

Impact of genetic variations of Polymerase γ gene (*POLG*) on breast cancer patients

Ieva Golubickaite^{1,*}, Rasa Ugenskiene^{1,2}, Erika Korobeinikova³, Jurgita Gudaitiene^{2,3}, Domas Vaitiekus³, Lina Poskiene^{4,5}, Elona Juozaityte^{2,3}

¹Institute of Biology Systems and Genetic Research, Lithuanian University of Health Sciences, ²Institute of Oncology, Lithuanian University of Health Sciences, ³Department of Oncology and Hematology, Hospital of Lithuanian University of Health Sciences, ⁴Department of Pathological Anatomy, Lithuanian University of Health Sciences, ⁵Department of Pathology, Hospital of Lithuanian University of Health Sciences

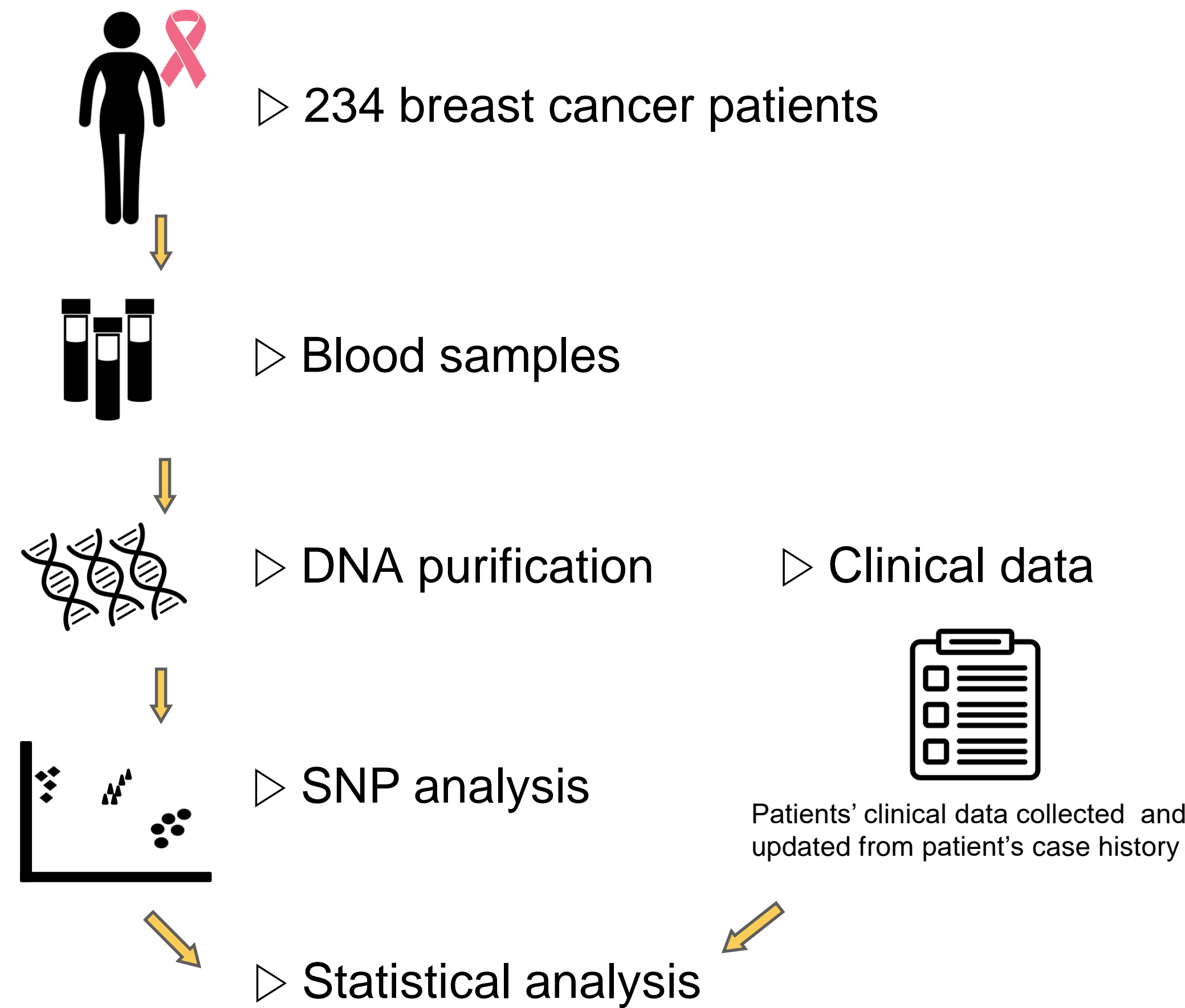
Background

- ▷ Breast cancer is the leading cause of cancer-related deaths among women.
- ▷ Variations in mitochondrial DNA (mtDNA) and nuclear DNA, coding for mitochondrial proteins or regulatory molecules, are seldom under consideration to be related to cancer.
- ▷ Polymerase gamma is important for mtDNA repair.
- ▷ It is also the only polymerase involved in mtDNA replication.

Aim

- ▷ Assess *POLG* gene variations rs3087374, rs2307441, rs2072267, rs976072 and associations with tumor phenotype, disease outcome for 234 breast cancer patients.

