

Inhibition of Fatty Acid Oxidation by the Peroxisome Proliferator Activated Receptor- α Antagonist TPST-1120 Elicits Tumor Regression Concomitant with Immune Activation

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OBJECTIVE

To assess TPST-1120 activities on tumor progression, FAO, and immune activation in preclinical models and clinical samples from patients with advanced solid tumors receiving TPST-1120 as monotherapy or in combination with immune checkpoint inhibitors

BACKGROUND

Peroxisome Proliferator Activated Receptor- α (PPAR α)

- Nuclear hormone receptor that functions as master regulator of fatty acid oxidation (FAO), upregulated in multiple cancers, and expressed by immune cells (Figure 1A)¹
- Represses multiple proinflammatory transcription factors (e.g., NF- κ B, STAT-1, STAT3, and AP-1), thereby limiting anti-tumor immunity (Figure 1B)¹
- Numerous tumors utilize FAO as a primary metabolic pathway, including renal cell carcinoma (RCC), cholangiocarcinoma (CCA), and hepatocellular carcinoma (HCC), making PPAR α an attractive target for therapeutic intervention²

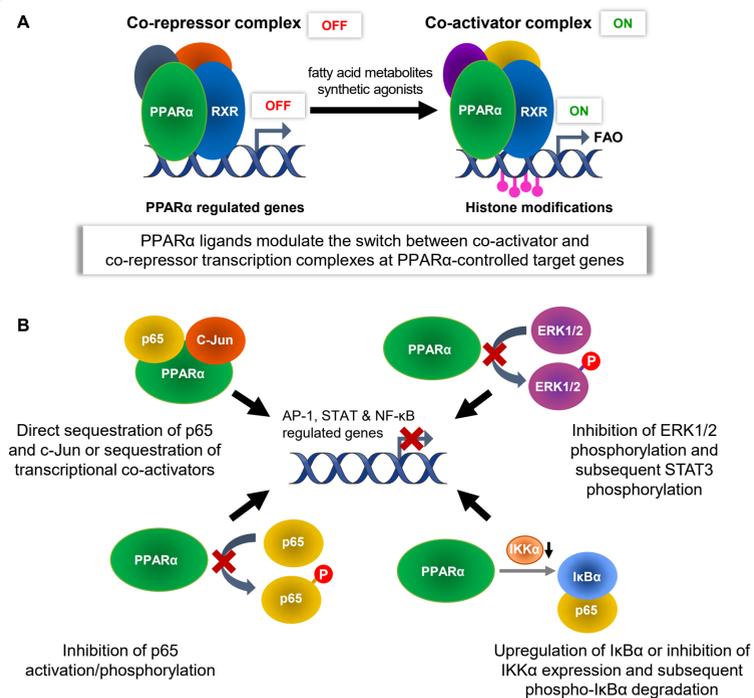


Figure 1. (A) PPAR α -Regulated Transcription; (B) PPAR α Transrepressive Activities

TPST-1120

- First-in-class, orally-bioavailable, potent, highly selective competitive antagonist of PPAR α
- Hypothesized to block tumor-promoting FAO and alleviate PPAR α -induced immune suppression
- Clinical experience:
 - Phase 1 trial in advanced solid tumors (NCT03829436):** 53% disease control rate as single agent and 30% objective response rate (ORR) in combination with nivolumab at the two highest TPST-1120 doses tested³
 - Phase 1b/2 randomized trial in 1L metastatic HCC (NCT04524871):** 30% confirmed ORR with TPST-1120 plus atezolizumab + bevacizumab vs. 13.3% with atezolizumab + bevacizumab alone; favorable PFS and OS hazard ratio in TPST-1120 arm vs. control arm⁴
 - Tolerable safety profile alone and in combination w/ nivolumab or atezolizumab + bevacizumab^{3,4}

METHODS

X-Ray Crystallography

- The ligand binding domain (LBD) of human PPAR α (amino acids 198-468) was expressed and purified from *E. coli*. The PPAR α /TPST-1120 co-crystals were grown at 4°C by sitting drop vapor diffusion. Data collection was performed at Spring8 BL45XU beamline, and the structure was determined by molecular replacement with existing structure PDB ID: 1K7L

