

عنوان مقاله:

Expression analysis of CDYF and CDFF transcripts in Iranian breast cancer patients

محل انتشار:

چهارمین کنگره بین المللی و شانزدهمین کنگره ملی ژنتیک (سال: 1399)

تعداد صفحات اصل مقاله: 2

نویسندگان: Ali Sattari - Department of Medical Genetics, Shahid Beheshti University of Medical Sciences, Tehran, Iran

Mohammad Taheri - Department of Medical Genetics, Shahid Beheshti University of Medical Sciences, Tehran, Iran

خلاصه مقاله:

Background and Aim: The importance of cancer stem cells (CSCs) in initiation and progression of breast cancer has been well established. This population of cells is characterized by high expression of CDFF and low expression of CDYF. However, the relative abundance of CDYF and CDFFtranscripts in breast cancer tissues and adjacent noncancerous tissues (ANCTs) has not been quantified yet. Methods: In the present investigation, we assessed expression of CDYF and CDFF at transcript level in breast cancer tissues and ANCTs in association with clinical determinants of patients' outcome and parameters that predict response to therapeutic options. Results: There was no significant difference in expression of CDYF and CDFF in breast cancer tissues compared with ANCTs (Expression ratios: 1.0 P and o.Af, P values: o.Af and o.f1, respectively). However, CDff expression was associated with tumor size in a way this gene was up-regulated in all of small sized (≤Y cm) tumors compared with the corresponding ANCTs (P value=o.of). Besides,CDff expression was significantly higher in tumors with extracapsular nodal extension compared with those without extension (P=∘.∘F). Expression of CDYF was higher in grade ™ tumors compared with grade Y tumors (P=o.of). Expression levels of CDYF and CDFF were correlated with each other in ANCTs but not in tumoral .tissues.Conclusion: The current study shows another aspect of CSC markers in the development of breast cancer

کلمات کلیدی: CD۴۴, CD۲۴, breast cancer

لینک ثابت مقاله در پایگاه سیویلیکا:

https://civilica.com/doc/1195567

