

Fig. 1A

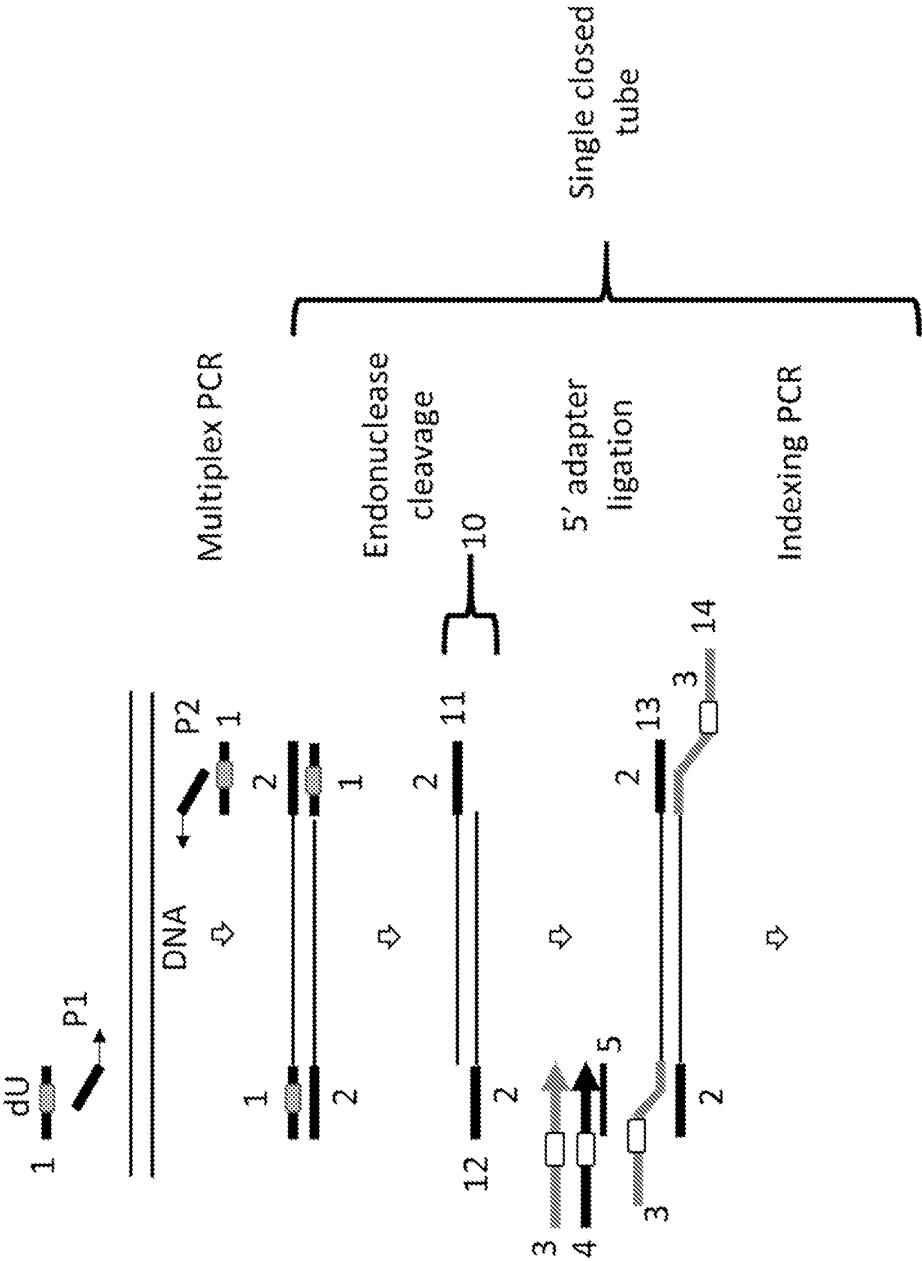


Fig. 1A (continued)

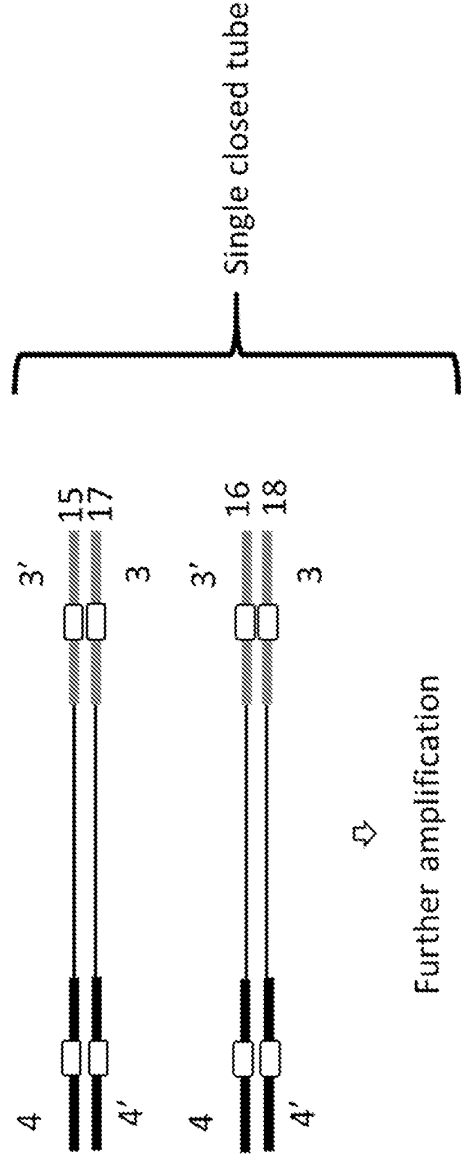


Fig. 1B

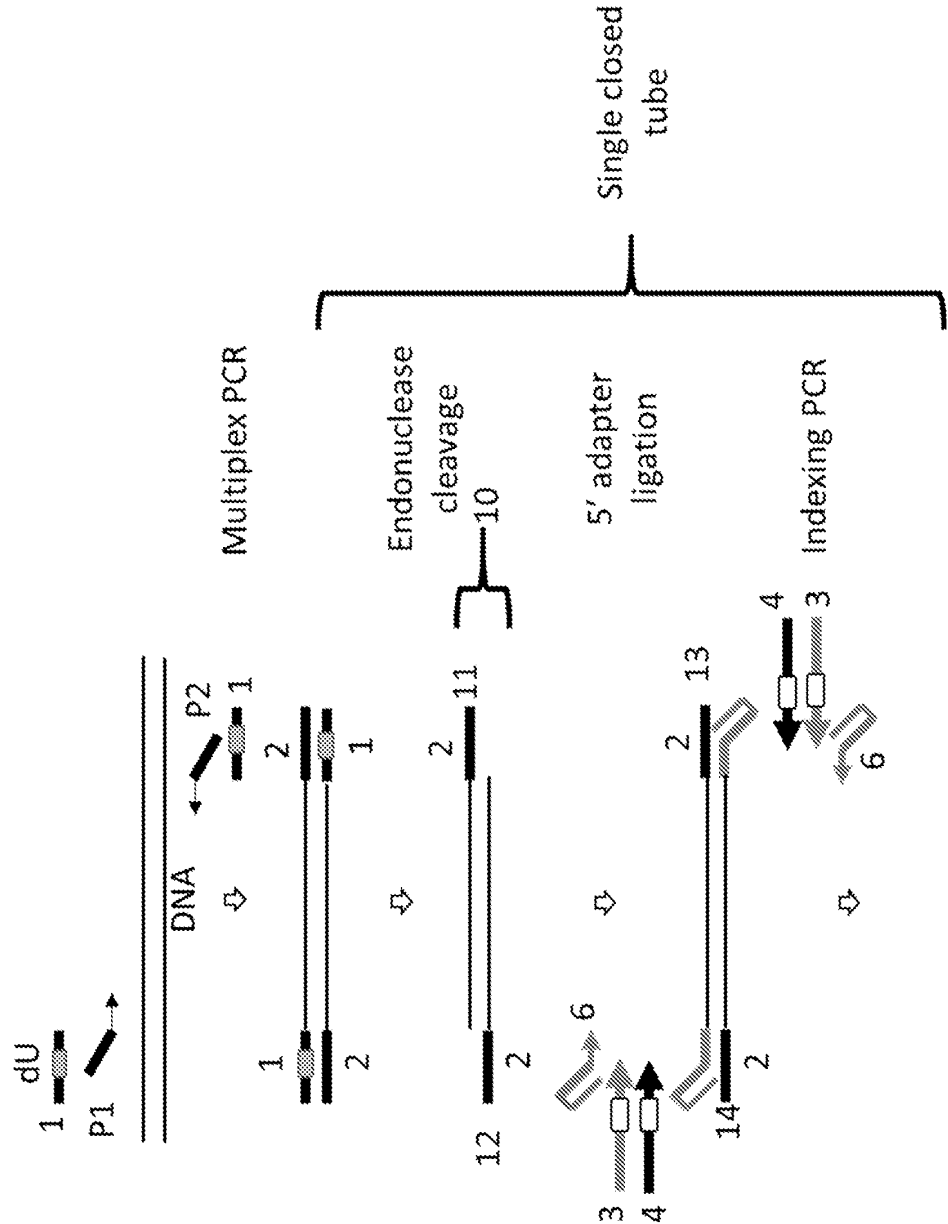


Fig. 1B (continued)