In Vitro PPAR α and Gene Expression Assays

- TPST-1120 inhibitory activity against PPAR α was determined in a CHO cell-based reporter cell line with a PPAR α response element (PPRE) linked to a luciferase gene
- TPST-1120-mediated effects on gene expression and lipid content in THP-1 and U937 myeloid cell lines were assessed using qPCR and flow cytometry

In Vivo Experiments: Pancreatic, Colon, and Liver Tumor Models

- BALB/c or C57BL/6 mice bearing PancOH7 (pancreatic), MC38 (colon), or Hepa 1-6 (liver) murine tumors were treated with TPST-1120 as monotherapy or in combination with mouse anti-PD-1
- Tumors were allowed to establish for 8 days (average of 171 \pm 5 mm³) prior to treatment with vehicle (n = 10), TPST-1120 at 30 mg/kg orally BID (n = 7), anti-PD-1 200 μ g IP Q3D (n = 10), or TPST-1120 + anti-PD-1 (n = 15). Animals were followed for tumor growth

- Animals in vehicle group (n = 10) were sacrificed on treatment Day 16 due to tumor growth (2024 \pm 153 mm³)

TPST-1120-01 Phase 1 Clinical Study Design (NCT03829436)

Key Eligibility Criteria	Part 1: TPST-1120 Monotherapy Dose Escalation	Endpoints
Inclusion <ul style="list-style-type: none"> Advanced/metastatic solid tumor ECOG PS 0-1 Adequate organ function No standard therapy available Exclusion <ul style="list-style-type: none"> Immunosuppressive meds Autoimmune disease Fibrates within 28 days of enrollment 	Solid Tumors 3+3 Design TPST-1120 100 mg up to 600 mg BID	<ul style="list-style-type: none"> Safety MTD and/or OBD of TPST-1120 Pharmacokinetics Preliminary efficacy
	Part 2: TPST-1120 Combination with anti-PD-1 Dose Escalation HCC, RCC, Cholangiocarcinoma 3+3 Design TPST-1120 200 mg up to 600 mg BID Full dose nivolumab (480 mg Q4W)	Translational Readouts <ul style="list-style-type: none"> Baseline tumor mutational status Whole blood gene expression Lipidomics

BID, twice daily; ECOG PS, Eastern Cooperative Oncology Group performance status; HCC, hepatocellular carcinoma; MTD, maximum tolerated dose; OBD, optimal biological dose; RCC, renal cell carcinoma; Q4W, once every 4 weeks.

Clinical Biomarker Assays

- Changes in whole blood gene expression and plasma lipid levels in patients enrolled in a phase 1 clinical trial (NCT03829436) were assessed using the Nanostring nCounter[®] PanCancer Immune Profiling panel and liquid chromatography-tandem mass spectrometry, respectively

RESULTS

TPST-1120 is a Competitive Antagonist of PPAR α

- X-ray co-crystal structures demonstrated that TPST-1120 binds to the PPAR α LBD, inducing a repressive conformation (Figure 2A)
- In an *in vitro* cell-based PPAR α assay, TPST-1120 potently competed against the PPAR α -specific synthetic agonist GW7647 and endogenous agonist oleoylethanolamide (OEA) (Figure 2B, 2C)

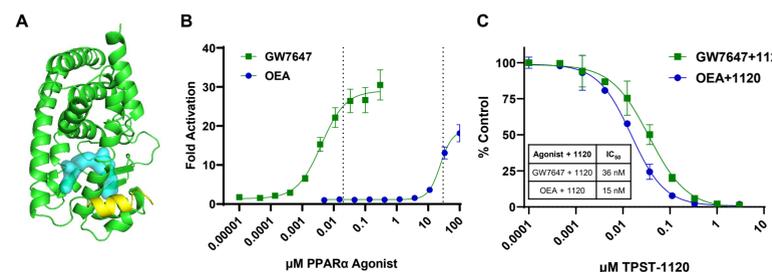


Figure 2. TPST-1120 is a Competitive Antagonist of PPAR α . (A) The TPST-1120 antagonist-bound ligand binding domain (green) shows that TPST-1120 (blue) situates the AF-2 activation helix (yellow) in an inactive conformation. The inactive conformation has a lower affinity to co-activator motifs and disfavors activation of PPAR α regulated genes. (B) Fold-activation of CHO cell-based reporter cell line following treatment with PPAR α -specific synthetic agonist GW7647 and endogenous agonist OEA. Vertical dotted lines correspond to agonist concentration tested in panel C. (C) Decrease in PPAR α -dependent activation by increasing concentrations of TPST-1120.

