

### US006143496A

# United States Patent [19]

Brown et al.

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METHOD OF SAMPLING, AMPLIFYING AND QUANTIFYING SEGMENT OF **NUCLEIC ACID, POLYMERASE CHAIN** REACTION ASSEMBLY HAVING NANOLITER-SIZED SAMPLE CHAMBERS, AND METHOD OF FILLING ASSEMBLY

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[51] **Int. Cl.**<sup>7</sup> ...... **G01N 33/543**; G01N 33/68

**U.S. Cl.** ...... 435/6; 435/287.1; 435/287.2; [52] 435/288.3; 435/288.7; 436/164; 436/172; 436/518; 436/524; 436/527; 436/531; 436/805; 436/809; 422/58

435/288.3, 288.7; 436/164, 172, 518, 524, 527, 531, 805, 809; 422/58

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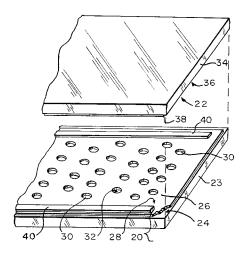
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#### **ABSTRACT** [57]

Methods of detecting and quantifying genomic nucleic acid molecule sequences are provided using the simultaneous amplification of a plurality of discrete nanoliter-sized samples. A miniaturized closed assembly is also provided for carrying out amplification of a nucleic acid molecule by polymerase chain reaction in multiple nanoliter-sized samples. Methods of filling miniaturized sample chambers are also provided as are methods for determining the number of template molecules in a sample by conducting replicate nucleic acid sequence amplification reactions on a set of terminally diluted samples and counting the number of positive amplification reactions. The methods can be used to detect a single starting nucleic acid target molecule.

### 17 Claims, 7 Drawing Sheets



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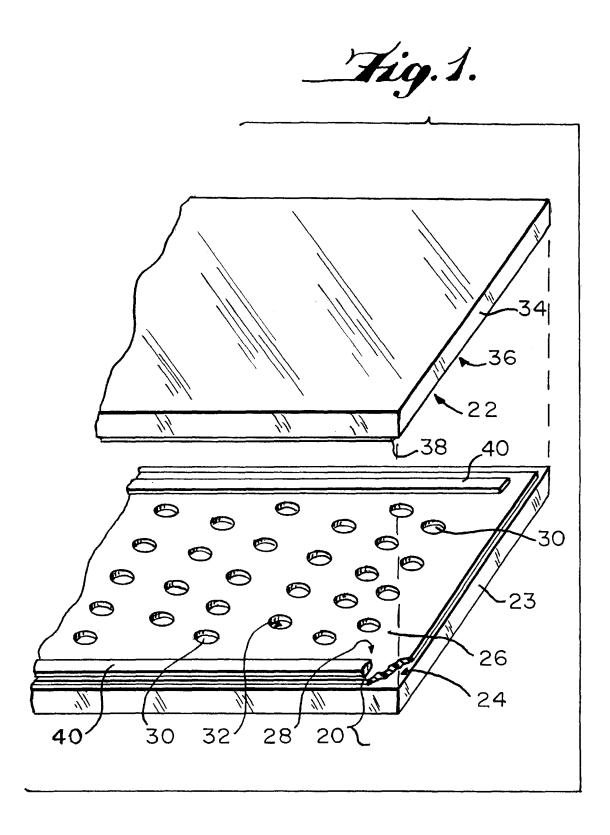
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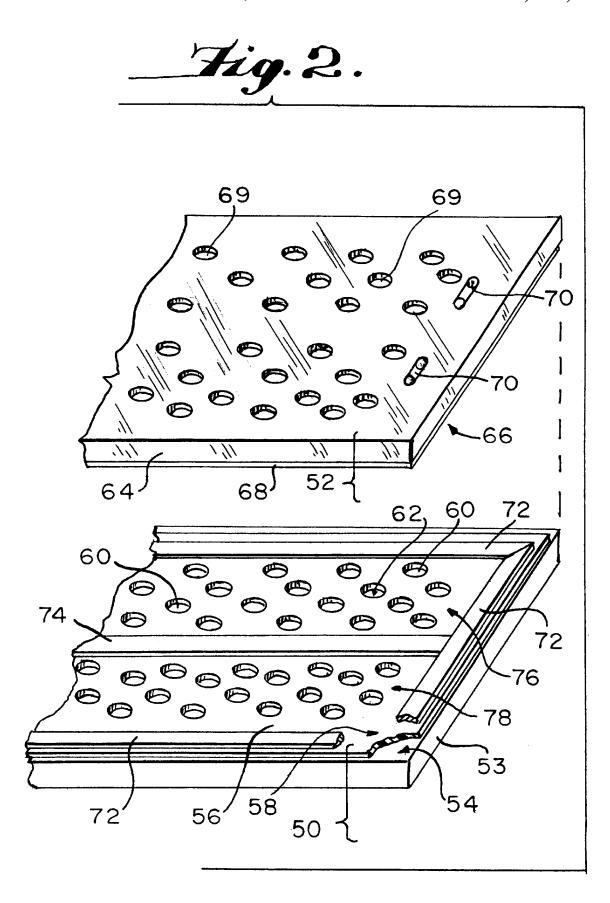
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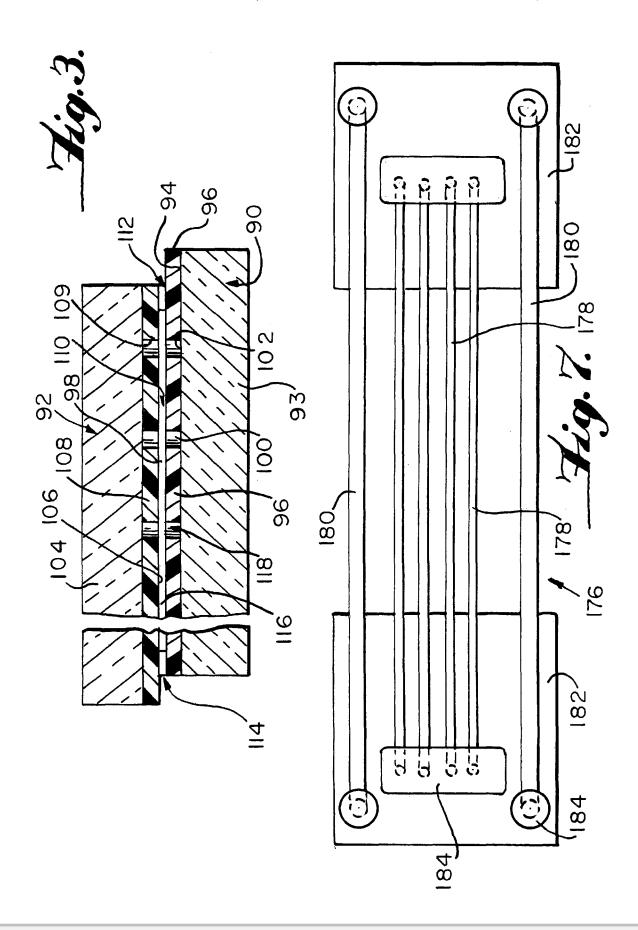
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