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HUANG, Andrew [US/US]; 4841 Moriah Hill Road,
Durham, NC 27707 (US).

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(74) Agents: **ZELANO, Anthony, J.** et al.; Millen, White, Ze-
lano & Branigan, P.C., 2200 Clarendon Boulevard, Suite
1400, Arlington, VA 22201 (US).

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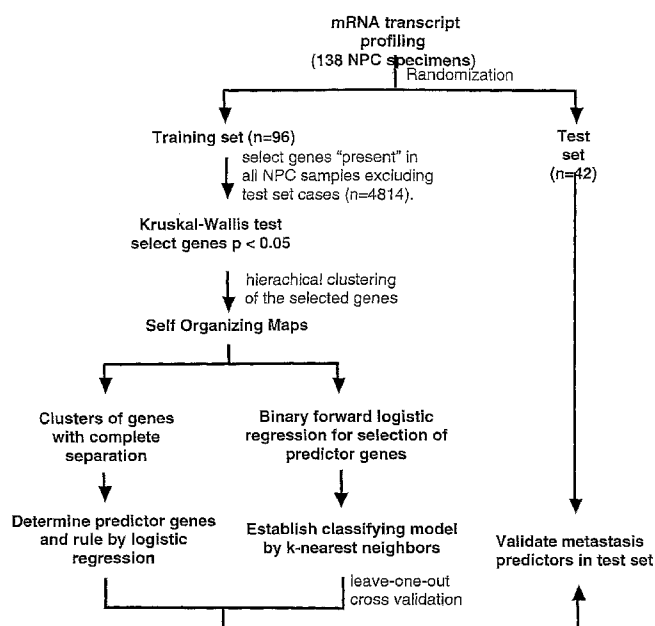
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SYNTHETIC RUBBER CORPORATION** [CN/CN];
7Th Floor, No. 113, Section 2, Zhongshan North Road,
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(72) Inventors; and

(75) Inventors/Applicants (for US only): **KAO, Kuo-jiang**

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(54) Title: GENE EXPRESSION PROFILING FOR IDENTIFICATION OF PROGNOSTIC SUBCLASSES IN NASOPHARYN-
GEAL CARCINOMAS

(57) Abstract: mRNA transcript profiling can be used to formulate molecular predictors of distant metastasis for primary NPCs. The predicted results are highly correlated with short metastasis-free and overall survival. Predictions are made using 52-genes based and 12- genes based predictors.

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