

# Engineered immunostimulatory cells can convert PBMCs from chronic lymphocytic leukemia (CLL) patients into potent tumor killing immune cells

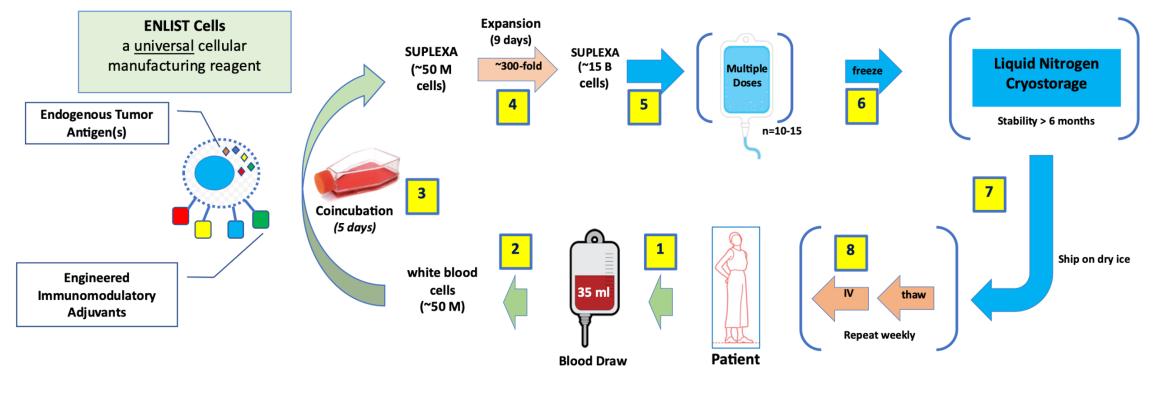
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## ABSTRACT #7517

#### Background

A unique autologous cellular therapeutic (**SUPLEXA**) has been developed from human PBMC. It is comprised of NK cells, NKT-like cells,  $\gamma\delta$  T cells and CD8+ T effector cells, capable of broadly lysing a variety of tumor cell lines in vitro. SUPLEXA cells are manufactured using an efficient 2 weeks xeno-free manufacturing procedure employing two proprietary engineered leukocyte stimulator cell lines (**ENLIST**) that express an array of immunomodulatory proteins. This process leads to a 300-fold expansion of NK cells, CD8+ T cells, NKT-like cells, and TCR $\gamma\delta$  T cells that are called SUPLEXA cells, which will be cryopreserved, and then transferred back into patients as an autologous immune cell therapy for cancer. In this study, PBMCs from CLL patients were used to generate SUPLEXA cells as a first approach to comparatively profile SUPLEXA cells from cancer patients and normal healthy volunteers (NHVs).

The SUPLEXA cell manufacturing process uses peripheral blood mononuclear cells (PBMCs) from cancer patients. PBMCs are stimulated with ENLIST cells for a 5-day induction period, which is then followed by a 9-day cytokine-induced expansion period. SUPLEXA cells are then cryopreserved to use as an autologous adoptive immunotherapy. A first-in-human clinical trial for this novel adoptive cellular immunotherapy for cancer is projected to begin later this year.



#### **Methods**

**ENLIST cells:** Engineered SK-MEL2 melanoma cell lines that express curated sets of > 20 different immunomodulatory proteins that are engineered for membrane expression.

**SUPLEXA cell generation:** Two million (M) PBMCs isolated from direct blood draws from 10 CLL patients or 5 NHVs were incubated with 0.4 M freeze/thaw killed ENLIST cells for 5 days in XVIVO-15 medium with 2% heat-inactivated human AB serum (XAB2) and then split 1:15 in XAB2 containing IL-7 and IL-15 to expand. After 9 days, SUPLEXA cells were harvested and cryopreserved.

**Mass Cytometry (CyTOF):** SUPLEXA cells were comprehensively characterized by mass cytometry (CyTOF) using a 47-marker antibody panel. CyTOF data analysis was done using an analysis workflow of dimensional reduction by PCA embedded opt-SNE using OMIQ

**Tumor Cell Killing Assay:** Tumor cytolytic activity was measured by flow cytometry using fluorescent tumor cell targets at 2:1, 1:1, and 1:2 effector:target cell ratios. M14 melanoma cells that express red fluorescent protein (RFP) were used as tumor cell targets for these studies. Cytolysis was measured at 48 hours.

