

## NEOANTIGEN-SPECIFIC STIMULATION OF T CELLS FOR EFFECTIVE CANCER ADOPTIVE CELL THERAPIES

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**Background** Targeting neoantigens by adoptive cell therapy (ACT) can effectively treat advanced solid tumors.<sup>1–5</sup> However, the conventional rapid expansion protocol (REP) for T-cell expansion can stimulate bystander cells<sup>6–8</sup> and cause differentiation and exhaustion of T cells.<sup>9–11</sup> This can lead to inadequate expansion of neoantigen-reactive T cells and hence ineffective ACT.

**Methods** We developed an *in vitro* culture method, termed NeoExpand, where T-cell receptor-engineered T cells (TCR-T) or neoantigen-reactive tumor infiltrating lymphocytes (neoTIL) were selectively expanded by neoantigen-specific stimulation. Briefly, T cells were co-cultured with antigen presenting cells or engineered cell lines loaded with neoantigens for ~2 weeks in the presence of interleukins 2 and 21.

**Results** When NeoExpand was used to expand TCR-T cells expressing previously identified CD8+ TCRs targeting shared p53 or KRAS neoantigens,<sup>4, 12</sup> selective expansion of TCR-expressing CD8+ T cells were observed when compared to REP [1.6 fold,  $p < 0.001$ ,  $n = 8$  (TCRs)]. Phenotypically, NeoExpand expanded CD39-CD69- cells, reportedly less differentiated T cells with stem-like features,<sup>7</sup> relative to REP (9.9 fold,  $p < 0.001$ ,  $n = 12$ ).

Next NeoExpand's ability to facilitate neoantigen-reactive TCR isolation was tested. From 25 TIL samples from tumors expressing p53 or KRAS mutations, the conventional screening<sup>13</sup> identified 14 neoTIL clonotypes (i.e., neoantigen-reactive TCRs) (3 CD4; 11 CD8), while NeoExpand enabled identification of 42 clonotypes (14 CD4; 28 CD8), indicating neoTIL's repertoire expansion during NeoExpand.

Next, we examined the effect of NeoExpand on expansion, phenotypes and functions of neoTIL. When 11 TIL samples from patients with p53-mutated or RAS-mutated gastrointestinal or breast cancer were tested, greater expansion of neoTIL with NeoExpand was noted relative to REP (4.0 fold,  $p = 0.02$ ). Single-cell transcriptome analysis revealed expansion of neoTILs with stem-like memory cell phenotypes uniquely in the NeoExpand conditions. These neoTILs expressed stem and memory markers, including CD62L, IL7R, and TCF1 and lacked exhaustion-associated gene expression, including CD39 and TIM3. Finally, TILs expanded through NeoExpand or REP were functionally compared using xenograft mouse models. Three TIL samples, one containing p53<sup>R175H</sup>-reactive TILs and two containing KRAS<sup>G12V</sup>-reactive TILs were expanded through NeoExpand or REP and were adoptively transferred to NSG mice engrafted with p53<sup>R175H+</sup> TYK-nu human ovarian cancer cells or KRAS<sup>G12V+</sup> patient-derived xenograft cancer cells. TILs expanded through NeoExpand led to significant tumor regression ( $p < 0.001$ ,  $n = 5$  mice/group).

**Conclusions** Collectively, NeoExpand selectively expands neoantigen-reactive T cells compared to REP and enables sensitive identification of neoantigen-reactive TCRs by expanding neoTIL repertoire. NeoExpand's ability to enhance phenotypes and functions of neoantigen-reactive T cells warrants its evaluation for clinical use.

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