

# Safety Outcomes by *UGT1A1* Status in the Phase 3 TROPiCS-02 Study of Sacituzumab Govitecan in HR+/HER2- Metastatic Breast Cancer

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## CONCLUSIONS

SG had a manageable safety profile (consistent with previous reports<sup>5,11</sup>) regardless of *UGT1A1* status (including neutropenia, diarrhea, and febrile neutropenia, which typically resolved within 10 days) in patients with HR+/HER2- mBC

Patients who were homozygous for *UGT1A1* \*28/\*28 experienced numerically higher rates of grade ≥ 3 TEAEs, TEAEs leading to discontinuation, any-grade anemia, and grade ≥ 3 diarrhea and neutropenia, although sample sizes were small

Because patients treated with SG are closely monitored for adverse events according to the prescribing information and per standard practice regardless of *UGT1A1* genotype, *UGT1A1* testing is not needed for SG use in pretreated HR+/HER2- mBC

Active monitoring and early intervention with routine strategies for AE management (and G-CSF treatment) is recommended for all patients being treated with SG



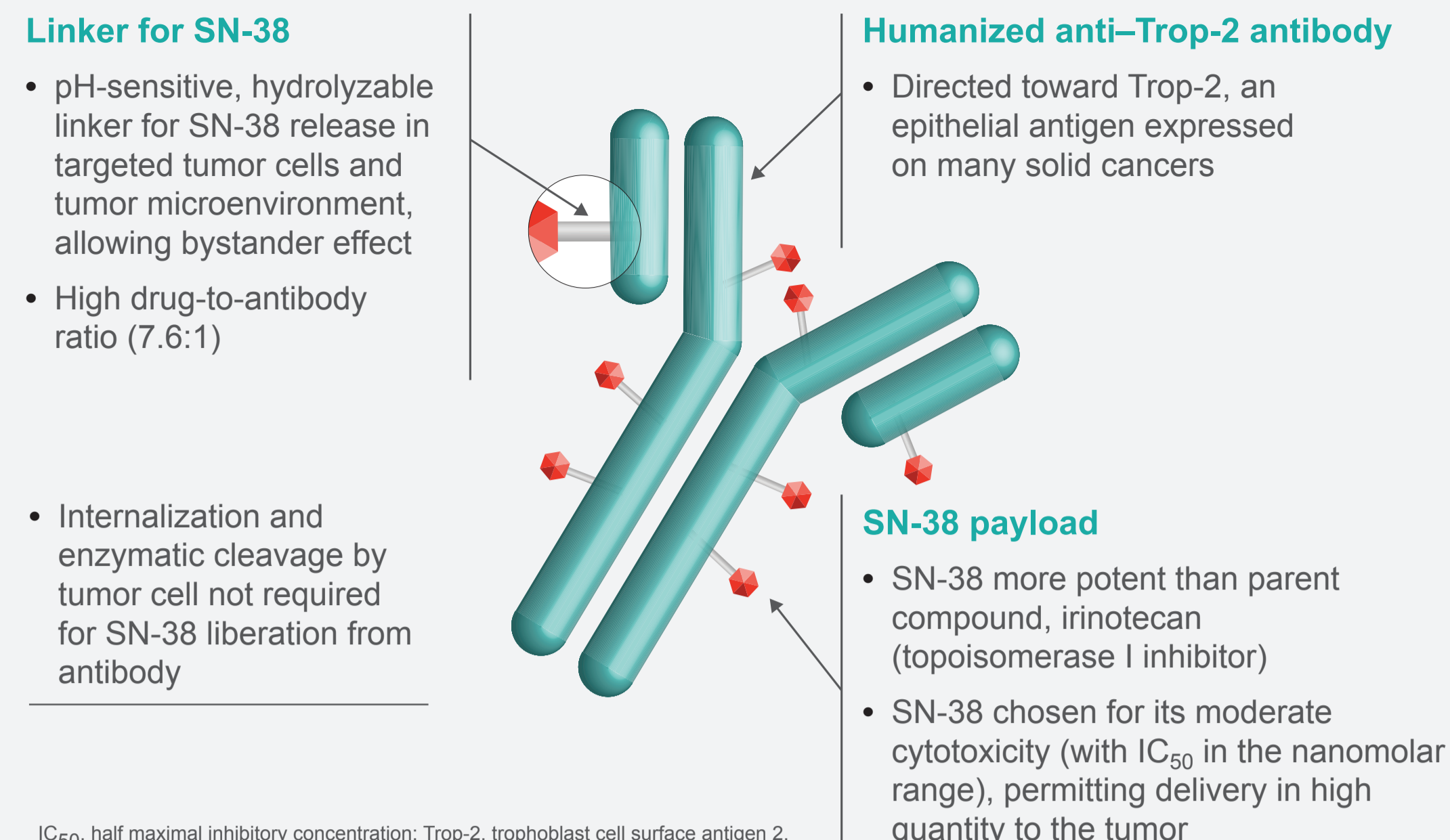
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## BACKGROUND

