Figure S1. Analysis of the whole-exome sequencing raw data (variant cell format files) using NextGENe® software. Compared with the reference sequence, the WES data suggest that there are two close variants in *TMTC3*, one is a missense variant (c.1123G>A) and the other is a four bases-deletion variant (c.1126_1129delCGAG). TMTC3, transmembrane O-mannosyltransferase targeting cadherins 3.