

Prognostic Significance Of Beclin1 and TGF- β 1 In Ovarian Cancer

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Abstract

Ovarian cancer is a gynecological malignancy with a high mortality rate. Autophagy is lysosomal degradation of damaged subcellular structures which is known as type II programmed cell death. Autophagy was initially thought to be a tumor-suppression mechanism and dysregulation of autophagy is suggested to be involved in tumor genesis. BECN1 is a tumor suppressor gene involved in the initiation of autophagy. It encodes Beclin-1 protein, which inhibits tumor growth, there is wide controversy about its role in initiation, promotion of tumor and prognostic importance of autophagic molecules. Transforming growth factor β 1 induce process of epithelial-mesenchymal transition (EMT), keeping epithelial cells more motile and invasive leading to cancer progression and metastasis.

Material and methods: Fifty Blocks of paraffin-embedded ovarian tissue were selected, representing cases diagnosed as ECO. The immunohistochemical staining procedure was done using Beclin 1 and TGF- β 1

Results: Parameters detect their expression and to correlate it with the different clinical parameters. Positive Beclin1 expression was observed in 54% and Positive TGF- β 1 staining was observed in 70% of patients tissue samples. Beclin1 significantly correlated with lower tumor grade ($P < 0.031$) and lower FIGO stage, $P = 0.01$, a significant association was observed between higher FIGO stage and TGF- β 1 expression. All metastatic cases were positive for TGF- β 1 versus 27.3% of metastatic cases positive for beclin.

Beclin1 showed significant correlation with non-recurring disease, $P = 0.005$ and was associated with less mortality $P = < 0.001$. TGF- β 1 was significantly associated with higher mortality rates and relapsing disease, $P = 0.015$, $OP = 0.005$.

Conclusion. Beclin1 protein could be considered a good prognostic factor in OC cases while TGF- β 1 considered adverse factor which could be of benefit in OC molecular targeting therapy