

Cobolimab with dostarlimab and docetaxel in patients with advanced non-small cell lung cancer (NSCLC): COSTAR Lung

Poster No. 1223

Background



Immune checkpoint inhibitors targeting PD-(L)1 are approved in the first line setting for the treatment of lung cancer.¹ However, primary and acquired resistance to anti-PD-(L)1 therapy is common in solid tumors and novel immunotherapy combinations are needed.



TIM-3 is an immune checkpoint receptor highly expressed on multiple immune cell types and is associated with suppressed anti-tumor responses.²



TIM-3 upregulation on PD-1 positive tumor infiltrating lymphocytes is associated with reduced proliferation and secretion of cytokines important for T cell-mediated anti-tumor activity.³⁻⁵ Concurrent TIM-3 and PD-1 blockade is more effective at reducing tumor growth than blocking either pathway alone.⁶

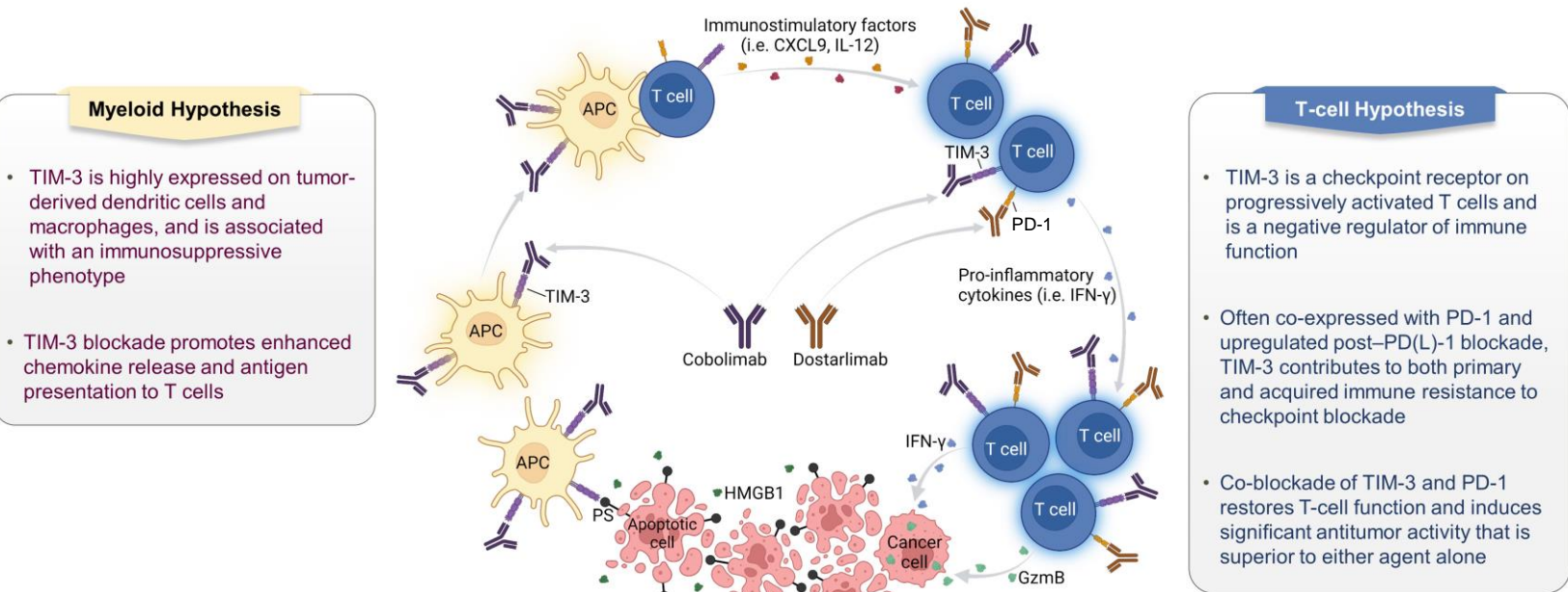


Cobolimab (TSR-022/GSK4069889), a novel first-in-class anti-TIM-3 mAb, activates immune cell function and induces significant anti-tumor activity when combined with anti-PD-1 agents (**Figure**).



