

## **Next Generation Direct-Acting Antivirals**

Glecaprevir (formerly ABT-493) pangenotypic NS3/4A protease inhibitor

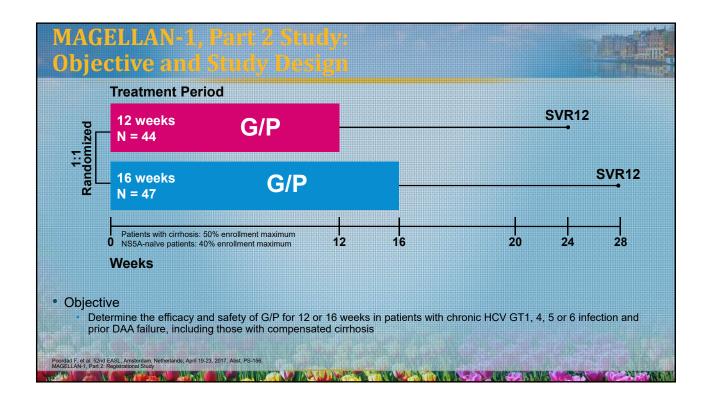
**Pibrentasvir** (formerly ABT-530) pangenotypic NS5A inhibitor

Coformulated: G/P

- · Once-daily oral dosing with food
- Potent against common NS3 and NS5A polymorphisms<sup>1</sup>
- Minimal metabolism, primary biliary excretion, and negligible renal excretion (<1%)</li>
  - Favorable safety profile: mostly mild AEs with few Grade ≥3 lab abnormalities²
  - Concomitant PPI use allowed
  - Phase 2 (12 weeks): 95% mITT SVR12 rate in patients with prior NS5A and/or PI failure<sup>3</sup>

G/P is coformulated and dosed once daily as three 100 mg/40 mg pills for a total dose of 300 mg/120 mg Glecaprevir was identified by AbbVie and Enanta

Ng TI, et al. Antimicrobial Agents and Chemotherapy, 2017. DuFour, J-F et al. EASL 2017. FRI-238 Poordad F, et al. Hepatology 2017. DOI: 10.1002/hep.29081. coordad F, et al. 52nd EASL, Amsterdam, Netherlands, April 19-23, 2017. Abst. PS-156



## MAGELLAN-1, Part 2 Study: Baseline Demographics and Clinical Characteristics

Characteristic	G/P 12 weeks N = 44	G/P 16 weeks N = 47
Male, n (%)	31 (70)	33 (70)
White race, n (%)	34 (77)	35 (75)
Black race, n (%)	9 (20)	11 (23)
Age, median years (range)	57 (22 – 67)	56 (36 – 70)
BMI, median kg/m² (range)	28 (21 – 41)	29 (20 – 52)
HCV RNA, median log <sub>10</sub> IU/mL (range)	6.1 (4.7 – 7.2)	6.3 (4.7 – 7.1)
Compensated cirrhosis, n (%)	15 (34)	12 (26)
HCV subtype*, n (%)		
1a	35 (80)	32 (71)
1b	8 (18)	10 (22)
1e	0	1 (2)
4	1 (2)	3 (6)
1 (not subtyped)	0	1 <sup>†</sup> (2)

\*Genotype and subtype determined by phylogenetic analysis for samples with available sequences †One patient had GT1, based on LIPA analysis, but was not subtyped

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MAGELLAN-1, Part 2 Study: Clinical Characteristics (Cont'd)

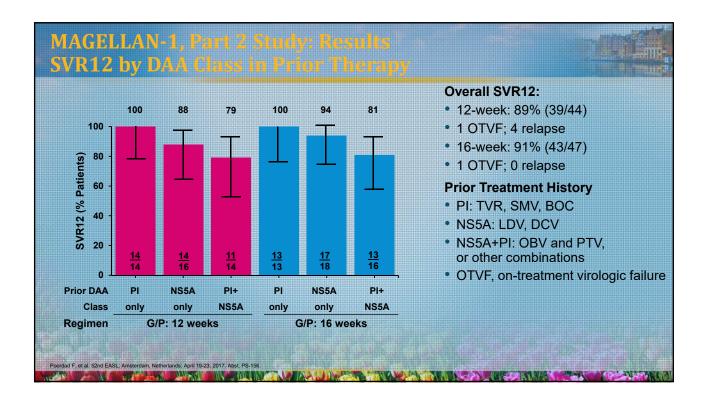
	Characteristic , n (%)	G/P 12 weeks N = 44	G/P 16 weeks N = 47
F	Previous DAA regimen class*		
	NS3/4A PI only (NS5A inhibitor-naïve)	14 (32)	13 (28)
	NS5A inhibitor only (PI-naïve)	16 (36)	18 (38)
	NS3/4A PI + NS5A inhibitor	14 (32)	16 (34)
F	Prior DAA treatment response		
	On-treatment failure	14 (32)	13 (28)
	Virologic relapse	30 (68)	34 (72)
F	Presence of key baseline substitutions <sup>†</sup>		
	None	13 (30)	13 (30)
	NS3 only	2 (5)	4 (9)
	NS5A only	24 (55)	23 (52)
	NS3 + NS5A	5 (11)	4 (9)

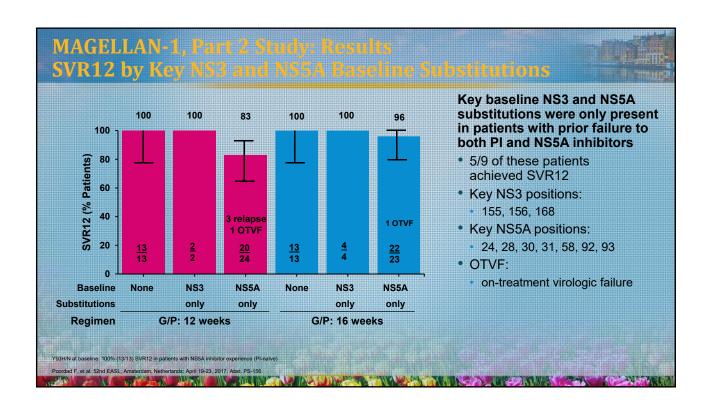
NS3 Pis included: PTV, SMV, ASV, TVR, or BOC; NS5A inhibitors included: DCV, LDV, or OBV

\*Sofosbury (NS56 inhibitor) could be included in any prior treatment regimen

†Only 44 patients and sequencing data available in each arm; percentages are based on N = 44; substitutions detected by next generation sequencing using 15% detection threshold at positions 155, 156, and 168 in NS3, and 24, 28, 30, 31, 58, 92, and 33 in NS5A.

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## MAGELLAN-1, Part 2 Study: Conclusions

- Patients with prior failure to PI containing regimens (NS5A inhibitornaïve):
  - 100% SVR12 with 12 or 16 weeks of G/P treatment
- Patients with prior failure to both PI- and NS5A inhibitor-containing regimens had lower SVR12 rates
- Patients with prior failure to NS5A inhibitors (i.e., LDV or DCV); NS3/4A PI-naïve:
  - 94% SVR12 with 16 weeks of G/P treatment with no relapse
  - No impact of baseline NS5A substitutions on SVR12
- G/P for 12 or 16 weeks was well tolerated; Grade 3 lab abnormalities were rare, with no discontinuations due to AEs, and no DAA-related serious AEs

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