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The Combination of Elbasvir and Grazoprevir for the Treatment of Chronic HCV Infection in Japanese Patients: A Randomized Phase II/III Study

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**Table S1** Prevalence of baseline NS3 and NS5A RAVs in parts 1 and 2

Population	Total number of patients	Number of patients with sequences available	Number of patients With RAVs		Baseline RAVs (number of patients with variants detected)
			RAVs not detected, <i>n</i> (%)	RAVs detected, <i>n</i> (%)	
<b>NS3 RAVs</b>					
Overall	321	321	218 (67.9)	103 (32.1)	D168D/E (3), D168E (2), M175L (2), M175L/M (1), Q80K (4), S122G (57), S122S/G (13), S122T/A (1), S122T/S/A/G (1), T54S (12), T54S/T (1), V107I (1), V107I/V (1), V158I (1), V170I/M/V (2), V170I/T (1), V170I/V (8), V36L (1), V55A (1)
<b>By genotype and subtype</b>					
GT1a	5	5	3 (60.0)	2 (40.0)	Q80K (2)
GT1b	316	316	215 (68.0)	101 (32.0)	D168D/E (3), D168E (2), M175L (2), M175L/M (1), Q80K (2), S122G (57), S122S/G (13), S122T/A (1), S122T/S/A/G (1), T54S (12), T54S/T (1), V107I (1),

					V107I/V (1), V158I (1), V170I/M/V (2), V170I/T (1), V170I/V (8), V36L (1), V55A (1)
<b>NS5A RAVs</b>					
Overall	321	321	263 (81.9)	58 (18.1)	L31L/I/M (1), L31L/M (1), L31M (11), L31V (1), M28M/V (1), R30H (1), R30Q/R (1), Y93H (22), Y93Y/C (2), Y93Y/H (19)
<b>By genotype and subtype</b>					
GT1a	5	5	3 (60.0)	2 (40.0)	M28M/V (1), Y93Y/C (1)
GT1b	316	316	260 (82.3)	56 (17.7)	L31L/I/M (1), L31L/M (1), L31M (11), L31V (1), R30H (1), R30Q/R (1), Y93H (22), Y93Y/C (1), Y93Y/H (19)

RAVs resistance-associated variants, *GT*, genotype