

Clinical Value of Testing for c-Met Protein Overexpression in Patients With Non-Small Cell Lung Cancer

Samuel Crawford^{1*}, Kellie Woll¹, Pam Martin², Tim Klein², Pallavi Mhaske¹, Andrea Wagner¹

¹AbbVie Inc., North Chicago, IL, USA;

²Medical Decision Modeling Inc, Indianapolis, IN, USA.

*Presenting author

OBJECTIVE

To evaluate the clinical value of testing patients with advanced/metastatic *EGFR* wild-type non-squamous non-small cell lung cancer (NSCLC) with MET immunohistochemistry (IHC) to identify those with c-Met protein overexpression (OE) or high OE, who would then be eligible for telisotuzumab vedotin (Teliso-V), an antibody-drug conjugate

CONCLUSION

The model findings suggest clinical value in implementing testing for c-Met protein OE among all eligible NSCLC patients as demonstrated by improved outcomes among patients when MET IHC testing is conducted

For additional information or to obtain a PDF of this poster



Scan QR code or to download an electronic version of this presentation and other AbbVie ISPOR 2025 scientific presentations
QR code expiration date: April 16, 2026
To submit a medical question, please visit www.abbvieinfo.com

Funding

AbbVie funded this study and participated in the study design, research, analysis, data collection, interpretation of data, reviewing, and approval of the publication. All authors had access to relevant data and participated in the drafting, review, and approval of this publication. No honoraria or payments were made for authorship. Medical writing support was provided by Joann Hettasch, PhD of Avalere Health, Ltd., and funded by AbbVie, Inc.

Disclosures

Samuel Crawford, Kellie Woll, Pallavi Mhaske, and Andrea Wagner are employees of AbbVie and may hold stock or stock options.

Pam Martin and Tim Klein are employees of Medical Decision Modeling Inc, which received funding from AbbVie, Inc. for the conduct of this study.

References

- American Cancer Society. Cancer Facts and Figures 2023. <https://www.cancer.org/research/cancer-facts-statistics/all-cancer-facts-figures/2023-cancer-facts-figures.html>; 2023 [accessed 14 March 2025].
- Hendriks LE, et al. *Annals of Oncology*. 2023;34(4):358–376.
- Leight NB, et al. *J Clin Oncol*. 2025;43:e17–e30.
- Bazhenova L, et al. *J Clin Oncol*. 2025;42:e72–e86.
- Bahrani A, et al. *J Cell Physiol*. 2017;232(10):2657–73.
- Camidge DR, et al. *Clin Cancer Res*. 2021;27(21):5781–92.
- De Miguel M, et al. *Annals of Oncology*. 2024;35 (suppl_2):S805–6.
- Gymnopoulos M, et al. *Mol Oncol*. 2020;14(1):54–68.
- Yang CY, et al. *Acta Pharmacol Sin*. 2019;40(7):971–9.
- Cappuzzo F, et al. *Annals of Oncology*. 2023;34 (suppl_4):S1671.

INTRODUCTION

- NSCLC accounts for 81% of all lung cancers in the US¹
- Guidelines recommend that patients with locally advanced or metastatic NSCLC receive targeted therapy or immunotherapy,^{2–4} thus comprehensive biomarker testing in these patients is essential for diagnosis and selection of appropriate treatment
- c-Met OE has emerged in recent years as a clinically relevant and potentially actionable protein biomarker in NSCLC⁵
- c-Met protein targeted agents currently being investigated in NSCLC include the antibody drug conjugates, Teliso-V,⁶ telisotuzumab adizutecan,⁷ RC108 (NCT05821933), TR1801,⁸ SHR-A1403,⁹ and the bispecific antibody MCLA-129,¹⁰ with Teliso-V at the most advanced stage in development
- These investigational c-Met protein targeted therapies may require tissue-based biomarker assessment by IHC prior to use as therapy
- Thus, it is important to understand the overall prevalence and prognostic value of c-Met OE as a biomarker

