

# Role of RIZ1 delP 704 polymorphism in metastatic breast cancer patients treated with exemestane

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**Background:** Estrogens sustain growth of estrogen receptor positive (ER+) breast cancer (BC) cells. ER activity is enhanced by several co-activators, including RIZ1, encoded by the tumor suppressor *PRMD2* gene. RIZ1 is a zinc finger protein representing a specific ER $\alpha$  co-activator. ER $\alpha$  function is strongly enhanced by RIZ1, which promotes optimal estrogen response in several tissues. Due to its critical role, polymorphisms on this gene, as well as on *ESR1* gene, coding for ER $\alpha$ , could affect estrogen activity. Estrogens are synthesized by aromatase, which is the target of several inhibitor drugs, including exemestane. Unfortunately, in a subset of ER+ MBC patients, exemestane is not effective and, in some cases, even toxic. Several germ-line single nucleotide polymorphisms (SNPs) have been described in *PRMD2* and *ESR1* genes. SNPs in this estrogens pathway, by affecting estrogens activity, could be responsible for the different treatment outcomes.

**Aims:** Our study aimed at investigating the predictive value of SNPs on *PMRD2* and *ESR1* genes in terms of Response Rate (RR) in ER+ MBC patients treated with exemestane as first-line hormone therapy.

**Methods:** This multicenter study enrolled 302 patients with ER+ MBC or locally-advanced BC with complete clinical data. For RR assessment, patients having a Complete Response (CR) or a Partial Response (PR) were compared with patients showing a Stable Disease (SD) or Progression Disease (PD) according to RECIST criteria. Multivariate statistical analyses for the associations between SNPs and clinical outcome were performed by logistic regression model. 95%CI and p-values underwent an internal validation by re-sampling through the application of bootstrap analysis with 1000 replications.

**Results:** In this study we found an association between RR and the insertion/deletion (indel) polymorphism RIZ1\_delP704 (rs2308040), which causes the insertion of a Proline residue at position 704 of the protein. Indeed, according to the recessive model, patients bearing the homozygous insertion (Pro/Pro) genotype had a higher probability of being responsive to exemestane therapy (OR<sub>DOM</sub>=2.23, 95%CI 1.25-4.00, p=0.007) than Del/Del or Del/Pro patients.

**Conclusions:** This study pointed out the predictive role of RIZ1\_delP704 (rs2308040) Pro/Pro genotype in the RR of MBC patients treated with the aromatase inhibitor exemestane. Once validated, this pharmacogenetic marker might be useful for treatment personalization.