**Cytokines:** A 33 cytokine Luminex panel was used to assess cytokine levels in tumor cell cytolysis supernatants.

|                 | <u></u>        | TOF Antibody Panel                             |
|-----------------|----------------|--|
| Marker          | Metal          | Role   |
| CD45            | 89Y            | Pan-leukocyte marker                           |
| CD172ab         | 111Cd          | Myeoloid cell marker                           |
| CD8a            | 111Cd          | CD8 T cell alpha-chain                         |
| CD20            | 112cu<br>113ln | B cell marker                                  |
| CD20            | 113m           | CD4 T cell marker                              |
| CD4<br>CD3      | 114Ca          |  |
| DNAM1           | 115m           | All T cell marker                              |
|                 |                | Adhesion molecule for NK cell activation       |
| Granulysin*     | 141Pr          | Effector cell tumor killing                    |
| KIR2DS1<br>PD-1 | 142Nd          | NK receptor that interacts with DAP12          |
|                 | 143Nd          | Activation and checkpoint inhibition           |
| DAP12           | 144Nd          | Signaling adaptor of NK cell activation        |
| CD19            | 145Nd          | B cell marker                                  |
| CD14            | 146Nd          | monocyte and macrophage                        |
| CD86            | 147Sm          | costimulation                                  |
| Granzyme A      | 148Nd          | Effector cell tumor killing                    |
| Granzyme K      | 1495m          | Effector cell tumor killing                    |
| CD26            | 150Nd          | Dipeptidyl peptidase                           |
| TCR-Vd1         | 151Eu          | gamma delta T cell receptor                    |
| CTLA-4          | 152Sm          | Checkpoint inhibition                          |
| CD69            | 153Eu          | Lympocyte activation marker                    |
| TCRgd           | 154Sm          | gamma delta T cell identification              |
| EOMES           | 155Gd          | TF for effector lymphocyte function            |
| CD8b            | 156Gd          | CD8 T cell beta-chain                          |
| CD16            | 157Gd          | NK marker                                      |
| CD39            | 158Gd          | ATP to Adenosine enzyme                        |
| PLZF            | 159Tb          | Transcription factor, regulates NK cells       |
| NKp30           | 160Gd          | NK cell marker                                 |
| PSGL1           | 161Dy          | Adhesion molecule for tissue infiltration      |
| CD56            | 162Dy          | Pan-NK cell marker                             |
| 4-1BB           | 163Dy          | T cell costimulation                           |
| GITR            | 164Dy          | Glucocorticoid-induced recetptor - Tregs       |
| NKG2A           | 165Ho          | Effector NK cell marker - negative             |
| CD107a          | 166Er          | Degranulation marker                           |
| Grz B           | 167Er          | Effector cell tumor killing                    |
| NKp46           | 168Er          | Activating receptor on NK cells                |
| TCRVa7.2        | 169Tm          | Invariant NK T cells                           |
| PD-L1           | 170Er          | PD-1 ligand for checkpoint inhibition          |
| TCR-Vd2         | 171Yb          | gamma delta T cell receptor                    |
| NKG2D           | 172Yb          | Effector NK cell marker - positive             |
| TCRab           | 173Yb          | alpha-beta T cell receptor                     |
| Ki67            | 174Yb          | Recently proliferated cell marker              |
| Tbet            | 175Lu          | TF for NK and effector lymphocyte subets       |
| Perforin        | 176Yb          | Effector cell tumor killing                    |
| CD57            | 194Pt          | T cell exhaustion marker                       |
| HLA-DR          | 195Pt          | Hiscompatability                               |
| SLAMF6          | 196Pt          | Trigger cytolytic activity of NK and NKT cells |
| CD38            | 198Pt          | NK adhesion to endothelium                     |
| TIGIT           | 209Bi          | Inhibitory receptor on Tregs                   |
| * Highlighted m | arkers a       | re stained for intracellular detection         |