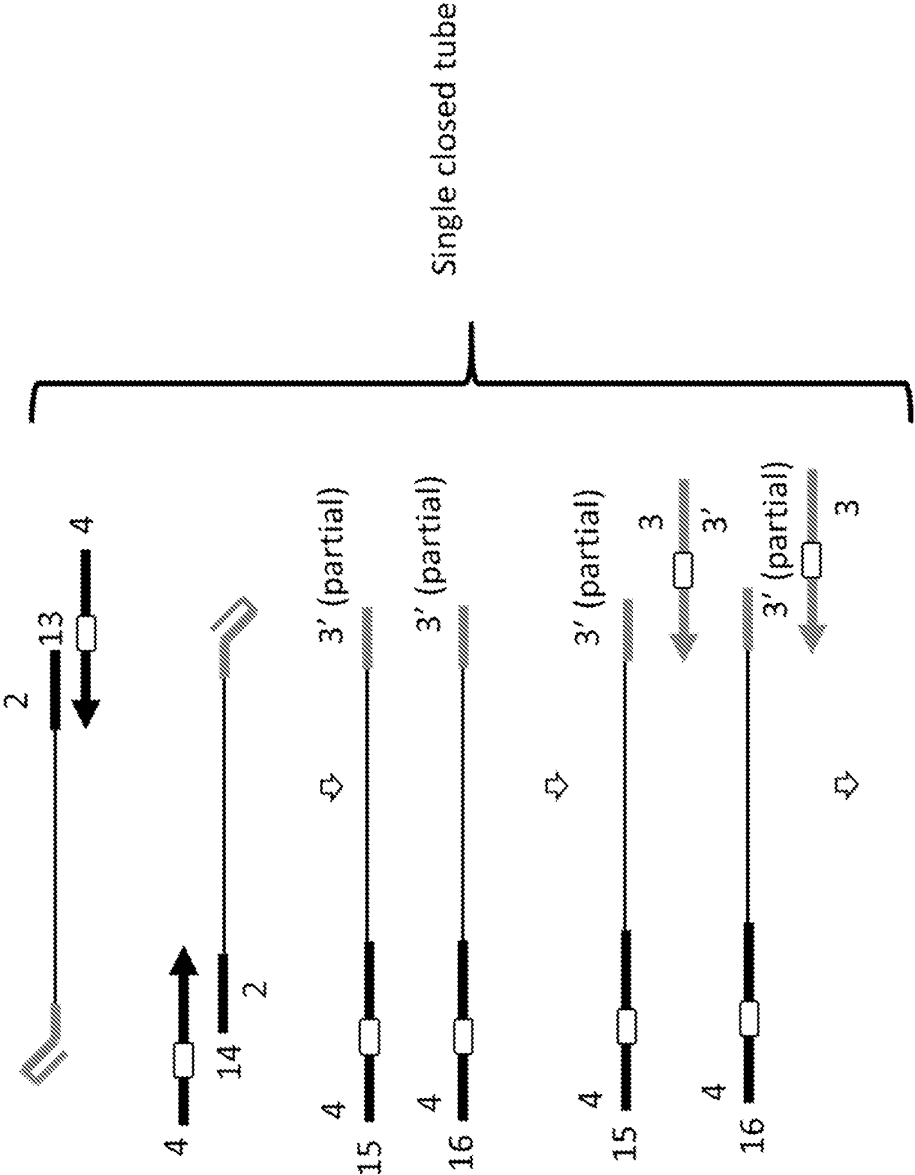
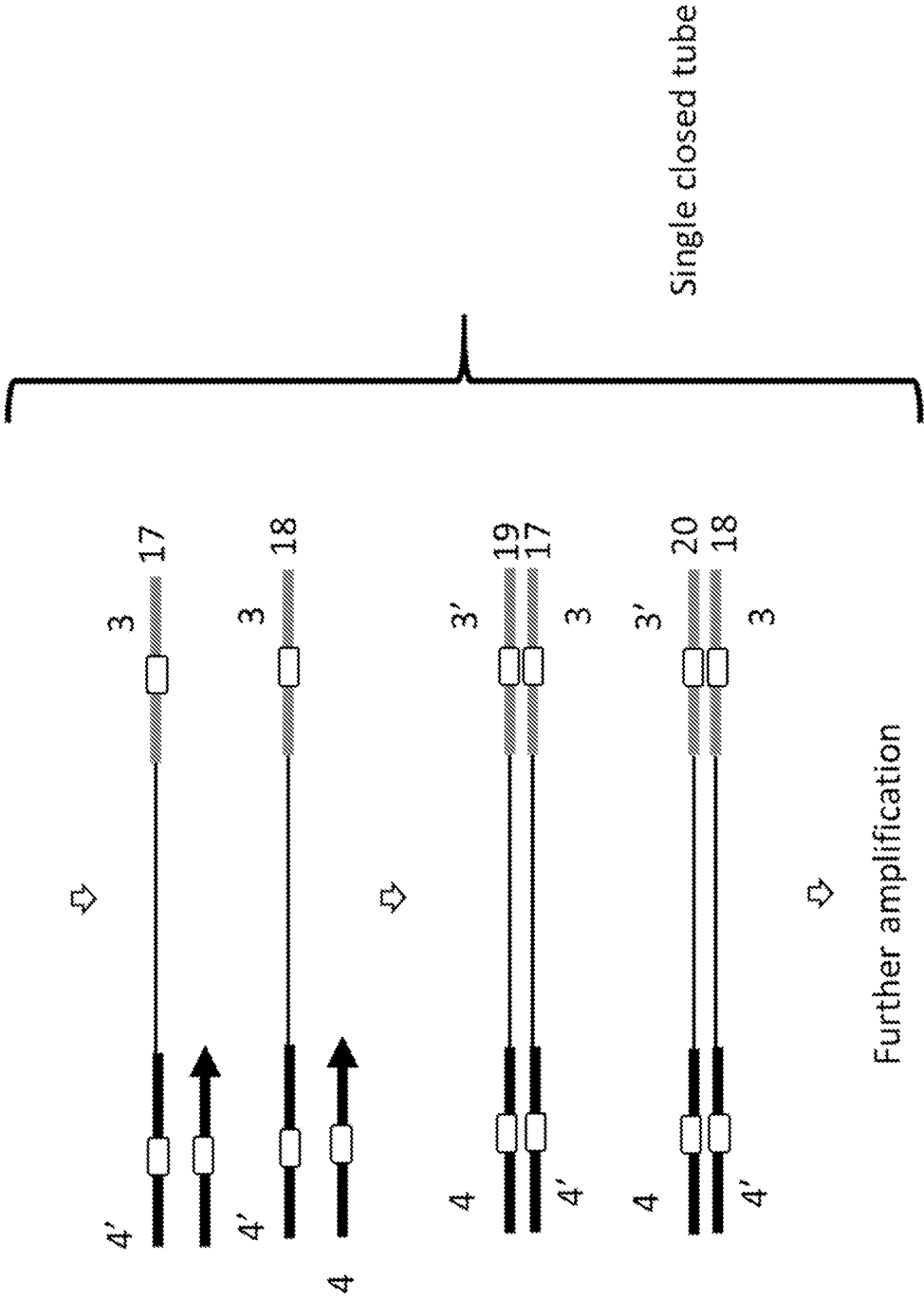
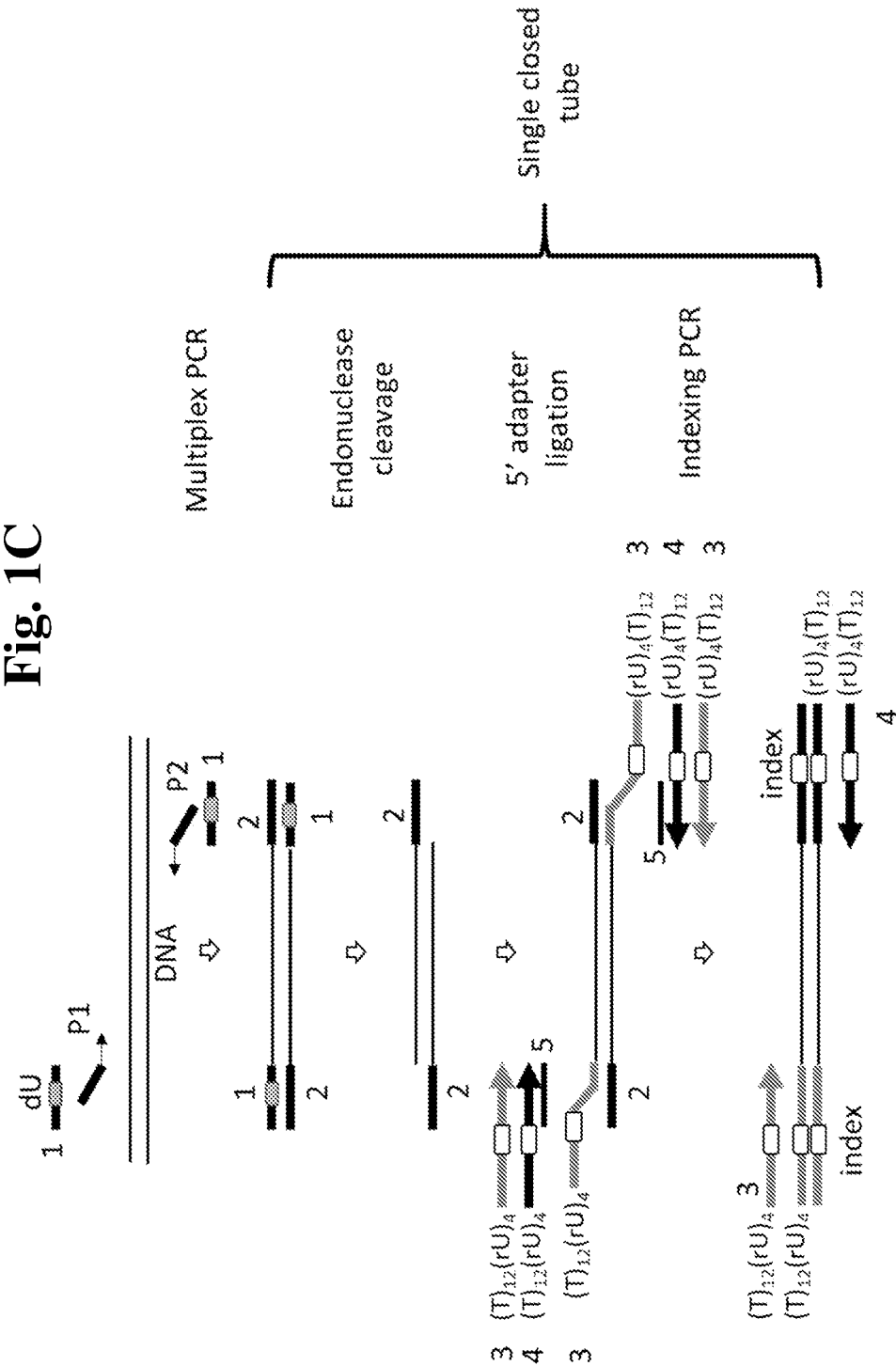
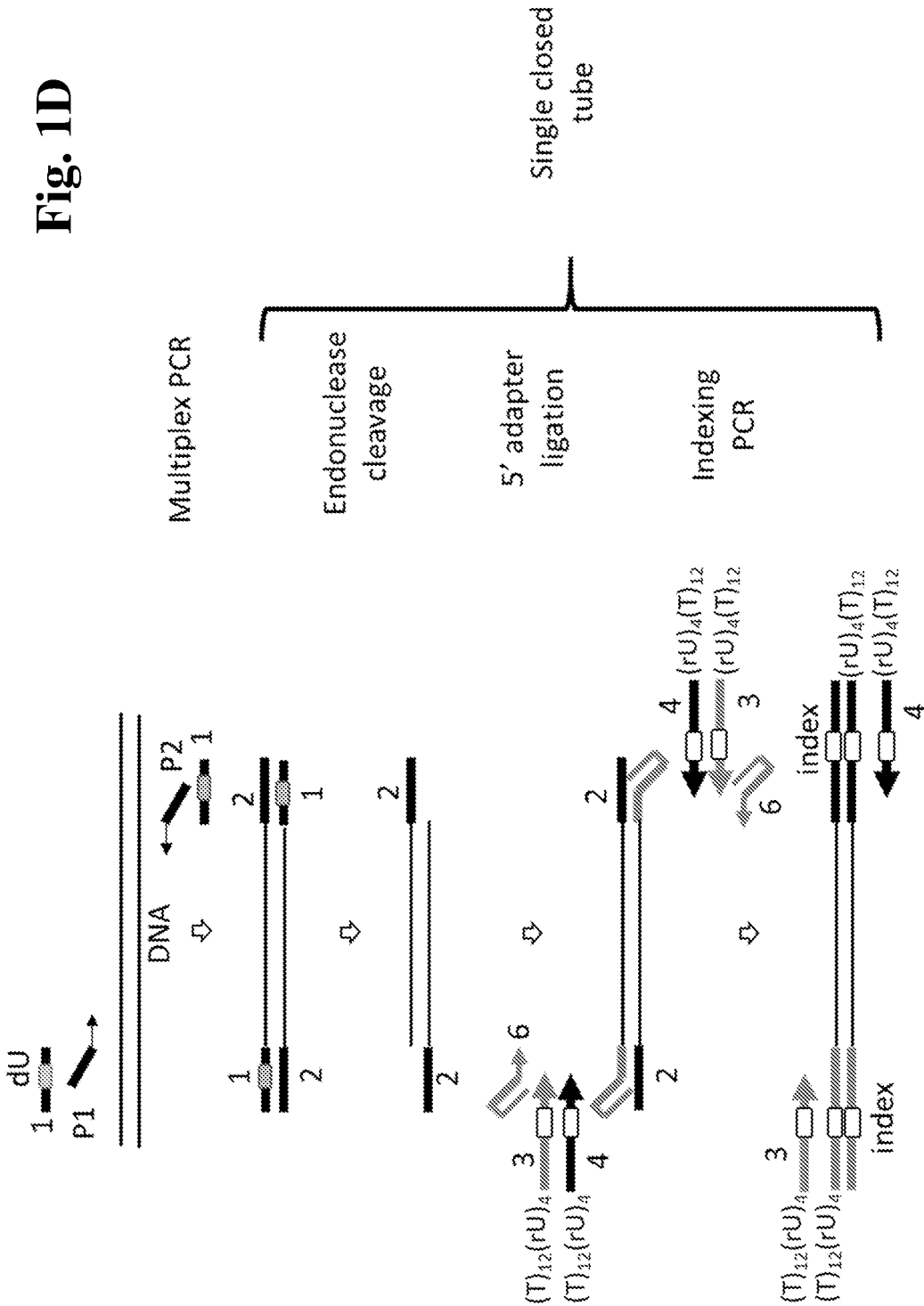
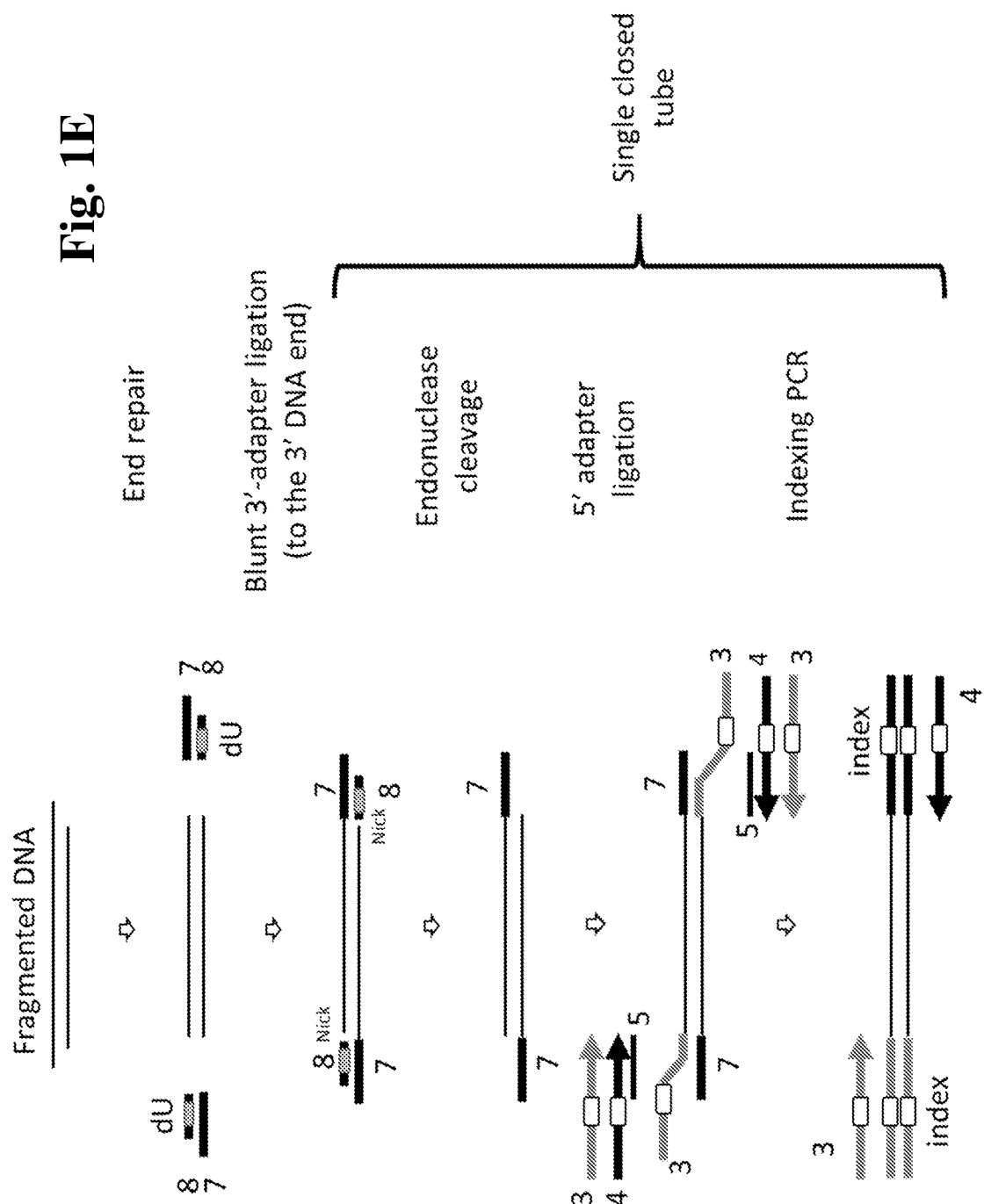


