Protein Profiling of *Escherichia coli* in response to Carbon source variation

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ABSTRACT

The response of *E.coli* to carbon source variation is well understood and it is commonly used as model biological system when validating an analytical method. We have characterized the proteins isolated from *E.coli* grown in glucose, lactose and acetate by using two dimensional polyacrylamide gel electrophoresis (2D-PAGE). The quantitative results obtained from our study (unpublished data) were comparable to other existing protein profiling and transcriptional profiling approaches.

Key words: Carbon metabolism, *E.coli*, Protein profiles.