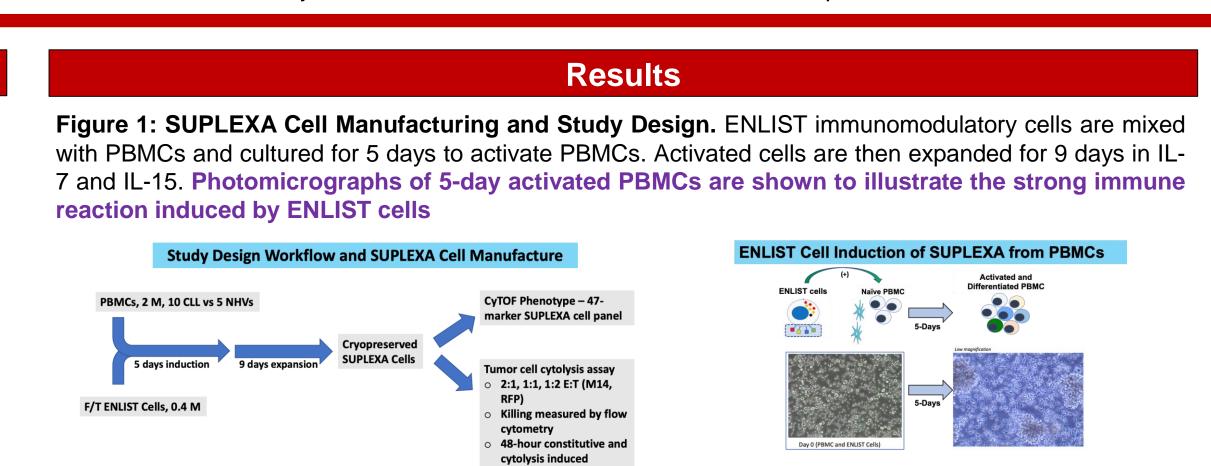
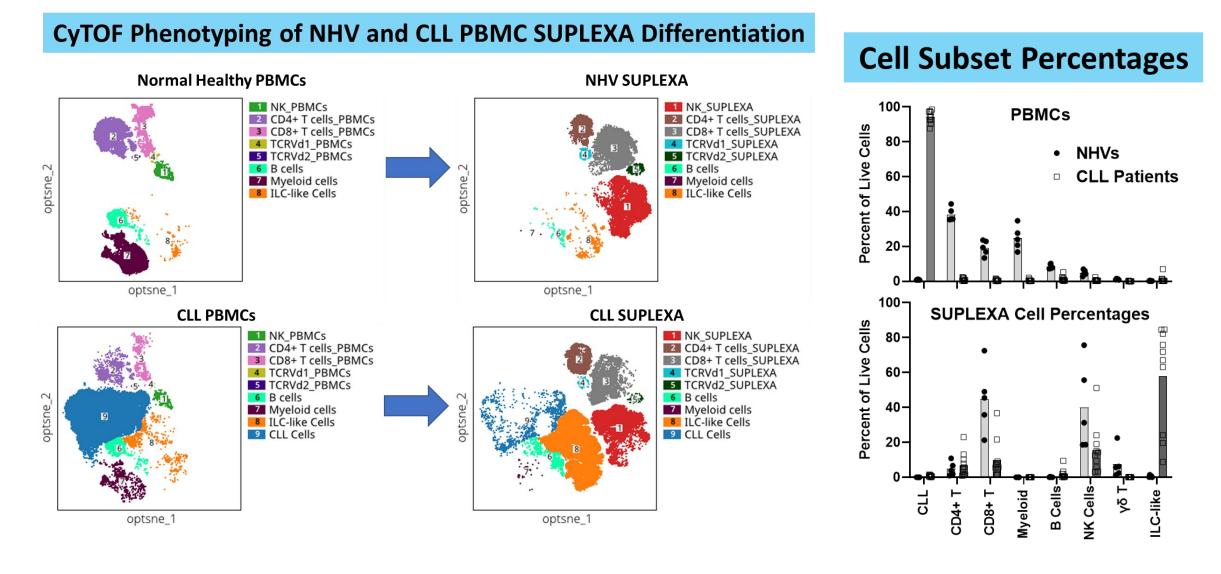
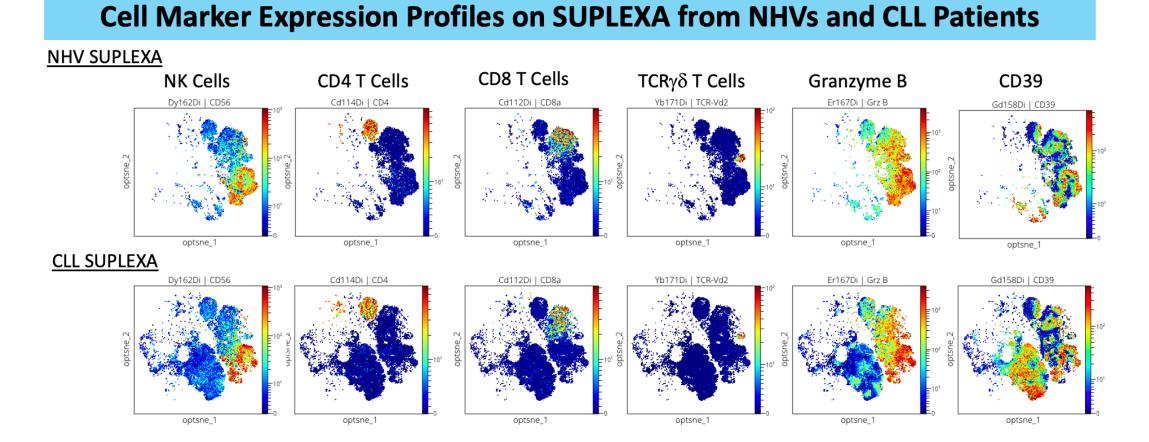


Figure 2: SUPLEXA Cell Phenotyping By CyTOF: SUPLEXA cells from CLL patients and NHVs were analyzed to compare single-cell phenotypes using a customized 47-marker CyTOF antibody panel to identify NK cell and T cell subsets as well as transcription factors, adhesion markers, and functional molecules. Our CyTOF data analysis workflow, cell subset overlay plots of SUPLEXA differentiation of PBMCs from NHVs vs. CLL patients, and cell marker expression profiles of SUPLEXA from NHVs and CLL patients for comparison are shown. Results indicate that SUPLEXA cells can be generated from CLL PBMCs that contain 90% CLL cells.







