



BACKGROUND & AIMS

- Glecaprevir/Pibrentasvir (G/P) is approved for adult patients with chronic HCV GT1-6 infection without cirrhosis or with compensated cirrhosis.
- \blacksquare G/P is also indicated for the treatment of adult patients with HCV GT1 infection who previously have been treated with a regimen containing an HCV NS5A inhibitor or an NS3/4A protease inhibitor, but not both.
- It is not recommended in patients with decompensated cirrhosis.
- We report real-world safety and efficacy of G/P in HCV-TARGET participants.

METHODS

- Patients enrolled were treated according to the local standards of care at academic (n=45) and community medical centers (n=19) in North America (n=60) and Europe (n=4).
- Detailed information on demographics, clinical course, and adverse events was abstracted from medical records into a unique centralized data core and independently monitored for completeness and accuracy.
- This analysis includes patients who started G/P before September 1st, 2018.
- Demographic, clinical, adverse events (AEs) and virological data were collected throughout treatment and post-treatment follow-up.
- Patient characteristics, Serious AEs and disposition are reported for all patients who started treatment.
- AEs are reported for patients whose treatment has concluded.
- When reporting SVR rates, SVR12 virological outcome was reported as available.
- Per Protocol Population (PP) consists of patients with available virological outcomes, excluding patients who discontinued early except for whom lack of efficacy was recorded.

DISCLOSURES

HCV-TARGET is an investigator-initiated study jointly sponsored by the University of Florida, Gainesville, FL (PI: Nelson) and the University of North Carolina at Chapel Hill, Chapel Hill, NC (PI: Fried). Funded in part by Janssen, Merck, AbbVie, Gilead, BMS, Kadmon, GSK, Genentech, and Vertex.

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For a listing of investigators and additional information about HCV-TARGET go to HCVTARGET.org ClinicalTrials.gov Identifier: NCT01474811

	8 wke	12 wke	16 wks	Other	τοται
	0 WK5	12 WK5	10 WK5	Omer	
N	430 (100)	184 (100)	25 (100)	30 (100)	726 (100)
		Demographics	N(%)		
Male	237 (55)	131 (71)	18 (72)	16 (53)	433 (60)
Age 60+	123 (29)	85 (46)	11 (44)	6 (20)	243 (34)
∋enotype: 1	294 (68)	132 (72)	18 (72)	22 (73)	512 (71)
2	59 (14)	20 (11)	1 (4)	3 (10)	88 (12)
3	63 (15)	21 (11)	5 (20)	1 (3)	96 (13)
4-6	13 (3)	10 (5)		2 (7)	25 (3)
Nos	1 (0)	1 (1)	1 (4)	2 (7)	5 (1)
x Experienced	36 (8)	28 (15)	17 (68)	3 (10)	88 (12)
Cirrhotic	17 (4)	83 (45)	12 (48)	4 (13)	120 (17)
iver Transplant		8 (4)	2 (8)		10 (1)
listory of Decomp.	3 (1)	9 (5)	1 (4)	1 (3)	14 (2)
Pialysis	6 (1)	9 (5)	1 (4)		16 (2)
rior SecondGen DAA	2 (1)	3 (2)	12 (48)	2 (7)	19 (3)
l Experience		2 (1)	2 (8)		4 (1)
IV co-infection	7 (2)	3 (2)	1 (4)		12 (2)
IS5A RAS Tested	44 (10)	20 (11)	6 (24)	5 (17)	78 (11)
RAS Present	11 (3)	7 (4)	2 (8)	3 (10)	23 (3)
	E	aseline Chemistry Medi	ian (Min-Max)		
Albumin (g/dL)	4.2 (1.5-5.3)	4 (1.8-4.9)	4.1 (2.8-4.7)	4.3 (2.9-5.2)	4.2 (1.5-5.3)
LT (IU/L)	42 (8-493)	55.5 (3-509)	52 (8-156)	39 (18-141)	47 (3-509)
Billirubin (mg/dL)	0.5 (0.2-3)	0.6 (0.2-2.2)	0.4 (0.2-2.1)	0.6 (0.2-1.3)	0.5 (0.2-3)
latelets (10 ³ /uL)	225.5 (29-581)	179 (40-443)	180 (57-371)	229 (35-575)	215.5 (29-581)
(ELD (amona cirrhotics)	7 (6-11)	8 (6-23)	7 (6-14)	7 (7-8)	7 (6-32)
CV RNA (log10 IU/mL)	6.2 (2.3-8.1)	6.3 (3.6-7.6)	6.4 (4.9-7.2)	5.9 (1.5-7.4)	6.2 (0.6-8.1)

