Tislelizumab Versus Docetaxel in Previously Treated Advanced Non-Small Cell Lung Cancer: Final Analysis of RATIONALE-303

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Objectives: In RATIONALE-303 (NCT03358875) tislelizumab significantly improved overall survival (OS) versus docetaxel in the intent-to-treat (ITT) population at the interim analysis (IA), based upon which tislelizumab was approved in China for treatment of advanced non-small cell lung cancer (NSCLC) patients with progressive disease after chemotherapy. Here, we report outcomes of the final analysis (FA) and post hoc biomarker analysis.

Methods: Patients ≥18 years with histologically confirmed, locally advanced or metastatic squamous or nonsquamous NSCLC were randomized (2:1) to intravenous (IV) tislelizumab 200 mg or IV docetaxel 75 mg/m² every 3 weeks. Co-primary endpoints were OS in the ITT and PD-L1 tumor cell (TC) ≥25% populations. The study had one planned IA only in the ITT population. The FA was conducted in the PD-L1 TC ≥25% population with secondary endpoints (investigator-assessed progression-free survival [PFS], objective response rate, and duration of response) tested sequentially once superiority of OS in the PD-L1 TC ≥25% population was demonstrated in the FA. Exploratory biomarker analyses included PD-L1 expression, tumor mutational burden (TMB), and gene expression profile.

Results: Between November 30, 2017, and April 8, 2020, 805 patients were randomized to tislelizumab (N=535) or docetaxel (N=270). The co-primary endpoint of OS (ITT) was met at IA (data cut-off August 10, 2020). At data cut-off CSCO 2022

(July 15, 2021), FA was conducted in the PD-L1 TC ≥25% population. Median follow-up times (reverse Kaplan-Meier method) were 30.9 months for tislelizumab and 27.5 months for docetaxel. In the ITT population, tislelizumab continued to improve OS versus docetaxel (median OS 16.9 months vs 11.9 months, respectively; hazard ratio [HR]=0.66). In the PD-L1 TC ≥25% population, tislelizumab showed a statistically significant OS benefit versus docetaxel (median OS 19.3 months vs 11.5 months; HR=0.53; P<0.0001). A consistent OS benefit was observed for almost all pre-defined subgroups. The study also met secondary endpoints at this FA. In the post hoc biomarker analysis, the association of TMB and genetic alterations including single-target gene mutation or pathway mutations with clinical outcomes was further explored. Compared with TMB, which was correlated with PFS benefit for tislelizumab versus docetaxel but was not correlated to OS benefit, except at the highest cutoff (≥14 mut/Mb), NOTCH1-4 mutations showed association with better tislelizumab efficacy, which was correlated with both PFS and OS benefit (Table). No new safety signals were identified.

Conclusions: Tislelizumab continued to improve OS versus docetaxel in pretreated advanced NSCLC regardless of PD-L1 expression at final analysis. Biomarker analysis implied the potential association of *NOTCH1-4* mutations with greater tislelizumab efficacy for both OS and PFS.

Table

	ITT Population		PD-L1 TC ≥25% Population		NOTCH1-4 Mut Population		NOTCH1-4 WT Population	
	TIS	D	TIS	D	TIS	D	TIS	D
	(N=535)	(N=270)	(N=227)	(N=116)	(N=26)	(N=15)	(N=218)	(N=101)
	365	206		87 (75.0)	13 (50.0)	13 (86.7)		
OS events, n (%)	(68.2)	(76.3)	141				152	79
[IA]	[275	[166	(62.1)				(69.7)	(78.2)
	(51.4)]	(61.5)]						
	16.9	11.9						
Median OS (95% CI), months [IA]	(15.2 <i>,</i> 19.1)	(9.6 <i>,</i> 13.5)	19.3 (16.5, 22.6)	11.5 (8.2, 13.5)	24.7 (14.2, NE)	7.7 (3.3, 14.3)	15.7 (13.9, 17.9)	12.9 (10.4, 14.9)
	[17.2 (15.3,	[11.9						
	20.0)]	13.9)]						
Stratified HR ^c (95% CI) [IA]	0.66 (0.56, 0.79) P<0.0001 ^{a,b} [0.64 (0.53, 0.78) P<0.0001 ^a]		0.53 (0.40, 0.70) P<0.0001 ^a		0.22 (0.10, 0.49) P=0.0002 ^{a,b}		0.75 (0.57, 0.99) P=0.0390 ^{a,b}	

PFS _{INV} events, n	451	208	177	94 (81.0)	14 (53.8)	14 (93.3)	187	83 (82.2)
(%)	(84.3)	(77.0)	(78.0)	54 (61.0)	14 (33.0)	14 (33.3)	(85.8)	03 (02.2)
Median PFS _{INV}	4.2	2.6	6.5	2.4	14.1	2.6	4.1	3.3
(95% CI), months	(3.9, 5.5)	(2.2, 3.8)	(6.2, 8.3)	(2.1, 4.1)	(6.2, NE)	(2.0, 4.1)	(2.2, 6.2)	(2.1, 4.1)
Stratified HR ^c (95% CI)	0.63 (0.53, 0.75)		0.37 (0.28, 0.49)		0.17 (0.08, 0.37)		0.72 (0.55, 0.95)	
ORR _{INV} , n (%)	121 (22.6)	19 (7.0)	85 (37.4)	8 (6.9)	-	-	-	-
Median DoR _{INV} , (95% CI), months	13.5 (8.5, 19.6)	6.0 (2.1, 7.2)	11.9 (8.3, 19.6)	4.2 (0.6, 6.1)	-	-	-	-

IA data cut-off: August 10, 2020.

FA data cut-off: July 15, 2021.

Abbreviations: CI, confidence intervals; D, docetaxel; DoR_{INV}, investigator-assessed duration of response; FA, final analysis; HR, hazard ratio; IA, interim analysis; ITT, intent-to-treat; mut, mutation; NE, not estimable; ORR_{INV}, investigator-assessed objective response rate; OS, overall survival; PD-L1, programmed death-ligand 1; PFS_{INV}, investigator-assessed progression-free survival; TC, tumor cell; TIS, tislelizumab; WT, wild type.

^a 1-sided stratified log-rank test.

^b Descriptive *P*-value.

^c Stratified by histology (squamous vs nonsquamous) and lines of therapy (second vs third).