Table S3. Somatic mutation found by WES of tumoral material of the TC0011. Mutations were validated by Sanger and ultra-deep sequencing

Chr	Start	End	Ref	Alt DP	Func.refge	neGene.refgene	ExonicFunc.: SOMATIC
chr12	113531016	113531016	5 C	G	353 exonic	DTX1	nonsynonynyes
chr2	223086090	223086090) C	T	308 exonic	PAX3	nonsynonynyes
chr7	140477790	140477790) -	CTGAGTACT	52 splicing	BRAF(NM_0043	33:e:NA yes