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| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Polymorphism | Author | Year | population | Sample size | | MAF(%)  (controls) | Influence on | References |
| cases | controls |
| rs187084  (A/G) | Chihab et al. | 2019 | Moroccan | 239 | 133 | 32.33 | The progression of HBV-related liver disease | [74] |
| rs5743836  (A/G) | Chihab et al. | 2019 | Moroccan | 239 | 133 | 14.40 | HBV DNA load | [74] |
| Wu et al. | 2012 | Taiwanese | 278 | - | - | HBV clearance | [64] |
| rs352140  (C/T)  (G/A) | He et al. | 2015 | Chinese Han | 1191 | 273 | - | Susceptibility to HBV infection | [76] |
| Gao et al. | 2015 | Chinese | 51 | 348 | 42.20 | Susceptibility to HBV intrauterine transmission | [43] |
| Abbreviation: MAF: minor allele frequency. | | | | | | | | |