# A Pivotal, Multicenter, Randomized Study Evaluating the Novel Antibody-Drug Conjugate Glembatumumab Vedotin (CDX-011; CR011-vcMMAE) in Patients with Metastatic, Triple-negative, gpNMB Over-expressing Breast Cancer

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# Glycoprotein NMB (gpNMB)

- An internalizable transmembrane glycoprotein over-expressed in ~40-60% of breast cancers as well as other tumors
- Expressed on epithelial tumor cells and supporting stromal cells
- Promotes migration, invasion, and metastases
- Shorter metastasis-free and overall survival have been noted in patients with high gpNMB-expressing tumors (including breast,<sup>1</sup> small cell lung cancer,<sup>2</sup> and glioblastoma<sup>3</sup>)

### Tumor Epithelial gpNMB Expression is Common in Triple Negative Breast Cancer (ER-/PR-/HER2-) and is Associated with Recurrence<sup>1</sup>



## Glembatumumab Vedotin

- Novel antibody-drug conjugate that delivers the potent cellular toxin monomethylauristatin E (MMAE) to gpNMB-expressing tumor cells
- Fully human gpNMB-specific monoclonal antibody (CR-011) conjugated to MMAE via a protease-sensitive peptide linker, designed to cleave upon cellular internalization
- Same linker-MMAE technology as that used successfully in Adcetris<sup>™</sup> (brentuximab vedotin; Seattle Genetics)



- Three completed clinical studies in melanoma and breast cancer
- MTD (Phase II dose) established as 1.88 mg/kg q3w
- Favorable ORR and PFS in heavily-pretreated populations
- Clinical benefit appears greatest in patients with triple-negative tumors and/or tumors expressing higher levels of gpNMB