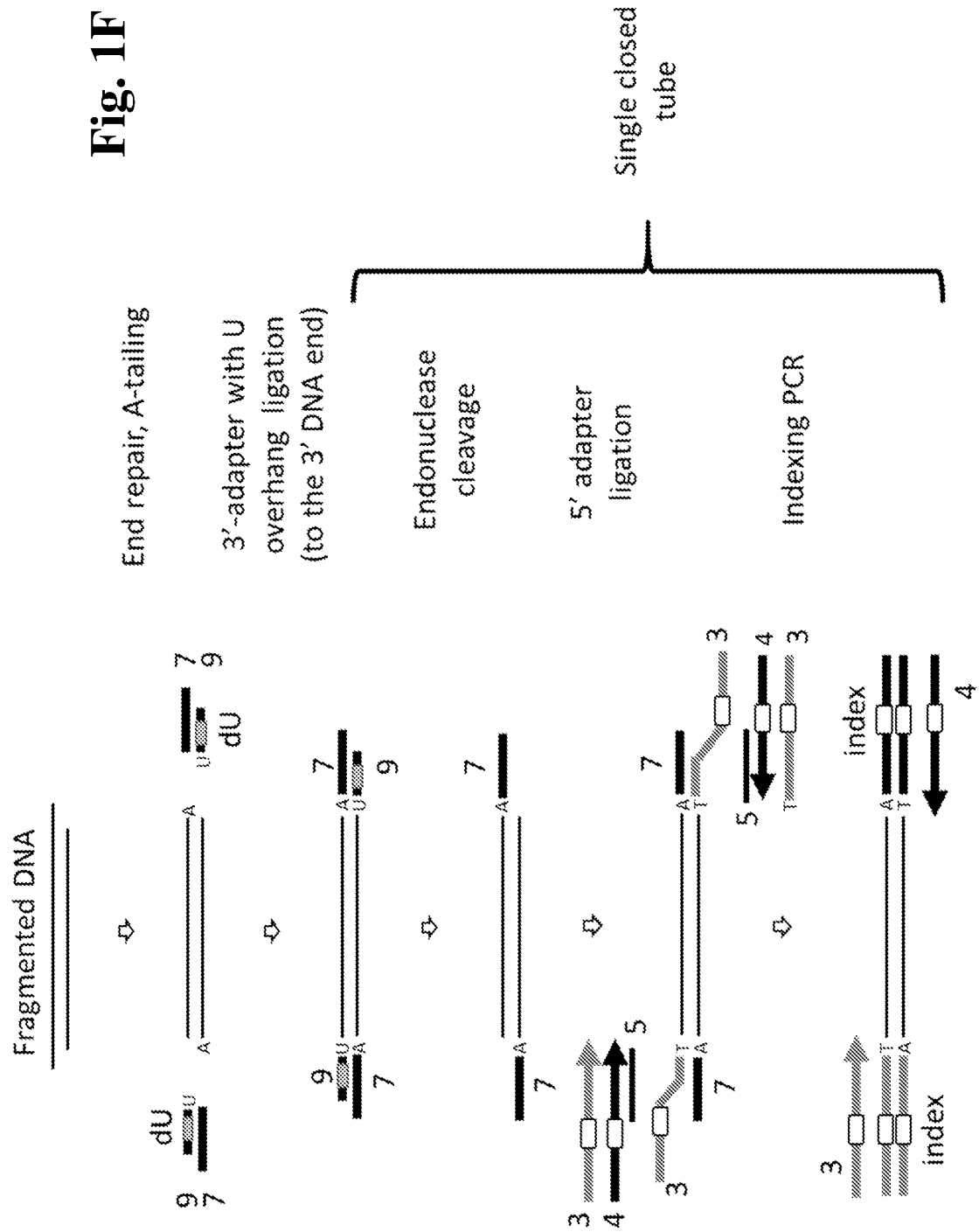
Fig. 1B (continued)

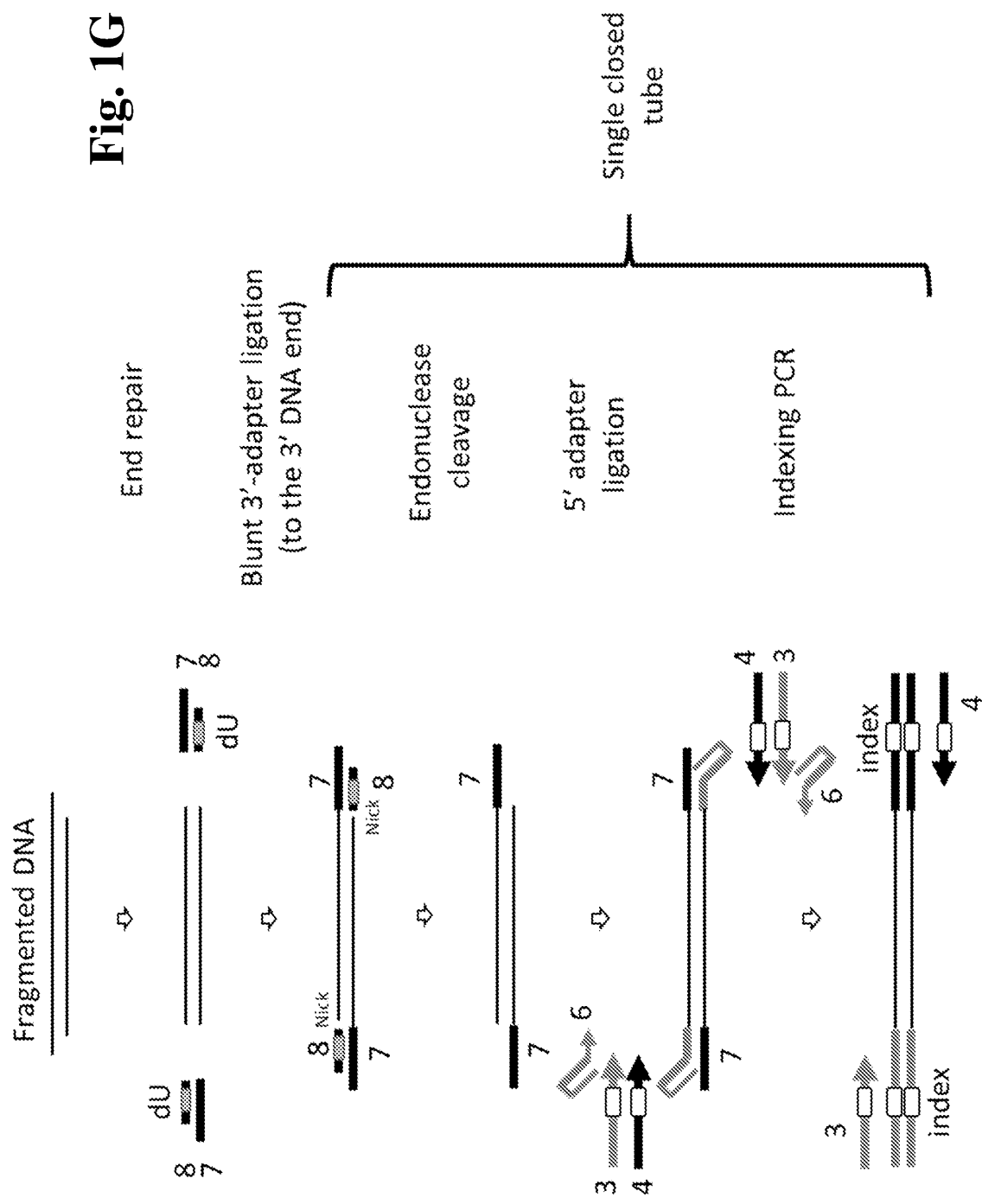












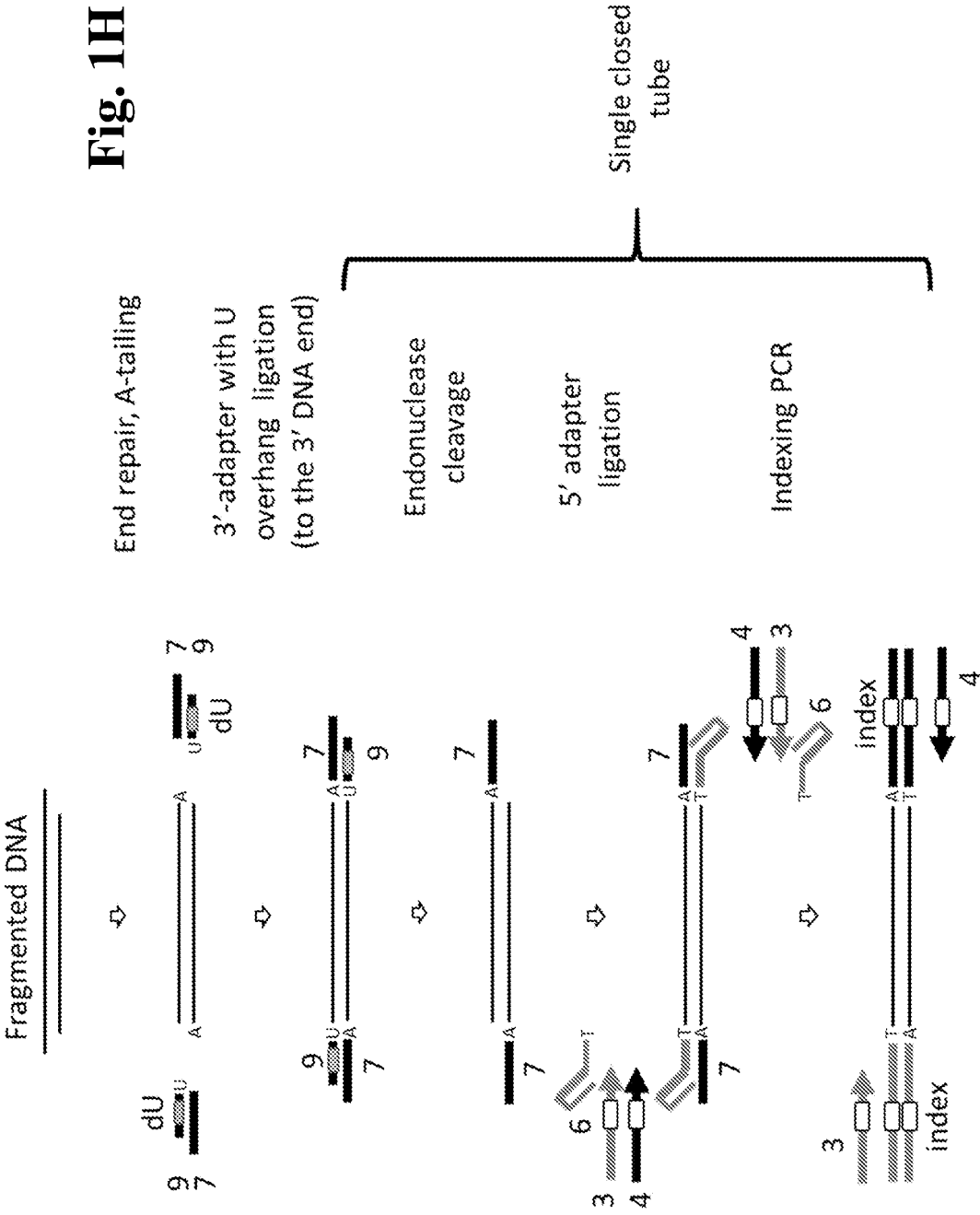
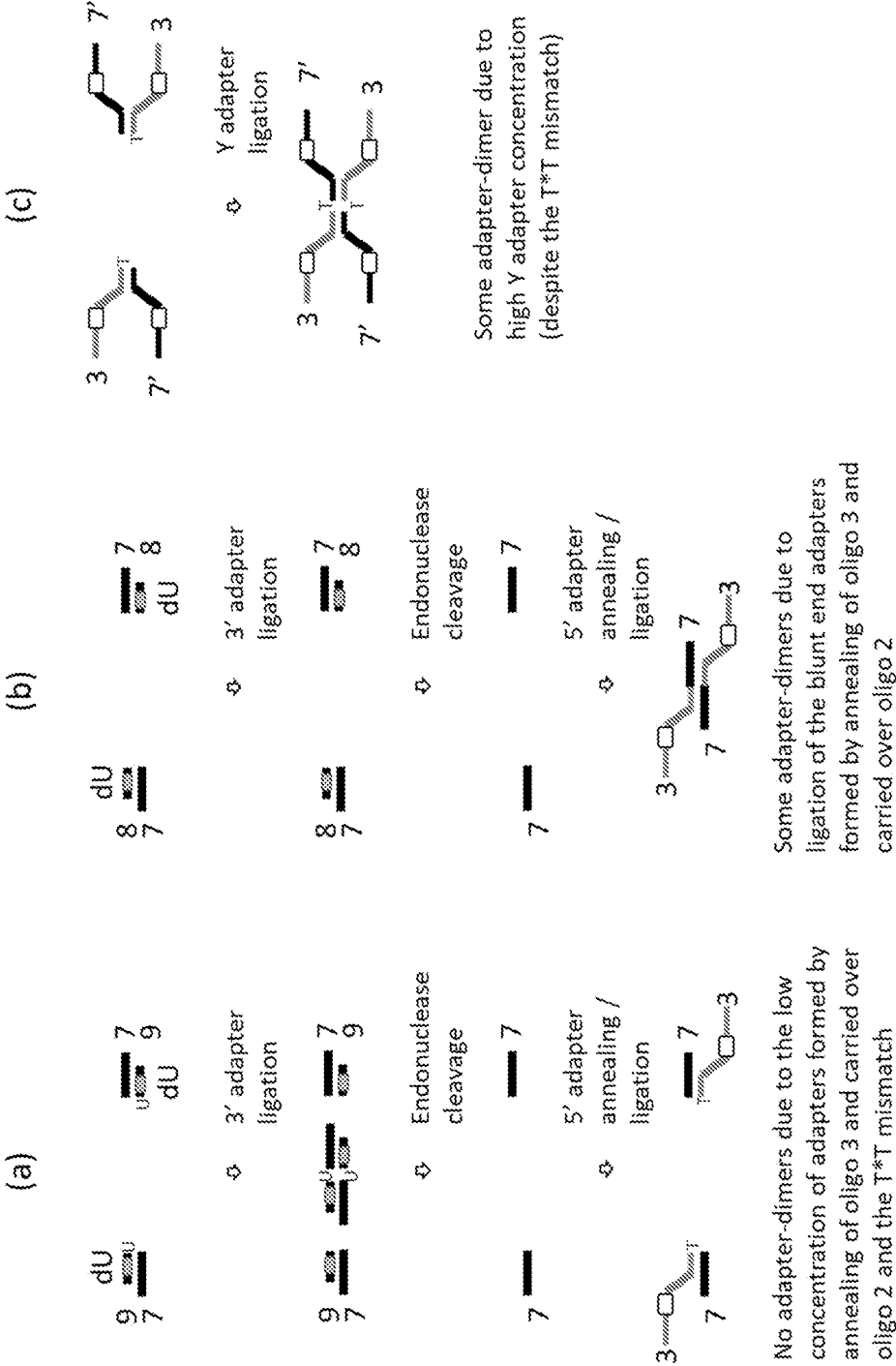


