

## A Phase 1 Evaluation of Tebotelimab, a Bispecific PD-1 x LAG-3 DART® Molecule, in Combination with Margetuximab in Patients with Advanced HER2+ Neoplasms

Patel MR¹, Luke JJ², Hamilton E³, Chmielowski B⁴, Blumenschein G⁵, Kindler H⁶, Bahadur S⁷, Sun J⁶, Chen F⁶, Zhang X⁶, Muth J⁶, Kaminker P⁶, Moore PA⁶, Sumrow BJ⁶, Ulahannan SV¹⁰

¹Florida Cancer Specialists/Sarah Cannon Research Institute, Sarasota, FL; ²UPMC Hillman Cancer Center, University of California Los Angeles, Los Angeles, CA; ¹Sarah Cannon Research Institute, Sarasota, FL; ²UPMC Hillman Cancer Center, University of California Los Angeles, Los Angeles, CA; ¹Sarah Cannon Research Institute, Sarasota, FL; ²UPMC Hillman Cancer Center, University of California Los Angeles, Los Angeles, CA; ¹Sarah Cannon Research Institute, Sarasota, FL; ²UPMC Hillman Cancer Center, University of California Los Angeles, Los Angeles, CA; ¹Sarah Cannon Research Institute, Sarasota, FL; ²UPMC Hillman Cancer Center, University of California Los Angeles, Los Angeles, CA; ¹Sarah Cannon Research Institute, Sarasota, FL; ²UPMC Hillman Cancer Center, University of California Los Angeles, Los Angeles, CA; ¹Sarah Cannon Research Institute, Sarasota, FL; ²UPMC Hillman Cancer Center, University of California Los Angeles, Los Angeles, CA; ¹Sarah Cannon Research Institute, Sarasota, FL; ²Sarah Cannon Research Institute, Sarasota, Sarasota, FL; ²Sarah Cannon Research Institute, Sarasota, Sar <sup>5</sup>Department of Thoracic Head & Neck Medical Oncology, Division of Cancer Medicine, WD Anderson Cancer Center, Houston, TX; <sup>6</sup>Division of Hematology/Oncology, Department of Medicine, University of Chicago, Ch 8Sidney Kimmel Comprehensive Cancer Center, Johns Hopkins School of Medicine, Baltimore, MD; 9MacroGenics, Inc., Rockville, MD; 10SCRI Nashville/OUHSC, Oklahoma City, OK

#### Background

#### **Tebotelimab (MGD013)**

- PD-1 and LAG-3 receptors are expressed on "exhausted" T cells
- Interactions with corresponding ligands negates anti-tumor T-cell activity
- Tebotelimab, an investigational DART protein, targets PD-1 and LAG-3 with a single molecule
- with IgG4 Fc - Greater synergistic T-cell activation (IFN-y) in vitro with tebotelimab compared with combination of individual constituents

Tetravalent (bivalent for each target) structure

- Ongoing tebotelimab Phase 1 study demonstrated safety up to 1200 mg Q2W, with evidence of monotherapy antitumor activity in various
- advanced solid tumors<sup>1</sup> Objective responses associated with increased baseline LAG-3 expression and IFN-y gene signature

# PD-1 x LAG-3 **Tetravalent Bispecific DART Molecule**

Fab region

Fc region

**Tebotelimab** 

#### Margetuximab

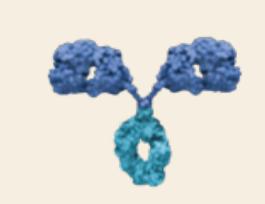
#### Investigational Fc-engineered anti-HER2 mAb

 Similar anti-HER2 properties as trastuzumab Enhanced Fc-mediated effector function<sup>2</sup>

Monotherapy RP2D defined as 600 mg

- Superior PFS to trastuzumab in Phase 3 SOPHIA study in mBC<sup>3</sup>
- Anti-tumor activity in advanced gastric cancer In combination with anti-PD-1<sup>4</sup>

#### Trastuzumab



- Binds HER2 with high specificity
- Disrupts signaling that drives cell proliferation and survival
- Wild-type immunoglobulin G1 (IgG1) immune effector domains Binds and activates immune cells