RESULTS (cont.)

TPST-1120 Inhibits Tumor Growth in Pancreatic, Colon, and Liver Tumor Models

- TPST-1120 elicited > 50% tumor growth inhibition in PancOH7, MC38, and Hepa1-6 models with enhanced inhibition observed in the latter two models when co-administered with anti-PD-1 (Figure 3)

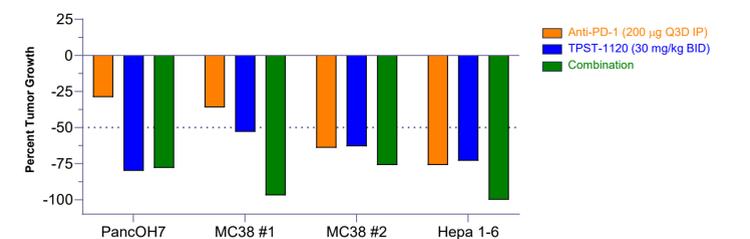


Figure 3. Percent Tumor Growth Inhibition with TPST-1120. The mean change in tumor volume from treatment initiation was calculated at end of study (approximately day 21) for each tumor model treated with indicated therapy. Results from two independent experiments with MC38 shown.

Mice Exhibiting Complete MC38 Tumor Regression are Protected from Tumor Re-challenge

- At treatment Day 16, treated MC38 tumors were significantly smaller compared to control group (TPST-1120 $P = 0.002$; anti-PD-1 $P = 0.002$; Combo $P < 0.00001$) (Figure 4A)
- Three animals in the combination group exhibited complete tumor regression (considered cured) which was maintained at Day 60; when cured mice were rechallenged with fresh autologous tumors, they remained protected from tumor growth while tumors grew as expected in the naïve, age-matched control mice (Figure 4B)

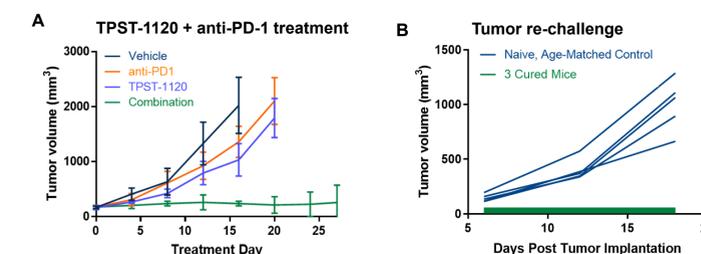


Figure 4. TPST-1120 + Anti-PD-1 Elicits Tumor Regression and Long-Standing Immune Memory in MC38 Colorectal Cancer Tumor Model in C57BL/6 Immunocompetent Mice. (A) C57BL/6 mice bearing 150 mm³ MC38 flank tumors treated with TPST-1120 30 mg/kg BID, 200 μ g anti-PD-1 Q3D or combination. (B) Three mice from combination therapy group with no observable tumors at day 60 or 5 age-matched control received MC38 tumors in flank and were monitored for 18 days.

TPST-1120 Induces Increased Neutral Lipid Content and Elevated Immune-Related Markers

- The human myelomonocytic cell line THP-1 treated with TPST-1120 in the presence of oleic acid demonstrated changes in lipid content (Figure 5A), immune gene expression (Figure 5B) and corresponding protein levels (Figure 5C) as a function of dose.