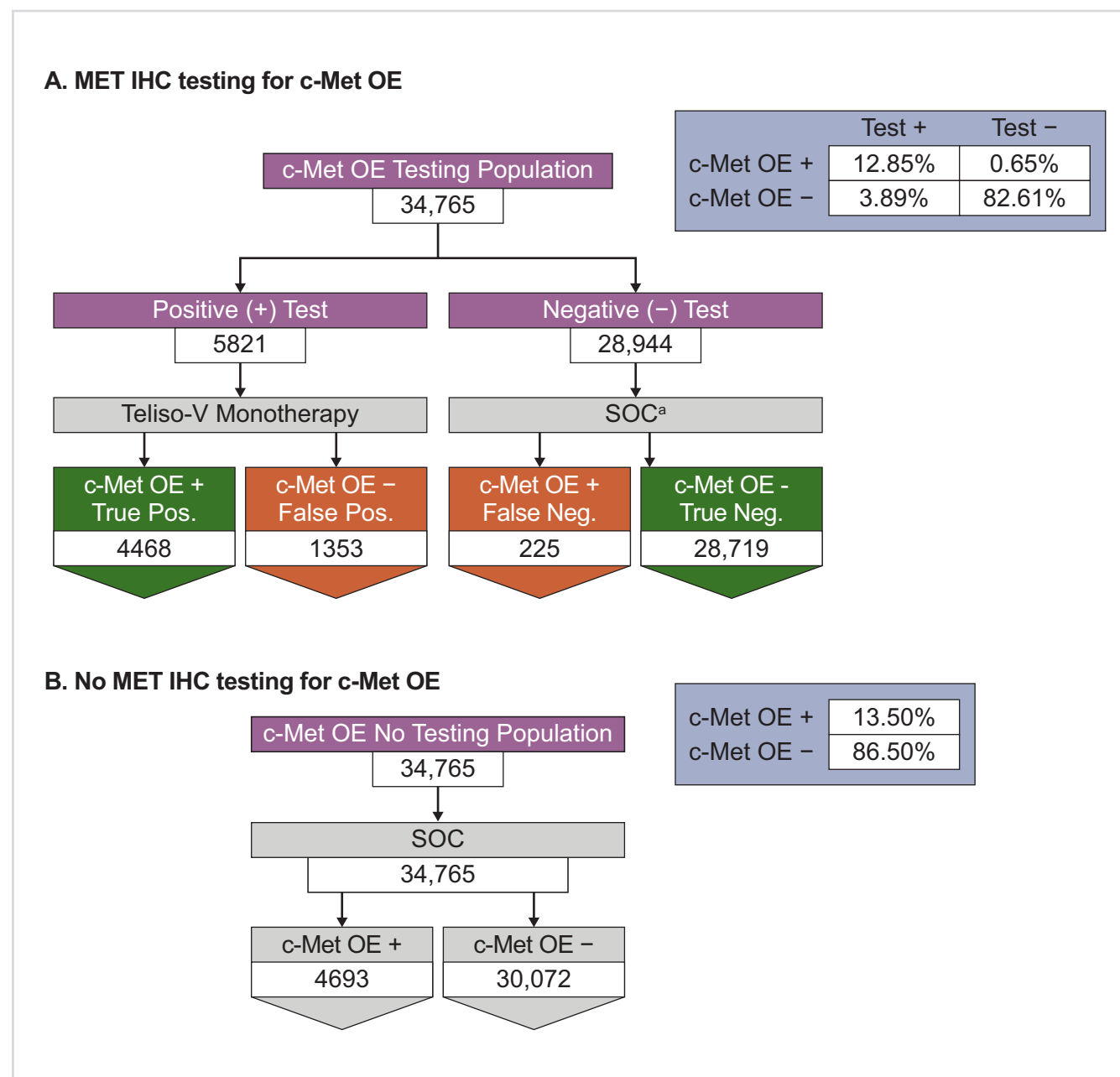
METHODS

- A deterministic decision-tree (for treatment and MET IHC testing allocations) converted into a stochastic partition-survival model (for efficacy-related outcomes) was used to evaluate the clinical utility of MET IHC testing
- The model simulated patients into two cohorts: MET IHC tested or not tested
 - Patients with no testing received a market basket of standard of care (SOC) regimens
 - Patients with testing and c-Met protein OE positivity received Teliso-V, while patients with c-Met low/no OE received SOC
 - Two scenarios were analyzed: c-Met OE or c-Met high OE, assessed by IHC with these thresholds:
 - c-Met high OE (≥50% tumor cells staining at 3+ intensity)
 - c-Met OE (≥25% tumor cells staining at 3+ intensity)
 - c-Met low/no OE (<25% tumor cells staining at ≤3 intensity)
- Prevalence of NSCLC and c-Met protein OE as well as clinical outcomes among patients with c-Met protein OE treated with SOC or Teliso-V were based on published literature (**Table 1**)
- Efficacy inputs (ie, overall survival, progression-free survival) were estimated based on exponential survival curves derived from published clinical and real-world data (**Supplemental Figures 1** and **2**)
- The model estimated expected and incremental life years (LYs) and quality-adjusted life years (QALYs)

RESULTS

- The deterministic decision-tree for c-Met high OE is illustrated in **Figure 1**, which demonstrates the testing vs no testing strategies and highlights the sensitivity and specificity of the MET IHC test to identify patients with c-Met high OE