- Breast cancer is the second leading cause of cancer death in women,<sup>1</sup> and hormone receptor-positive (HR+)/human epidermal growth factor receptor 2-negative (HER2-) cancers make up approximately 70% of breast cancers<sup>2</sup>
- Sacituzumab govitecan (SG) is a trophoblast cell surface antigen 2 (Trop-2)-directed antibody-drug conjugate (ADC) (Figure 1) approved for triple-negative breast cancer (TNBC) in multiple countries and HR+/HER2- metastatic breast cancer (mBC) in the US<sup>3,4</sup>
- In the phase 3 randomized TROPiCS-02 study, SG versus treatment of physician's choice (TPC) demonstrated significantly improved median progression-free survival (PFS; 5.5 vs 4.0 months; hazard ratio [HR], 0.66; P = .0003) and median overall survival (OS; 14.4 vs 11.2 mo; HR, 0.79; P = .020), with a manageable safety profile in patients with pretreated, endocrine-resistant HR+/HER2- mBC<sup>5</sup>
- Polymorphisms in uridine diphosphate glucuronosyltransferase 1A1 (*UGT1A1*) have been associated with increased incidence of known adverse events, such as neutropenia, febrile neutropenia, anemia, and diarrhea, following treatment with some systemic anticancer agents, including SG, irinotecan, pazopanib, sunitinib, and nilotinib due to reduced rate of SN-38 glucuronidation<sup>6-10</sup>
- We present safety analyses of SG versus TPC by *UGT1A1* genotype status from TROPiCS-02

Figure 1. Sacituzumab govitecan antibody-drug conjugate



## Results

### Patients

- Baseline characteristics were balanced between the treatment arms (Table 1)
- Of 543 patients enrolled, 517 were included in the safety population (SG, n = 268; TPC, n = 249)
  - 9 patients (3%) in the SG arm and 2 (1%) in the TPC arm remained on treatment at data cutoff
  - The most common reason for treatment discontinuation was progressive disease (SG, 80%; TPC, 73%)

Table 1. Demographics and baseline characteristics

	SG (n = 272)	TPC (n = 271)
<b>Female, n (%)</b>	270 (99)	268 (99)
<b>Median age at study entry (range), y</b>	57.0 (29-86)	55.0 (27-78)
<b>Race,<sup>a</sup> n (%)</b>		
White	184 (68)	178 (66)
Asian	11 (4)	5 (2)
Black	8 (3)	13 (5)
<b>ECOG performance status, n (%)</b>		
0	115 (42)	126 (46)
1	157 (58)	145 (54)
<b>Metastatic disease, n (%)</b>	262 (96)	264 (97)
<b>Number of prior chemotherapies, n (%)</b>		
2-3	127 (47)	119 (44)
> 3	144 (53)	152 (56)
<b>Median prior systemic regimens,<sup>b</sup> n (range)</b>	7.0 (3-17)	7.0 (3-16)
<b>Setting of prior systemic therapies,<sup>c</sup> n (%)</b>		
Adjuvant	186 (68)	206 (76)
Neoadjuvant	67 (25)	62 (23)
Metastatic	272 (100)	271 (100)
Other	8 (3)	7 (3)
<b>BRCA1/2 mutational status, n (%)</b>		
Negative	109 (40)	114 (42)
Positive	21 (8)	11 (4)
Unknown	142 (52)	146 (54)

BRCA, breast cancer gene; ECOG, Eastern Cooperative Oncology Group; SG, sacituzumab govitecan; TNBC, triple-negative breast cancer; TPC, treatment of physician's choice.  
<sup>a</sup>Race was not reported for 69 (25%) patients in the SG arm and 70 (26%) patients in the TPC arm, and 5 (2%) patients in the TPC arm were reported as other.  
<sup>b</sup>Anticancer regimens refer to any treatment regimen that was used to treat breast cancer in any setting.  
<sup>c</sup>Patients may have received prior systemic therapy in more than 1 setting.