Fig. 11



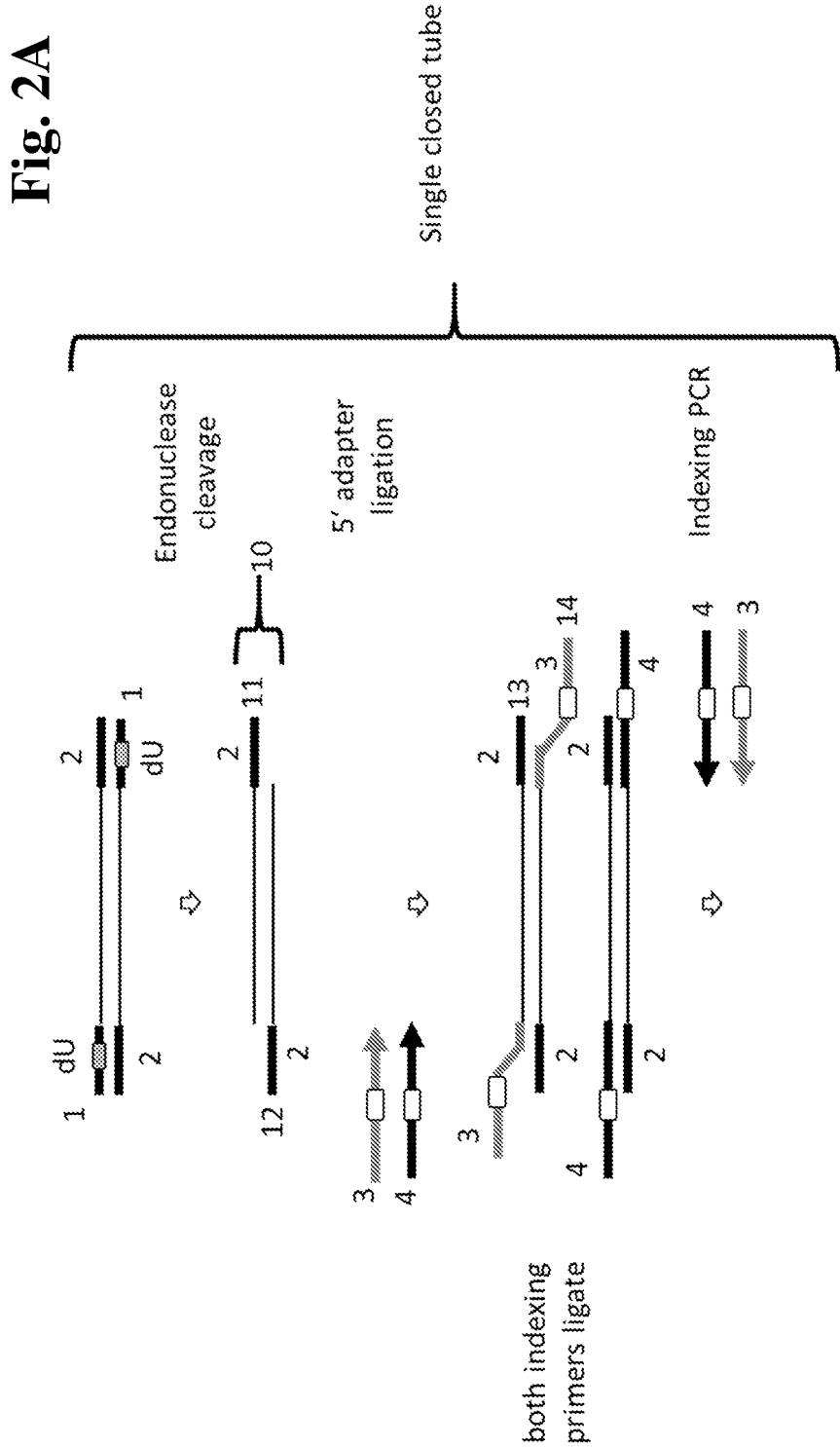


Fig. 2A
(continued)

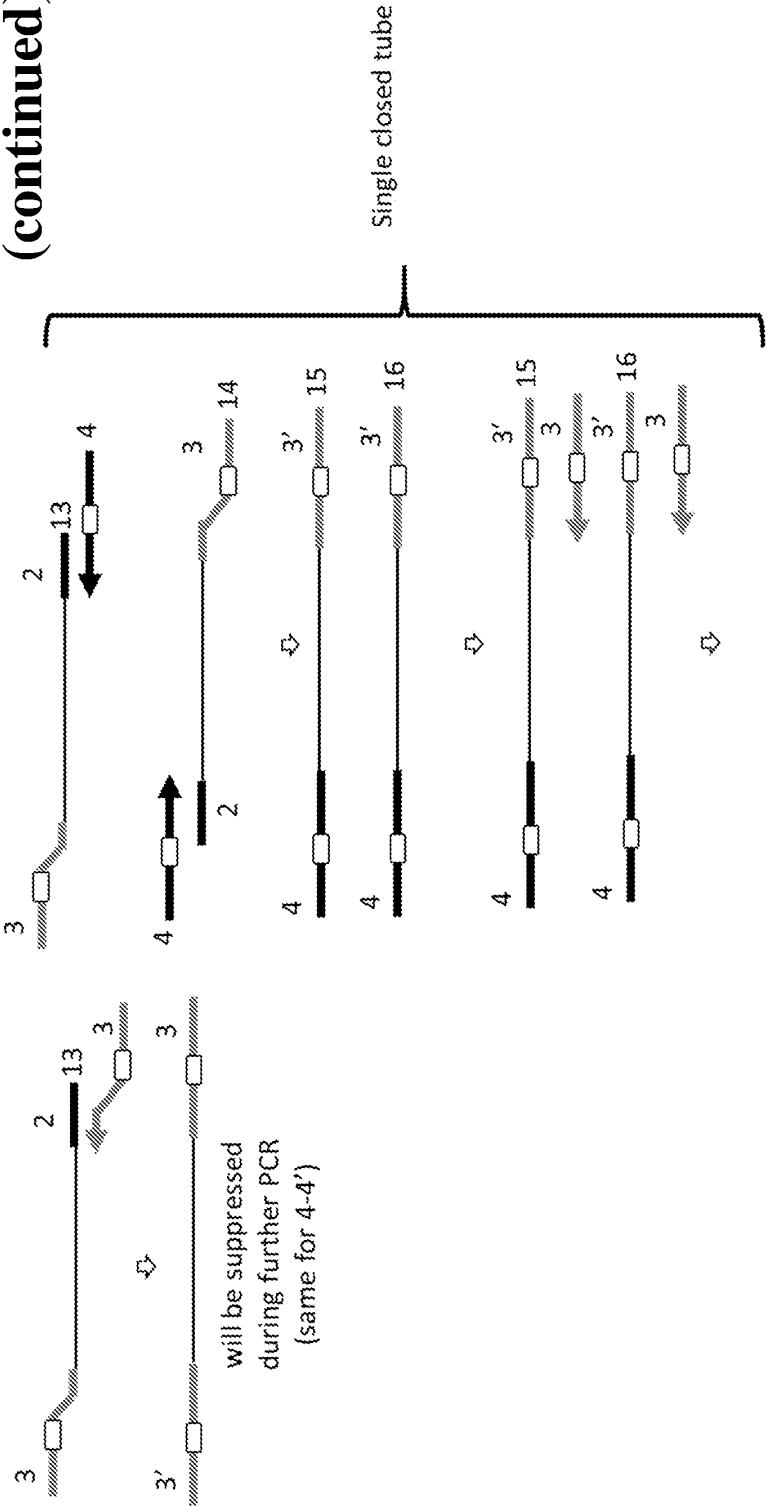


Fig. 2A
(continued)

