**Table S2. Associations of** **IFIH1-DHX58 SNPs with HCV infection outcomes in dominant and additive models of multivariable analyses**

|  |  |  |  |
| --- | --- | --- | --- |
| **SNPs** | **Dominant model** | **Dominant model** | **Additive model‡** |
| ***P*** | ***PFDR*** | ***P*** | ***PFDR*** | ***P*** | ***PFDR*** |
| Rs10930046 (T>C) | 0.259 | 0.389 |  <0.001 |  <0.001 | 0.005 | 0.008 |
| Rs2074158 (A>G) | 0.001 | 0.003 |  <0.001 |  <0.001 |  <0.001 |  <0.001 |
| Rs2074160 (G>A) | 0.748 | 0.748 | 0.115 | 0.115 | 0.410 | 0.410 |

 *PFDR* is expressed as the P value after multiple corrections by the FDR method