TABLE 155: VIRAL LOAD < THRESHOLD GENOTYPE 4 TREATMENT-EXPERIENCED: RELATIVE RISKS AND RISK DIFFERENCE FOR ALL TREATMENT COMPARISONS — RANDOM EFFECTS MODEL

| TREATMENT | REFERENCE | RR (95% CRI) | RD % (95% CRL) |
|-------------------------|-------------------------------|-------------------------|------------------------|
| SOF24 + RBV24 | SOF12 + RBV12 | 1.02 (0.46 to 1.48) | 1.57 (-47.73 to 30.03) |
| DCV24 + ASU24 + PR24 | | 1.05 (0.61 to 1.55) | 4.79 (-35.24 to 33.60) |
| DCV24 + ASU24 + PR24 | SOF24 + RBV24 | 1.03 (0.58 to 2.40) | 3.10 (-39.92 to 56.32) |
| | | | |
| Random effect model | Residual deviance | 3.341 vs. 4 data points | |
| | Deviance information criteria | 12.853 | |
| Fixed effect model | Residual deviance | 3.262 vs. 4 data points | |
| | Deviance information criteria | 12.716 | |

ASU = asunaprevir; CrI = credible interval; DCV = daclatasvir; 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.