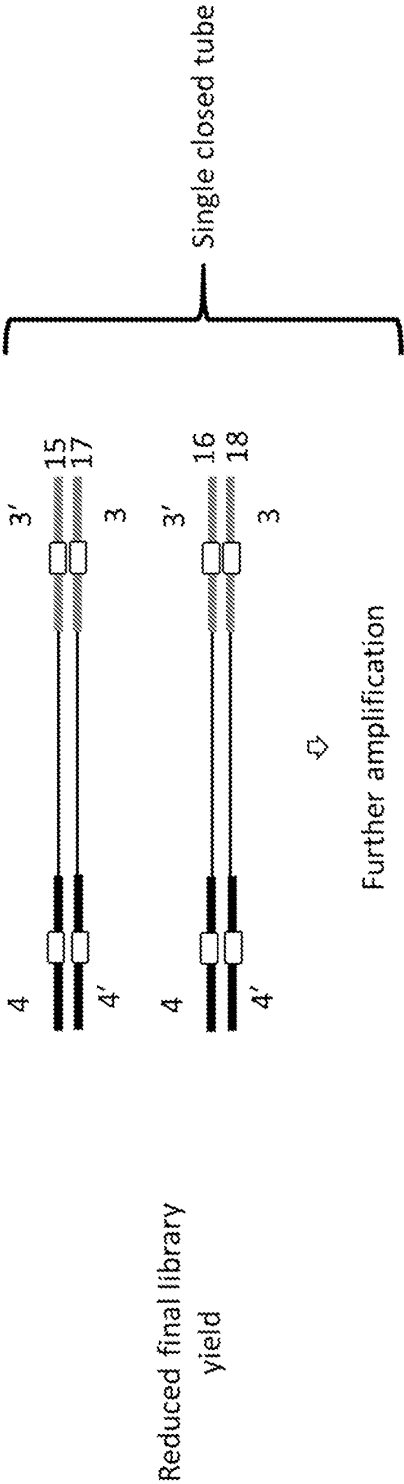
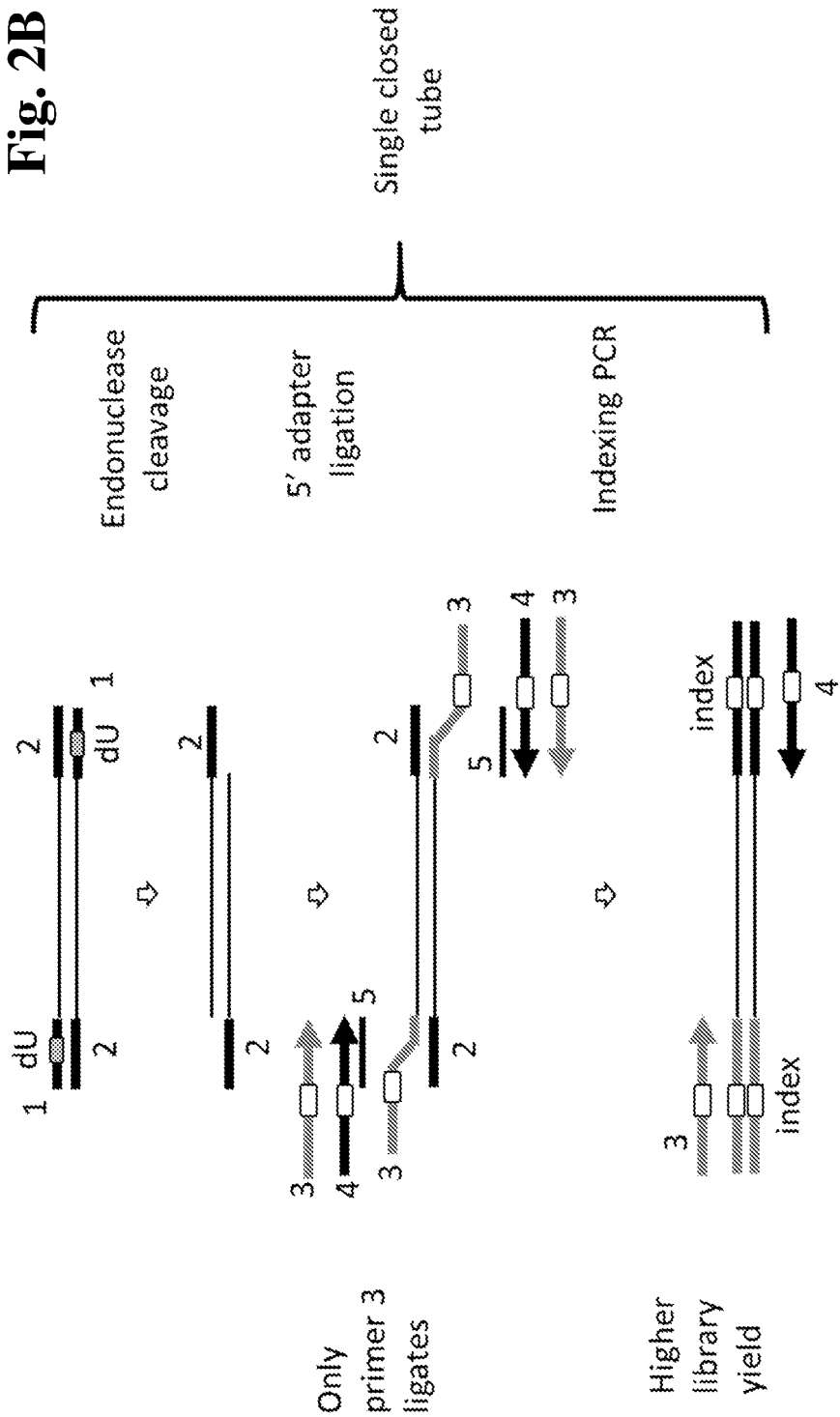
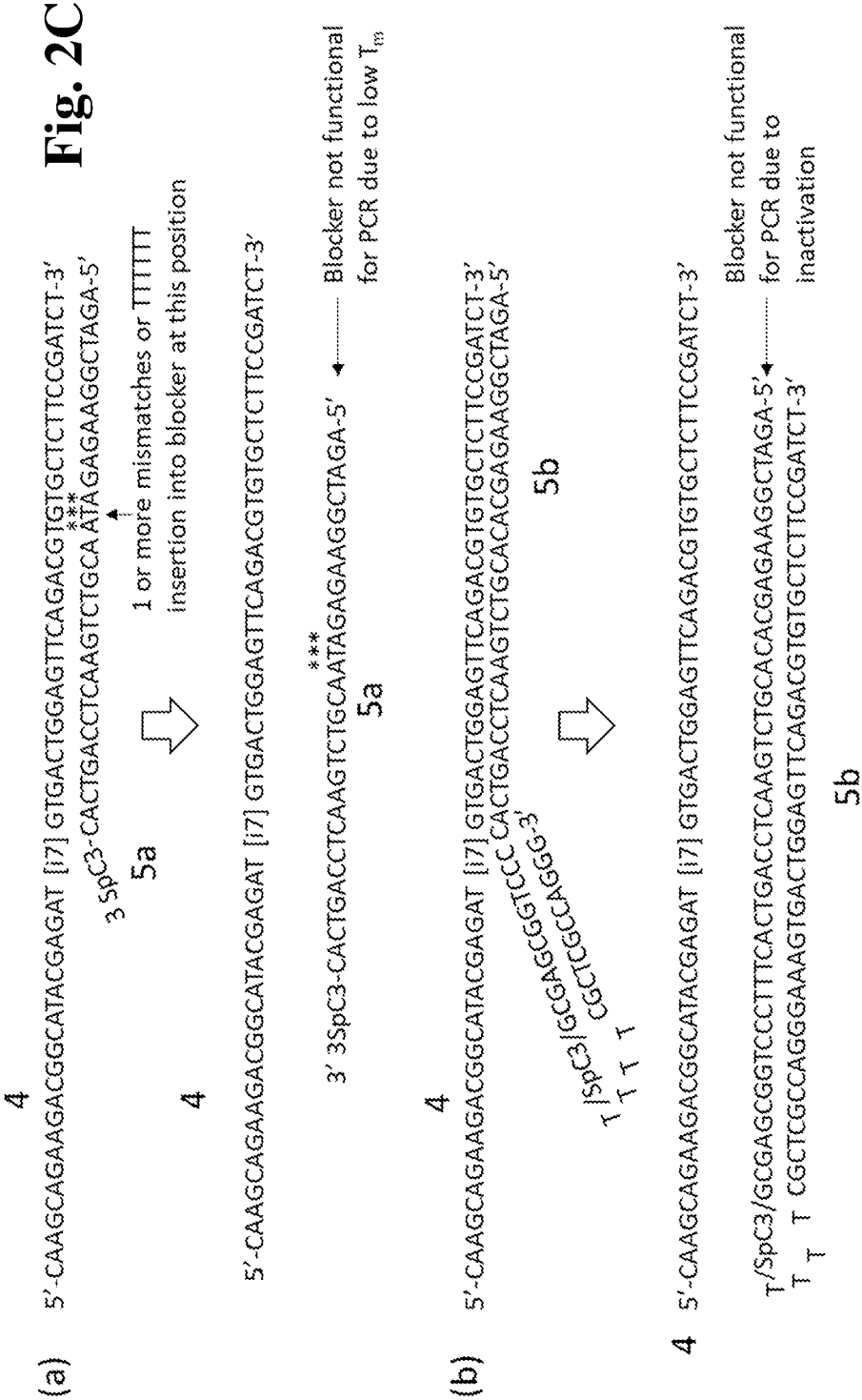