1. Rose, et al. CCR, 2010 2. Li, et al. APMIS 2013 3. Kuan, et al. CCR 2006

# BACKGROUND





<b>Completed Phase II Study in Patients wi</b>	th Advanced Breast Car	ncer: "EMERGE"		
Study designed to examine whether anti-cancer	<ul> <li>Population:</li> </ul>			
activity of glembatumumab vedotin is dependent upon distribution/intensity of gpNMB expression	<ul> <li>gpNMB+ breast cancer (≥5 cells positive by centralized</li> </ul>	% of epithelial or stroma I IHC)	Triple-negative breast cancer (TN overexpressing gpNMB *	
<ul> <li>Treatments (2:1 randomization):</li> <li>Glembatumumab vedotin (1.88 mg/kg q3w)</li> <li>"Investigator's Choice" (IC) single-agent</li> </ul>	<ul> <li>Refractory/resistant to approved therapies (taxane, anthracycline, capecitabine; and if HER2+, trastuzumab and lapatinib)</li> <li>Progression within 6 months of last regimen</li> </ul>		N=300	
<ul> <li>chemotherapy</li> <li>Cross-over from IC to glembatumumab vedotin</li> <li>permitted at progression</li> </ul>	- 98% with metastatic disease		* Central analysis of tumor obtained after	
	- Median of 6 prior lines of anticancer therapy (4 lines		of locally advanced/metastatic disease	
Tissue Screening for gpNMB Expression	of cytotoxic therapy for adv	anced disease)	<ul> <li>gpNMB overexpression: ≥ 25% of tumo epithelial cells positive by IHC</li> </ul>	
100%			- 40% of INBC expected to meet crite	
90% 80%	All Screened Breast Cancer Patients		(EIMERGE experience)	
70% 60% 50% 40% 20% 10% 0%	<ul> <li>99% of screened patients gpNMB expression</li> <li>Epithelial gpNMB expression</li> </ul>	s met eligibility for sion was more frequent	<ul> <li>TNBC (centrally confirmed):</li> <li>ER/PR: &lt; 1% of cells positive by IHC</li> <li>HER2: IHC staining of 0 or 1+, FISH copies or ratio &lt; 1.8</li> </ul>	
25% 210% 225% 25% 210% 225% % gpNMB Expressing % gpNMB Expressing Tumor cells Stromal cells				
	S	afoty	Additional Eligibility Criteria	
Response to Glemabatumumab Vedotin Glembatumumab Vedotin	, but not to IC stigator's Choice	Glembatumumab vedotin well-tolerated	<ul> <li>0 or 1 chemotherapy-containing regime advanced breast cancer</li> </ul>	
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	<ul> <li>Below threshold</li> <li>Above threshold</li> <li>Above threshold</li> <li>Treatment-related toxicity: Rash, neutropenia, fatigue, nausea vomiting</li> </ul>		<ul> <li>Progression/recurrence-free interval &gt; completion of neoadjuvant or adjuvant chemotherapy</li> </ul>	
10%     10%     10%     10%     10%     10%	p = 0.7 p = 0.9	alopecia, decreased appetite and peripheral neuropathy	<ul> <li>Resistant to taxane therapy</li> <li>If progression-free interval &gt;12 montonic or adjuvant taxane, must receive tax</li> </ul>	
<b>Bai</b> 5% % %		Less hematologic	advanced disease	
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	241228810%25%Expression Thresholdial cells expressing gpNMB by IHC)	toxicity than Investigator's Choice	<ul> <li>If taxane administered for advanced demonstrate progression during or v months of completing therapy</li> </ul>	
Activity: Tumor Response, Progression-Fr	ee Survival (PFS) and Over	all Survival (OS)	<ul> <li>Receipt of anthracycline-containing che any setting, with no further anthracyclin indicated</li> </ul>	
All Patients Triple-Neg	ative High gpNMB Expression <sup>§</sup>	Triple Negative & High gpNMB <sup>§</sup>	- i.e., contraindicating cardiac condition	
GV (n=96) IC (n=41) GV (n=31) I	C (n=11) GV (n=27) IC (n=11)	GV (n=12) IC (n=6)	intolerance, or cumulative dose ≥ 24 doxorubicin or equivalent	
Partial Response (PR) 13 (16%) 5 (14%) 5 (19%)	0 (0%) 8 (32%) 1 (13%)	4 (33%) 0 (0%)	Maggurable disease by DECIST 1.1 or	
Confirmed PR 8 (10%) 3 (8%) 2 (7%)	0 (0%) 4 (16%) 1 (13%)	1 (8%) 0 (0%)		
Stable Disease or Better 46 (57%) 19 (53%) 18 (67%)	3 (33%) 16 (64%) 3 (38%)	9 (75%) 1 (25%)	<ul> <li>ECOG Performance Status 0 or 1</li> </ul>	
Median PFS (months) 2.1 2.0 2.3	1.6 2.7 1.5	3.0 1.5	<ul> <li>Neuropathy Grade ≤ 1</li> </ul>	
p=0.38 p=0.43 p=0.14 p=0.008*		p=0.008*	No investigational therapy within 4 week	
Median OS (months) 7.5 7.4 6.9	6.5 10.0 5.7	10.0 5.5	treatment	
p=0.24 p=0.3	0 p=0.18	p=0.003*		

GV, Glembatumumab vedotin; IC, Investigator's Choice Therapy.  $\S \ge 25\%$  of tumor epithelial cells expressing gpNMB by IHC \* Statistically significant Patients who received Investigator's Choice and subsequently crossed over to receive glembatumumab vedotin (n=15) are included in the tumor response and PFS analyses for each treatment received, but are assigned to the Investigator's Choice arm only for OS analysis. Analysis of best response (RECIST 1.1) excludes patients who discontinued from study without evaluable post-baseline radiographic imaging (n=15 for GV arm; n=5 for IC arm).





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• Type I error rate ( $\alpha$ ) of 0.05 allocated between the co-primary endpoints

- Hypothesized HR = 0.64- Hypothesized median PFS: 4 months for capecitabine 6.25 months for glembatumumab vedotin