#### Margetuximab<sup>5,6</sup>



- Same specificity and affinity Similarly disrupts signaling
- ↑ Affinity for activating FcyRIIIA Activating - ↓ Affinity for inhibitory FcyRIIB

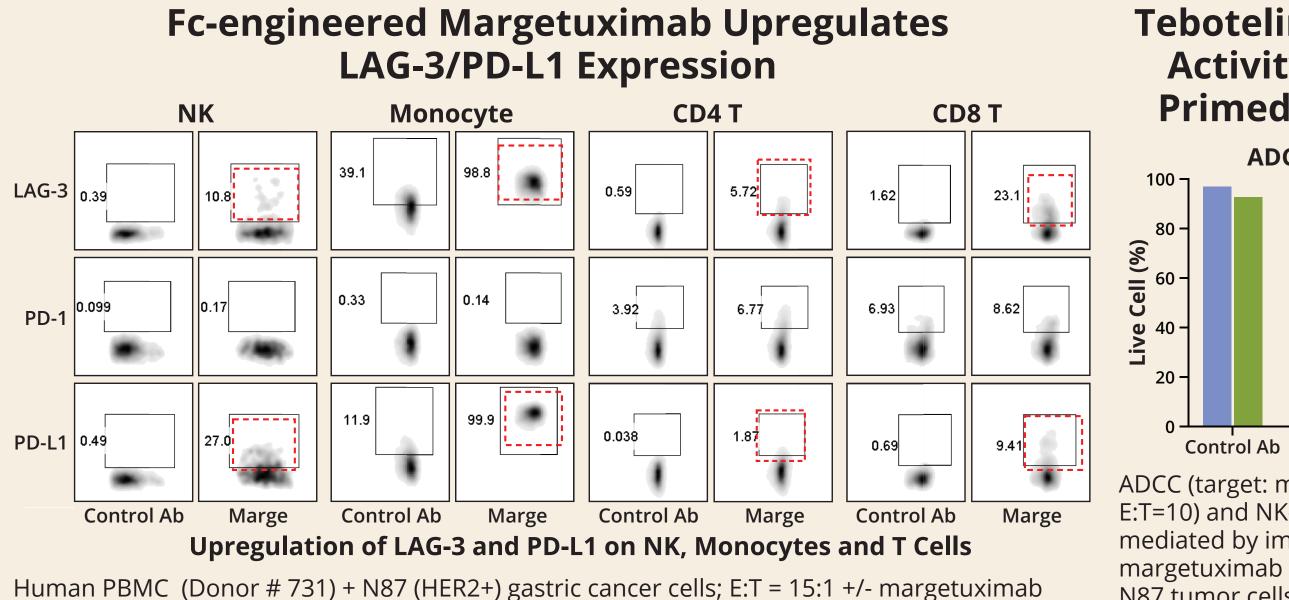
# • Fc engineering:

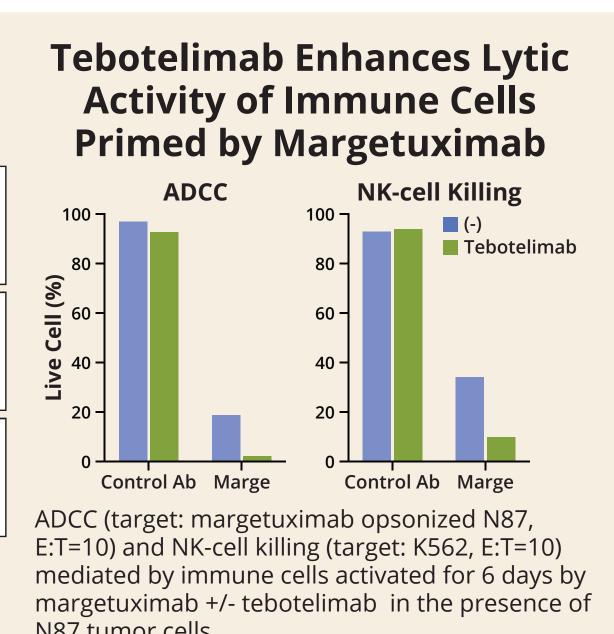
# **Margetuximab Binding to FcyR Variants:**