Second Gen DAA legimens: SOF/Siviv, SOF/LDV, SOF/VEL, SOD/DCV, OF EBR/GZR Containing legimen

DISPOSITION &

Started Treatment **Ongoing Treatment**

Completed Treatment Lost to on Treatment Follo **Discontinued Prematurely Adverse Event** Non Compliance with Study Drug Other Death

Virological Outcome Avail Lost to Post Tx Follow-up In Post Tx Follow-up EOT Records Pending **Discontinued Prematurely** * AEs that caused discontinuation: ABD * 3 patients died after completing the treatment

ADVERSE EVENT

			8 wks	12 wks	16 wks	Other	TOTAL
		Adv	verse Events N(%) – List of 10 m	ost common ev	ents	
Patients	with <i>i</i>	AE	211 (49)	100 (54)	14 (56)	12 (40)	337 (46)
Fatigu	е		67 (19)	30 (19)	1 (5)		98 (18)
Heado	ache		48 (13)	25 (16)	4 (19)		77 (14)
Nause	a		35 (10)	12 (8)	4 (19)		51 (10)
Diarrh	oea		24 (7)	12 (8)			36 (7)
Pruritu	S		11 (3)	6 (4)	1 (5)		18 (3)
Influer	nza lil	ke illness	6 (2)	10 (6)			16 (3)
Dizzine	ess		9 (3)	5 (3)			14 (3)
Abdor	minal	pain upper	5 (1)	5 (3)			10 (2)
Dyspn	oea		5 (1)	4 (3)	1 (5)		10 (2)
Vomiti	ng		8 (2)	2 (1)			10 (2)
		Serie	ous Adverse Ev	ents – List incluc	des all reported	SAEs	
8 wks	4	Aortic dissed	ction (1); Chror	nic obstructive	pulmonary dise	ase (2); Dyspn	oea (2); Rhi-
		novirus infec	tion (1); Tibia fr	acture (1);			
12 wks	7	Cardiac arre	st (1); Cellulitis	(2); Haemarthro	osis (1); Hyperte	nsive crisis (1);	Kidney trans-
		plant rejection	on (1); Mental s	tatus changes ((1); Respiratory	failure (1);	
16 wks	2	Acute myoc	ardial infarctior	n (1); Myastheni	a gravis (1);		

PATIENTS WHO EXPERIENCED VIROLOGICAL FAILURE

Pt	GT	Race*	TxWks	Outcome
Pt 1	1a	AA	8	RELAPSE
Pt 2	1a	AA	8	RELAPSE
Pt 3	1a	AA	9	RELAPSE
Pt 4	1a	W	8	ON TREATMENT FAILURE
Pt 5	1a	W	8	RELAPSE
Pt 6	1a	W	8	RELAPSE
Pt 7	1a	W	16	ON TREATMENT FAILURE
Pt 8	1b	AA	12	RELAPSE
Pt 9	1b	W	8	ON TREATMENT FAILURE
Pt 10	2b	W	8	ON TREATMENT FAILURE
Pt 11	2b	W	8	ON TREATMENT FAILURE
Pt 12	3	0	9	RELAPSE
Pt 13	3a	0	8	RELAPSE
* AA: Afri	can Am	erican; W: W	, Vhite: O: Otl	her.

HCV-TARGET

Safety and Efficacy of Glecaprevir/Pibrentasvir for the Treatment of HCV Genotype 1-6: Results of the HCV-TARGET Study

Sterling, RK; Zeuzem, S; Welzel, T; Manns, M; Reddy, RK; Terrault, N; Ben-Ari, Z; Vodkin, I; Toor, A; Vainorius, M; Akuschevich, L; Fried, MW; Nelson, DR; Sulkowski, M

OUTCOM	IE – ALL	PTS WF	IO STA	RTED T	X
	8 wks	12 wks	16 wks	Other	TOTAL
	Disposition N(%)			
	430 (100)	184 (100)	25 (100)	30 (100)	726 (100)
	N/A	N/A	N/A	N/A	57 (8)
	420 (98)	181 (98)	25 (100)		626 (86)
owup	7 (2)	1 (1)	•	23 (77)	31 (4)
y .	3 (1)	2 (1)		7 (23)	12 (2)
•	2(1)	•		4 (13)	6(1)
Study Drug	1 (0)	1 (1)		•	2 (0)

				2 (7)	2 (0)
		1 (1)		1 (3)	2 (0)
Treatment	Outcome St	atus N(%)			
lable	326 (76)	139 (76)	18 (72)	4 (13)	487 (67)
	40 (9)	18 (10)	3 (12)		61 (8)
	52 (12)	23 (13)	4 (16)		79 (11)
	N/A	N/A	N/A	N/A	57 (8)
without Outcome	12 (3)	4 (2)		26 (87)	42 (6)
DOMINAL PAIN (1) HEPATIC	CIPPHOSIS (1).		ANT PE IECTION (· PRIIRITIIS (2)·

IN (1); REPAIL CIRRECOID (1); RIDNET IRANOPLANT REJECTION (1); MIGRAINE (1); PRURITUD (2)

TxExp/RAS

Naive/.

