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T cell receptor pharmacodynamics associated with survival and response to tremelimumab (T) in combination with durvalumab (D) in patients (pts) with unresectable hepatocellular carcinoma (uHCC)

Patricia McCoon,¹ Young S. Lee,² R. Kate Kelley,³ Violeta Beleva Guthrie,² Song Wu,² Stephanie A. Bien,⁴ Alejandra Negro,⁵ Philip He,⁵ John Kurland,⁵ Carl Barrett,¹ Fernanda Pilataxi,⁵ Steven Ching,⁵ Ghassan K. Abou-Alfa⁶

¹Translational Medicine, AstraZeneca, Waltham, MA, USA; ²Translational Medicine, AstraZeneca, Gaithersburg, MD, USA; ³Hematology/Oncology, University of California, San Francisco, CA, USA; ⁴Adaptive Biotechnologies, Seattle, WA, USA; ⁵AstraZeneca, Gaithersburg, MD, USA; ⁶Memorial Sloan Kettering Cancer Center, New York, NY, USA

Objective

- We present an exploratory molecular analysis of peripheral blood T cell receptors following treatment with two tremelimumab (T) plus durvalumab (D) combinations, including a novel combination regimen containing a single, priming dose of T (T300+D), and T and D as monotherapies, and the potential association with objective response rate (ORR) and overall survival (OS), in patients with unresectable hepatocellular carcinoma

Conclusions

- A greater number of expanded T cell clones at Day 29 was associated with longer survival and a greater likelihood of a clinical response
- The number of expanded T cell clones increased with higher doses of tremelimumab, and a greater proportion of patients had above-the-median number of expanded T cell clones in the T300+D and T arms versus the T75+D arm
- These results, taken together with the clinical safety data,¹ support the clinical benefit of the T300+D regimen
- Further work is needed to understand the relative contributions of CD4 and CD8 T cell clonal expansion on the improved ORR and OS observed with T300+D and T monotherapy
- T300+D and D are being evaluated in the Phase 3 HIMALAYA study (NCT03298451) in unresectable hepatocellular carcinoma versus sorafenib²

Plain language summary

Why did we perform this research? Immunotherapy helps the body's own immune system to recognize and kill cancer cells. Tremelimumab and durvalumab are immunotherapies that are being developed for the treatment of hepatocellular carcinoma, a type of liver cancer. In Study 22, a single, priming dose of tremelimumab in combination with durvalumab (T300+D) appeared to help patients live longer and help their tumors shrink in size, compared with using multiple lower doses of tremelimumab, or using either treatment on its own¹

How did we perform this research? Blood samples were taken from patients before and after treatment to determine whether changes in an aspect of the immune response correlated with improved outcomes for the patient

What were the findings of this research and what are the implications? We found that this feature of the immune response measured in this research appeared to be dependent on the dose of tremelimumab, with the largest immune response seen in patients who received the higher doses of tremelimumab. Patients with an increased immune response tended to have better outcomes; patients lived longer and their tumors decreased in size. The novel T300+D combination is now being investigated, versus sorafenib, in more patients in a larger clinical study called HIMALAYA

Where can I access more information?
Clinical trial registry: <https://clinicaltrials.gov/ct2/show/NCT02519348>
QR code materials: <https://bit.ly/2RsoTIU>

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References: 1. Kelley RK, et al. J Clin Oncol 2020;38(15_suppl). Abs 4087
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Introduction

- Tremelimumab (T), an anti-CTLA-4 antibody, and durvalumab (D), an anti-PD-L1 antibody, are under investigation as monotherapies and in combination for the treatment of multiple tumor types, including hepatocellular carcinoma (HCC)^{1,3-6}
- In this Phase 2 study in unresectable HCC (uHCC; Study 22, NCT02519348), a novel, single, priming dose of tremelimumab plus durvalumab (T300+D) combination regimen has shown favorable clinical activity and safety versus D or T monotherapy or another combination (T75+D)¹
- Additionally, an expansion of proliferative CD8+ lymphocytes at Day 15 was associated with improved response and was most profound for the T300+D-treated patients¹
- Changes in peripheral blood T cell receptor (TCR) clonality have been associated with clinical benefit from anti-CTLA-4 therapy in other indications⁷
- Here, we present an exploratory molecular analysis of peripheral blood TCRs, evaluating changes in TCR clonality in patients receiving different doses of T and D

