

CDX3379-04: Phase 2 Evaluation of CDX-3379 in Combination with Cetuximab in Patients with Advanced Treatment-Refractory Head and Neck Squamous Cell Carcinoma

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BACKGROUND

- ErbB3 signaling may contribute to resistance to targeted therapies such as cetuximab
- CDX-3379 is a fully human anti-ErbB3 monoclonal IgG1 λ antibody with extended serum half-life and a potent inhibitor of ErbB3 activation via a unique mechanism of action that locks ErbB3 in an inactive configuration¹
- CDX-3379 antitumor activity has been observed in patients with head and neck squamous cell carcinoma (HNSCC) patients in previous clinical studies:
 - In a phase 1b study, a durable complete response (CR) with CDX-3379 and cetuximab in a patient with cetuximab-refractory HPV(-) HNSCC²
 - In a window of opportunity study, an exceptional clinical response in a patient with HPV(-) HNSCC³

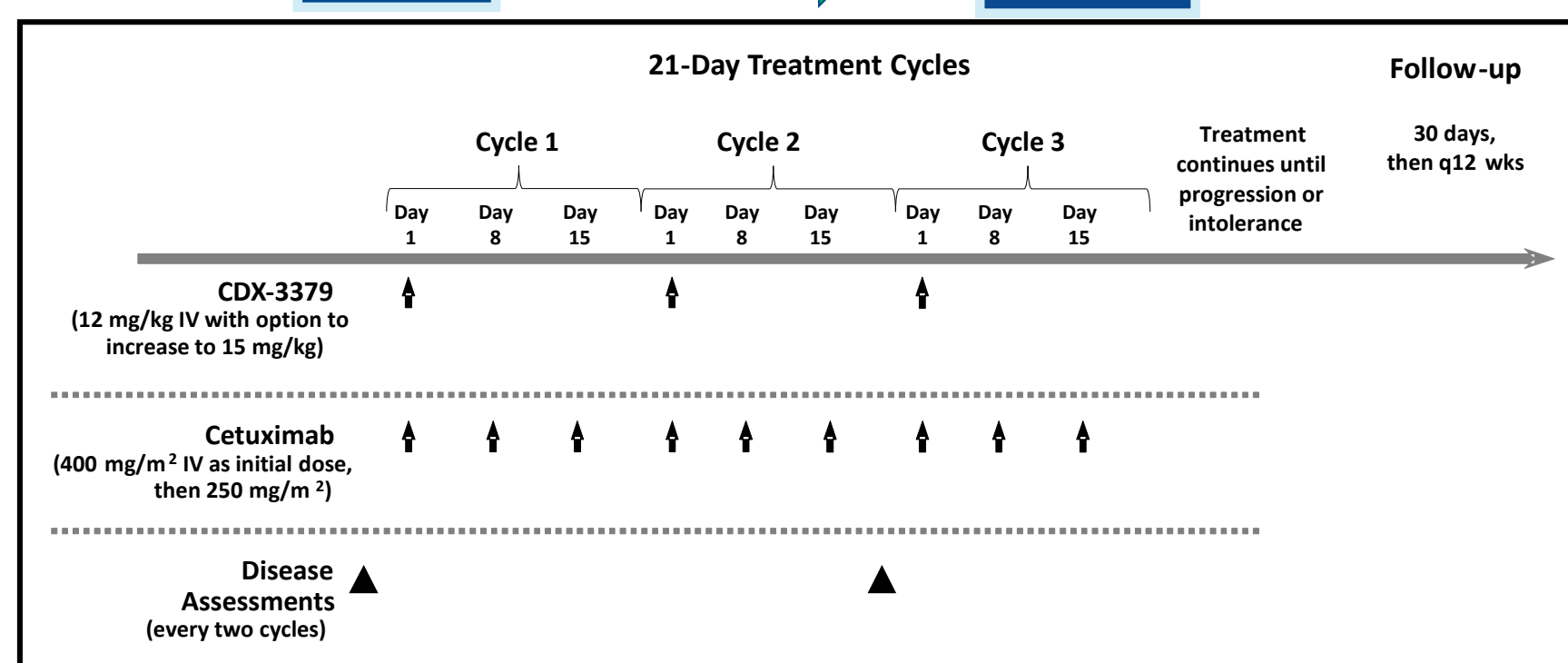
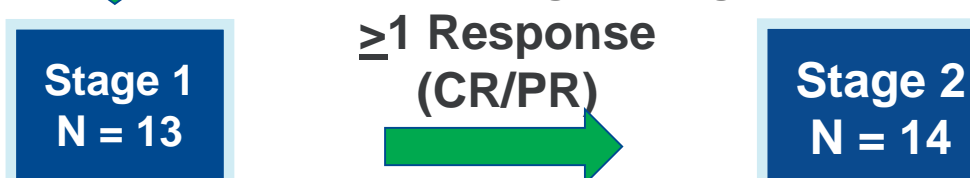
CDX3379-04 STUDY DESIGN

