

Novel Small-Molecule ROR γ Agonist Immuno-oncology Agent LYC-55716: Tumor Selection and Evaluation of Renal Cell and Bladder Cancer for Inclusion in Phase 2a Expansion

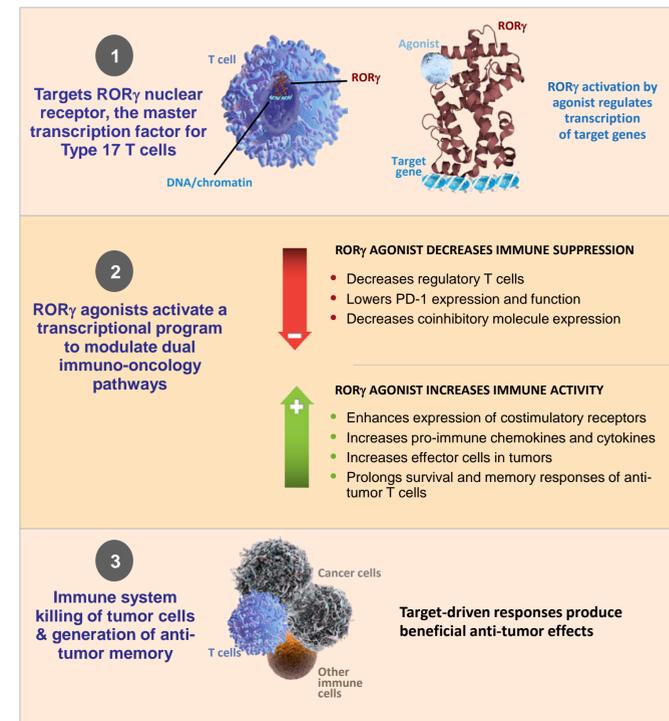
Abstract #67

Xiao Hu,¹ Xikui Liu,¹ Hongxiu Li,¹ Elizabeth Zawadzka,¹ Garry Weems,² Yilin Gao,¹ Brian Fox,³ H. Jeffrey Wilkins,² Laura Carter¹
¹Lycera Corp., Ann Arbor, MI; ²Lycera Corp., Plymouth Meeting, PA; ³Celgene Corp., Seattle, WA.

BACKGROUND

- Retinoic acid receptor-related orphan receptor γ (ROR γ) is the master transcription factor responsible for type 17 effector T cell differentiation and function.
- Synthetic ROR γ agonists can modulate immune cell gene expression by enhancing type 17 helper (Th17) and cytokine (Tc17) T cells, stimulating a potent antitumor response that includes increased immune activity and decreased immune suppression based on preclinical models (Figure 1).¹
- ROR γ agonists have shown promise as monotherapy and combination therapy in syngeneic tumor models.
- LYC-55716 is a first-in-class, oral, small-molecule investigational agent that selectively activates ROR γ .
- Phase 1 clinical testing of LYC-55716 identified a pharmacodynamically active dose and demonstrated that this agent was well tolerated in cancer patients.
- Preclinical and bioinformatics analyses were conducted to support the inclusion of patients with renal cell carcinoma (RCC) and bladder cancer (BC) in a Phase 2A expansion trial (NCT02929862).

Figure 1. ROR γ agonist as a novel immuno-oncology approach

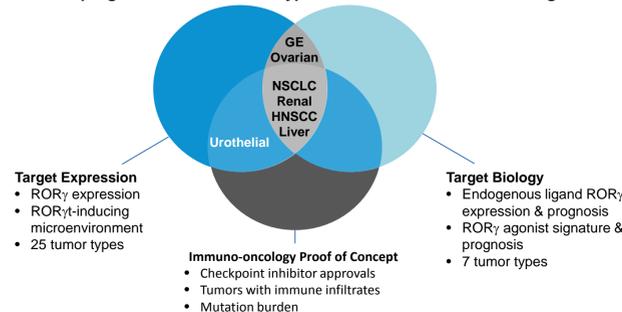


METHODS

- Bioinformatic analyses were conducted using data on patients with RCC and BC from The Cancer Genome Atlas (TCGA). The resulting dataset was evaluated for (Figure 2):
 - Expression of ROR γ and ROR γ -inducing cytokines;
 - Signature genes associated with ROR γ biology, biomarkers for endogenous ROR γ ligands, and correlations with patient survival rates; and
 - Tumor microenvironment immune profiles.