- Of patients treated with SG, 38% had wild-type, 44% had heterozygous, and 9% had homozygous *UGT1A1* genotypes (Table 2)
- The number of patients with a homozygous genotype was low, potentially limiting interpretation of these analyses

Table 2. Summary of *UGT1A1* status

	SG (n = 268)		
	*1/*1 Wild-type (n = 103)	*1/*28 Heterozygous (n = 119)	*28/*28 Homozygous (n = 25)
<b>Total, n (%)<sup>b</sup></b>	103 (38)	119 (44)	25 (9)
White	71 (39)	82 (45)	16 (9)
Asian	7 (64)	2 (18)	0
Black	2 (29)	3 (43)	1 (14)

SG, sacituzumab govitecan; TEAE, treatment-emergent adverse event.  
<sup>a</sup>Patients (1%) treated with SG had other genotypes, one each of \*1\*36, \*1\*37, and \*28\*36; 2 of these patients were white, and 1 did not have race reported.  
<sup>b</sup>Racial subgroups were calculated as percentage of total racial group in each genotype category.

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## ACKNOWLEDGMENTS

- We thank the patients and their caregivers for helping us realize the possibilities of this research
- We thank the dedicated clinical trial investigators and their devoted team members for participating in TROPiCS-02
- This study is sponsored by Gilead Sciences, Inc.
- Medical writing and editorial support was provided by Ben Labbe, PhD of Parexel and funded by Gilead Sciences, Inc.

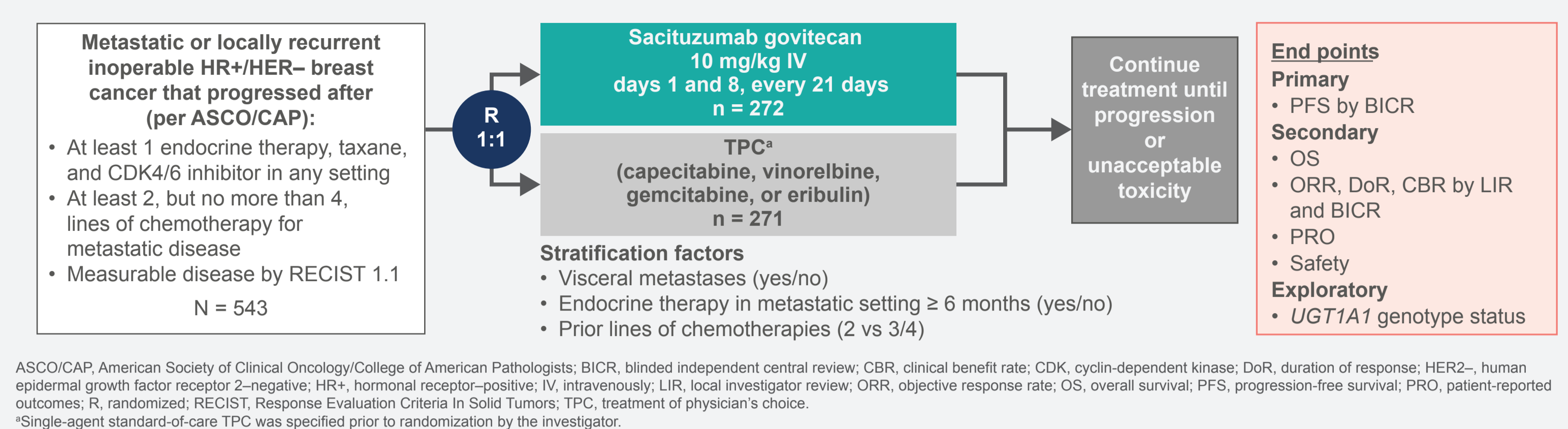
## DISCLOSURES

- Dr. Frederik Marmé reports research funding from Roche, Novartis, AstraZeneca, GSK/tesaro, MED, Clovis, Vaccibody, Gilead Sciences, and Eisai; consultancy/advisory roles with AstraZeneca, GSK/tesaro, Pfizer, Eisai, Gilead Sciences, Vaccibody, and Genomic Health; honoraria from AstraZeneca, Clovis, GSK/tesaro, Eli Lilly, Novartis, Pfizer, Roche, Myriad Genetics, PharmaMar, Eisai, MSD, Immunomics/Gilead Sciences, Pierre-Fabre, Agendia, Genomic Health, and Seattle Genetics; and travel accommodations/expenses from Pfizer, Roche, and AstraZeneca.
- Dr. Peter Schmid reports research funding from AstraZeneca, Genentech, Novartis, Oncogenes, Roche, and Medivation; and advisory roles with Amgen, AstraZeneca, Bayer, Boehringer Ingelheim, Celgene, Eli Lilly, Merck, Novartis, Pfizer, Puma, Roche, Gilead Sciences, Inc., Eisai, MSD, and Seagen.