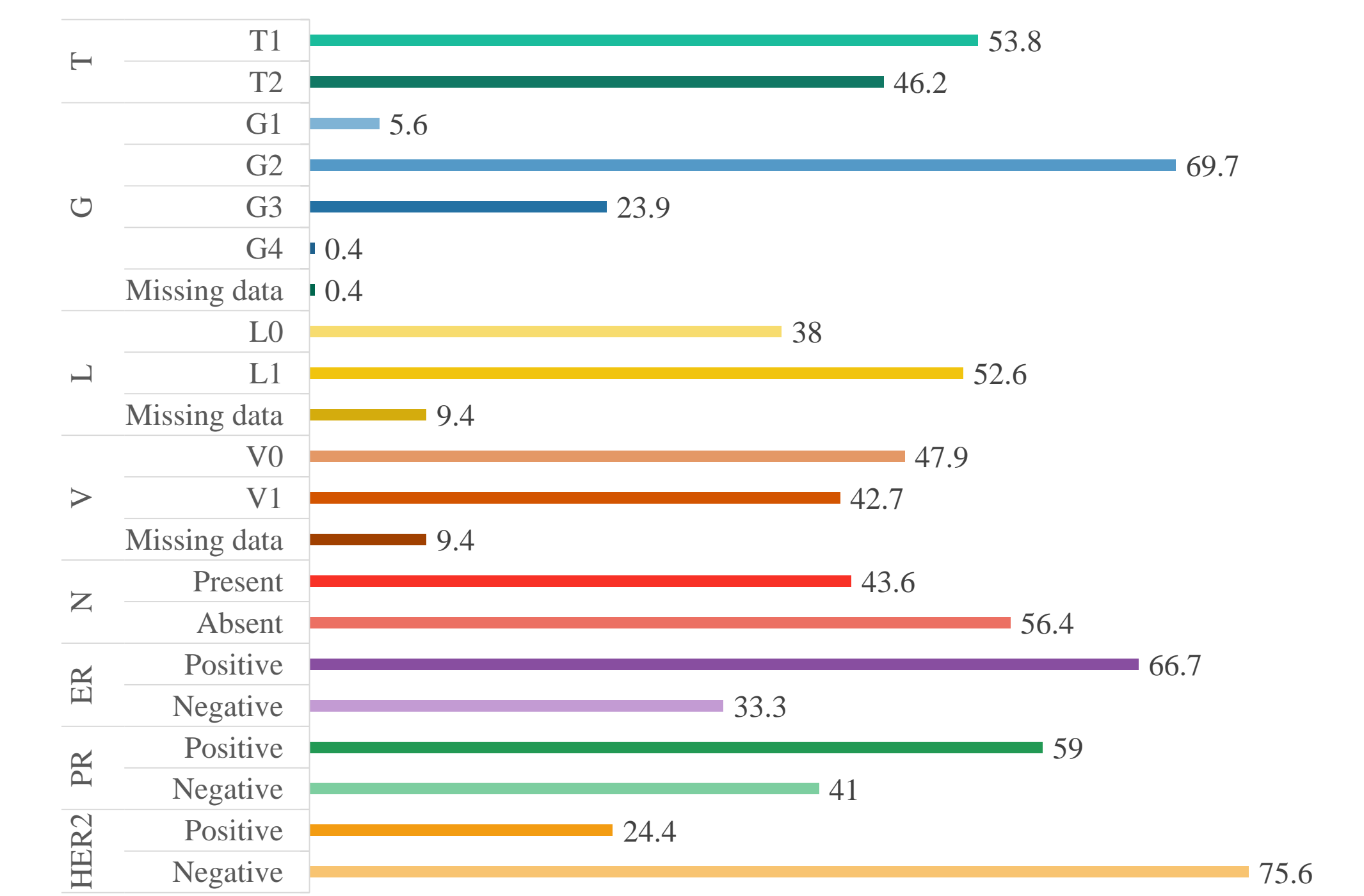
Methods



Results

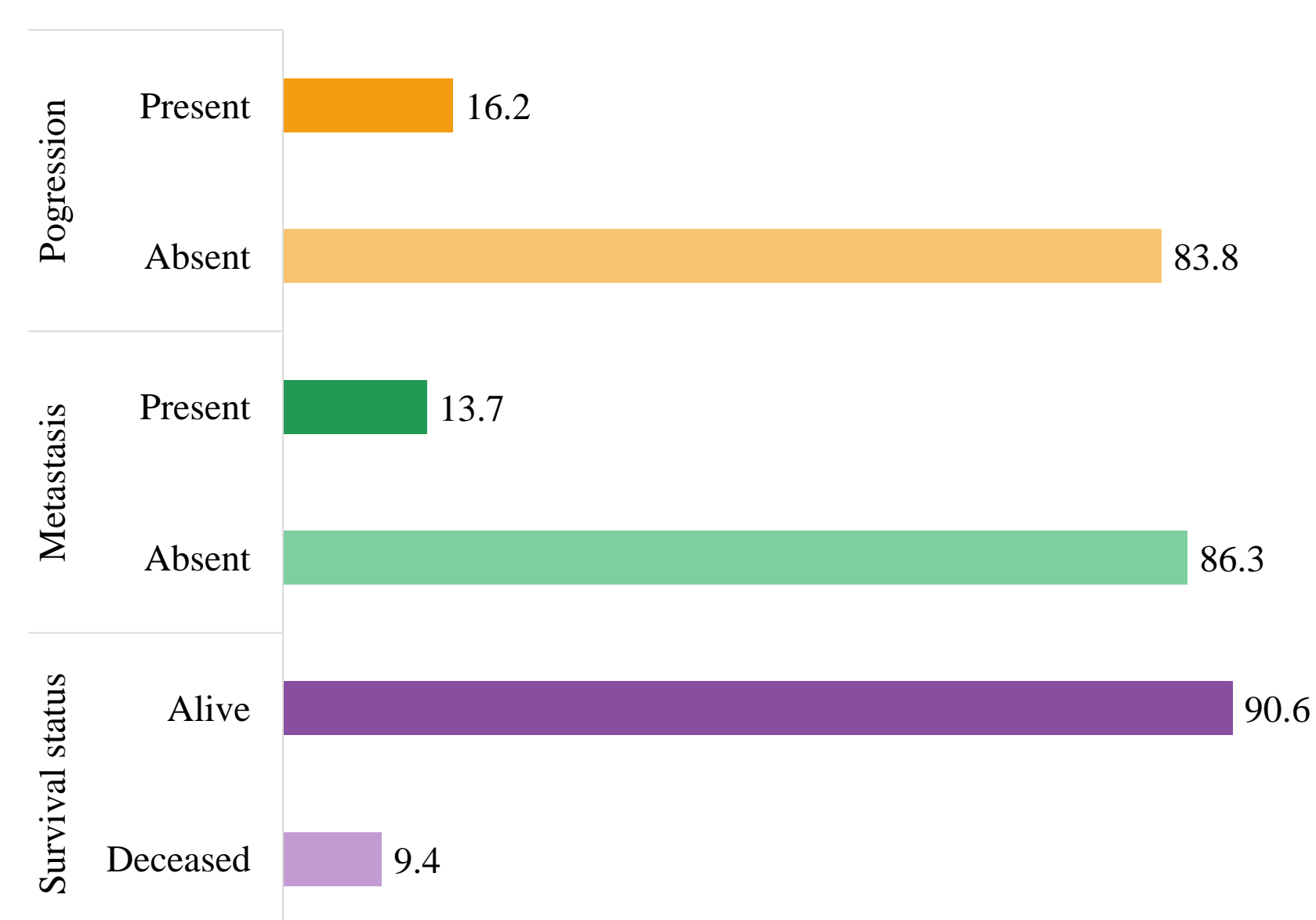
- ▷ Patients with rs2307441 TT and CT genotypes had a lower probability for tumor vascular invasion than CC genotype (OR=0.075; 95% CI 0.017-0.323; p=0.001)
- ▷ The carriers of the T allele in rs2307441 had a lower probability of tumor vascular invasion than non-carriers (OR=0.064; 95% CI 0.015-0.273; p=0.000)
- ▷ Patients with rs2072267 AG genotype were predisposed for disease progression compared with GG genotype (OR=3.386; 95% CI 1.262-9.086; p=0.015)
- ▷ The carriers of the A allele of rs2072267 had a higher probability of disease progression than the non-carriers (OR=3.197; 95% CI 1.237-8.261; p=0.016)