#### CD16A Genotype May Predict Anti-HER2 Antibody Benefit

- Improved responses to trastuzumab observed in patients with HER2+ breast cancer<sup>7</sup> or gastric cancer<sup>8</sup> who are homozygous for CD16A-158V (V/V), which encodes the higher-affinity Fcy receptor when compared to CD16A-158F carriers (V/F, F/F)
- V/V present in approximately 15% of Caucasian, African American, and Asian populations
- Lower affinity F/F and intermediate affinity V/F carriers comprise the remainder of the population (~85%) Hypothesis: Greater margetuximab benefit in lower-affinity CD16A-158F carriers (e.g. V/F, F/F) - Increased affinity of margetuximab for CD16A-158F over trastuzumab (wild-type IgG1)

#### Tebotelimab Plus Margetuximab: Combinatorial Biology





#### Phase 1 Study Design **Cohort Expansion (Ongoing)** 3 + 3 Dose Escalation (Complete) (Advanced HER2+ solid tumors) **HER2+ Breast Cancer** Tebotelimab 600 mg Margetuximab 15 mg/kg HER2+ GC/GEJ Cancer Tebotelimab 300 mg Margetuximab 15 mg/kg Other HER2+ Cancers

#### **Study Objectives**

Both study drugs administered Q3W.

### Primary

- Characterize safety of combination in patients with HER2+ locally advanced or metastatic cancers Secondary
- Characterize PK and immunogenicity of combination administered Q3W
- Assess preliminary antitumor activity of combination using RECIST v1.1 and irRECIST

#### Exploratory

- Explore relationships between PK/PD, patient safety, and antitumor activity of combination Determine relationship between clinical response and expression of PD-L1, HER2, and/or LAG-3 on tumor cells and immune cell infiltrates
- Investigate immune-regulatory activity of combination in vivo, including various measures of T-cell activation in peripheral blood and/or tumor biopsy specimens
- Explore relationships among Fcy receptor allelic variation in CD16A, CD32A, and CD32B and clinical response

#### **Key Entry Criteria**

Age ≥ 18 years

Data cut-off: 05Oct2020

- ECOG performance status 0 or 1
- Life expectancy ≥ 12 weeks
- Measurable disease per RECIST v1.1 Acceptable laboratory parameters
- Cancer must have progressed following standard therapy, or during/after HER2-directed therapy if approved and available for patients with HER2+ gastric, GEJ, or breast cancer
- History of HER2 positivity defined as (1) IHC 3+ or IHC 2+ with ISH positivity or (2) HER2 amplification by NGS

#### Patient Demographics and Disposition

#### Demographics **HER2+ Tumor Types Treated** Salivary gland, Cholangiocarcinoma 63 (29, 86) Median age (range), years Gender, n (%) Endometrial, 4 17 (41.5) 24 (58.5) Female ECOG PS, n (%) 19 (46.3) 22 (53.7) Esophageal, 4 **Median prior lines of therapy (range) Prior Checkpoint Inhibitor** 9 (22.0) 32 (78.0) Ovarian, 5 Colorectal, 6 **Prior HER2-directed therapy** 28 (68.3) 13 (31.7)

End of Treatment Disposition			
	Tebotelimab + Margetuximab		
Patients treated	41		
Response-evaluable patients	28 (68.3)		
Median duration of therapy, weeks (min, max)	9.8 (1.7, 64.4)		
Continuing treatment, n (%)	11 (26.2)		
Treatment discontinuation, n (%)	30 (73.2)		
Reasons for discontinuation, n (%) Disease progression Adverse event Patient/physician decision/withdrawal Unknown	21 (51.2) 6 (14.3) 2 (4.9) 1 (2.4)		

### **Preliminary Evidence of Anti-tumor Activity** Cholangiocarcinom Esophageal adenocarcinoma Cervical cancer Tebotelimab 300 mg + margetuximab 15 mg/kg Q3W Tebotelimab 600 mg + margetuximab 15 mg/kg Q3W + Prior CPI therapy \* Ongoing # Prior HER2-directed therapy D-L1 Combined Positive Score (CPS) calculated as follows: Number of PD-L1+ cells (tumor cells, lymphocytes and macrophages)/total number of Breast cancer Endometrial cancer Cholangiocarcinoma Esophageal adenocarcinoma Cervical cancer Urethral carcinoma First new lesion Treatment ongoing **Weeks Since Treatment Initiation**

