HER2 Testing in the **MOUNTAINEER Trial: Analysis of** Treatment Response Based on **Central HER2 Assessment Using IHC/ISH and NGS**

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Objectives

- To explore correlations between tissue and blood-based biomarkers and clinical outcomes in the chemo-refractory HER2+ RAS WT mCRC patient population
- To evaluate HER2 testing methodologies (tissue and blood) that can be used to identify patients with HER2+ mCRC who may benefit from treatment with TUC + Tras

Conclusions

The percent agreement of HER2 status was high across the three different platforms (IHC/FISH, Tissue NGS, and Blood NGS), with higher percent agreement between tissue-based assays

Treatment response to TUC + Tras was predicted by a HER2+ result from any of the three testing platforms

- Patients in which HER2 amplification was not detected by ctDNA NGS may have HER2 amplification by a tissue-based assay and may benefit from treatment with a HER2-directed therapy
- cORR in IHC2+/ISH+ was numerically lower than IHC3+ but remained clinically relevant

These data support the use of both tissue and blood-based methods to identify HER2+ mCRC in patients that may benefit from treatment with TUC + Tras until an FDA-approved assay is available

Abbreviations

AE, adverse event; BICR, blinded independent central review; BID, twice a day; BRAF, proto-oncogene B-Raf; C1D1, cycle 1 day 1; CE, Conformité Européenne; CISH, chromogenic in situ hybridization; CLIA, Clinical Laboratory Improvement Amendments; cORR, confirmed objective response rate; CR, complete response; ctDNA, circulating tumor DNA; DOR, duration of response; FDA, United States Food and Drug Administration; FISH, fluorescence in situ hybridization; HER2, human epidermal growth factor receptor 2; IHC, immunohistochemistry; ISH, in situ hybridization; ISO, International Organization for Standardization; mAb, monoclonal antibody; mCRC, metastatic colorectal cancer; mDOR, median duration of response; mOS, median overall survival; mPFS, median progression-free survival; NA, not available; NGS, next-generation sequencing; OS, overall survival; PD, progressive disease; PFS, progression-free survival; PO, orally; PR, partial response; Q3W, every 3 weeks; QC, quality control; R, randomization; RAS, rat sarcoma virus; RECIST, Response Evaluation Criteria in Solid Tumors; SD, stable disease; TKI, tyrosine kinase inhibitor; Tras, trastuzumab; TUC, tucatinib; VEGF, vascular endothelial growth factor

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- HER2 overexpression and/or amplification (HER2+) occurs in 3%-5% of patients with metastatic colorectal cancer (mCRC)¹⁻⁵
- Rates of HER2+ may be as high as ~10% in patients with RAS/BRAF wild-type mCRC tumors
- Established regional guidelines recommend HER2 testing and HER2-directed treatment options for mCRC^{1,6}
- Tucatinib is a highly selective HER2-directed TKI approved by the FDA in combination with trastuzumab for treatment of patients with RAS wild-type HER2+ unresectable or mCRC that has progressed following treatment with fluoropyrimidine-, oxaliplatin-, and irinotecan-based chemotherapy
- FDA approval was based on the primary analysis of the MOUNTAINEER trial (NCT03043313)

MOUNTAINEER Summary

• Results from the MOUNTAINEER primary analysis showed clinically meaningful activity and demonstrated tucatinib in combination with trastuzumab was well tolerated⁷

Efficacy (n=84)	
cORR per BICR	38.1%
mDOR	12.4 months
mPFS	8.2 months
mOS	24.1 months

Safety (n=86)

- No deaths due to AEs Diarrhea was most common AE • Grade 1 or 2: 60.5% • Grade 3: 3.5%
- Low discontinuation rate due to AEs: 5.8%

Samples Available for Central HER2 Testing MOUNTAINEER Cohorts A+B: Response Assessment by Testing Platform Patients enrolled in all cohorts per ≥1 local testing method^a: 114 Cohort C **Cohort A** Cohort B 30 patients 39 patients 45 patients Cohorts A + B: Samples submitted for central testing^b with evaluable results Blood NGS: Tissue NGS: IHC/FISH

^aPatients were enrolled after locally testing tissue by at least one of the approved testing methods (IHC, ISH,

^bTissue samples were retrospectively analyzed for HER2 status at the sponsor-designated central laboratory ^cVariability observed in sample size for each assay is due to differences in QC/acceptance criteria of each assay and collection/availability of samples

Percent Agreement Between Three Central HER2 Testing Modalities

 High level of percent agreement of HER2 status observed across multiple central HER2 testing modalities (all cohorts included; pairwise comparisons)