Early clinical studies show durable responses with an acceptable safety profile when cobolimab was administered in combination with an anti-PD-1 mAb, dostarlimab, in prior-treated or treatment-naïve advanced or metastatic NSCLC.¹

Figure: Cobolimab Mechanism of Action



Objective

COSTAR Lung (NCT04655976) aims to compare the efficacy and safety of cobolimab plus dostarlimab plus standard of care chemotherapy (docetaxel; Arm A) versus dostarlimab plus docetaxel (Arm B) and docetaxel alone (Arm C) in patients with PD-1/PD-L1 relapsed/refractory NSCLC.

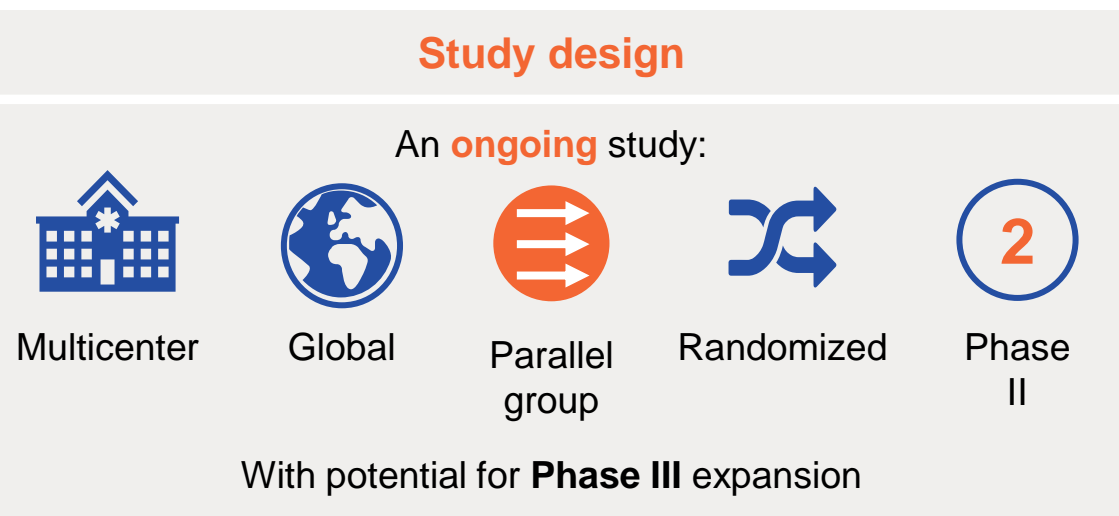
Abbreviations

AE, adverse event; ALK, anaplastic lymphoma kinase; APC, antigen-presenting cell; CTLA-4, cytotoxic T-lymphocyte-associated protein 4; CXCL9, C-X-C Motif Chemokine Ligand 9; DoR, duration of response; ECOG PS, Eastern Cooperative Oncology Group performance status; EGFR, epidermal growth factor receptor; EORTC-QLQ-C30, European Organisation for Research and Treatment of Cancer Quality of Life Questionnaire; EORTC-QLQ-C13, EORTC-QLQ-C13; I-Score, Lung Cancer Module; FFPE, formalin-fixed paraffin embedded; Gy, gray; GzmB, Granzyme B; HMGB1, high mobility group box 1; IA, interim analysis; IFN, interferon; iAE, immune-related AE; RECIST v1.1, immune-related RECIST version 1.1; mAb, monoclonal antibody; NSCLC, non-small cell lung cancer; ORR, objective response rate; OS, overall survival; PBCT, platinum-based chemotherapy; PD-1, programmed cell death protein-1; PD-L1, programmed cell death ligand 1; PFS, progression-free survival; RECIST v1.1, Response Evaluation Criteria in Solid Tumors version 1.1; ROS-1, ROS proto-oncogene 1; RT, radiotherapy; SAE, serious adverse event; TEAE, treatment-emergent AE; TIM-3, T-cell immunoglobulin and mucin-domain containing-3; TPS, tumor proportion score; TTD, time to treatment discontinuation.

