

Genotype 1a

TABLE 108: SVR GENOTYPE 1A TREATMENT-EXPERIENCED WITH PRIOR RELAPSE: RELATIVE RISKS AND RISK DIFFERENCE FOR ALL TREATMENT COMPARISONS — RANDOM EFFECTS MODEL

TREATMENT	REFERENCE	RR (95% CRI)	RD % (95% CRL)
SIM12 PR24-48 RGT	PR48	2.62 (1.85 to 3.60)	45.63 (26.75 to 60.65)
PAR/RIT12 + OMB12 + DAS12 + RBV12		3.30 (2.51 to 4.36)	64.94 (48.52 to 73.97)
PAR/RIT12 + OMB12 + DAS12 + RBV12	SIM12 PR24-48 RGT	1.26 (0.98 to 1.66)	19.08 (-2.04 to 37.74)
Random effect model	Residual deviance	5.624 vs. 6 data points	
	Deviance information criteria	33.823	
Fixed effect model	Residual deviance	5.68 vs. 6 data points	
	Deviance information criteria	33.668	

Cri = credible interval; DAS = dasabuvir; OMB = ombitasvir; PAR = paritaprevir; PR = pegylated interferon plus ribavirin; RBV = ribavirin; RD = risk difference; RGT = response-guided therapy; RIT = ritonavir; RR = relative risk; SIM = simeprevir; vs. = versus.

Note: Please refer to Treatment Regimen Nomenclature table for description of dosages.