Fig. 2B





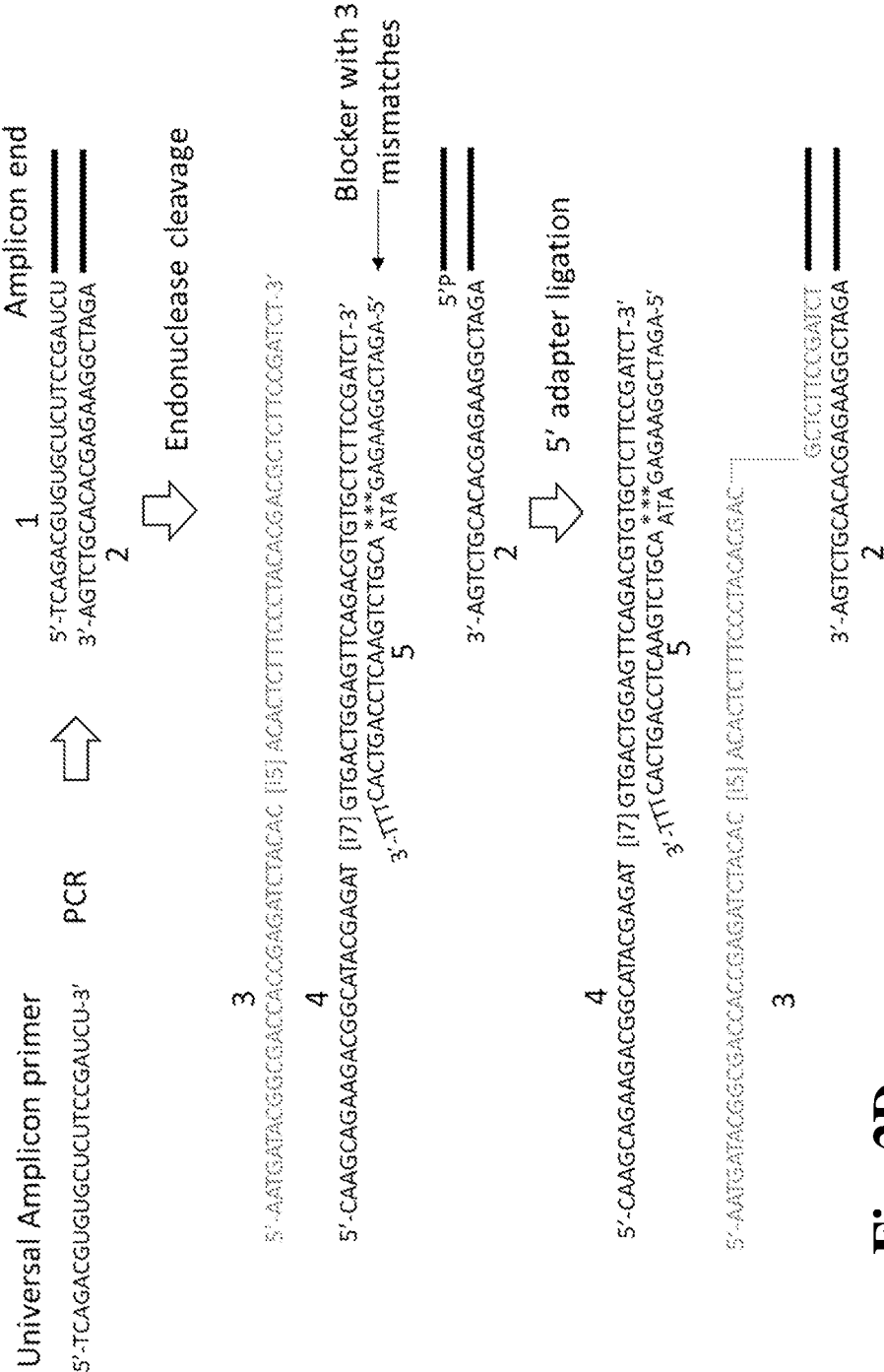


Fig. 2D

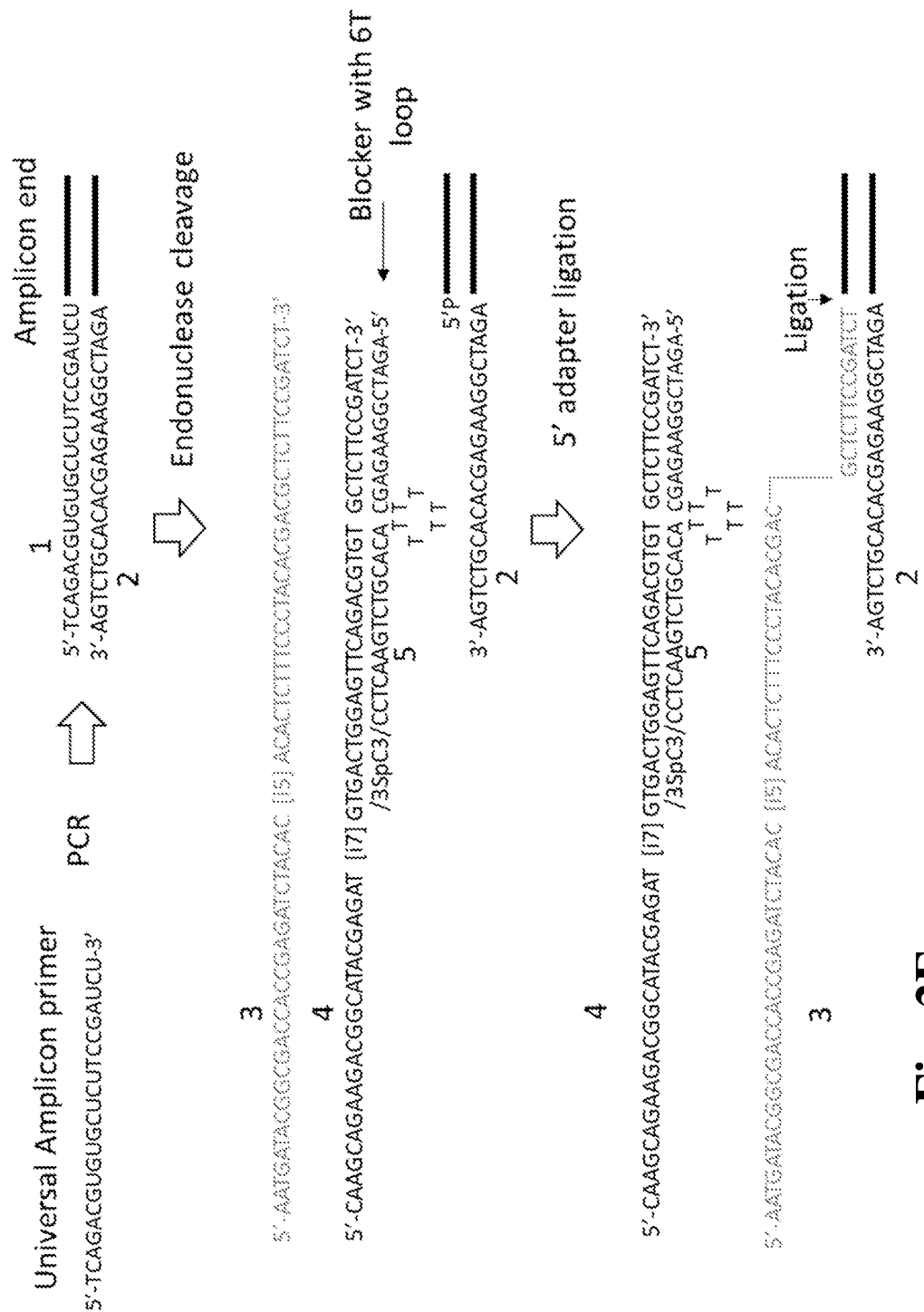


Fig. 2E

Fig. 2F

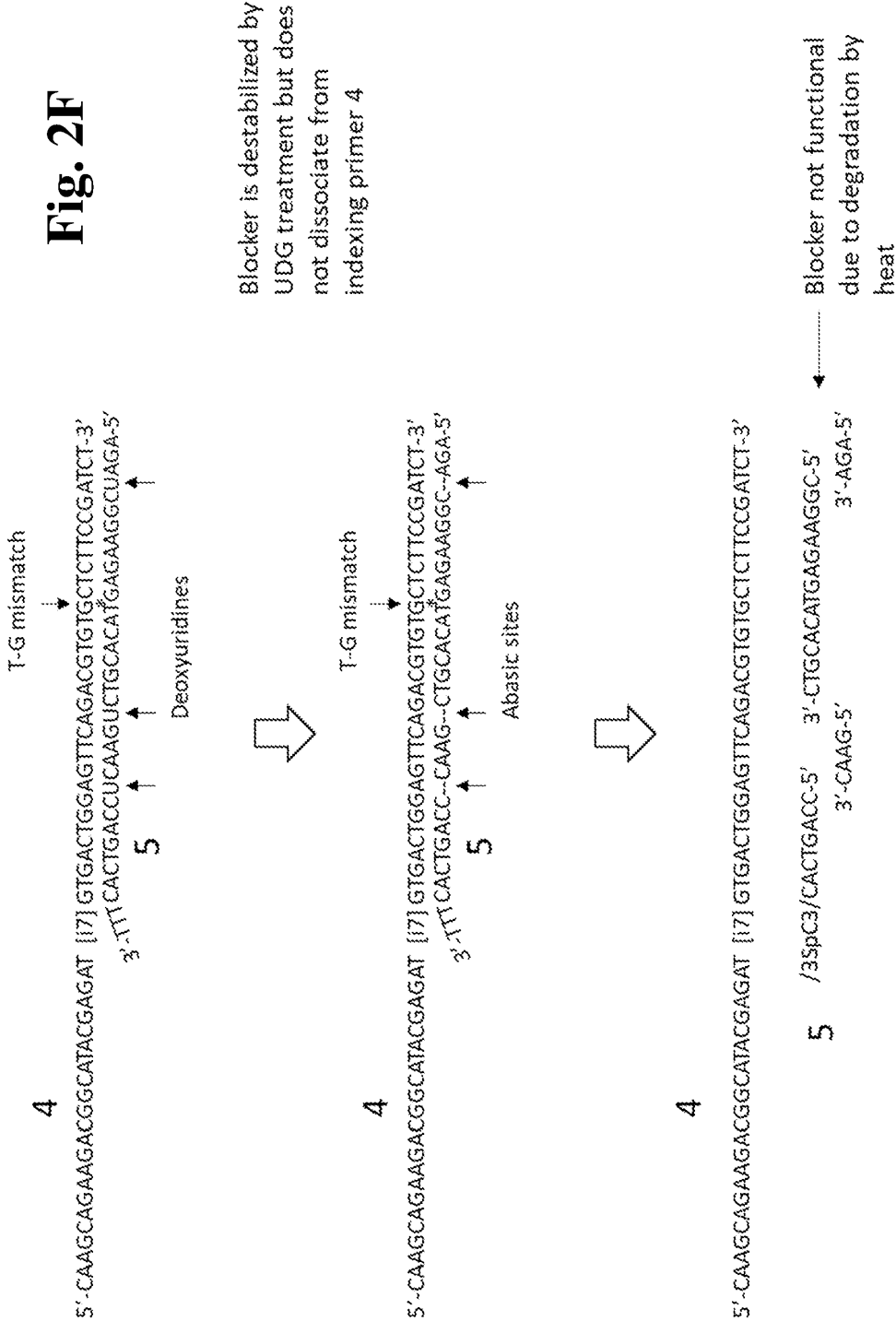


Fig. 2G

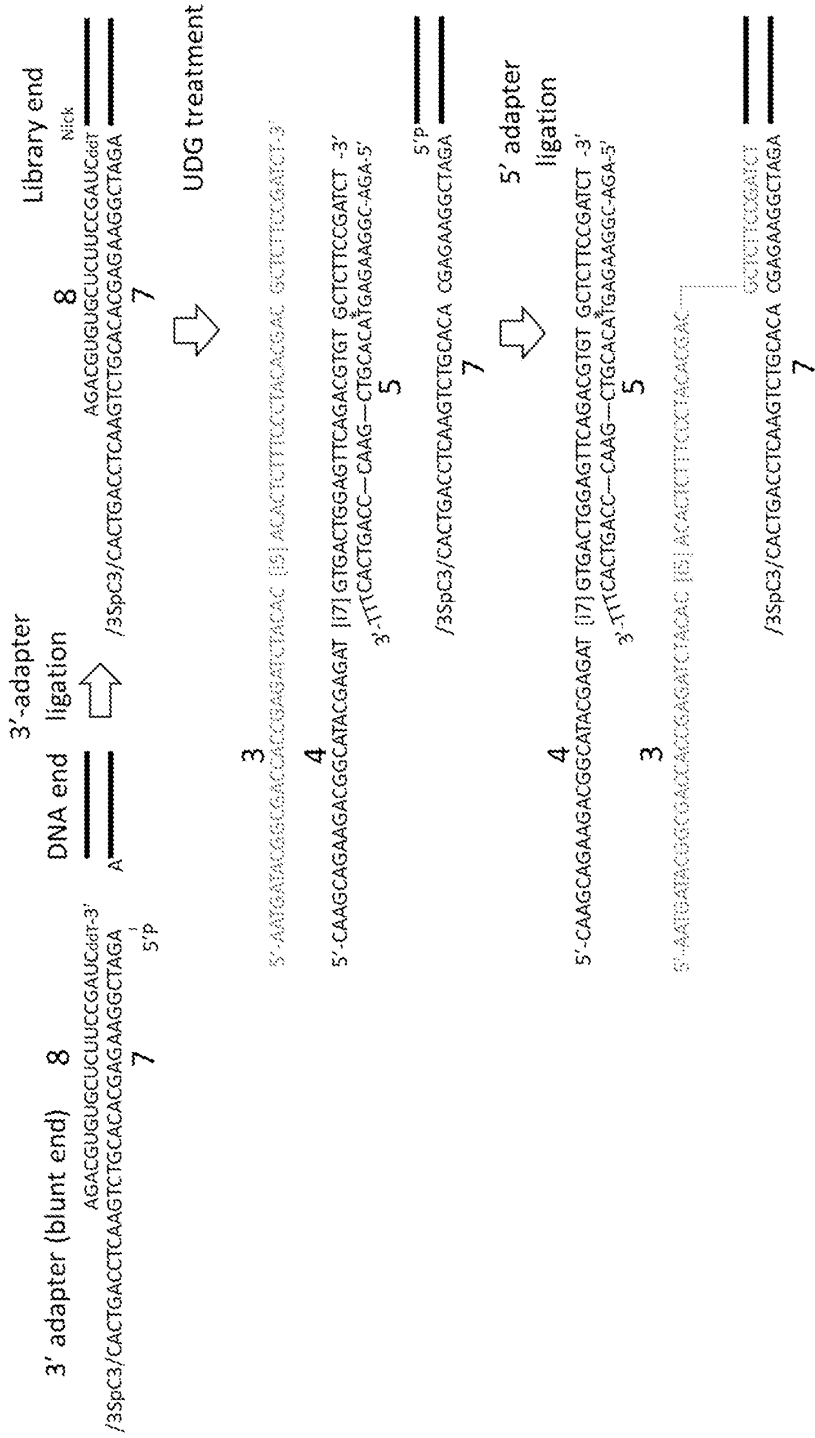