**Results** 

Figure 3: Tumor Cell Cytolysis Assay and Killing Activity. SUPLEXA cells from NHVs and CLL patients were compared for tumor cell cytolytic activity against fluorescent tumor target cells using a flow cytometry method. Figures show representative killing of M14-RFP target cells and comparative tumor cytolysis activity SUPLEXA from 2 NHVs and 10 CLL patients. Results show potent and similar tumor cytolytic activity between NHV and CLL SUPLEXA cells.

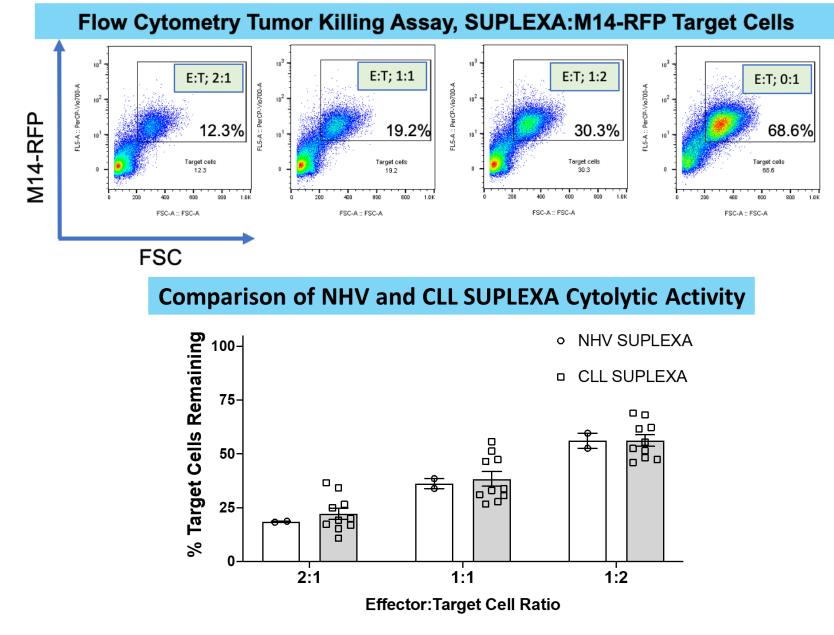
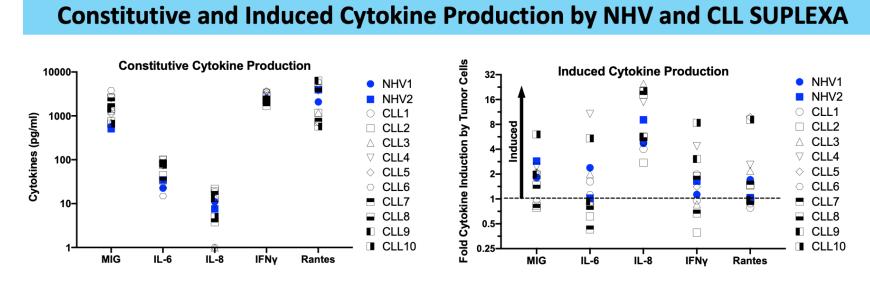


Figure 4: Cytokine Production by SUPLEXA During Tumor Cytolysis. SUPLEXA cells were incubated without or with M14 tumor cells for 48 hours. Supernatants were tested for cytokines by 33-plex Luminex. Plots of constitutive and tumor cell (M14) induced cytokine production by SUPLEXA from NHVs and CLL patient are shown. Results show comparable levels of cytokine production between CLL and NHV SUPLEXA cells.



### Conclusions

- 1. We show consistent ENLIST cell induced generation of SUPLEXA cells from CLL patients and CLL cells die or are killed during the SUPLEXA manufacturing process
- 2. CyTOF single-cell phenotyping of SUPLEXA cells from CLL patients and NHVs showed overlapping phenotypes, but also some important differences:
  - $\circ \qquad \text{No TCR} \gamma \delta \text{ T cell expansion from CLL PBMCs}$
  - Expansion of ILC-like cells in CLL patients CD45+,CD39+,CD69+, Tbet+, GranzymeB and SLAMF6+ cells without T, B, myeloid, or NK markers
- SUPLEXA cells from NHV and CLL patients showed identical levels of potent tumor cell killing activity and cytokine production profiles with normal distribution of heterogeneity Funding: Alloplex Biotherapeutics, Inc and NCI R01CA258924 (PI Brown)