NCT03254927 is a phase 2, multicenter, open-label, single-arm clinical trial to evaluate safety and efficacy of CDX-3379 in combination with cetuximab in patients with advanced HNSCC

Key Eligibility Criteria

- Recurrent/metastatic HPV(-) HNSCC, not curable with local treatment (eg, surgery, radiation)
- Cetuximab resistance (progression within 6 months)
- Prior PD-1 targeted checkpoint inhibition, unless not a candidate
- RECIST 1.1 measurable disease
- ECOG 0 or 1; life expectancy \geq 12 weeks
- No active brain metastases
- No nasal, paranasal sinus, or nasopharyngeal WHO Type III carcinoma

Simon's 2-stage design



STUDY ASSESSMENTS

- Tumor response (MRI/CT): every 6 weeks during treatment
- Tumor biopsy: Screening, Cycle 2, and at progression
- Safety and toxicity assessments
- CDX-3379 pharmacokinetics and immunogenicity

STUDY OBJECTIVES

- Primary Objective:** Objective Response Rate (ORR): CR or PR (RECIST 1.1)
- Secondary Objectives:** Clinical benefit rate, duration of response, progression-free survival, overall survival, safety, pharmacokinetics, immunogenicity
- Exploratory Objectives:** Biomarker analysis

STAGE 1 RESULTS

Patient Demographics (N=15)

Median Age, years (range)	61 (46-68)
Male:	11 (73%)
Race:	White 13 (87%) Black 1 (7%) Asian 1 (7%)
ECOG at Baseline:	0 1 (7%) 1 14 (93%)
Smoking or Tobacco Use:	Prior Smoker 13 (87%)
Site of Primary Tumor:	Oral Cavity 8 (53%) Pharynx 5 (33%) Larynx 1 (7%) Other 1 (7%)
P16 Status:	Negative 7 (47%) (only required for pts with oropharyngeal tumors) Positive 1 (7%) Not Done 7 (47%)
# Prior Regimens, median (range)	3 (2-6)
Prior Treatment with Cetuximab	14* (93%)
Prior Checkpoint Inhibitor	15 (100%)

Exposure to Study Drug

CDX-3379 Median # doses (min, max)	3 (1, 16)
Cetuximab Median # doses (min, max)	6 (1, 46)
Duration of treatment	< 3 months 12 (80%) 3 - 6 months 2 (13%) \geq 12 months 1 (7%)
Dose Reductions, Delays and/or Holidays	10 (67%)

Reasons for Discontinuing Study Treatment	Disease Progression 7 (47%) Clinical Progression 4 (27%) Adverse Event 2 (13%) Death 1 (7%)
Ongoing Treatment	1 (7%)

