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(54) Title: GENE SIGNATURES FOR USE WITH HEPATOCELLULAR CARCINOMA

(57) Abstract: The present invention provides a method for predicting prognosis of hepatocellular carcinoma patients based on measurement of the relative level of expression of a combination of 15 immune genes of interest, or a subset thereof, in the tumors of such patients. Tumor material can come from surgical resection or biopsy. The relative gene expression information may be combined in an algorithm. The signature can be used by itself or in combination with other information such as stage information.



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