Fig. 3A

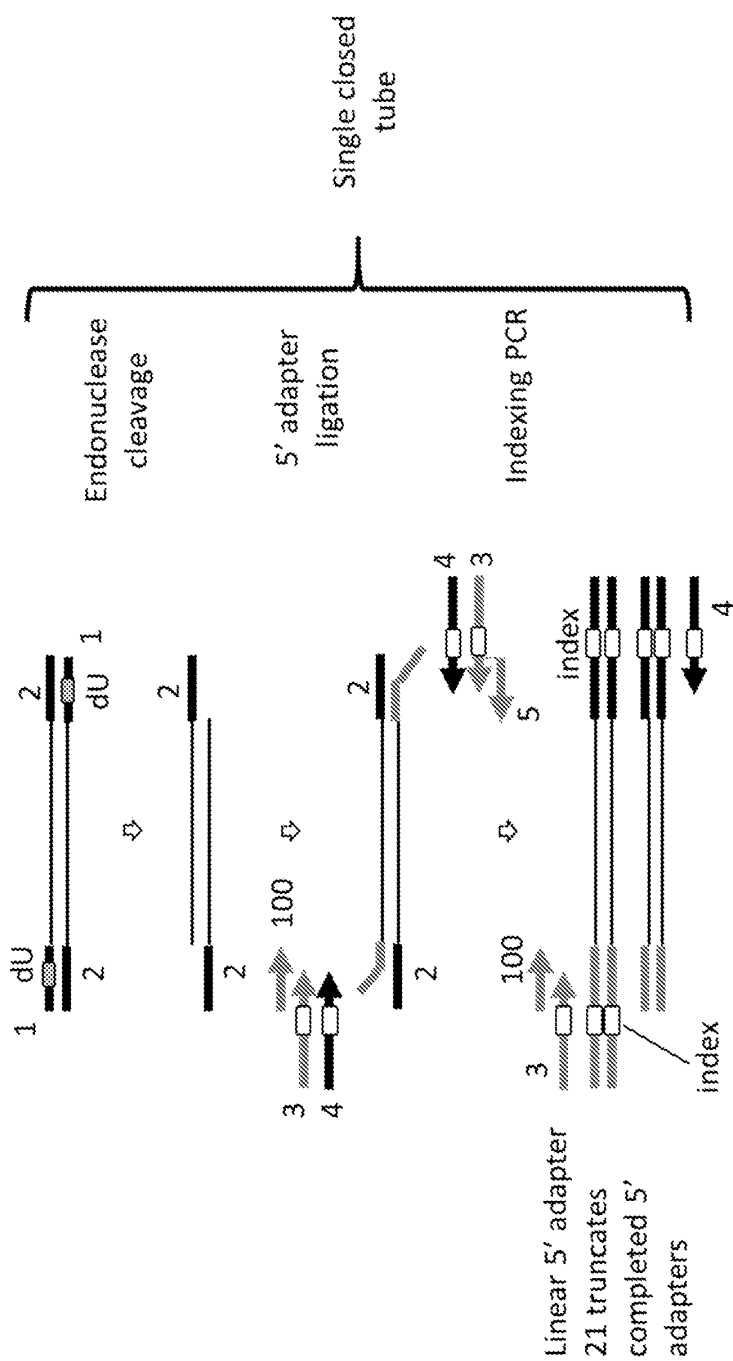


Fig. 3B

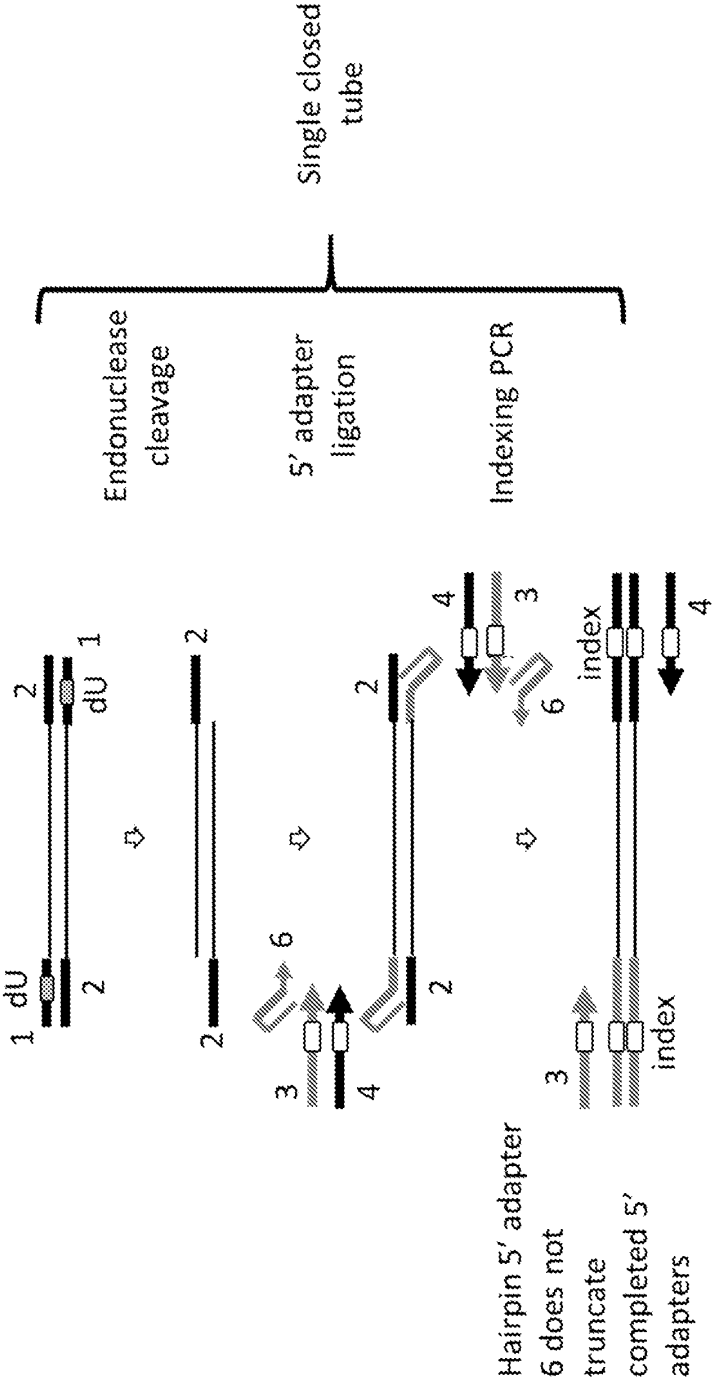
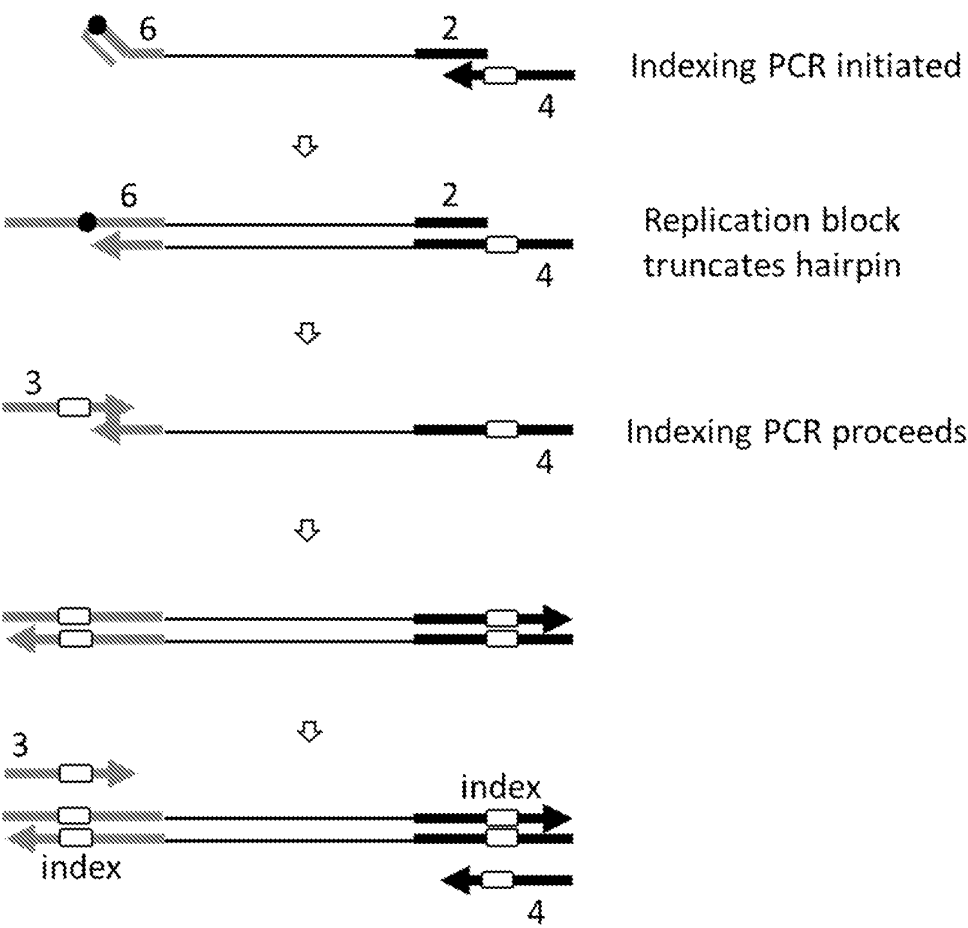
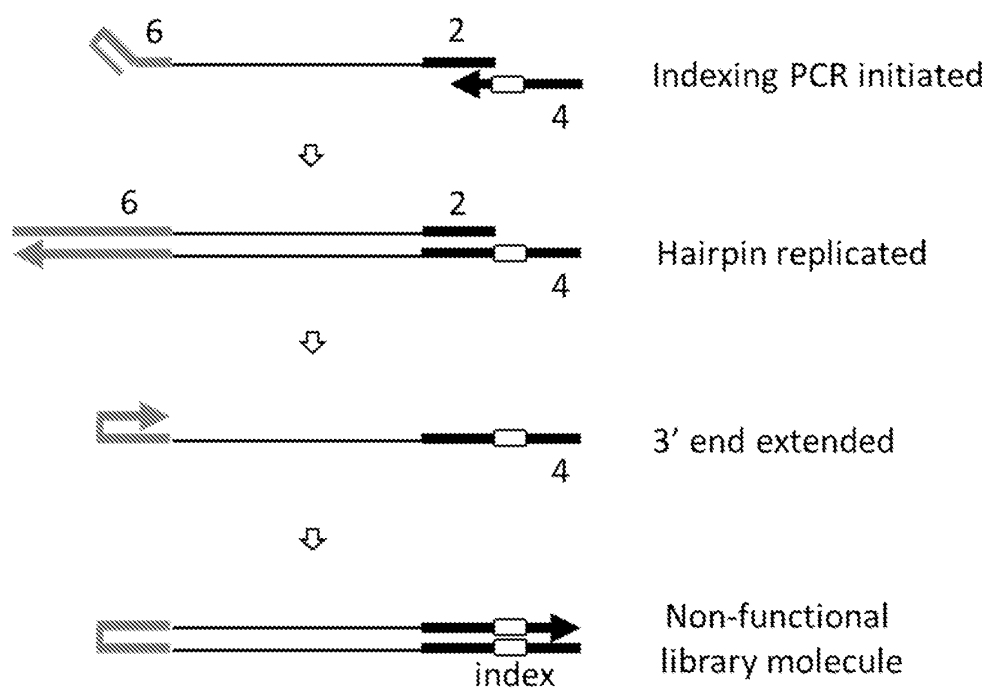


