

Two Gene Prognostic Model with ERCC1 and BRCA1 in NSCLC Patients Treated With Platinum Doublets

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Background/Aims: Among genes involving DNA repair pathway, we found that germline variation of ERCC1 codon 118 or a haplotype of BRCA1 was associated with differences in outcome in NSCLC Pts. treated with platinum doublets. Approach with multiple genes could be able to generate a more robust and useful model for clinicians to potentially utilize as a guide in patient assignment. **Methods:** The Lung Cancer Cohort of Inha University Hospital has constructed a database system that includes clinical information and matched peripheral blood DNA. We chose 300 patients with advanced-stage disease who were treated with more than two cycles of platinum-based chemotherapy as a first-line treatment, who underwent full follow-up at our hospital, and whose peripheral blood lymphocytes were available for analysis. These 300 consecutive NSCLC patients chosen gave their written informed consent and agreed to the purposes of the study. **Results:** The median age was 63 years. Histologically, 139 (46.3%) of the patients had squamous cell carcinomas and 137 (45.7%) had adenocarcinomas. Median survival time of Pts. was 13.0 months. We observed no significant association between ten SNPs (Q504K, IVS5-425C/T, IVS5+33C/A, IVS4+86A/G, N118N, and -284G/T for ERCC1; S1613G, IVS13-1893A/C, IVS12-1207 C/T, and IVS12112C/A for BRCA1) and overall survival. In gene-gene interaction analyses between ERCC1 and BRCA1, Pts. with variant genotype of ERCC1 N118N and wild-type genotypes of four tagging SNPs of BRCA1 showed statistically significant shorter survival in both analyses of Kaplan Meier and Cox's proportional hazard model (adjusted HR, 95% CI: 1.984, 1.103 to 3.569 for N118N+1613S/S; 1.889, 1.089 to 3.277 for N118N+IVS13-1893A/A; 2.205, 1.077 to 4.517 for N118N+IVS12-1207C/C; 2.248, 1.095 to 4.615 for N118N+ IVS12112C/C). In haplotype to haplotype analyses, Pts. with variant type haplotype (ATG) of ERCC1 and wild-type haplotype (AACC) of BRCA1 showed statistically significant shorter survival (adjusted HR, 95% CI: 2.643, 1.416 to 4.932). **Conclusions:** This result suggests that the model of tagging SNPs of ERCC1 and BRCA1 can give clinicians to useful guide in treating NSCLC Pts. with platinum doublets.

