The impact of DNA damage response gene polymorphisms on therapeutic outcomes in late stage ovarian cancer.

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Legends to Supplementary Figures

Supplementary Figure 1. Overall survival and Progression Free Survival curves by ATM genotypes.

Supplementary Figure 2. Overall survival and Progression Free Survival curves by ATR genotypes.

Supplementary Figure 3. Overall survival and Progression Free Survival curves by CHK1 (upper panels) and CHK2 (lower panels) genotypes.

Supplementary Table 1. SNPs evaluated.

Gene	Chromosome	Most severe consequence	Ref. SNP ID n°	MFA
ATM	11	T to C transition: intron variant	rs664677	0,35 (C)
	11	G to A transition: 3 prime UTR variant	rs664143	0,37 (A)
ATR	3	C to T transition: missense variant	rs2229032	0,10 (T)
	3	G to A transition: missense variant	rs2227928	0,40 (A)
CHEK1	11	G to A transition: intron variant	rs521102	0,44 (A)
СНЕК2	22	T to C transition: intron variant	rs2267130	0,31 (C)
CDK12	17	G to A transition: 3 prime UTR variant	rs1054488	0,22 (A)

MFA= minor allele frequency





