

DIFFERENCE GEL ELECTROPHORESIS (DIGE) OF SERUM FROM PATIENTS WITH GRADE IV ASTROCYTOMAS AND CONTROL SUBJECTS

Fischer, J.S.G.¹, Carvalho, P.C.², Carvalho, M.G.C.³, Fonseca, C.O.⁴, Neves-Ferreira, A.G.C.⁵, Albuquerque, L. M.⁵, Trugilho, M.R.O.⁵; Perales, J.⁵; Domont, G. B.¹

¹Department of Biochemistry – Institute of Chemistry – Federal University of Rio de Janeiro (UFRJ) and the Proteomics Network of Rio de Janeiro, Rio de Janeiro, Brazil; ²Systems Engineering and Computer Science Program – UFRJ, Rio de Janeiro, Brazil; ³Institute of Biophysics – UFRJ, Rio de Janeiro, Brazil; ⁴Institute of Biology – Fluminense Federal University, Rio de Janeiro, Brazil; ⁵Department of Pharmacodynamic and Physiology, Oswaldo Cruz-Fiocruz Institute, Rio de Janeiro, Brazil.

More than 60% of primary brain tumors are of glial origin, occurring at any age. Grade IV astrocytoma is the most malignant type and can quickly spread to other brain parts. The identification of new biomarker panels holds promise for early detection of this cancer being crucial for a successful treatment. Our aim was to use differential in gel electrophoresis (DIGE) to search for differences in protein expression level between seven control subjects and seven astrocytoma patients. The Ettan DIGE System and CyDye fluors were used to simultaneously separate three samples in each 2-D gel. The gels were scanned with the Typhoon™, and the DeCyder software used to locate putative biomarkers. The paired Student t test indicated 21 differentially expressed proteins ($p < 0.01$) that are being identified by mass spectrometry.

Acknowledgements: FAPERJ, CNPq, CAPEs, Ary Frauzino Foundation and Genesis Molecular Biology Laboratory

Key words: DIGE, astrocytoma, cancer, biomarker