



Fig. S2. The coverage across target regions. The coverage in the Y axes (values between 0–200) and the target region (exons 1–6, 7–27b, 28–49) in the X axes for both sample sets A and B. In addition to the exon 1, there were no other regions with constantly low coverage. Black arrows indicate positions of confirmed mutations, red arrows are positions of false negative and asterisk indicate positions of false positive mutations.