	Central IHC + FISH (n=70)		
Response	Positive (IHC3+) (n=45)	Positive (IHC2+/ISH+) (n=15)	Negative (n=10)
CR	3	0	0
PR	18	3	1
SD ^a	17	5	4
PD	7	6	5
NA	0	1	0
cORR, n (%) (95% CI)	21 (46.7%) (31.7, 62.1)	3 (20.0%) (4.3, 48.1)	1 (10.0%) (0.3, 44.5)
mDOR, months (95% CI)	16.4 (*	10.6, 25.5)	_
mPFS, months (95% CI)	10.1 ((4.2, 15.2)	2.8 (1.2, 6.3)

^aIncludes non-CR/non-PD

Response

SD ^a PD NA CORR, n ((95% CI) mDOR, m mPFS, mo	PR
PD NA CORR, n ((95% CI) mDOR, m mPFS, mo alncludes nor	SD ^a
NA cORR, n ((95% CI) mDOR, m mPFS, ma alncludes nor	PD
cORR, n ((95% CI) mDOR, m mPFS, mo alncludes nor	NA
mDOR, m mPFS, mo ^a Includes nor	cORR, n ((95% CI)
^a Includes nor	mDOR, m
^a Includes nor	mPFS, mo
	^a Includes nor

PR SD^{a}

PD NA cORR, n (95% CI) mDOR, m mPFS, mo

^aIncludes non-CR/non-PD

Background

MOUNTAINEER: Global, Open-Label, Phase 2 Trial



^aTucatinib dose: 300 mg PO BID, trastuzumab dose: 6 mg/kg Q3W (loading dose 8 mg/kg C1D1); each treatment cycle is 21 days ^bPatients remained on therapy until evidence of radiographic or clinical progression, unacceptable toxicity, withdrawal of consent, or study closures

^cStratification: left sided tumor primary vs other

Patients were allowed to cross over and receive tucatinib and trastuzumab if they experienced radiographic progression at any time point or if they had not achieved a PR or CR by week 12

^eEfficay assessed in patients who received any anount of study treatment and had HER2+ tumors fRECIST v1.1 per BICR

Results

• cORR per BICR was 40.0% to 47.7% for patients identified as having HER2+ disease by any of the three assays • DOR per BICR was 12.4-16.4 months for patients identified as having HER2+ disease by any of the three assays • Tumors confirmed as HER2+ by central IHC/ISH had a mDOR of 16.4 months (95% CI: 10.6, 25.5) and mPFS of 10.1 months (95% CI: 4.2, 15.2)

PGDx tissue NGS HER2 amplification HER amplification detected not detected n=44) 20 0 (0%) (0, 45.9) 21 (47.7%) (32.5, 63.3) 15.3 (8.9, 25.5) nonths (95% CI) 2.1 (1.3, -) 10.9 (7.0, 20.7) onths (95% CI) on-CR/non-PD



	Guardant ctDNA			
e	HER amplification detected (n=56)	HER2 amplification not detected (n=15)		
	1	1		
	22	2		
	18	7		
	14	4		
	1	1		
%)	23 (41.1%) (28.1, 55)	3 (20.0%) (4.3, 48.1)		
onths (95% CI)	12.4 (6.2, 38.3)	_		
onths (95% CI)	8.1 (3.1, 10.2)	10.9 (2.0, 18.4)		





^aConfirmed HER2-positive mCRC tested at CLIA-certified or ISO-accredited laboratory: IHC: HER2 3+ IHC by an FDA-approved or CE-marked HER2 IHC test following interpretational manual for breast cancer, ISH: HER2 2+ IHC with amplification by an FDA-approved or CE-marked HER2 in situ hybridization assay (FISH or CISH) following interpretational manual for breast cancer NGS: HER2 amplification by CLIA-certified or ISO-accredited NGS sequencing assay

^bHER2 status was evaluated in tissue and blood samples using multiple assays. Tissue samples were analyzed by IHC (Ventana HER2 (4B5) IHC Assay), FISH (Dako iQFISH pharmDx assay) per the interpretational manual for breast cancer. Tissue NGS was performed using the PGDx elio tissue complete assay. NGS analysis of ctDNA was done using the Guardant 360 assay

- There are currently no established best practices for HER2 testing and/or interpretation in mCRC
- Previously presented analysis of MOUNTAINEER showed that there is a very high level of concordance between breast and gastric algorithms, suggesting either algorithm (with appropriate training) can be used to identify patients with HER2+ mCRC⁸
- Here we present clinical outcomes stratified by HER2 testing method for patients with mCRC treated with TUC + TRAS



The one responder that was HER negative by central IHC/FISH was HER2 amplified by other central methods