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Elbasvir/Grazoprevir Plus Sofosbuvir ± Ribavirin in Treatment-Naive And Treatment-Experienced People With Hepatitis C Virus Genotype 3 Infection and Compensated Cirrhosis: SVR24 Results of the C-ISLE Study

Background

- Genotype (GT)3 is a common hepatitis C virus (HCV) genotype, accounting for 30% of infections globally¹
 GT3 infection is associated with high rates of hepatocellular carcinoma and rapid progression to cirrhosis
- Despite recent advances, HCV GT3 infection and cirrhosis are challenging to treat with all-oral regimens
 Treatment-experienced people or those with baseline NS5A resistance-associated substitutions (RASs) are particularly challenging
- Elbasvir (EBR) is a once-daily NS5A inhibitor and grazoprevir (GZR) is a once-daily HCV NS3/4A protease inhibitor (Figure 1)
- The combination of EBR and GZR is approved in Europe, the United States, Canada, and other countries worldwide²
- Broad activity vs most HCV genotypes in vitro³⁻⁵
- Efficacious in treatment-naive and -experienced people, cirrhotic and noncirrhotic people,
- HIV/HCV co-infected people, and those with chronic kidney disease⁶⁻⁹
 Sofosbuvir (SOF) is an NS5B inhibitor indicated for the treatment of chronic HCV infection as a component of a combination antiviral treatment regimen (Figure 1)¹⁰
- The C-SWIFT study evaluated the safety and efficacy of EBR/GZR + SOF for 4-12 weeks in treatment-naive participants with HCV GT1 or 3 infection¹¹
- High sustained virologic response (SVR) rates (>90%) in HCV GT3-infected, treatment-naive participants receiving EBR/GZR + SOF for 12 weeks
- Similar efficacy in cirrhotic participants treated for 12 weeks and noncirrhotic participants treated for 8 or 12 weeks

Figure 1. EBR/GZR plus SOF



Aim

 To evaluate the regimen of EBR/GZR + SOF ± ribavirin (RBV) in cirrhotic participants with HCV GT3 infection for durations ranging from 8-16 weeks

Participants and Methods

Study Design

- C-ISLE (NCT02601573; Protocol PN083-02) was a randomized, open-label, UK-based clinical trial in
- HCV GT3-infected participants with compensated cirrhosis (Figure 2)
 Adults with chronic HCV GT3 infection were included
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 Compensated liver cirrhosis defined by liver biopsy (METAVIR F4) or transient elastography (>12.5 kPa)
 Treatment-naive, experienced to peginterferon/RBV, monoinfected or HIV co-infected
- All participants received EBR 50 mg/GZR 100 mg + SOF (400 mg as per prescribing information) ± RBV 800-1400 mg/day
- Treatment-naive participants were treated for 8 or 12 weeks
- Treatment-experienced participants were treated for 12 or 16 weeks
- Randomization of treatment-experienced participants was stratified based on prior relapse vs nonrelapse (partial, null, interferon-intolerant)
- Target enrollment was 25 participants per arm
- The primary endpoint was SVR 12 weeks after completion of therapy (SVR12, HCV RNA <15 IU/mL [cobas[®] TaqMan[®] v2.0])
- SVR 24 weeks after completion of therapy (SVR24) was assessed as a secondary endpoint
 SVR12 data were presented at The Liver Meeting[®] 2016¹²; this presentation provides the final
- SVR12 data were p
- Efficacy analysis population
- Full analysis set (FAS) included all participants who received ≥1 dose of study drug
- Modified FAS (mFAS) excluded participants who discontinued for reasons unrelated to the treatment regimen, including discontinuation for non-drug-related AEs
- Resistance analysis was performed using next-generation sequencing: detection limit for variants of ≈1% or ≈15% prevalence
- Resistance analysis population included all participants who had baseline sequencing available and a treatment outcome of either SVR12 or virologic failure
- NS3 polymorphisms at positions 36, 54, 55, 56, 80, 107, 122, 132, 155, 156, 158, 168, 170, or 175
- NS5A polymorphisms at positions 24, 28, 30, 31, 32, 38, 58, 62, 92, or 93
 NS5B polymorphisms at positions 06, 142, 150, 282, 280, 240, 290, 57, 201
- NS5B polymorphisms at positions 96, 142, 159, 282, 289, 316, 320, or 321

Figure 2. Study design



[†]The primary endpoint was HCV RNA <15 IU/mL at FW12 (SVR12). SVR24 was assessed as a secondary endpoint.