Figure 1. Decision tree for c-Met high OE scenario



*SOC includes a market basket of immunotherapies (eg, atezolizumab, pembrolizumab, nivolumab), chemotherapies (eg, docetaxel with or without nintedanib), and targeted therapies (eg, ramucirumab + docetaxel).
OE, overexpression; SOC, standard of care.

METHODS (CONTINUED)

Table 1. Model features and parameters

Parameter	Source/input
Prevalence and population	
Prevalence NSCLC among adults	SEER*StatDatabase: Incidence - SEER Research Plus Data, 18 Registries. 2016; BIM
Patients initially diagnosed with locally advanced disease in 1 year	SEER*StatDatabase: Incidence - SEER Research Plus Data, 18 Registries. 2017; BIM
Proportion with advanced/metastatic NSCLC including recurrent disease	Huntzinger 2021 <i>J Thor Onc</i> 2021;16(3): S317-S318.
Patients with 2+ lines of therapy	Nadler E, et al. <i>J Cancer Res Clin Oncol</i> . 2021; 147(3):671-690.
Percent EGFR wild-type	Dogan S, et al. <i>Clin Cancer Res</i> . 2012;18(22):6169-6177.
Population cohort	N = 34,765
Biomarker prevalence: % c-Met OE, % c-Met high OE	Camidge DR, et al. <i>J Clin Oncol</i> . 2024;42(25):3000-3011.
Efficacy inputs	
Progression-free survival	Camidge DR, et al. <i>J Clin Oncol</i> . 2024;42(25):3000-3011. Le 2024 ESMO
Overall survival	Camidge DR, et al. <i>J Clin Oncol</i> . 2024;42(25):3000-3011. Le 2024 ESMO
Model analysis year costs	2025 US dollars
Length of analysis	120 months
Comparators	Teliso-V, SOC

BIM, budget impact model; OE, overexpression; SOC, standard of care; teliso-V, telisotuzumab vedotin.