Figure 2. Factors considered in tumor selection

Clinical program identified tumor types that meet at least 2 of 3 categories



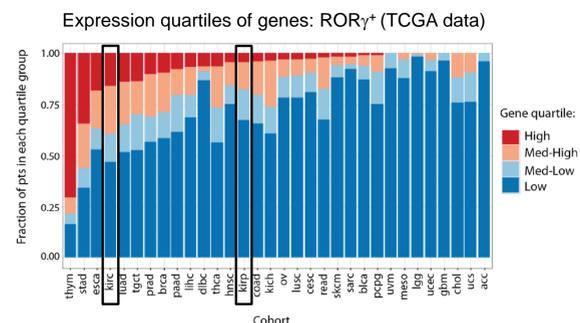
GE, gastroesophageal; HNSCC, head & neck squamous cell carcinoma; NSCLC, non-small-cell lung cancer.

- ROR γ expression and the in vitro effects of a ROR γ agonist on peripheral blood mononuclear cells (PBMCs) were assessed from patients with genitourinary (GU) cancer.

TARGET EXPRESSION

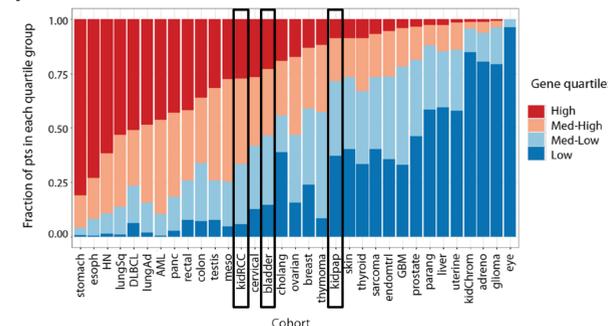
- RNA sequencing analysis identified ROR γ t expression in a significant fraction of RCC and BC samples (Figure 3).
 - >40% RCC patients and 20% of renal papillary patients have high ROR γ t expression, implying they are infiltrated by Type 17 T cells.
 - A low percentage of BC patients express high ROR γ t.

Figure 3. ROR γ t expression across tumor types



- ROR γ -inducing cytokines (IL-1 β , IL-6 and IL-23) that support type 17 differentiation were highly expressed in both RCC and BC.
 - >50% of RCC and BC patients have expression of genes that would support the formation of ROR γ t T cells (Type 17 cells) (Figure 4).

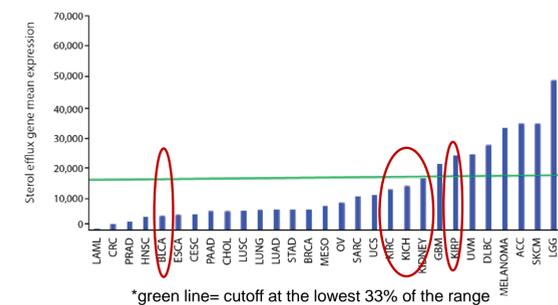
Figure 4. Expression of genes that support ROR γ t expression



TARGET BIOLOGY

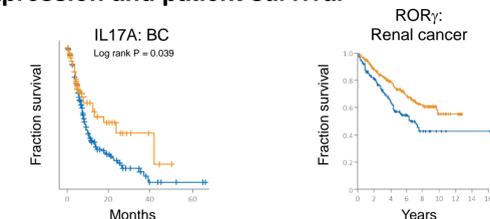
- Certain sterols can function as ROR γ ligands and partially activate ROR γ ,² but sterol levels are likely low in exhausted T cells and can be influenced by cancer cells in the tumor microenvironment.
 - TCGA analysis of sterol efflux gene expression (a surrogate for endogenous ligand levels) revealed differential expression across tumor types (Figure 5).
 - Low expression of sterol efflux genes in BC tumors suggested low levels of endogenous ROR γ ligands in the tumor microenvironment.
 - Medium levels of sterol efflux genes were found in kidney renal cell (KIRC), kidney chromophobe (KICH), and kidney renal papillary cell (KIRP) carcinomas.

Figure 5. Expression of sterol efflux genes across tumor types



- Analysis revealed a positive correlation between patient survival and expression of ROR γ t (or the ROR γ signature gene IL17A).
 - IL17A expression is associated with statistically significant improvements in survival in BC (Figure 6).
 - High expression of ROR γ (both isoforms) is also associated with better survival in patients with KIRC, KICH, and KIRP.