Tumor pathomorphological characteristics



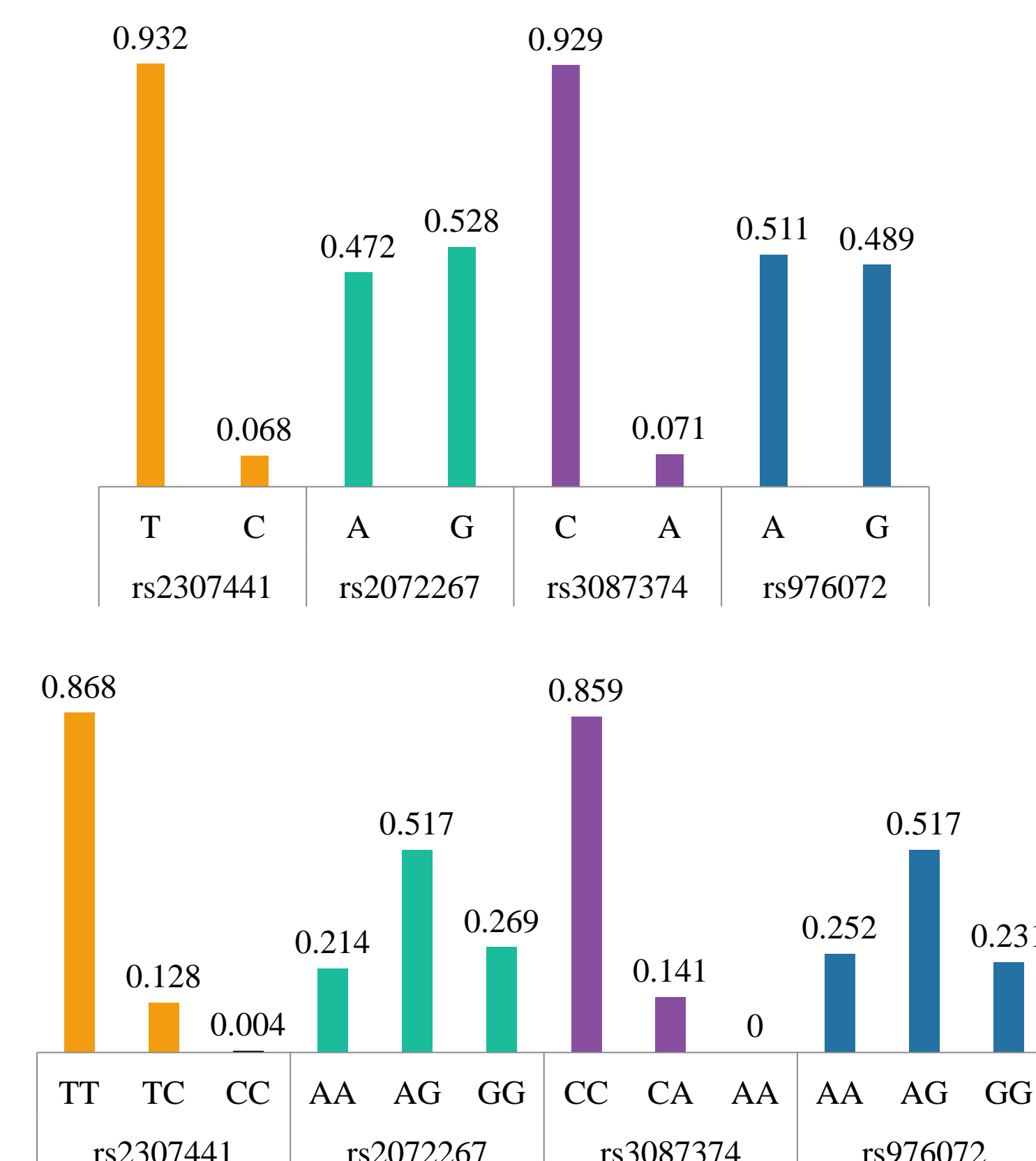
T - Pathological tumor size; G - Tumor differentiation grade; L - Lymphatic invasion; V - Vascular invasion; N - Lymph node involvement; ER - Estrogen receptors; PR - Progesterin receptors; HER2 - Human epidermal growth factor receptor 2.