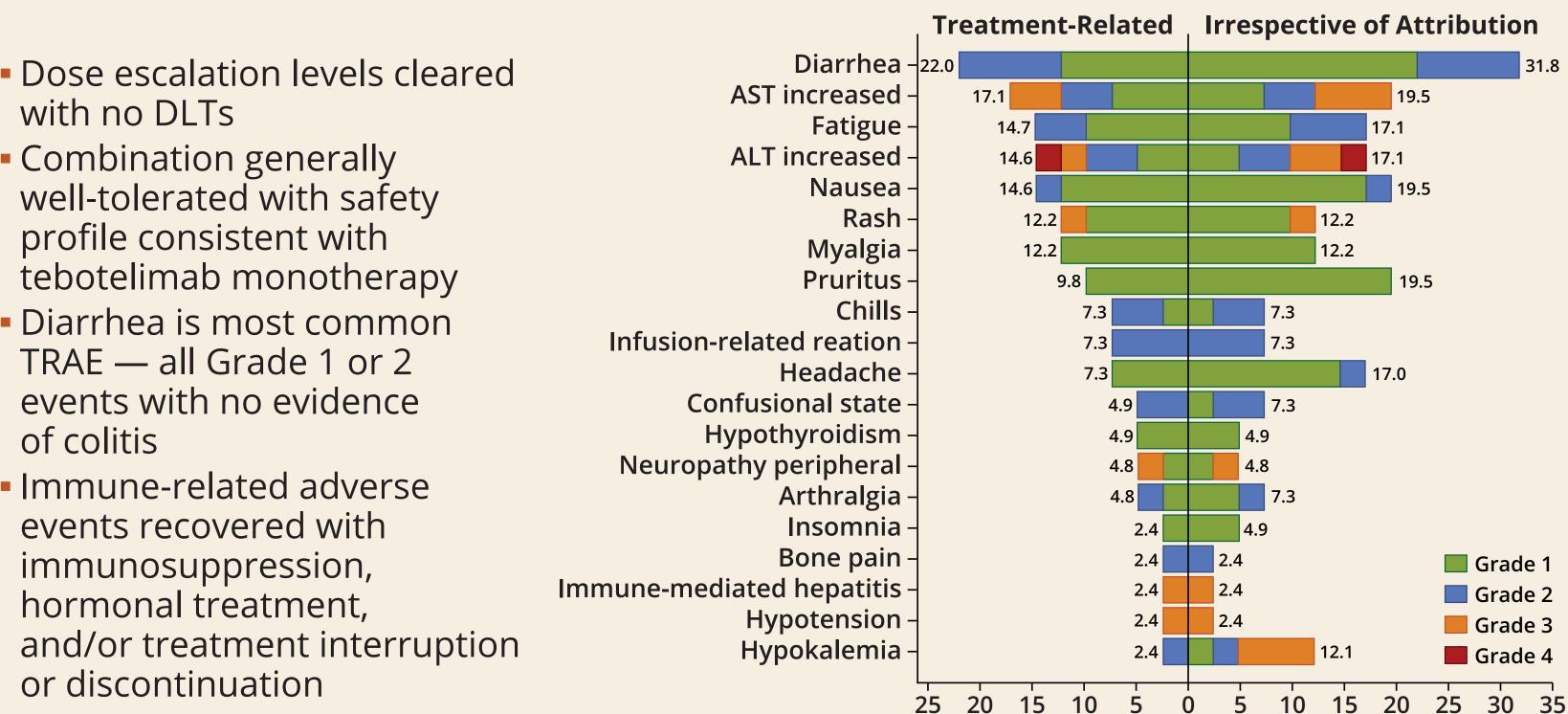
## **Safety Overview**

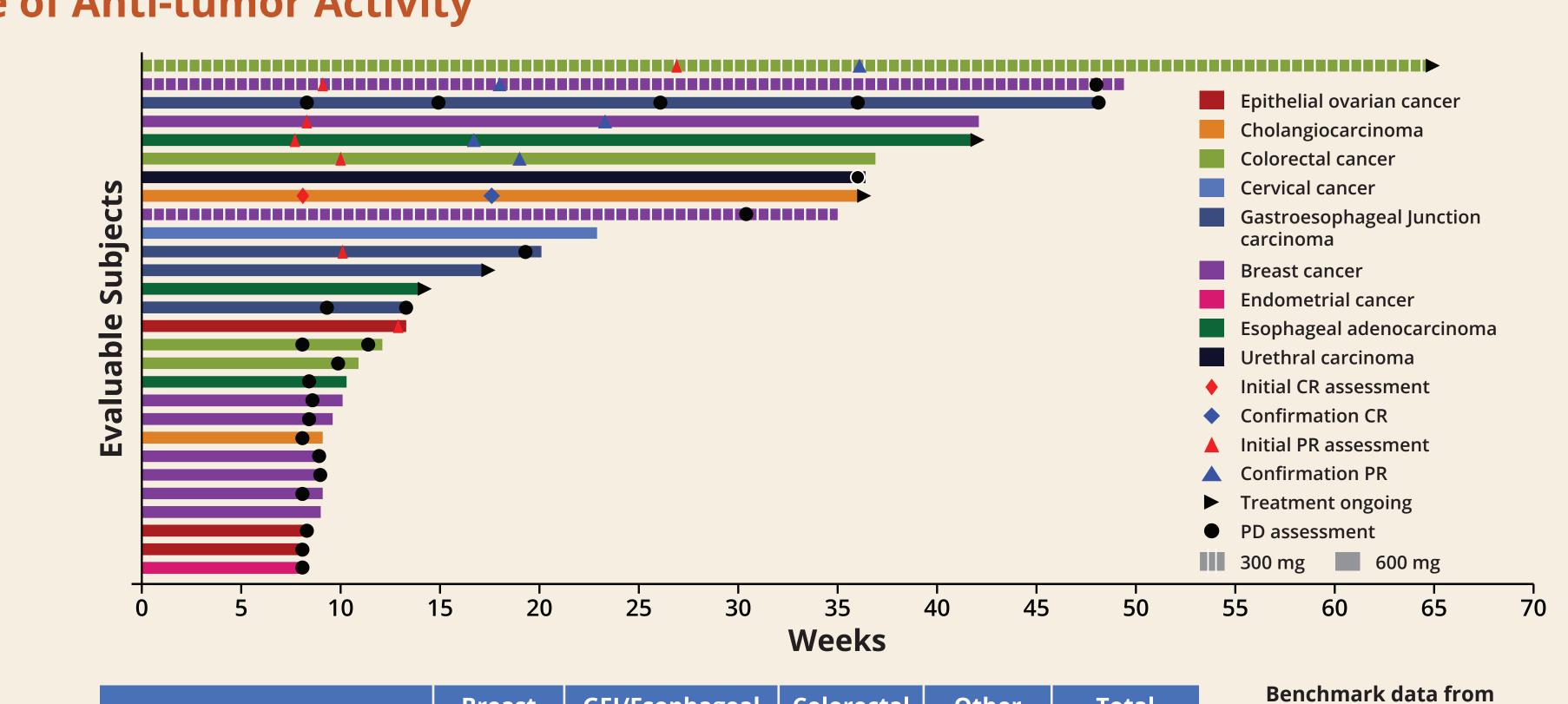
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Overall AE Totals	All Grades (N=41)	≥ Grade 3 (N=41)	
AE (irrespective of causality)	37 (90.2)	17 (41.5)	
Treatment-related AE	30 (73.2)	8 (19.5)	
SAE (irrespective of causality)	7 (17.1)	6 (14.6)	
Treatment-related SAE	2 (4.9)	2 (4.9)	
AE leading to treatment discontinuation	6 (14.6)	4 (9.8)	
Adverse Events of Special Interest (AESIs)			
Infusion-related reaction	3 (7.3)	0 (0)	
Arthralgia	2 (4.9)	0 (0)	
Immune-mediated hepatitis	1 (2.4)	1 (2.4)	
Left ventricular dysfunction	1 (2.4)	1 (2.4)	
Pancreatitis	1 (2.4)	1 (2.4)	
Hyperthyroidism	1(2.4)	0 (0)	
	Adverse Events  Treatment-Related   Irrespective of Attribution		
<ul> <li>Dose escalation levels cleared with no DLTs</li> </ul>	Diarrhea - 22.0  AST increased - 17.1  Fatigue - 14.7	19.5 17.1	