Disclosures

HRK is a consultant and on advisory board for Merck, Bristol Myers Squibb (BMS), Ono, Takeda and AstraZeneca. HRK receives payment or honoraria for speaker's bureau from Merck, BMS, Ono, Takeda and AstraZeneca. CG served at a speaker's bureau and advisory board and received honoraria from AstraZeneca, BMS, Merck Sharp & Dohme, and Roche. DK declares no conflicts of interest. AT declares no conflicts of interest. EF declares advisory board participation for Abbvie, Amgen, AstraZeneca, Bayer, Beigene, Bluebird bio, Bristol Myers Squibb, Boehringer Ingelheim, BMS, Lilly, GlaxoSmithKline (GSK), Janssen, Medical Trends, Merck KGaA, Merck Sharp & Dohme, Novartis, Pemetrex, Pfizer, Puma Biotechnology, Regeneron, Roche, Sanofi, Genzyme, Syneos Health, Takeda, and Grifols; speaker bureau participation for AstraZeneca, Boehringer Ingelheim, BMS, Lilly, Merck Sharp & Dohme, Novartis, Pemetrex, Pfizer, prime Oncology, Roche, Takeda, Touchstone, and CME Outfitters; research funding from Fundación Merck Salud and a grant for oncology innovation; and is an independent member of the Board at Grifols. VV has received fees for consulting or serving on advisory boards for BMS, Merck, GSK, Foundation Medicine, AstraZeneca, EMD Serono, Novartis, and Novocure. YJK declares no conflicts of interest. TOG declares no conflicts of interest.

Methods



Cobolimab dose of 300 mg is determined based on the Phase 1 study AMBER (GSK Study 213348; NCT02817633)⁷

Participants will be stratified according to:

- Prior lines of therapy: 1 prior line of therapy vs 2 prior lines of therapy
- PD-L1 status: tumor proportion score $\geq 50\%$ vs $< 50\%$
- Histology: squamous vs non-squamous

This study will be completed in three phases: screening, treatment, and follow-up

Key inclusion/exclusion criteria:



Key inclusion criteria

- ≥ 18 years old
- Pathologically-proven advanced or metastatic NSCLC, and only squamous or non-squamous cell carcinoma
- Received ≤ 2 lines of therapy for advanced/metastatic disease
- Measurable disease per RECIST v1.1
- Documented radiographic disease progression on prior PBCT and on prior anti-PD-(L)1 therapy
- Agrees to submit an archival FFPE tumor tissue specimen that was collected on or after diagnosis
- Documented PD-L1 status
- ECOG PS of 0 or 1
- Life expectancy of at least 3 months and adequate organ function
- Participant has recovered from any prior treatment-related toxicities



Key exclusion criteria

- Previous treatment with an anti-PD(L)-1 agent that resulted in permanent discontinuation due to AE
- Previous treatment with an anti-TIM-3 or anti-CTLA-4 agent or docetaxel
- Documented sensitizing EGFR, ALK, or ROS-1 mutation
- Radiologic or clinical disease progression ≤ 8 weeks after initiation of prior anti-PD-1/PD-L1 antibody treatment
- Received > 30 Gy RT to the lung within 6 months of first dose of study treatment
- New or progressive brain metastases and/or leptomeningeal metastases
- Active autoimmune disease or active infection
- Current interstitial lung disease, current pneumonitis, or history of pneumonitis

Acknowledgments

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Figure: COSTAR Lung Study Schema