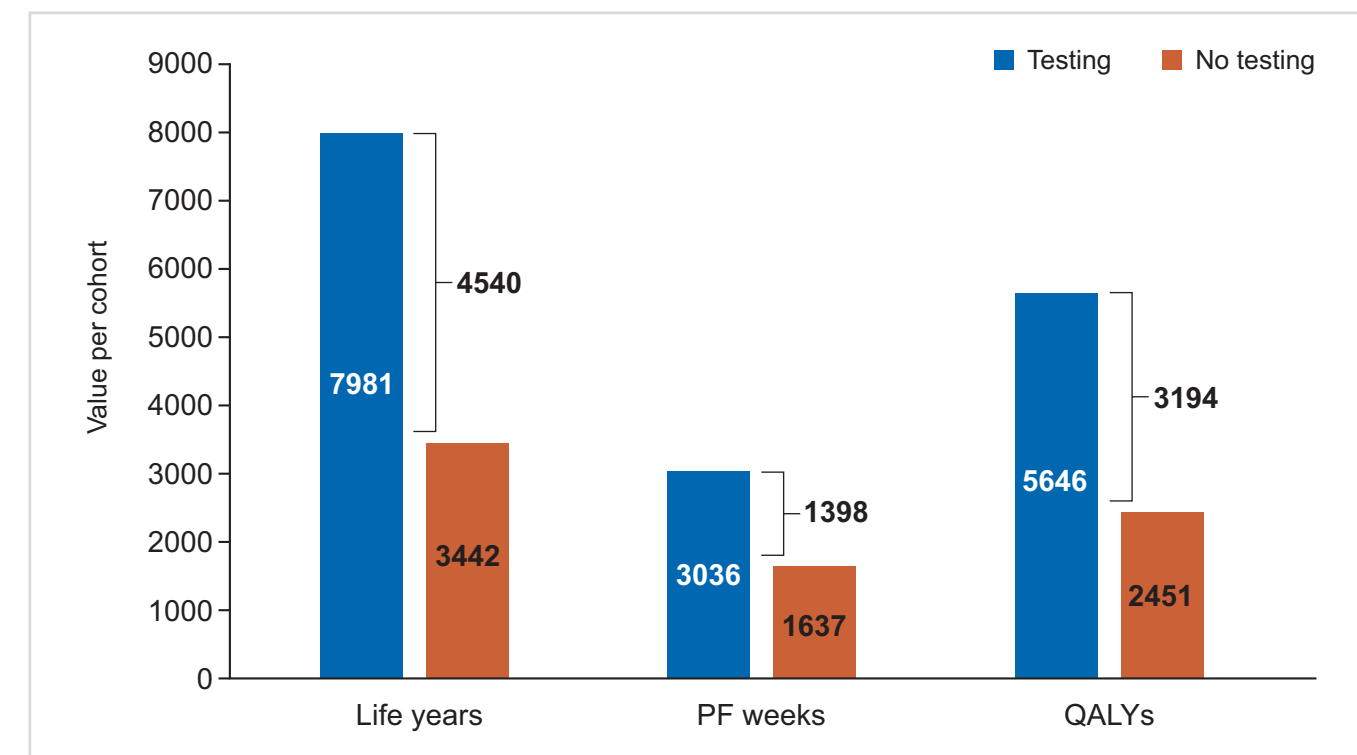
c-Met high OE scenario

- Based on results obtained from the model, the c-Met high OE tested cohort had an incremental improvement of 4540 LYs and 3194 QALYs relative to patients in the not tested cohort, among all eligible patients in the US (**Figure 2**).
- Among patients who would have tested positive for c-Met high OE in both the tested and non-tested cohorts, those in the tested cohort had a 131.9% and 130.3% incremental improvement in LYs and QALYs, respectively

c-Met OE scenario

- Similar trends were observed for patients in the c-Met OE scenario:
 - Among all eligible patients in the US who would have tested positive for c-Met protein OE in the tested cohort, an incremental improvement of 6154 LYs and 4363 QALYs was observed relative to the non-tested cohort
 - Among patients who would have tested positive for c-Met protein OE in both the tested and non-tested cohorts, those in the tested cohort had a 79.0% and 79.9% incremental improvement in LYs and QALYs, respectively

Figure 2. Incremental improvement in life years, progression-free survival, and QALYs in tested (and presumed c-Met high OE positive) vs not tested cohorts



c-Met high OE positive for the testing cohort is the sum of patients on Teliso-V who tested positive and those who tested negative but were actually c-Met high OE positive and received SOC.
PF, progression-free; QALYs, quality-adjusted life years.