

Differential expression of E-cadherin gene in human neuroepithelial tumors

F.J.N. Motta^{1,2}, E.T. Valera³, A.K.B. Lucio-Eterovic³, R.G.P. Queiroz³, L. Neder⁴, C.A. Scrideli³, H.R. Machado⁵, C.G. Carlotti-Junior⁵, S.K.N. Marie⁶ and L.G. Tone³

¹Departamento de Genética, Faculdade de Medicina de Ribeirão Preto, Universidade de São Paulo, Ribeirão Preto, SP, Brasil
²Universidade Federal do Piauí, Campus Ministro Reis Veloso, Parnaíba, PI, Brasil
³Departamento de Puericultura e Pediatria
Faculdade de Medicina de Ribeirão Preto, Universidade de São Paulo, Ribeirão Preto, SP, Brasil
⁴Departamento de Patologia, Faculdade de Medicina de Ribeirão Preto, Universidade de São Paulo, Ribeirão Preto, SP, Brasil
⁵Departamento de Anatomia e Cirurgia, Faculdade de Medicina de Ribeirão Preto, Universidade de São Paulo, Ribeirão Preto, SP, Brasil
⁶Departamento de Neurologia, Faculdade de Medicina, Universidade de São Paulo, São Paulo, SP, Brasil

Corresponding author: F.J.N. Motta E-mail: motta@ufpi.edu.br

Genet. Mol. Res. 7 (2): 295-304 (2008) Received December 18, 2007 Accepted February 8, 2008 Published April 8, 2008

ABSTRACT. Cadherins are cell-to-cell adhesion molecules that play an important role in the establishment of adherent-type junctions by mediating calcium-dependent cellular interactions. The *CDH1* gene encodes the transmembrane glycoprotein E-cadherin which is important in maintaining homophilic cell-cell adhesion in epithelial tissues. E-cadherin interacts with catenin proteins to maintain tissue architecture. Structural defects or loss of expression of E-cadherin have been reported as a common feature in several human cancer types. This study aimed to evaluate the expression of E-cadherin and their correlation with clinical features in microdissected brain tumor samples from 81 patients, divided into 62 astrocytic tumors grades I to IV and 19 medulloblastomas, and from 5 white matter non-neoplasic brain tissue samples. E-cadherin (CDH1) gene expression was analyzed by quantitative real-time polymerase chain reaction. Mann-Whitney, Kruskal-Wallis, Kaplan-Meir, and logrank tests were performed for statistical analyses. We observed a decrease in expression among pathological grades of neuroepithelial tumors. Non-neoplasic brain tissue showed a higher expression level of CDH1 gene than did neuroepithelial tumors. Expression of Ecadherin gene was higher in astrocytic than embryonal tumors (P = 0.0168). Low-grade malignancy astrocytomas (grades I-II) showed higher CDH1 expression than did high-grade malignancy astrocytomas (grades III-IV) and medulloblastomas (P < 0.0001). Non-neoplasic brain tissue showed a higher expression level of CDH1 gene than grade I malignancy astrocytomas, considered as benign tumors (P = 0.0473). These results suggest that a decrease in E-cadherin gene expression level in high-grade neuroepithelial tumors may be a hallmark of malignancy in dedifferentiated tumors and that it may be possibly correlated with their progression and dissemination.

Key words: Cancer; Neuroepithelial tumors; *CDH1* expression; Real-time polymerase chain reaction

Genetics and Molecular Research 7 (2): 295-304 (2008)