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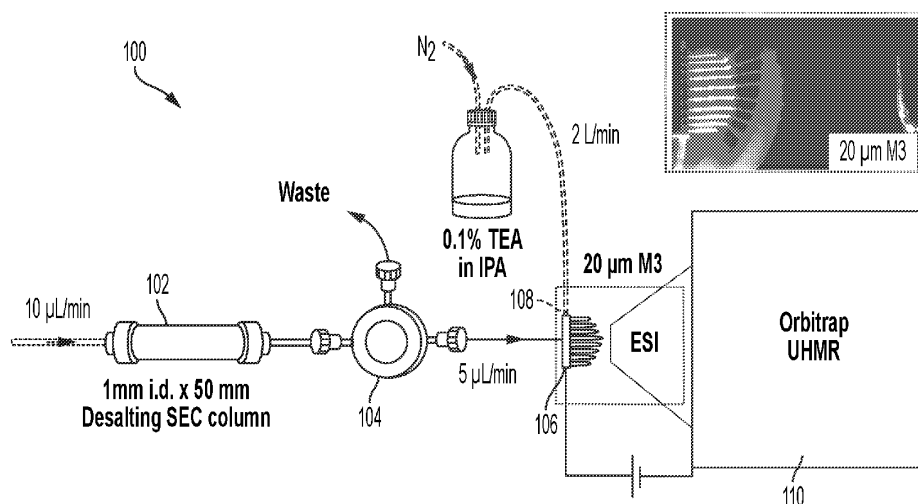


FIG. 2

(57) Abstract: Methods for determining the relative abundance of viral capsid components in a sample of viral particles are disclosed. In embodiments, methods for determining the relative abundance of empty capsids, partially-full capsids and full capsids (e.g., containing a heterologous nucleic acid molecule) of adeno-associated virus are disclosed.

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