



Fig. S3. Mutations found in sequencing with Roche 454 GS Junior as illustrated in Integrative Genomics Viewer (IGV). Inserts: Sanger sequencing of the corresponding area. (A) E579 intronic delT, a homopolymer-related false positive. Confirmed mutations: (B) E396 c.3911T>G, (C) S47 c.4922G>A, (D) E71 c.499_502delTGTT.