Safety

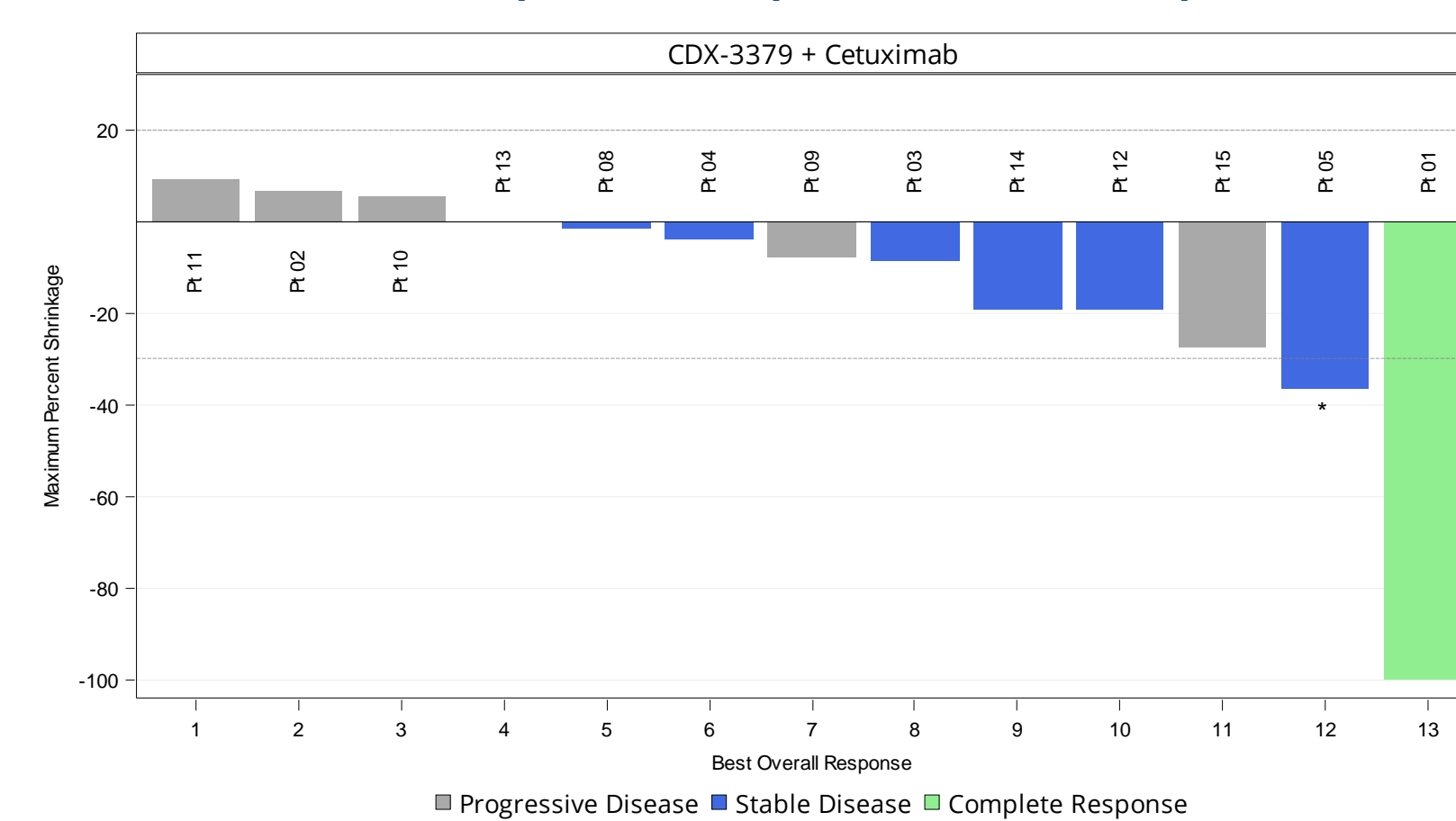
CDX-3379 Treatment-Related AEs (N=15) \geq 10% or \geq G3 in \geq 2 patients	Overall N (%)	Grade 3-5 N (%)
Diarrhea	10 (67)	4 (27)
Fatigue	2 (13)	-
Hypokalemia	4 (27)	3 (20)
Hypomagnesemia	3 (20)	2 (13)
Electrocardiogram QT prolonged	2 (13)	1 (7)
Mucositis	3 (20)	2 (13)
Palmar-plantar erythrodysesthesia syndrome	2 (13)	-
Rash	2 (13)	1 (7)

AEs, adverse events. Three patients experienced serious treatment-related AEs including diarrhea, QT prolongation and mucosal inflammation.

