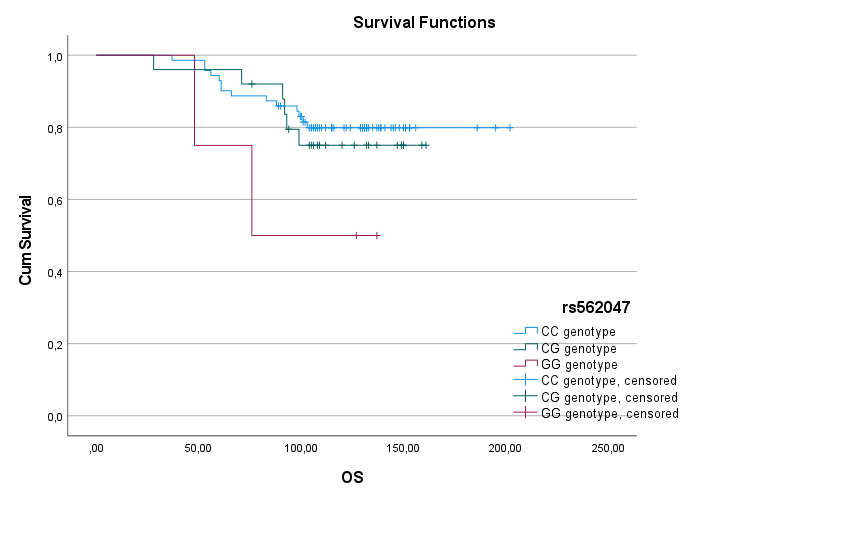
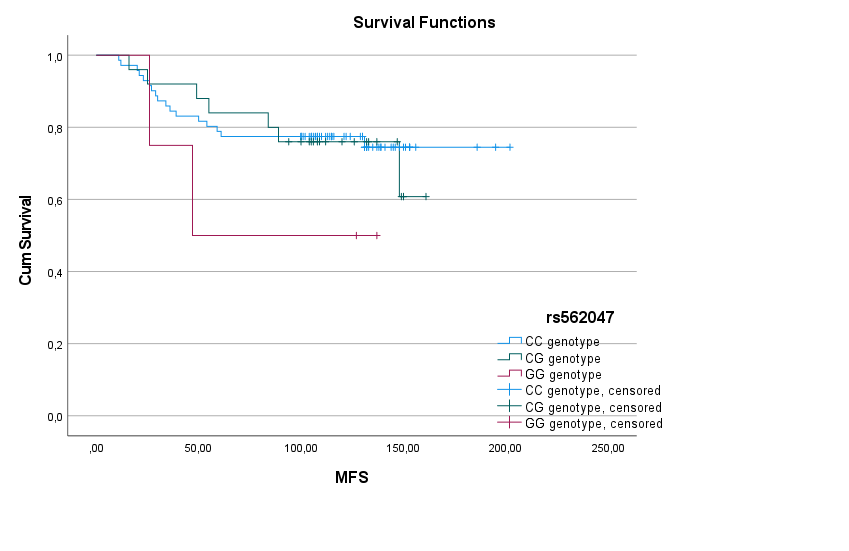
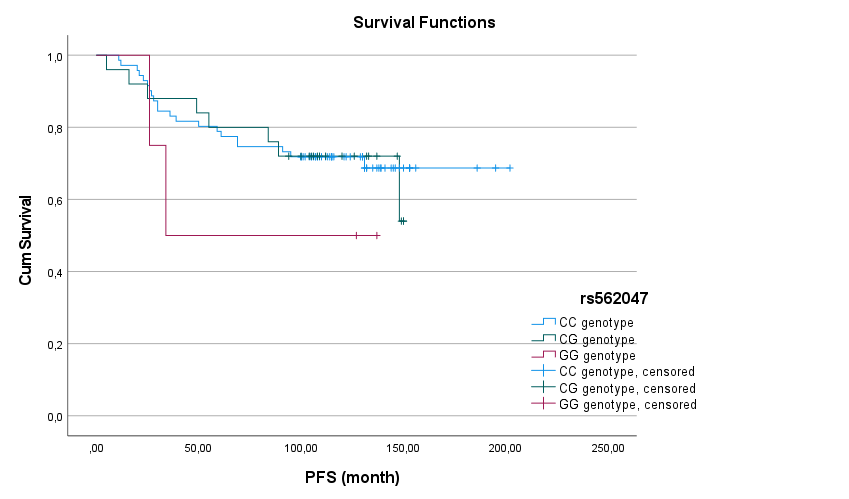
* + 1. ***HSPA1A rs562047***

The possible associations between *HSPA1A rs562047* polymorphism and BC patient survival were assessed (Kaplan-Meier analysis, Log Rank test). In our study, no significant link between this SNP in genotype model and PFS (p=0.587), MFS (p=0.587), OS (p=0.226) was determined. In allelic model there was also no-significant correlation found with *PFS* (p=0.308, p=0.651), MFS (p=0.198, p=0.560) and OS (p=0.095, p=0.379) and C or G alleles, respectfully. Due to non-significant results Cox survival analysis was not further performed. The representative PFS, MFS and OS curves, generated with Kaplan–Meier, are presented in Figure 7.



***Figure 6. Kaplan-Meier curves for PFS, MFS and OS for different HSPA1A rs562047 genotypes***