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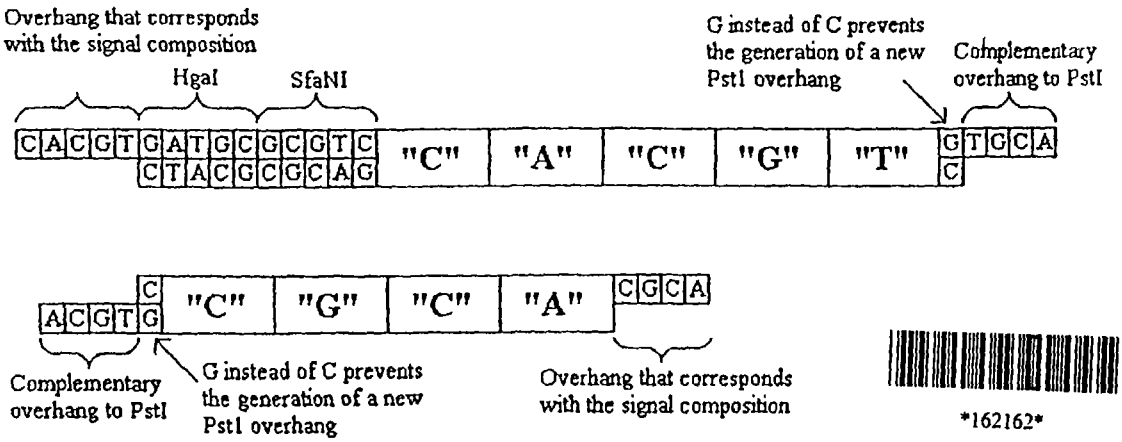
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(57) Abstract

The present invention provides a method of sequencing all or part of a target nucleic acid molecule by determining the sequence of a portion of said nucleic acid molecule in addition to information regarding the position of said portion and in particular provides a new method of sequencing involving the magnification of one or more bases of said bases to aid identification.