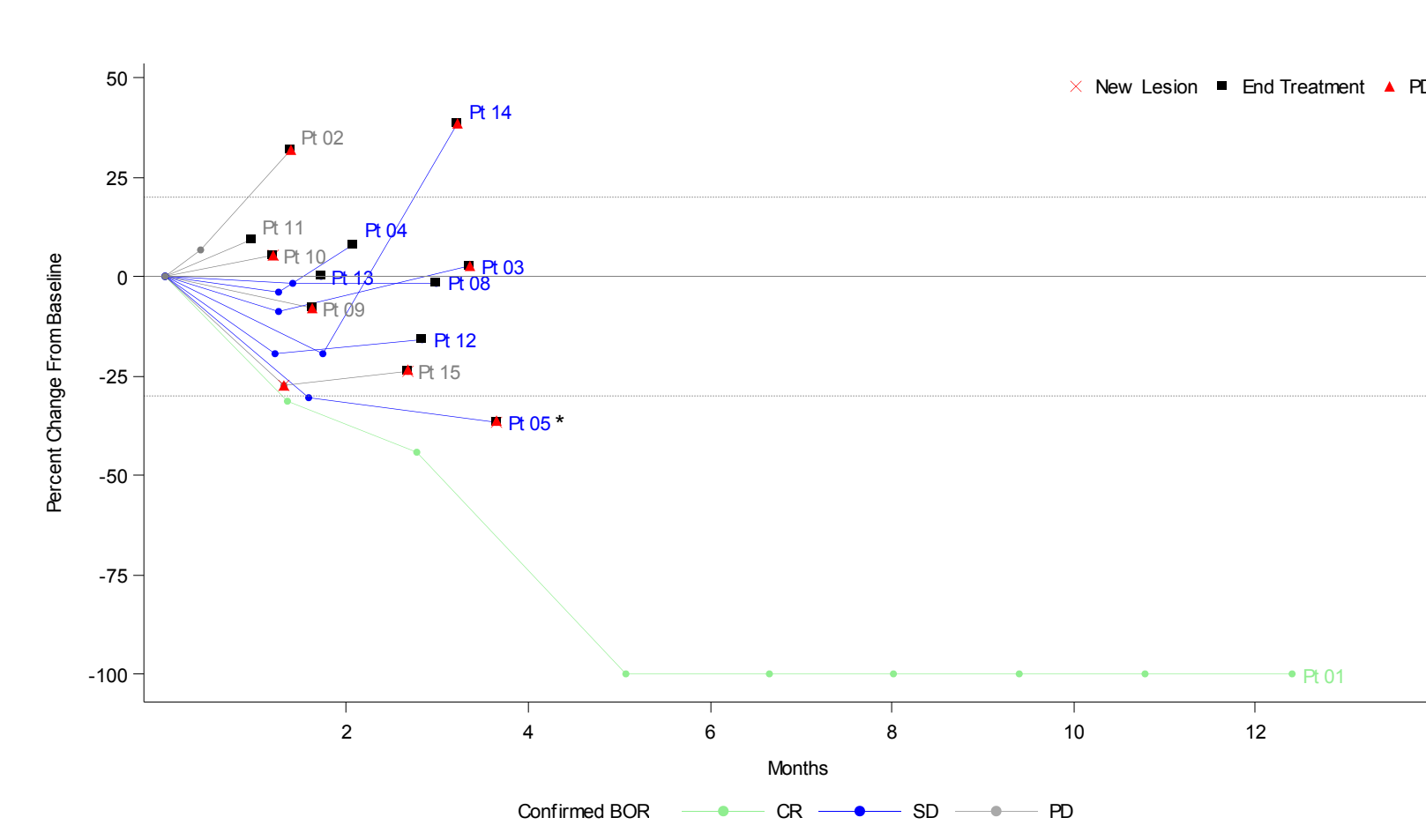
Efficacy

Best Overall Response (BOR) (N=15)	n (%)
Complete Response (CR)	1 (7%)
Partial Response (PR)	0 (0%)
Stable Disease (SD)	7 (47%)*
Progressive Disease (PD)	6 (40%)
Not Evaluable (NE)	1 (7%)
Clinical Benefit Rate (N=14)	
SD \geq 12 weeks	3 (21%)
Clinical Benefit Rate (Objective Response or SD \geq 12 wks)	4 (29%)

Best Overall Response – Response Evaluable Population**



Spider Plot of Tumor Burden – Response Evaluable Population**



* Includes one cetuximab-naïve patient, who experienced a single-timepoint PR
**Two patients did not have a post-baseline disease assessment

