**Supplementary Table 2. Impact of virological markers on HBsAg relapse**

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| --- | --- | --- | --- |
|  | **HBsAg relapse**  ***n*=7** | **HBsAg non-relapse**  ***n*=41** | ***p-*value** |
| cccDNA, log10 copies/cell |  |  | 0.166 |
| +(*n*=35) | 7 (20%) | 28 (80%) |  |
| −(*n***=**13) | 0 (0%) | 13 (100%) |  |
| HBcrAg, mean (SD) | 3.81 (0.46) | 3.53 (0.54) (*n***=**29） | 0.207 |
| HBsAb, IU/mL |  |  |  |
| ≥ 100 | 0 | 19 | 0.033 |
| < 100 | 7 | 22 |  |
| HBsAg immunohistochemistry (+/−) | 4/3 | 8/33 | 0.055 |

cccDNA, covalently closed circular DNA; HBcrAg, hepatitis B core related antigen; HBsAb, hepatitis B surface antibody; HBsAg, hepatitis B surface antigen; SD, standard deviation.