Clinical disease characteristics

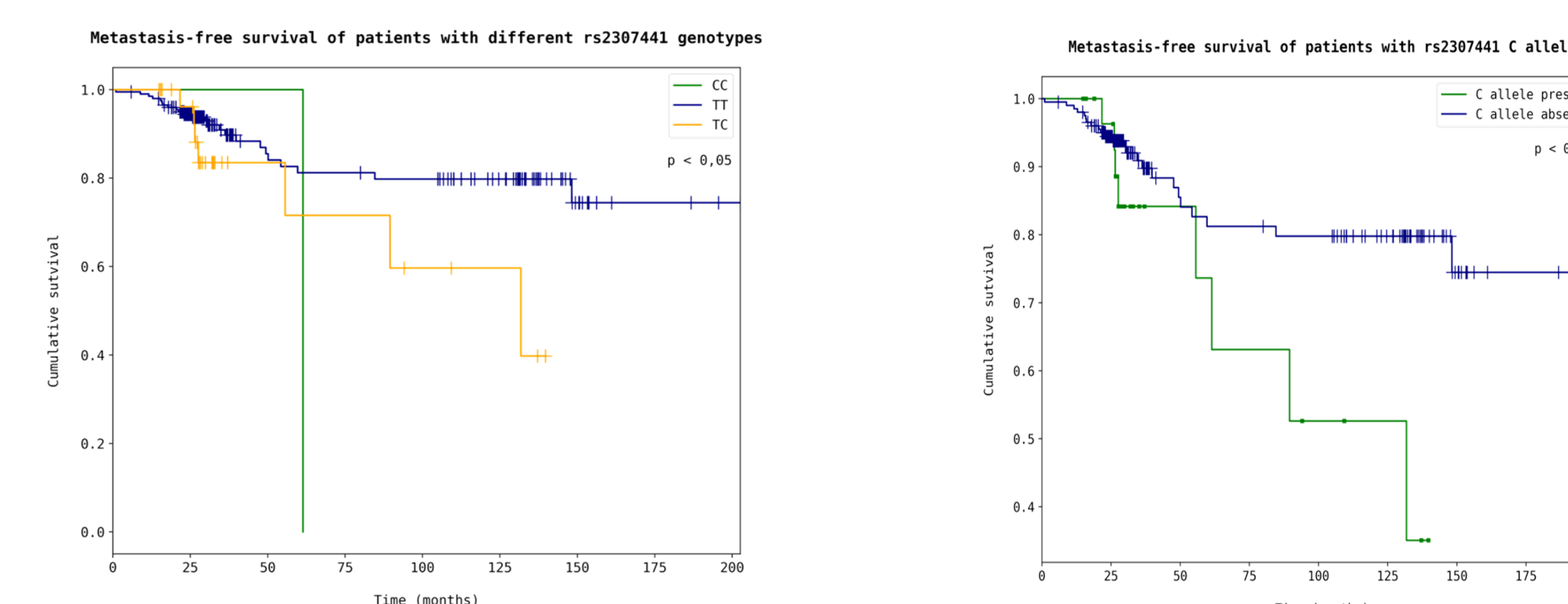


- ▷ Most of the patients presented without progression or metastasis and with deaths for only 9.4% of all patients.

POLG genotypes and alleles distribution



Survival analysis



- ▷ Patients with rs2307441 CT genotype presented with a tendency of shorter MFS when compared to the patients with TT genotype (p=0.051)
- ▷ The C allele holders had approximately 2.5 higher probability of shorter MFS than those without C allele (HR = 2.495; CI 1.114-5.587; p=0.026)

Conclusions

Our data indicate that nuclear DNA variations in *POLG* (rs2307441, rs2072267) gene play an important role for in breast cancer patients

Key words

POLG, SNV, breast cancer, tumor phenotype, outcome