## Treatment-Experienced Patients All Patients

## TABLE 125: SVR GENOTYPE 2 TREATMENT-EXPERIENCED: RELATIVE RISKS AND RISK DIFFERENCE FOR ALL TREATMENT COMPARISONS — RANDOM EFFECTS MODEL

| TREATMENT           | REFERENCE                     | RR (95% CRI)            | RD % (95% CRL)          |
|---------------------|-------------------------------|-------------------------|-------------------------|
| SOF16 + RBV16       | SOF12 + RBV12                 | 0.86 (0.63 to 1.02)     | -12.21 (-33.63 to 1.88) |
| SOF12 + PR12        |                               | 1.07 (0.93 to 1.15)     | 6.57 (-6.25 to 12.59)   |
| SOF12 + PR12        | SOF16 + RBV16                 | 1.23 (1.00 to 1.70)     | 18.27 (0.20 to 40.32)   |
|                     |                               |                         |                         |
| Random effect model | Residual deviance             | 6.58 vs. 8 data points  |                         |
|                     | Deviance information criteria | 34.221                  |                         |
| Fixed effect model  | Residual deviance             | 6.609 vs. 8 data points |                         |
|                     | Deviance information criteria | 34.152                  |                         |

Crl = credible interval; PR = pegylated interferon plus ribavirin; RBV = ribavirin; RD = risk difference; RR = relative risk; SOF = sofosbuvir; SVR = sustained virologic response; vs. = versus.

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