Results

Demographics and Characteristics

• 100 participants with HCV GT3 infection and compensated cirrhosis were enrolled (Table 1)

Table 1. Participant demographics

	Cirrhotic GT3-infected participants (n = 100)
Male, n (%)	68 (68)
Race, n (%)	
Asian	29 (29)
White	69 (69)
Other	2 (2)
Age, years, mean (SD)	53.4 (8.7)
BMI ≥30 kg/m², n (%)	28 (28)
Cirrhosis diagnosis method	
Liver biopsy, n (%)	16 (16)
FibroScan [®] , n (%)	84 (84)
FibroScan [®] score, kPa, mean (SD)	25.4 (12.1)
Prior treatment history, n (%)	
Naive	47 (47)
PR-experienced	53 (53)
HCV RNA log ₁₀ , IU/mL, mean (SD)	6.2 (0.7)
<i>IL28B</i> CC, n (%)	50 (50)
Albumin, g/dL, mean (SD)	3.6 (1.2)
ALT IU/L, median (range)	94 (21-389)
Total bilirubin, mg/dL, mean (SD)	0.7 (0.4)
Platelets × 10 ³ cells/µL, median (range)	138 (46-396)
Platelet count <100 × 10 ³ cells/µL, n (%)	24 (24)

ALT, alanine aminotransferase; BMI, body mass index; PR, peginterferon/ribavirin; SD, standard deviation.

Virologic Response

In the FAS, SVR24 was achieved by 91% and 88% of treatment-naive participants treated for 8 and

12 weeks, respectively (Figure 3)

 SVR24 was achieved by 94%, 94% and 83% of treatment-experienced participants treated for 12 or 16 weeks

- Two participants experienced virologic failure, both relapses in the 8-week treatment arm, with virologic recurrence prior to follow-up week (FW)12. There were no virologic failures after FW12
 One participant discontinued treatment due to the drug-related adverse event (AE) of cellulitis
- Seven participants were lost to follow-up: 2 prior to FW12 and 5 between FW12 and FW24



Figure 3. SVR24 in the FAS population

EOT, end of treatment

[†]All lost to follow-up or consent withdrawn, except 1 participant[‡] who discontinued treatment due to a drug-related AE.

 In the mFAS, SVR24 was achieved by 100% of treatment-naive and -experienced participants receiving EBR/GZR + SOF for 12 weeks (Figure 4)



[†]All lost to follow-up or consent withdrawn.

• Two participants relapsed (**Table 2**)

At baseline, participant 1 had Y93H, P58S, and S62T present in 44%, 99% and 62% of the viral population, suggesting these 3 RASs were linked within the same virus

Table 2. Relapse participants

	Treatment		Time of	NS5A RAVs		
Participant	history	Treatment regimen	relapse	Baseline	Failure	
1	Naive	EBR/GZR + SOF + RBV (8 weeks)	FW4	Y93H, P58S, S62T	P58S, S62T	
2	Naive	EBR/GZR + SOF + RBV (8 weeks)	FW8	WT	WT	

Nonvirologic Failures

• Eight participants had nonvirologic failure (**Table 3**)

Table 3. Nonvirologic failures

Participant	Treatment history	Treatment regimen	Reason for discontinuation	
Prior to FW	/12			
1	Naive	EBR/GZR + SOF (12 weeks)	LTFU after TW2	
2	Experienced	EBR/GZR + SOF + RBV (12 weeks)	Withdrew consent after day 7	
3	Experienced	EBR/GZR + SOF (16 weeks)	Discontinued after day 7 due to cellulitis	
Between FW12 and FW24				
4	Naive	EBR/GZR + SOF (12 weeks)	LTFU/withdrew consent	
5	Naive	EBR/GZR + SOF (12 weeks)	LTFU/withdrew consent	
6	Experienced	EBR/GZR + SOF (12 weeks)	LTFU/withdrew consent	
7	Experienced	EBR/GZR + SOF (16 weeks)	LTFU/withdrew consent	
8	Experienced	EBR/GZR + SOF (16 weeks)	LTFU/withdrew consent	

LTFU, lost to follow-up.

