TABLE 149: 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	1.25 (1.08 to 1.45)	19.04 (6.66 to 29.79)
Random effect model	Residual deviance	6.754 vs. 8 data points	
	Deviance information criteria	32.185	
Fixed effect model	Residual deviance	6.769 vs. 8 data points	
	Deviance information criteria	32.023	

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.