

Table S16. Associations between VDR haplotypes and progression free survival in 127 CRC patients.

VDR Haplotypes	Estimation	St Error	T-Test	HRR (IC95%)	p-value
TCA	-0.970	0.661	-1.466	0.378 [0.103 - 1.386]	0.142
TAA	0.032	0.543	0.060	1.033 [0.356 - 2.997]	0.951
CCA	-0.253	0.200	-1.262	0.775 [0.523 - 1.150]	0.206
CCG	-0.571	0.920	-0.621	0.564 [0.092 - 3.430]	0.534
CAA	0.068	0.239	0.286	1.071 [0.669 - 1.714]	0.774
CAG	0.037	0.852	0.043	1.038 [0.195 - 5.520]	0.964
Ref. haplotype = TAG -2x Log-likelihood (with covariates) =460.739477 -2 x Log-likelihood (without covariates) = 466.690810 (df = 6) p-value = 0.429					