Resistance analyses

The presence of baseline NS3 or NS5A RASs had no impact on SVR12 (Figure 5)

Figure 5. Prevalence and impact on SVR12 of NS3 and NS5A RASs



Next-generation sequencing: detection limit for variants of 15% prevalence.

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- Based on next-generation sequencing with a detection limit of 1%, 9 participants had Y93 RASs present at baseline (Table 4)
- Four participants had Y93 present in ≥11% of the circulating viral population

Table 4: Presence of Y93 RASs at baseline (next-generation sequencing 1%)

Participant	Treatment history	Treatment	Y93H in viral population (%)	Other NS5A RASs (% in total viral population)	Treatment outcome
1	TN	EBR/GZR + SOF + RBV for 8 weeks	1	A30V (2)	SVR12
2	TN	EBR/GZR + SOF + RBV for 8 weeks	44	P58S (99); S62A (3) S62T (62)	Relapse
3	TN	EBR/GZR + SOF for 12 weeks	87	A30L (7); A30S (5) A30V (2); S62T (90) S621T (62)	SVR12
4	TN	EBR/GZR + SOF for 12 weeks	11	A30M (14)	SVR12
5	TE	EBR/GZR + SOF for 12 weeks	2	S62T (99)	SVR12
6	TE	EBR/GZR + SOF + RBV for 12 weeks	1	A30V (1)	SVR12
7	TE	EBR/GZR + SOF + RBV for 12 weeks	28	A30F (17); A30V (75) S62T (11); S62L (7)	SVR12
8	TE	EBR/GZR + SOF + RBV for 12 weeks	2	P58T (1)	SVR12
9	TE	EBR/GZR + SOF for 16 weeks	7	P58H (1); P58T (1)	SVR12

TE, treatment-experienced; TN, treatment-naive.

Adverse events

• AEs are shown in **Table 5**

Table 5. Adverse events

	EBR/GZR + SOF + RBV 8 weeks (n = 23)	EBR/GZR + SOF 12 weeks (n = 41)	EBR/GZR + SOF + RBV 12 weeks (n = 18)	EBR/GZR + SOF 16 weeks (n = 18)
Drug-related AEs, n (%)				
Fatigue	3 (13.0)	9 (22.0)	9 (50.0)	5 (27.8)
Nausea	4 (17.4)	4 (9.8)	6 (33.3)	3 (16.7)
Skin rash/pruritus	2 (8.7)	3 (7.3)	8 (44.4)	2 (11.1)
Serious AE (SAE), n (%) [†]	0	1 (2.4)	3 (16.7)	1 (5.6)
Discontinued study medication due to AE, n (%) [‡]	0	0	0	1 (5.6)
Hemoglobin <10 g/dL, n (%) [¶]	0	1 (2.4)	2 (11.1)	0
Total bilirubin >5× baseline, n (%)	0	0	0	0
ALT/AST >5× ULN, n (%)	0	0	0	0

AST, aspartate aminotransferase; ULN, upper limit of normal.

[†]SAEs of lung infection, creatinine increased, chest pain, opiate overdose, and cellulitis were reported.

[‡]One participant had a drug-related SAE of vomiting on day 4 and then subsequently discontinued treatment at day 7 due to drug-related cellulitis.

[¶]Lowest hemoglobin level was 8.9 g/dL.

Conclusions

- High efficacy of EBR/GZR was demonstrated in treatment-naive and -experienced cirrhotic HCV GT3-infected participants
- In the mFAS analysis, SVR12 was 100% among participants receiving EBR/GZR + SOF for 12 weeks
- There were no virologic failures among participants receiving EBR/GZR + SOF ± RBV for 12 or 16 weeks
- Treatment duration longer than 12 weeks is not needed
- High efficacy regardless of presence of baseline NS5A RASs or participant characteristics
- Generally safe and well tolerated

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