TABLE 148: VIRAL LOAD ≥ THRESHOLD GENOTYPE 2 TREATMENT-NAIVE: RELATIVE RISKS AND RISK DIFFERENCE FOR ALL TREATMENT COMPARISONS — RANDOM EFFECTS MODEL

TREATMENT	REFERENCE	RR (95% CRI)	RD % (95% CRL)
SOF12 + RBV12	PR24	1.17 (1.07 to 1.28)	13.25 (5.60 to 20.55)
Random effect model	Residual deviance	6.838 vs. 8 data points	
	Deviance information criteria	38.161	
Fixed effect model	Residual deviance	6.96 vs. 8 data points	
	Deviance information criteria	37.961	

CrI = credible interval; PR = pegylated interferon plus ribavirin; RBV = ribavirin; RD = risk difference; RR = relative risk; SOF = sofosbuvir; vs. = versus.
Note: Please refer to Treatment Regimen Nomenclature table for description of dosages.