## METHODS

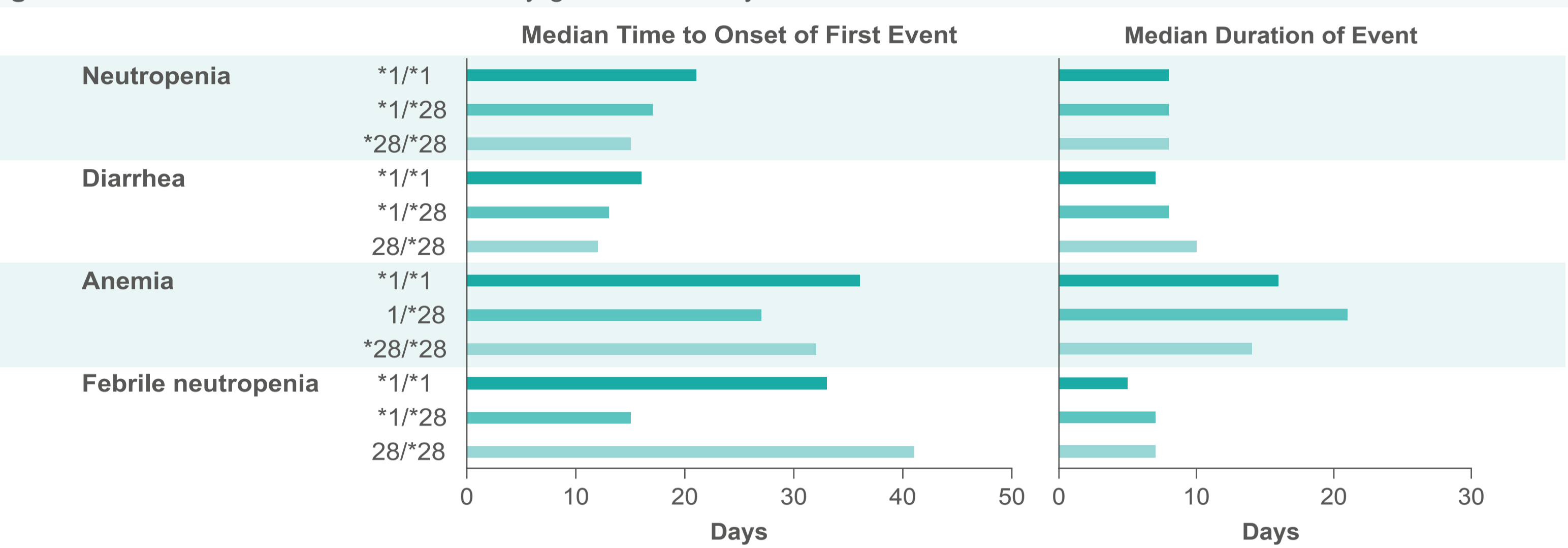
- TROPiCS-02 is a phase 3, randomized, open-label study of SG versus TPC (Figure 2)
- Exploratory safety analyses by *UGT1A1* status were performed
- The data cutoff for the analysis was July 1, 2022

Figure 2. TROPiCS-02 (NCT03901339): A phase 3 study of SG in HR+/HER2- locally recurrent inoperable or metastatic breast cancer<sup>11</sup>



ASCO/CAP, American Society of Clinical Oncology/College of American Pathologists; BICR, blinded independent central review; CBR, clinical benefit rate; CDK, cyclin-dependent kinase; DoR, duration of response; HER2-, human epidermal growth factor receptor 2-negative; HR+, hormone receptor-positive; IV, intravenously; LIR, local investigator review; ORR, objective response rate; OS, overall survival; PFS, progression-free survival; PRO, patient-reported outcomes; R, randomized; RECIST, Response Evaluation Criteria in Solid Tumors; TPC, treatment of physician's choice.  
<sup>11</sup>Single-agent standard-of-care TPC was specified prior to randomization by the investigator.

Figure 3. Time to onset and duration of any-grade TEAEs by *UGT1A1* status



SG, sacituzumab govitecan; TEAE, treatment-emergent adverse event. In patients treated with SG. Assessed in the safety population.