No. (%) of Patients

Percentage of Patients with Treatment-Related

**AEs and AEs Irrespective of Attribution** 





**Preliminary Results** 

Evaluable patients

Disease control rate

ORR (confirmed)

Association of Baseline LAG-3/PD-1 Expression with Clinical Response				
Dual <i>LAG3/PDCD1</i> Expression at Baseline Associates with	baseline LAG-3 and PD-1	Levels Inversely Correlate		
Objective Response with Best % Change in Target Lesions				
100 CR/PR SD	100 - CR/PR - SD	100 - CR/PR - SD		
80 - PD	80 - PD	80 - PD		
<b>ESH</b> 40-	F 60-	60 - 40 -		
3 40				
20 –	Correlation Pearson r=-0.6333	Correlation Pearson r=-0.3815		
0 20 40 60 80 100	-80 -60 -40 -20 0 20 40 60	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$		
PDCD1	% Change in Target Lesion(s)	% Change in Target Lesion(s)		

- Patients demonstrating objective responses exhibit higher expression of both LAG-3 and PD-1 (PDCD1) mRNA in baseline biopsy samples Expression of LAG3 and PDCD1 mRNA in baseline tumor biopsy inversely correlates with best change in
- Highest correlation observed with LAG-3 expression •ROC analyses of LAG-3 expression and objective responses (confirmed and unconfirmed) indicate a 75%
- response rate in LAG-3 biomarker high patients vs 9% in biomarker low patients Further analyses ongoing and will be extended to additional patients
- to further define potential enrichment biomarker component(s)
- The NanoString PanCancer IO 360™ assay was used to interrogate gene expression, including the abundance of 14 immune cell types and 32 immuno-oncology signatures, from archival biopsies of 19 HER2+ advanced solid tumor cohorts treated with margetuximab and tebotelimab. **Left:** Normalized expression scores (standardized 0-100) for *LAG3* were plotted against *PDCD1*. **Right:** Correlation of standardized *LAG3* and *PDCD1* expression levels to best percent change in tumor lesions from baseline, respectively. Receiver Operating Characteristic (ROC) analyses were performed on the 19 patient data set using Youden Index and Distance methods.

#### Conclusions

- Tebotelimab + margetuximab combination generally well tolerated
- Safety profile consistent with tebotelimab monotherapy
- Evidence of antitumor activity observed among refractory patients with various HER2+ tumor types - Objective responses (n=8; 6 confirmed) observed in multiple advanced HER2+ tumor types
- ORR (including unconfirmed responses) = 28.6% (Compares favorably to benchmark mBC data from PANACEA study)
- 64.3% of response-evaluable population with decrease of target lesion tumor burden - Duration of response (n=6 confirmed responders): 4.21–8.97 months (3 patients ongoing)
- Majority of responding patients with baseline PD-L1 expression of ≤ 1
- All responding patients carry less favorable CD16A-158F allotype (i.e. V/F or F/F)
- Baseline LAG-3 and PD-1 mRNA expression associated with clinical response Analyses ongoing to define patient enrichment biomarker
- Enrollment in HER2+ tumor-specific cohorts ongoing

#### References

1. Luke JJ, et al. ASCO 2020. 2. Nordstrom, et al., 2011 Breast Cancer Research, 13: R123. 3. Rugo, et al., ASCO 2019, Chicago, IL. 4. Catenacci, et al., ASCO GI 2019, San Francisco, CA / Catenacci et al. 2020 Lancet Oncology, in press. 5. Nordstrom JL, et al. Breast Cancer Res. 2011;13(6):R123. 6. Stavenhagen JB, et al. Cancer Res. 2007;67(18):8882-8890. 7. Musolino A, et al. J Clin Oncol. 2008;26(11):1789-1796. 8. Wang DS, et al. Onco Targets Ther. 2017;10:5065-5076.