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# **EXHIBIT "A"**

The GeneMarkS software

# **Description:**

GeneMark.hmm is a genomic data analysis tool designed to predict the presence of protein coding regions in raw prokaryotic DNA sequence text. The program consists of a single executable program. Parameters of the statistical models that represent the statistical characteristics of protein coding and non-coding regions of an individual prokaryotic genome are determined by self-training algorithm of GeneMarkS.

(Although GeneProbe does not take obligation of the GeneMarkS support or maintenance, we will do our best to address your questions, please see http://www.geneprobe.net/contactus.htm for faster communication regarding academic distribution use Fax: +1 (404) 603–8875 or e-mail custserv@genepro.com)