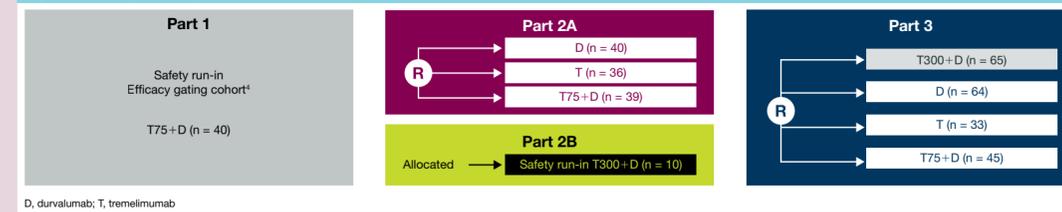
Methods

Study design

- Immune checkpoint therapy-naïve patients with uHCC who had progressed on, were intolerant to, or refused, sorafenib were randomized to receive one of the following dose regimens (Figure 1):

- T300+D: a single, priming dose of T 300 mg + D 1500 mg, then D 1500 mg Q4W
- D monotherapy: D 1500 mg Q4W
- T monotherapy: T 750 mg Q4W for a total of 7 doses, then Q12W
- T75+D: T 75 mg + D 1500 mg Q4W for a total of 4 doses, then D 1500 mg

Figure 1. Study 22 design

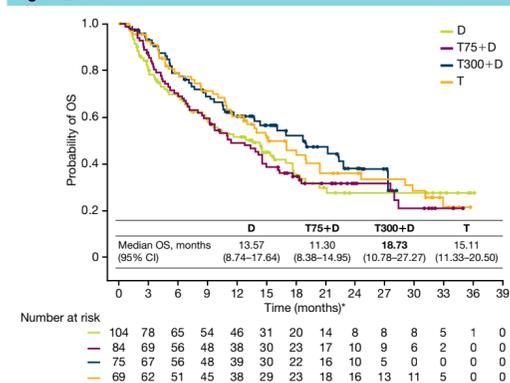


Results

Overall survival

- The longest median OS was observed with T300+D (Figure 2)¹

Figure 2. OS

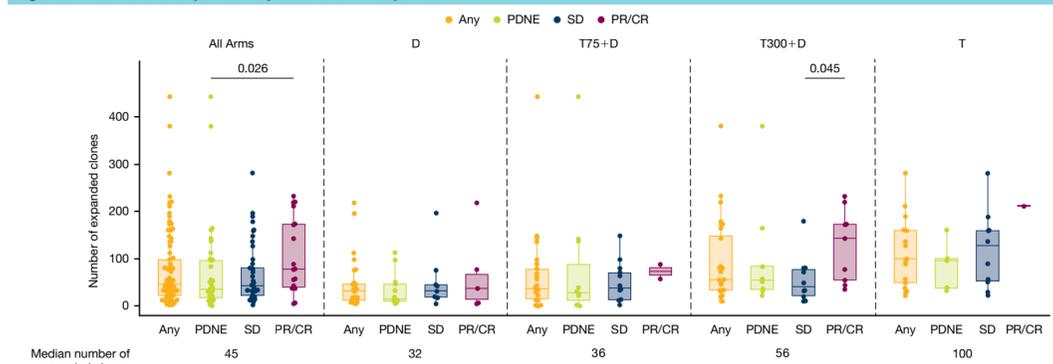


*Time from randomization (Part 2A, 3) or first dose (Part 2B)
CI, confidence interval; D, durvalumab; OS, overall survival; T, tremelimumab

Table 1. Response and clonality for clinically-evaluable and immunoSEQ-evaluable patients