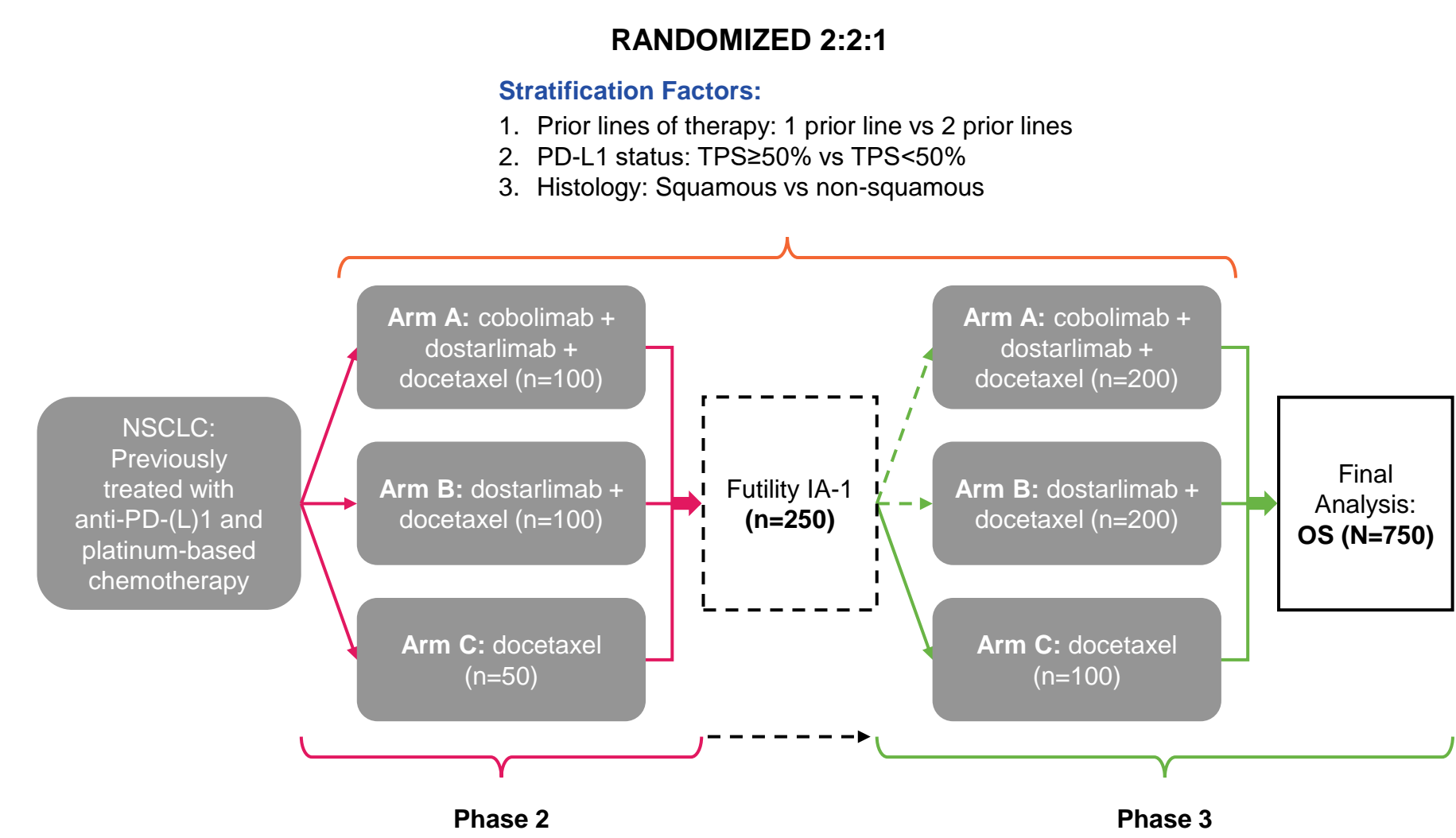
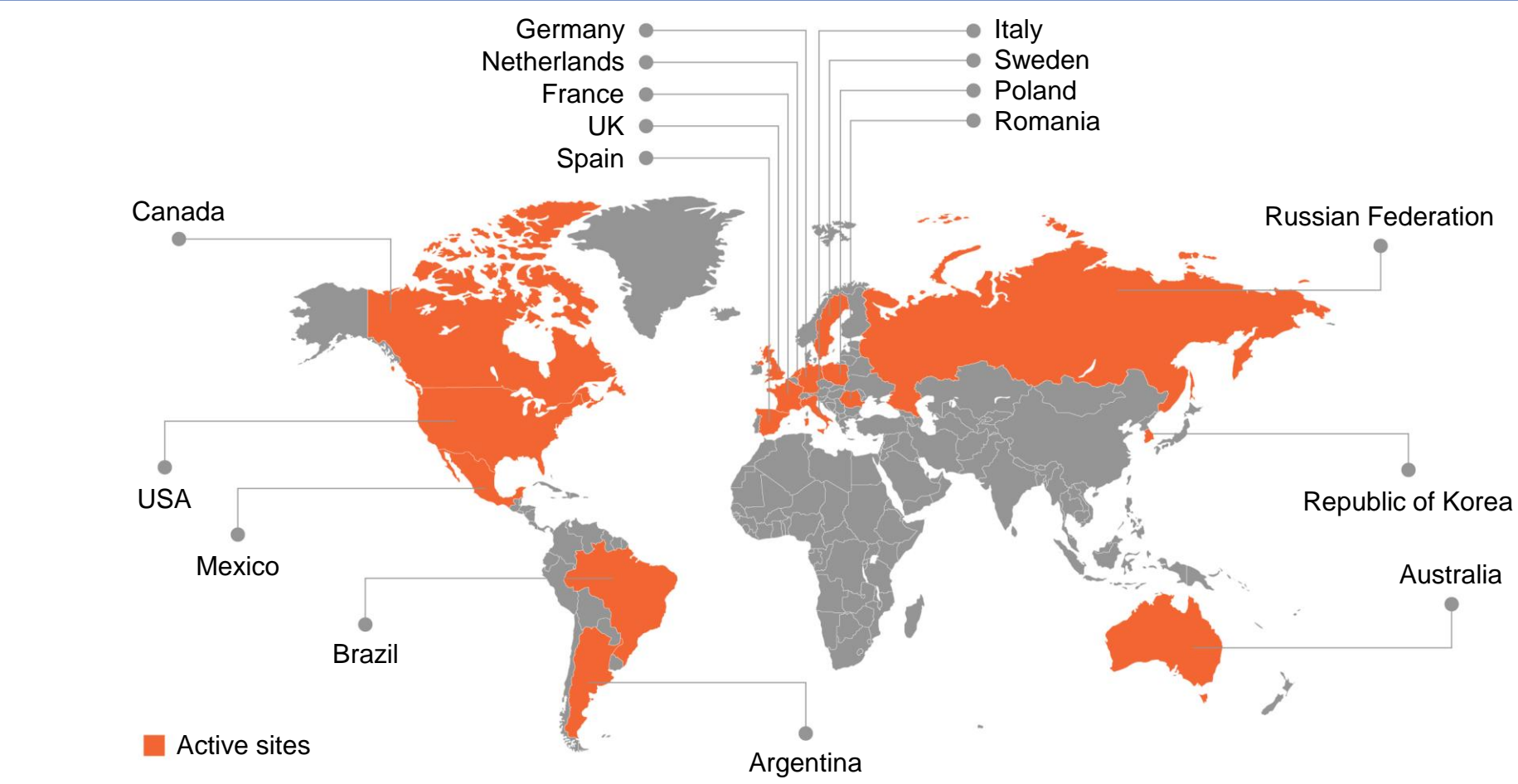