Figure 6. Correlation between ROR γ and IL17A expression and patient survival

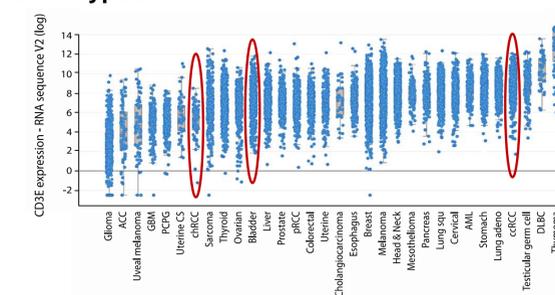


- A literature review also indicated that IL-17 is implicated in the efficacy of Bacillus Calmette-Guérin immunotherapy for bladder cancer.³

IMMUNE PROFILE

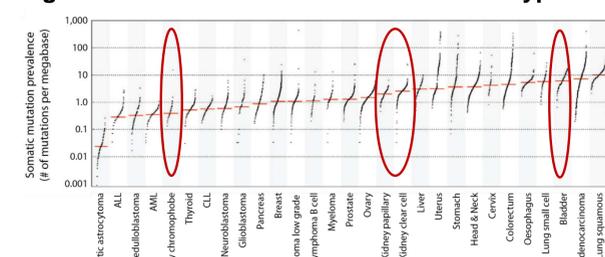
- Studies have shown a positive correlation between immunotherapy efficacy and infiltration of T cells (eg, CD3e, CD4, CD8a) in the tumor microenvironment.^{4,5}
 - Analysis of TCGA RNA sequencing data indicated a high expression of T cell markers such as CD3e in RCC and BC (Figure 7).

Figure 7. Tumor-infiltrating lymphocytes across tumor types



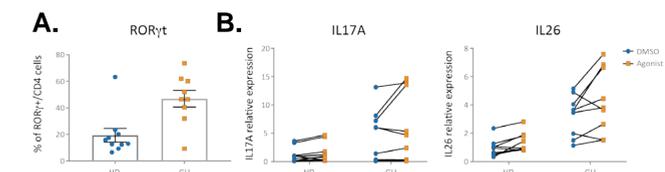
- The immune system may recognize tumors with a high mutational burden; patients with such tumors have shown improved response to immunotherapy and longer progression-free survival.^{6,7}
 - Analysis of RCC and BC tumors indicated a high mutation burden (Figure 8), which are associated with immunotherapy efficacy.

Figure 8. Mutational burden across tumor types



- PBMCs from patients with GU cancer (RCC, n=5; urothelial, n=4) showed increased expression of ROR γ t mRNA by qPCR compared with healthy donors (HD) (Figure 9A).
- Consistent with higher basal levels of ROR γ protein, PBMC from 6/9 GU patients express high levels of ROR γ target genes (IL-17A, IL-26) prior to treatment with ROR γ agonist compound. In most patients, the addition of ROR γ agonist further enhances target gene expression (Figure 9B).
- These data suggest that immune cells from GU patients are responsive to ROR γ agonist treatment.

Figure 9. PBMC expression of (A) ROR γ t and (B) target genes IL17A and IL26



CONCLUSIONS

- RCC and BC express ROR γ and ROR γ -supportive genes, and this expression is correlated with improved patient survival.
- ROR γ biology suggests RCC and BC are likely to respond to ROR γ agonist therapy.
- These translational and bioinformatics studies of ROR γ expression, biology, and tumor immune profiles support the inclusion of RCC and BC (Urothelial carcinoma) patients in an ongoing Phase 2a expansion trial of LYC-55716 (Table 1).

Table 1. Tumors selected for Phase 2a Expansion

Tumor Types
Non-small-cell lung cancer
Gastroesophageal cancer
Head and neck squamous cell carcinoma
Ovarian cancer
Renal cell carcinoma
Urothelial carcinoma

REFERENCES

- Hu X, et al. *Oncoimmunology*. 2016;5:e1254854.
- Hu X, et al. *Nat Chem Biol*. 2015;11:141-7.
- Takeuchi A, et al. *Eur J Immunol*. 2011;246-51.
- Alexandrov LB, et al. *Nature*. 2013;500:415-21.
- Lawrence MS, et al. *Nature*. 2013;499:214-8.
- Chen J, et al. *Int J Biol Sci*. 2011;7:53-60.
- Lawrence MS, et al. *Nature*. 2013;499:214-8.

DISCLOSURE: LYC-55716 is an investigational agent not yet approved by FDA: the safety and efficacy of LYC-55716 have not been established in patients. Copies of this poster obtained through Quick Response (QR) Code are for personal use only and may not be reproduced without permission from ASCO® and the author of this poster.

CLINICAL TRIAL CONTACT: wilkins@lycera.com

