their simple generation and their safety profile.

INTRODUCTION

T cell therapy has demonstrated great clinical and preclinical successes in treatment of hematological malignancies (1, 2). Despite these encouraging results, treatment of solid tumors with T cells yields limited therapeutic benefits (3). Most studies have focused on the coadministration of proinflammatory cytokines or immune checkpoint inhibitors with T cells to boost efficacy (4–8). However, unsatisfactory results and severe side effects observed in some patients necessitate the development of effective and safe combination therapies (9, 10). Previous studies have demonstrated that the suppressive metabolic state of the oxygen- and nutrient-deprived tumor microenvironment impedes T cell infiltration, survival, and effector function, which likely compromises the therapeutic benefit of solid tumor T cell therapy (11, 12). T cell metabolism involves multiple diverse pathways, offering a breadth of potential intervention targets (12). For example, T cell function is dependent on the amount of cholesterol on cell membrane to cluster T cell receptors (TCRs) and form an immunological synapse (13–15). Therefore, modulation of cholesterol metabolism in combination with T cell therapy holds potential for improving solid tumor immunotherapy.

Avasimibe (Ava), an inhibitor of the cholesterol-esterification enzyme acetyl-CoA acetyltransferase 1 (ACAT1), elevates plasma membrane cholesterol concentrations, which, in turn, promote TCR clustering and thus improve effector function of T cells (16). We therefore hypothesized that combining Ava with T cell therapy would boost solid tumor immunotherapy. However, the pharmacokinetics and biodistribution of Ava are different from those of T cells, which imposes a challenge on optimizing the two as a com-

bination therapy (17–20). Thus, there is a need to develop promising combinatorial technologies that maximize both individual therapies.

Genetically engineered T cells can serve as a living factory to produce designed protein drugs (21–23). However, heterogeneous expression of engineered proteins, combined with potential for toxicity, reduces the efficacy of this strategy (24, 25). In addition, small molecular drugs cannot be manipulated in this genetic manner. An alternative to genetic engineering can be accomplished by backpacking nanoparticulated drugs onto the T cell surface via chemical conjugation or ligand-receptor biorecognition. This strategy has been shown to augment T cell function and widen the therapeutic window of combined drugs (26–28). Of note, backpacking strategies may impair the physiological functions of T cells. These effects can be due to long-term occupation of the functional biomolecules on the T cell membrane or due to changes to the glycometabolism of the T cell (27–31). Thus, technology to backpack nanoparticulated drugs onto the T cell surface can be further improved to reduce the impact of backpacking on the function of T cells.

Here, inspired by glycosylphosphatidylinositol-anchored proteins on the plasma membrane and click chemistry (32–34), we developed an alternative strategy to backpack drugs on the T cell surface by T cell-surface anchor-engineering technology. Specifically, we introduced functional tetrazine (Tre) groups onto the T cell surface via lipid insertion in the cell membrane. Next, liposomal Ava containing bicyclo [6.1.0] nonyne (BCN) groups was clicked onto the cell surface without disturbing the physiological functions of engineered T cells. We show that the liposomal Ava was retained on the T cell surface during circulation as well as extravasation and locally released to increase cholesterol in the T cell membrane. The increased cholesterol promoted rapid TCR clustering and sustained T cell activation. Last, we found that engineering TCR transgenic CD8⁺ T cells and chimeric antigen receptor T cells (CAR T cells) to carry liposomal Ava showed superior antitumor efficacy in mouse models of melanoma and glioblastoma.

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51. E. Blanco, H. Shen, M. Ferrari, Principles of nanoparticle design for overcoming biological barriers to drug delivery. *Nat. Biotechnol.* **33**, 941–951 (2015).
52. S. T. Hess, T. J. Gould, M. V. Gudheti, S. A. Maas, K. D. Mills, J. Zimmerberg, Dynamic clustered distribution of hemagglutinin resolved at 40 nm in living cell membranes discriminates between raft theories. *Proc. Natl. Acad. Sci. U.S.A.* **104**, 17370–17375 (2007).
53. S. T. Hess, M. Kumar, A. Verma, J. Farrington, A. Kenworthy, J. Zimmerberg, Quantitative electron microscopy and fluorescence spectroscopy of the membrane distribution of influenza hemagglutinin. *J. Cell Biol.* **169**, 965–976 (2005).
54. Y. Han, H. Pan, W. Li, Z. Chen, A. Ma, T. Yin, R. Liang, F. Chen, Y. Ma, Y. Jin, M. Zheng, B. Li, L. Cai, T cell membrane mimicking nanoparticles with bioorthogonal targeting and immune recognition for enhanced photothermal therapy. *Adv. Sci.* **6**, 1900251 (2019).
55. C.-M. J. Hu, R. H. Fang, K.-C. Wang, B. T. Luk, S. Thamphiwatana, D. Dehaini, P. Nguyen, P. Angsantikul, C. H. Wen, A. V. Kroll, C. Carpenter, M. Ramesh, V. Qu, S. H. Patel, J. Zhu, W. Shi, F. M. Hofman, T. C. Chen, W. Gao, K. Zhang, S. Chien, L. Zhang, Nanoparticle biointerfacing by platelet membrane cloaking. *Nature* **526**, 118–121 (2015).
56. C. E. Brown, B. Aguilar, R. Starr, X. Yang, W.-C. Chang, L. Weng, B. Chang, A. Sarkissian, A. Brito, J. F. Sanchez, J. R. Ostberg, M. D'Apuzzo, B. Badie, M. E. Barish, S. J. Forman, Optimization of IL13R α 2-targeted chimeric antigen receptor T cells for improved anti-tumor efficacy against glioblastoma. *Mol. Ther.* **26**, 31–44 (2018).

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