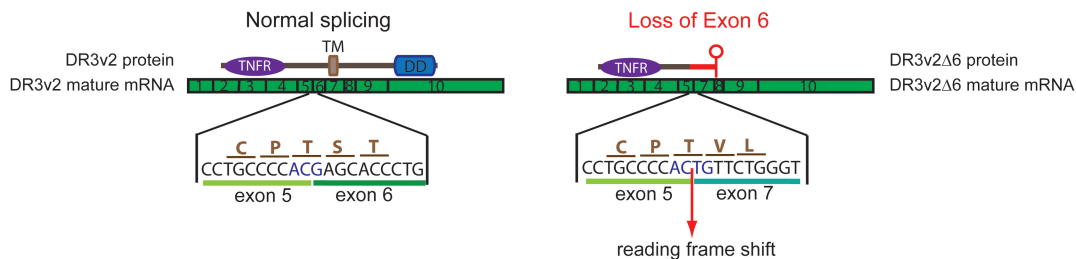


A

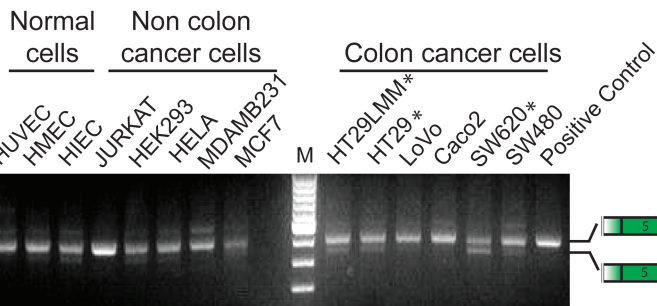


B

DR3v2	NP_003781.1	MEQRPRGCAA	VAAALLLVLL	GARAQGGTRS	PRCDCAGDFH	KKIGLFCCRG	CPAGHYLKAP	CTEPCGNSTC	70
DR3v2Δ6		MEQRPRGCAA	VAAALLLVLL	GARAQGGTRS	PRCDCAGDFH	KKIGLFCCRG	CPAGHYLKAP	CTEPCGNSTC	70
DR3v2	NP_003781.1	LVCPQDTFLA	WENHHNSECA	RCQACDEQAS	QVALENC SAV	ADTRCGCKPG	WFVECVQSQC	VSSSPFYCQP	140
DR3v2Δ6		LVCPQDTFLA	WENHHNSECA	RCQACDEQAS	QVALENC SAV	ADTRCGCKPG	WFVECVQSQC	VSSSPFYCQP	140
DR3v2	NP_003781.1	CLDCGALHRH	TRLLCSRRDT	DCGTCLPGFY	EHGDGCVSCP	TSTLGSCPER	CAAVCG---W	RQM-FWVQVL	206
DR3v2Δ6		CLDCGALHRH	TRLLCSRRDT	DCGTCLPGFY	EHGDGCVSCP	T-VLG--PGA	PGWPCGPPAW	GHPDLHIPPL	207
DR3v2	NP_003781.1	LAGLVVPLLL	GATLTYTYRH	CWPHKPLVTA	DEAGMEALTP	PPATHLSPLD	SAHTLLAPPD	SSEKICTVQL	276
DR3v2Δ6		LA-----	-----	-----	-----	-----	-----	-----	209
DR3v2	NP_003781.1	VGNSWTPGY P	ETQEALCPQV	TWSWDQLPSR	ALGPAAAPT L	SPESPAGSPA	MMLQPGPQLY	DVMDAVPARR	346
DR3v2Δ6		---SQAPGY---	---CR---	-----	-----	-----	-----	-----	217
DR3v2	NP_003781.1	WKEFVRTLGL	REAEIEAVEV	EIGRFRDQY	EMLKRWQQQ	PAGLGAVYAA	LERMGLDGCV	EDLRSRLQRG	416
DR3v2Δ6		-----	-----	-----	-----	-----	-----	-----	217

DR3v2 NP\_003781.1 P 417  
DR3v2Δ6 - 217

C



D