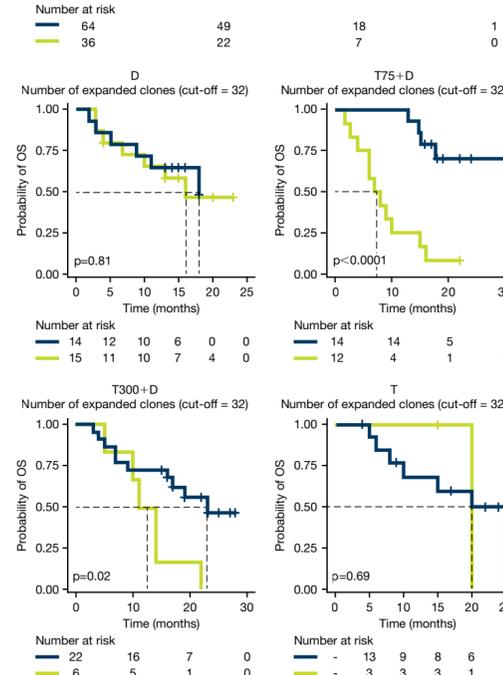
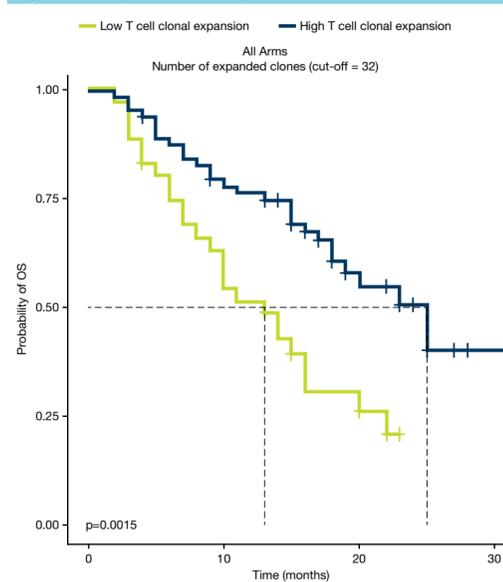
	D (n = 104)	T75+D (n = 84)	T300+D (n = 75)	T (n = 69)
Data for clinically-evaluable patients				
ORR, %	10.6	9.5	24.0	7.2
Median OS, months (95% CI)	13.6 (8.7-17.6)	11.3 (8.4-15.0)	18.7 (10.8-27.3)	15.1 (11.3-20.5)
Data for immunoSEQ-evaluable patients				
Patients with samples, n	30	26	28	17
ORR, %	23.0	7.7	32.0	5.9
Median OS, months (95% CI)	NA (10-NA)	16 (9-NA)	19 (11-NA)	21 (10-NA)
Median number of expanded T cell clones at Day 29 (95% CI)	32 (13-46)	36 (22-70)	56 (36-84)	100 (50-160)
Median baseline Simpson clonality (95% CI)	0.079 (0.046-0.890)	0.054 (0.040-0.090)	0.079 (0.046-0.110)	0.076 (0.036-0.880)
Median baseline richness, 10 ³ clones (95% CI)	59 (47-88)	62 (42-79)	51 (47-66)	60 (41-67)

Figure 3. T cell clonal expansion by best overall response



CR, complete response; D, durvalumab; PDNE, progressive disease and non-evaluable data plotted together; PR, partial response; SD, stable disease; T, tremelimumab

Figure 4. OS by T cell clonal expansion status



D, durvalumab; OS, overall survival; T, tremelimumab

Objectives and assessments

- Primary endpoint: safety
- Key secondary endpoints:
 - Overall survival (OS)
 - Objective response rate (ORR)
 - Duration of response

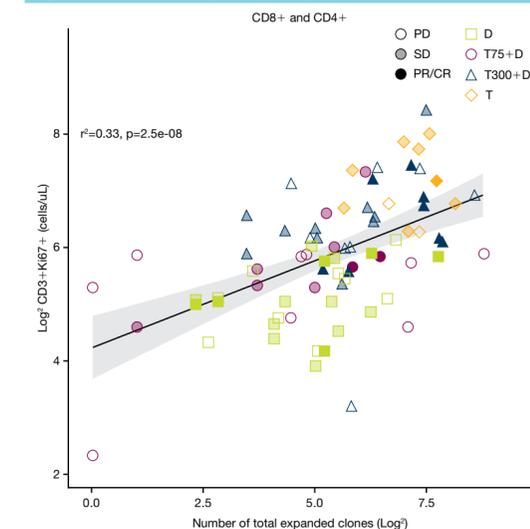
T cell receptor analysis

- DNA was isolated from PAXgene®-preserved whole blood collected at baseline and on Day 29, corresponding to 4 weeks after the single, initial dose of D and/or T from evaluable samples. Paired samples were available from a subset of clinically-evaluable patients
- CDR3 sequences of TCR β-chain were sequenced using the immunoSEQ® Assay (Adaptive Biotechnologies, Seattle, WA)
- Associations with ORR and OS were evaluated. Statistical measures of biomarker data are descriptive only and considered exploratory

- The increased number of expanded T cell clones correlates with increased proliferating T cells on treatment (Figure 5 and Supplementary Figure 3)

- Previously, we presented an association of response to T300+D with greater CD8+ T cell proliferation
- Here, we see a correlation between proliferating T cells and the number of expanded T cell clones
- Taken together, these data suggest that both expanding the number of unique T cell clones and increasing T cell proliferation are needed to optimize the chances for clinical response

Figure 5. T cell clonal expansion and proliferation



T cell proliferation was measured by flow cytometry on whole blood samples collected on Day 15 CD, cluster of differentiation; CR, complete response; D, durvalumab; PD, progressive disease; PR, partial response; SD, stable disease; T, tremelimumab

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Disclosures

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