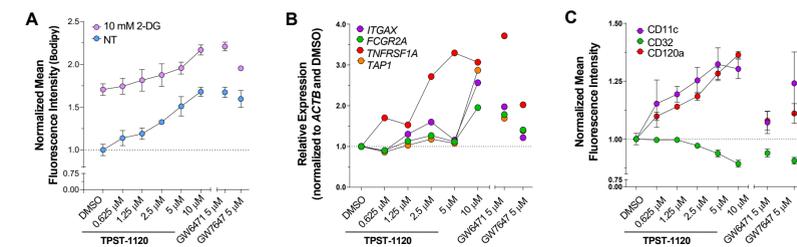


Figure 5. TPST-1120 Induces Increased Neutral Lipid Content and Elevated Immune-Related Genes in THP-1 Cells. (A) THP-1 cells were incubated for 24 hr in the presence of oleic acid (30 μ M) with or without 2-deoxyglucose (2-DG) in increasing concentrations of TPST-1120 after which Bodipy staining was analyzed. Similar results observed with U937 cells (data not shown); (B) Expression levels of listed genes in THP-1 cells incubated with increasing TPST-1120; (C) Cell surface levels of listed proteins on THP-1 cells in increasing TPST-1120. 2-DG: glucose analog that inhibits glycolysis; GW6471: PPAR α -selective antagonist; GW7647: PPAR α -selective agonist.

Biomarker Analysis of Gene Expression and FFA Supports TPST-1120 Mechanism of Action

- In the phase 1 trial, statistically significant, exposure-dependent elevations in expression levels of multiple immune-related genes including *FCGR2A* (CD32), *ITGAX* (CD11c), *TAP1*, and *TNFRSF1A* (CD120a) were observed, and patients exhibiting objective responses displayed increased free fatty acids (FFA) in plasma (Figure 6)

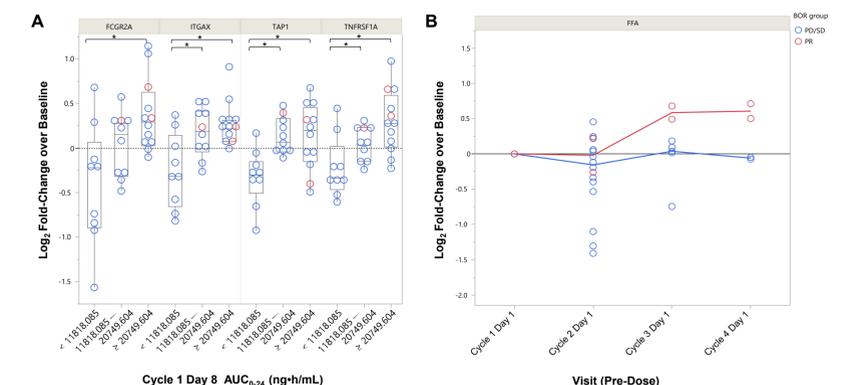


Figure 6. TPST-1120 Induces Increased Expression of Immune-Related Genes and Elevated Free Fatty Acids. (A) Genes differentially expressed on treatment day 8 vs. baseline, as a function of TPST-1120 exposure. Median elevation magnitudes in highest tertile were statistically increased above baseline values ($*P < 0.05$ by Wilcoxon's pair-wise analysis vs. first tertile [< 11818.05]). (B) Log₂-fold-changes in baseline normalized free fatty acids (FFA) in patients enrolled in Part 2, presented by response.

CONCLUSIONS

- TPST-1120 is a potent competitive antagonist of PPAR α that inhibits FAO and induces immune activation consistent with its proposed mechanism of action:
 - Co-crystal structure of TPST-1120 and the PPAR α ligand-binding domain show a repressive conformation suggesting reduced transcription of target genes
 - TPST-1120 potently competed against PPAR α -specific agonists in a cell-based assay, demonstrating specificity for the PPAR α isoform
 - Tumor growth was inhibited > 50% in pancreatic, colon, and liver tumor models in mice, with immune memory induction in the colon cancer model
 - Dose-dependent increases in lipid content and immune gene expression support PPAR α inhibition by TPST-1120
 - In a phase 1 study, patients with advanced solid tumors receiving TPST-1120 showed significant exposure-dependent elevation of several immune-related genes and increased plasma FFA associated with clinical response
- Clinical response and biomarker findings demonstrate that inhibition of PPAR α is an effective therapeutic strategy for the treatment of cancer and has been a basis for the ongoing late-stage clinical development of TPST-1120

REFERENCES: 1. Bougarne N, et al. *Endocr Rev*. 2018;39:760–802. 2. Nath A and Chan C. *Sci Rep*. 2016;6:18669 2. 3. Yarchoan M, et al. *J Clin Oncol*. 2022; 40(16): Abstract 3005; 4. Tempest Therapeutics Press Release, October 11, 2023. <https://ir.tempestx.com/news-releases/news-release-details/tempest-releases-new-data-demonstrating-superiority-tpst-1120>.

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