Naive/

Naive/N

Naive/

Naive/.

Naive/

Exp./Y

Naive/

Naive/

Naive/

Naive/

Naive/

Naive/

- Treatment experienced patient had prior DAA: LDV;SOF.
- All patients who failed had completed their assigned treatment duration and were not cirrhotic.

SVR 12 RATES AND 95% CONFIDENCE INTERVALS

Se

8 weeks

All	(312/323	8)		
Tx Naive	(281/292	2)		
Tx Experienced	(31/31)			H
Non-Cirrotic	(305/316	5)		
Cirrhotic	(7/7)			
Compensated	(7/7)			
Genotype 1	(206/213	3)		
Genotype 2	(44/46)			
Genotype 3	(48/50)			
Genotype 4-6	(13/13)			
econdGen DAA experienced	(2/2)			
HIV co-infected	(6/6)			

12 weeks

All	(138/139)
Tx Naive	(115/116)
Tx Experienced	(23/23)
Non-Cirrotic	(70/71)
Cirrhotic	(68/68)
Compensated	(60/60)
Decompensated	(8/8)
Genotype 1	(99/100)
Genotype 2	(16/16)
Genotype 3	(14/14)
Genotype 4-6	(8/8)
SecondGen DAA experienced	(2/2)
PI experienced	(1/1)
HIV on infactod	(1/1)
HIV CO-INTECTED	

16 weeks



* SVR rates are shown for Per Protocol population (N=480). 95% CIs were calculated by Clopper-Pearson method.

PREDICTORS OF SVR, PER PROTOCOL

- N OR LCL UCL Age 60+ 480 0.861 0.289 2.567 480 0.298 0.075 1.188 Male 8 wks 480 0.294 0.053 1.642 480 1.404 0.250 7.878 Tx Experience 5.778 0.334 99.84 Cirrhosis 471 0.694 0.034 14.02 Decompensatio Baseline ALB >=3.5 g/d 442 0.525 0.029 9.508 HCV RNA Quantifiable(3-6wks) 269 0.221 0.062 0.783 480 1.258 0.313 5.057 Use of PPI DAA Experience 480 0.285 0.046 1.756 480 0.520 0.089 3.040 Genotype 3 => Favors SVR 0.001 0.01 0.1 1 10 100
- Note that the number of failures is only 13. Hence univariable analysis was performed.
- Logistic regression adjusted by Firth's penalty. Limited to patients with treatment duration of 8/12/16 weeks.
- HCV RNA Quantifiable at 3-6 weeks is the only predictor for SVR that shows statistical significance at the 0.05 level.

SVR12 96.6 96.2 100.0 96.5 100.0 96.7 95.7 96.0 100.0 100.0 100.0	LCL 96.0 93.4 88.8 93.9 59.0 59.0 93.3 85.2 86.3 75.3 15.8 54.1	UCL 97.1 98.1 100.0 98.2 100.0 98.7 99.5 99.5 100.0 100.0 100.0
99.3 99.1 100.0 98.6 100.0 100.0 99.0 100.0 100.0 100.0 100.0 100.0	98.8 95.3 85.2 92.4 94.7 94.0 63.1 94.6 79.4 76.8 63.1 15.8 2.5 2.5	99.6 100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0
94.4 100.0 92.9 90.9 100.0 100.0 92.9 100.0 92.9 100.0 91.7 100.0	90.7 39.8 66.1 58.7 59.0 54.1 2.5 66.1 2.5 29.2 61.5 15.8	97.0 100.0 99.8 99.8 100.0 100.0 99.8 100.0 99.8 100.0 99.8 100.0

100

CONCLUSIONS

- Glecaprevir/Pibrentasvir (G/P) is safe and highly efficacious in this heterogeneous HCV-Genotype 1-6 infected population in a real-world setting.
- Predictors of SVR should be viewed with caution and not used in making clinical decisions regarding treatment continuation due to very low number of treatment failures in this population.