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(54) Title: BREAST CANCER EXPRESSION PROFILING

(57) Abstract: The present invention relates to a method for analyzing cancer.e.g., breast cancer comprising detection of differential expression of at least one of the 16 genes encoding serine/threonine kinases listed in Table 1, or of said 16 genes, and to a polynucleotide library comprising at least one said 16 genes. This finds use in the development of novel applications, in particular in the development of prognosis or diagnostic of breast cancer or for monitoring the treatment of a patient with a breast cancer.

WO 2009/083780 A8