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Conservation and Polymorphism of EBV *RPMS1* gene in EBV-Associated Tumors and Healthy Individuals from Endemic and Non-Endemic Nasopharyngeal Carcinoma Areas in China

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Highlights

- According to three meaningful mutations (g155325t, g155326a and c155389t) in our experiment, RPMS-1 was classified into RPMS1-A, RPMS1-B, RPMS1-C, RPMS1-D.
- There was significantly difference between EBV-associated tumors

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