- Granulocyte colony-stimulating factor (G-CSF) was initiated for management of neutropenia for 54% of patients treated with SG; G-CSF treatment was initiated less frequently in wild-type patients and median time to G-CSF initiation was shortest in heterozygous patients (Table 5)

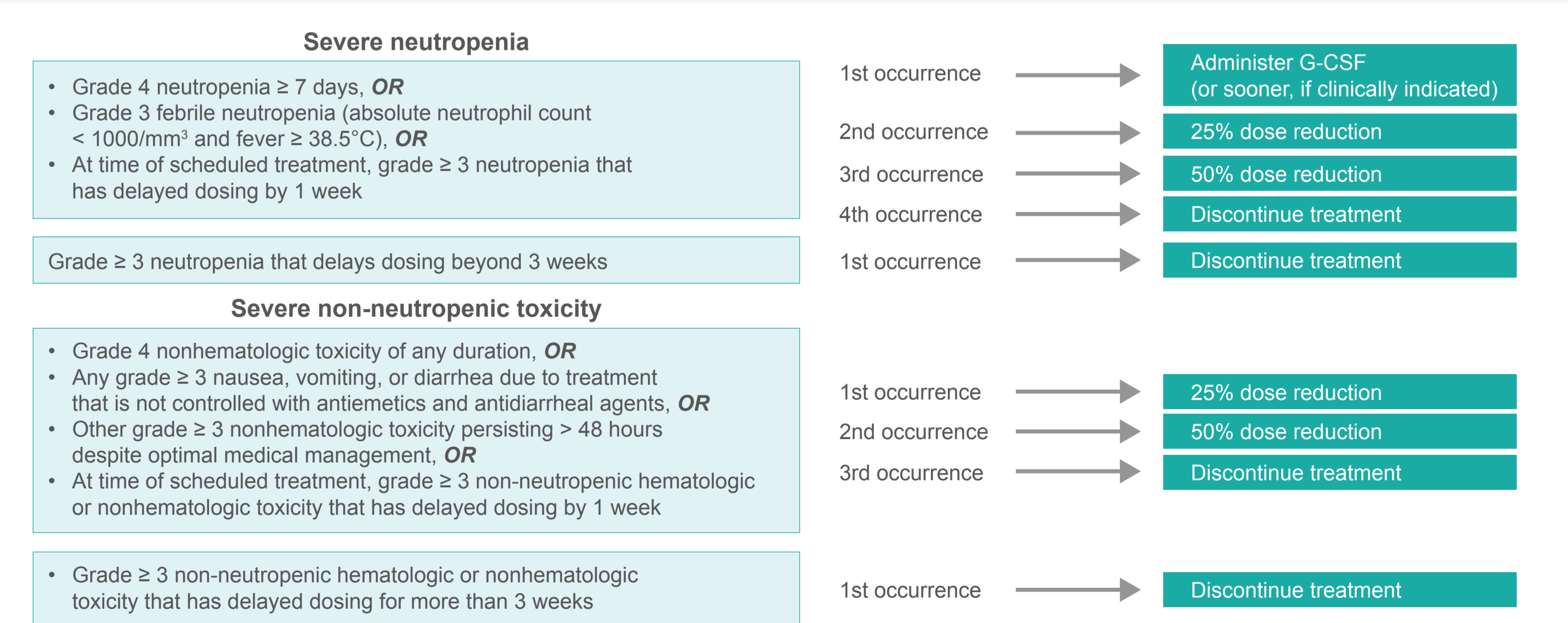
Table 5. G-CSF initiation

	SG (n = 268)			Total (n = 268)
	*1/*1 Wild-type (n = 103)	*1/*28 Heterozygous (n = 119)	*28/*28 Homozygous (n = 25)	
<b>G-CSF initiated, n (%)</b>	48 (47)	71 (60)	16 (64)	144 (54)
<b>Median time to initiation of G-CSF,<sup>a</sup> mo</b>	0.69	0.49	0.62	0.61

G-CSF, granulocyte colony-stimulating factor; SG, sacituzumab govitecan.  
<sup>a</sup>Date of G-CSF initiation was defined as the first date of G-CSF medication that started from first SG dose date to 30 days after last SG dose date.

- Severe TEAEs were managed using a well-defined treatment algorithm (Figure 4)

Figure 4. TEAE management strategies<sup>9</sup>



G-CSF, granulocyte colony-stimulating factor; TEAE, treatment-emergent adverse event.