Figure: Trial Status



Study endpoints and assessments

Primary endpoints:

- OS for Arms A or B vs C
 - OS defined as survival from the date of randomization to the date of death by any cause

Secondary endpoints:

- Arms A vs B:
 - OS
- Arms A vs B, A vs C, and B vs C:
 - Confirmed ORR per investigator assessment based on RECIST v1.1
 - PFS per RECIST v1.1
 - DoR per RECIST v1.1
 - TTD: defined as time from randomization to meaningful deterioration on a composite endpoint of dyspnea, chest pain, and cough (from EORTC-QLQ-LC13 domains)
 - Change from baseline as assessed by EORTC-QLQ-C30 and EORTC-QLQ-LC13 domains
- Arms A and B vs C:
 - Safety and tolerability per incidence of TEAEs, SAEs, irAEs, TEAEs leading to death, and AEs leading to discontinuation

Exploratory endpoints:

- Confirmed ORR per investigator assessment and PFS (iRECIST)
- Pharmacokinetics
- Immunogenicity
- Biomarkers of response
- Patient-reported efficacy and tolerability
- Healthcare resource utilization

Statistical analyses

Futility analysis to assess ORR across the 3 arms in the Phase 2 portion.

Conducted after 250 participants are enrolled in total (Arm A and B: 100 each, Arm C: 50), and have been followed-up for a minimum period, have withdrawn from the study, or died.

An additional 500 patients may be included in the Phase 3 portion and final analysis will be conducted on OS primary endpoint.

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Presenting author email: nobelg@yuhs.ac

