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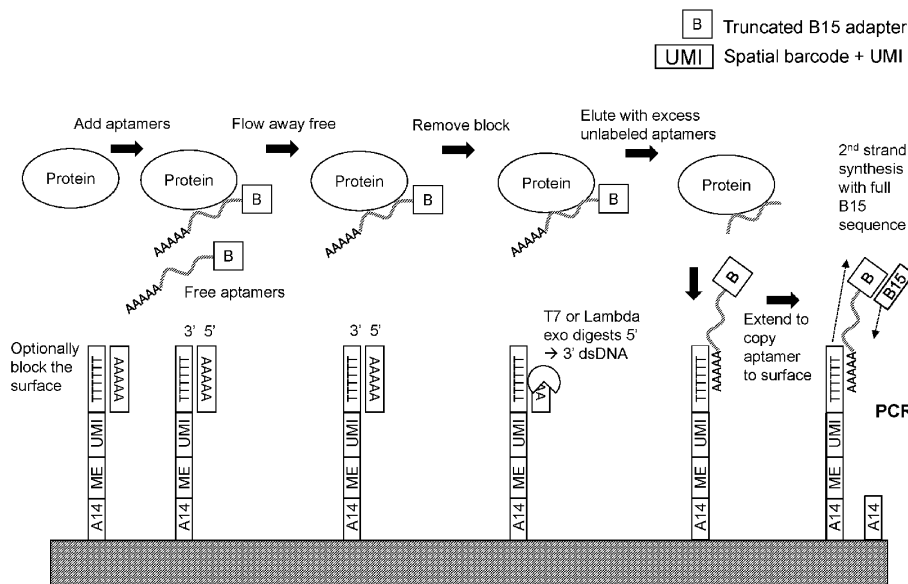


FIG. 1

(57) Abstract: The present disclosure relates, in general, to methods of preparing a spatial proteome and/or transcriptome sequencing library. The spatial proteome and/or transcriptome sequencing library from a biological sample is useful, in some aspects, to determine a genetic profile and help diagnose a subject who has or is at risk of having a disorder, and improve treatment of the subject.

[Continued on next page]

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