EXPLORATORY BIOMARKER ANALYSIS

- Next-generation sequencing was performed targeting >1400 cancer-related genes
- The analysis included 18 HNSCC tumor samples from three CDX-3379 clinical studies:
 - CDX3379-01 Phase 1b (CDX-3379 + cetuximab): one patient with cetuximab-refractory, HPV(-) HNSCC with a durable complete response
 - CDX3379-02 (CDX-3379 monotherapy; window of opportunity study): 3 patients, including a patient with an exceptional clinical response (92% tumor shrinkage assessed by physical examination after 2 doses of CDX-3379 every 2 weeks)³
 - CDX3379-04 (CDX-3379 + cetuximab): 14 patients, one ongoing, durable complete response (11+ months) and one unconfirmed partial response

Study	Responders																	
	3379-01 Pt A	3379-04 Pt 01	3379-04 Pt 05*	3379-02 Pt B	3379-04 Pt 04	3379-04 Pt 08	3379-04 Pt 12	3379-04 Pt 14	3379-04 Pt 03	3379-04 Pt 13	3379-04 Pt 02	3379-04 Pt 09	3379-04 Pt 10	3379-04 Pt 15	3379-04 Pt 11	3379-04 Pt 06	3379-02 Pt C	3379-02 Pt D
Primary Tumor	Oral Cavity	Oral Cavity	Oral Cavity	Oral Cavity	Oro-pharynx	Oral Cavity	Hypo-pharynx	Oral Cavity	Larynx	Oro-pharynx	Naso-pharynx	Oral Cavity	Oral Cavity	Oral Cavity	Hypo-pharynx	Oral Cavity	Oral Cavity	Pharynx
HPV Status	Negative	Negative	Not tested	Negative	Negative	Not tested	Not tested	Negative	Not tested	Negative	Not tested	Negative	Negative	Not tested	Negative	Positive	Negative	Positive
Best Response	CR	CR	uPR	Exceptional Response	SD (\geq 12 wks)	SD (\geq 12 wks)	SD (\geq 12 wks)	SD	SD	SD	PD	PD	PD	PD	PD	PD	0% Shrinkage	-16% Shrinkage
TP53																		
PIK3CA																		
FAT1																		
NOTCH1																		
NOTCH2																		
NOTCH3																		
NOTCH4																		

CR: complete response; uPR: unconfirmed partial response; SD: stable disease; PD: progressive disease
*Tumor samples from the CDX3379-01 and CDX3379-02 studies are italicized
†Per protocol, surgery was conducted following 2nd dose of CDX-3379

- FAT1 mutations were identified in all patients considered responders in this analysis
- NOTCH1, NOTCH2 or NOTCH3 mutations also appeared to be enriched in patients in whom clinical activity was observed
- Inactivating mutations in the FAT1 and NOTCH genes have been identified in 32% and 26% of HPV(-) HNSCC tumors, respectively⁴
- Loss of FAT1 function results in activation of the transcriptional cofactor YAP1⁵; YAP1 has been shown to upregulate components of ErbB signaling pathways^{6,7}, including the ErbB3 ligand NRG1⁸
- Preclinical studies investigating the association of CDX-3379 sensitivity and inactivating mutations of FAT1 and other genes are ongoing

CONCLUSIONS AND FUTURE DIRECTIONS

- Clinical activity has been observed in the CDX3379-04 study, including a durable complete response (11+ months) and tumor shrinkage, in patients with advanced, recurrent, HPV(-) HNSCC, where treatment options are limited
 - Clinical Benefit Rate of 29% was achieved
 - Dose reductions and/or delays to the combination therapy in the majority of patients may have impacted the magnitude of antitumor activity; dose modifications are being considered for future studies
 - CDX-3379 in combination with cetuximab was associated with the expected target-mediated adverse events of diarrhea and rash
- Across the CDX-3379 studies, mutations in FAT1 and NOTCH1, NOTCH2, or NOTCH3 and primary tumor site of oral cavity were associated with clinical activity in HNSCC
 - All 4 responding patients had FAT1 mutations and primary tumor site of oral cavity
 - Three of the responding patients also had NOTCH1, NOTCH2, or NOTCH3 mutations
- Future CDX-3379 development will focus on the clinical utility of biomarker-driven patient selection

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