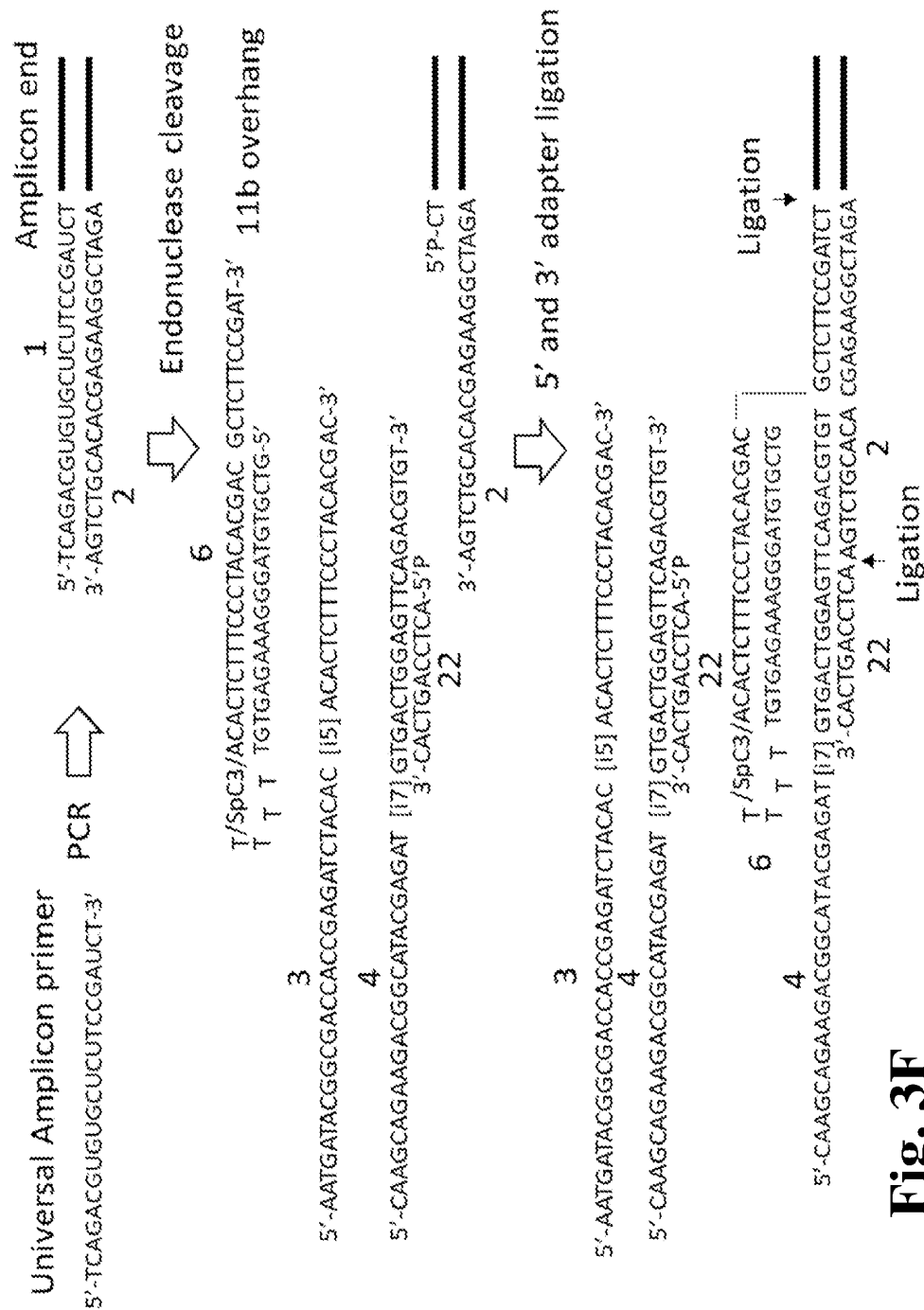
Fig. 3C



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Fig. 3D

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**Fig. 3F**

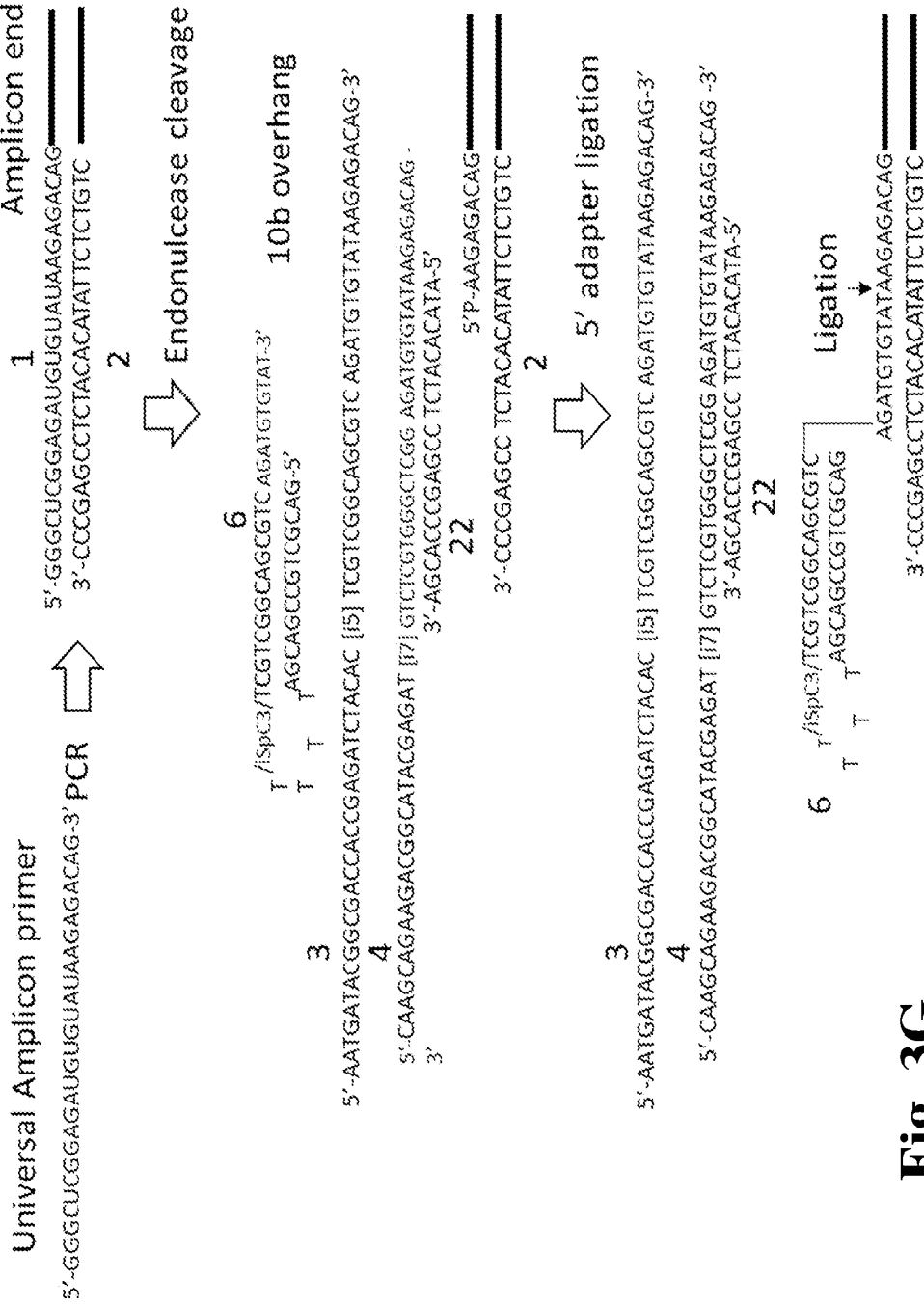


Fig. 3G

Fig. 3I

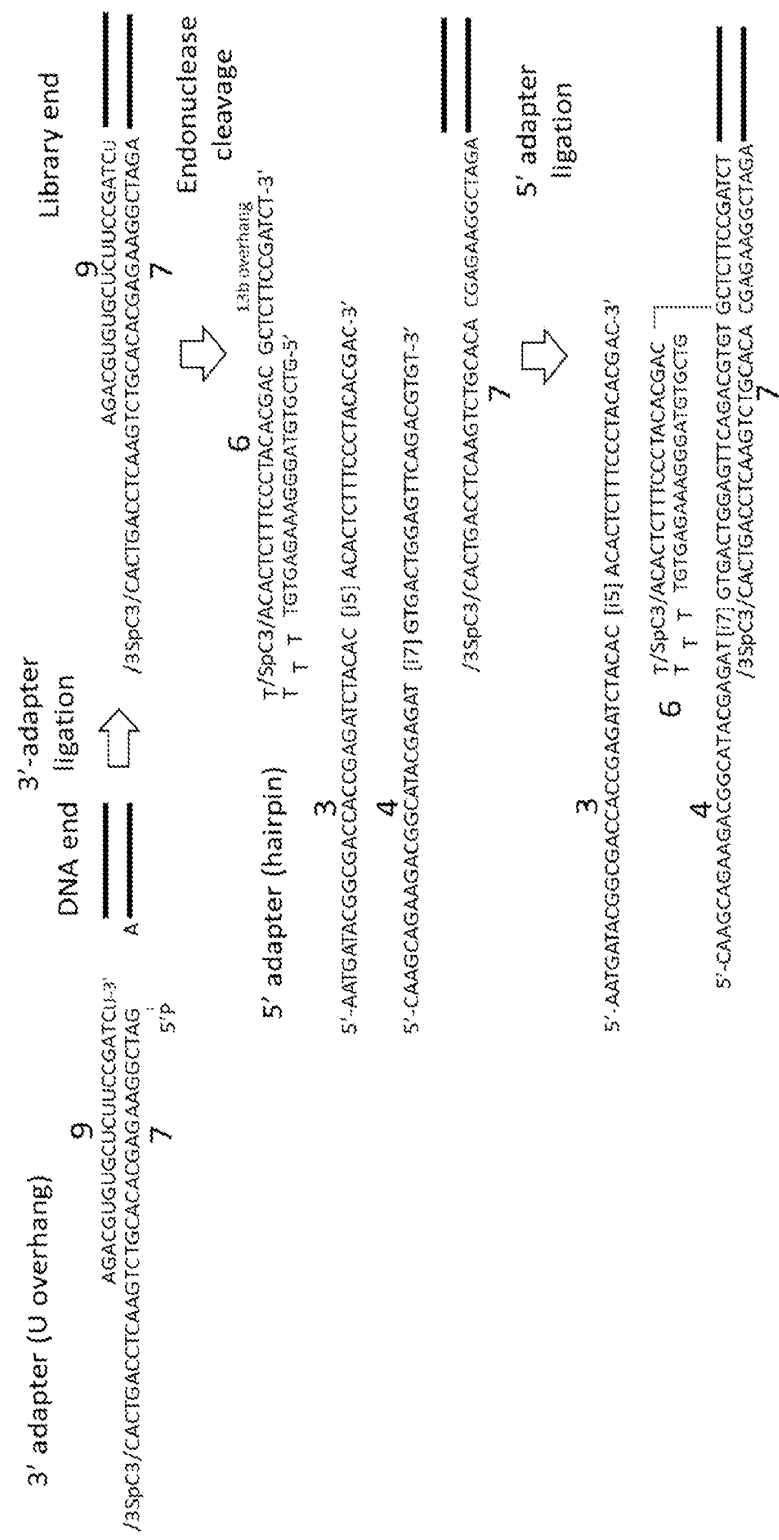


Fig. 3J

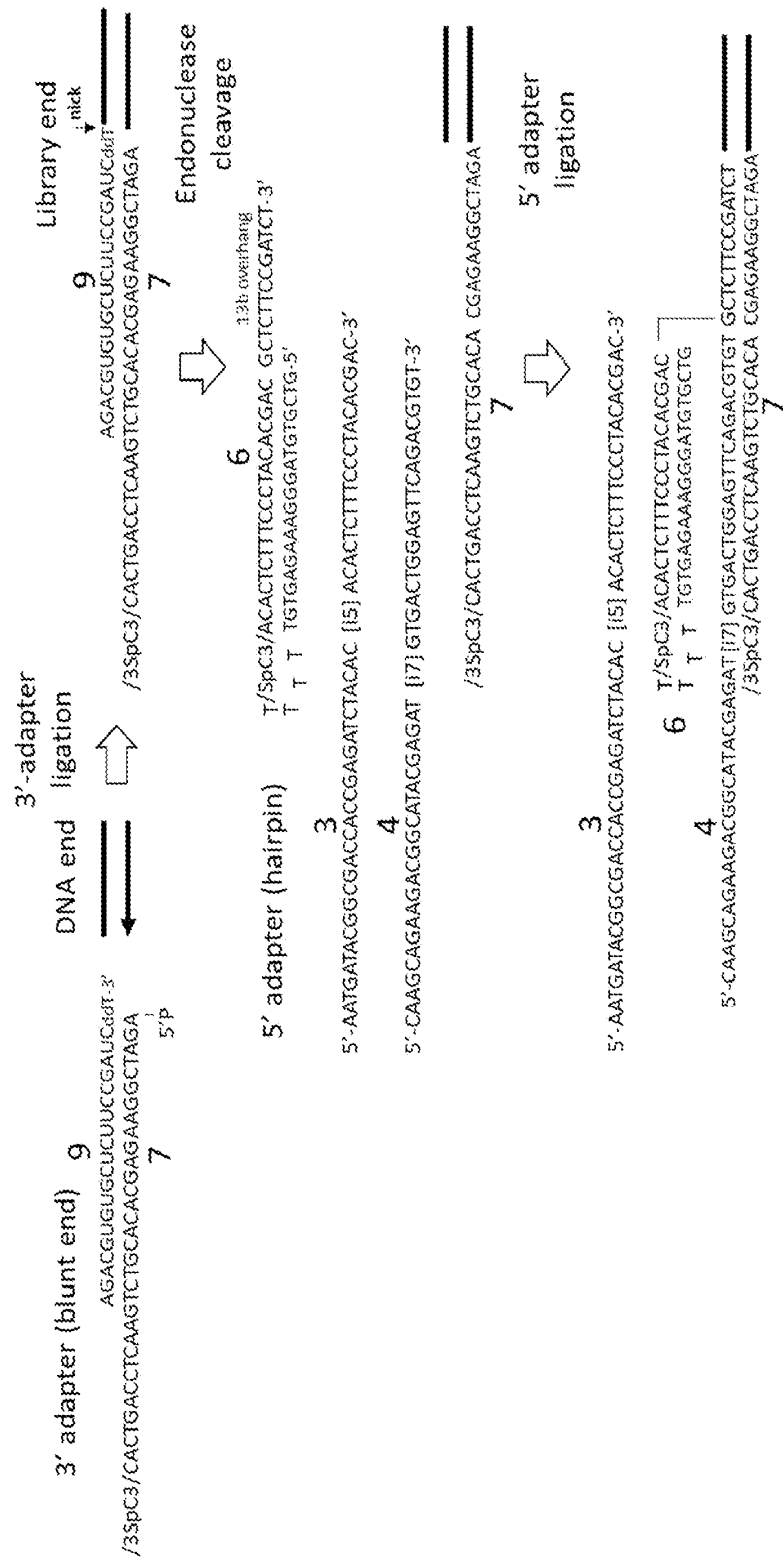


Fig. 3K

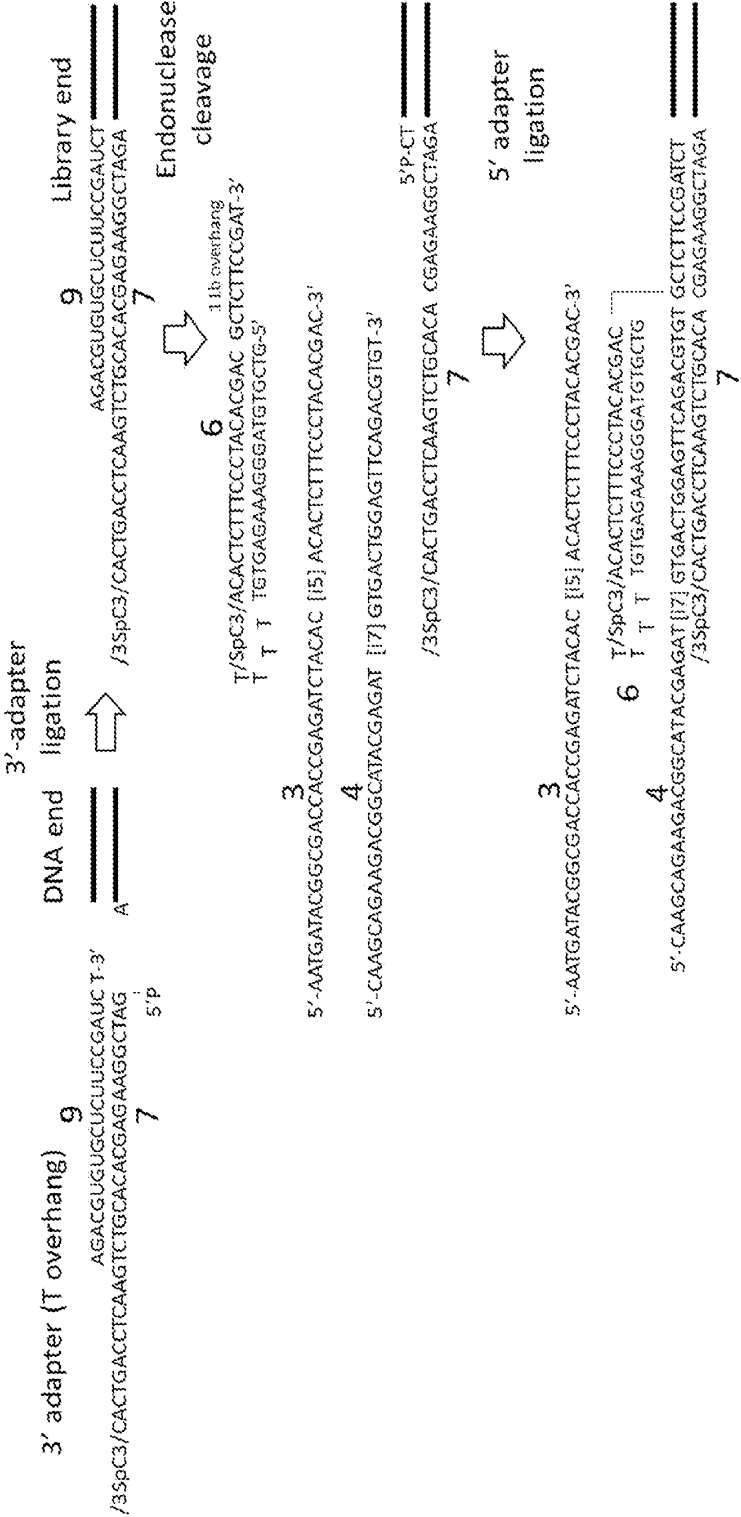
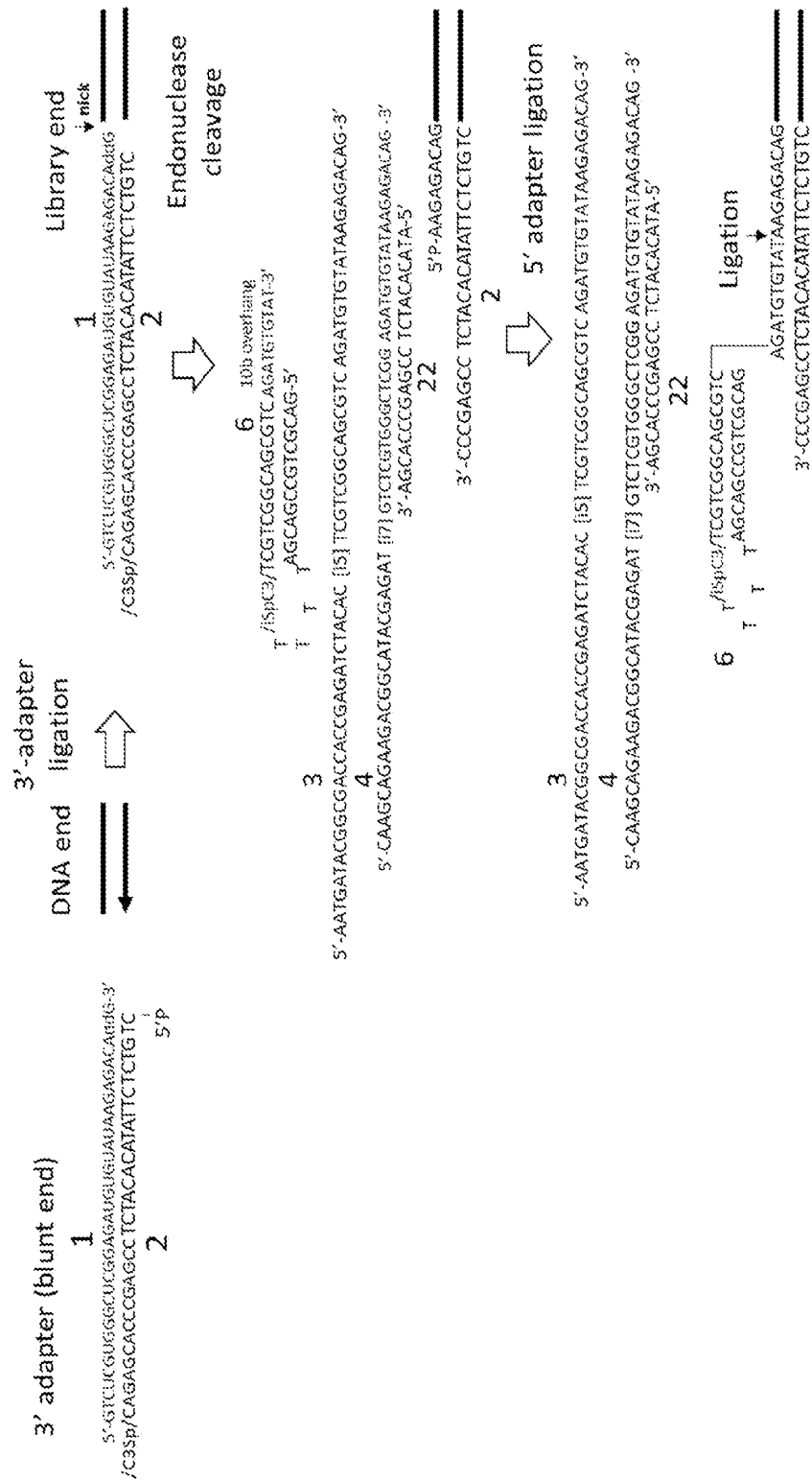
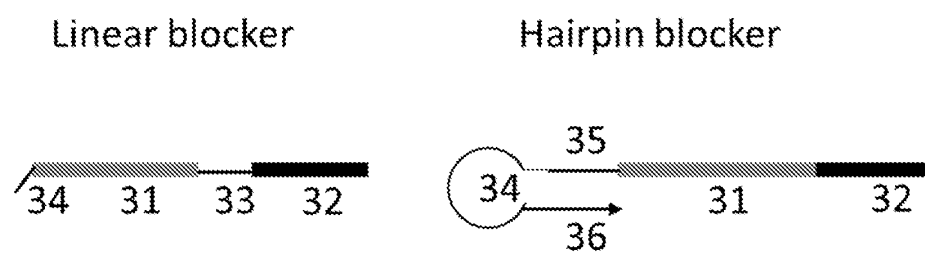


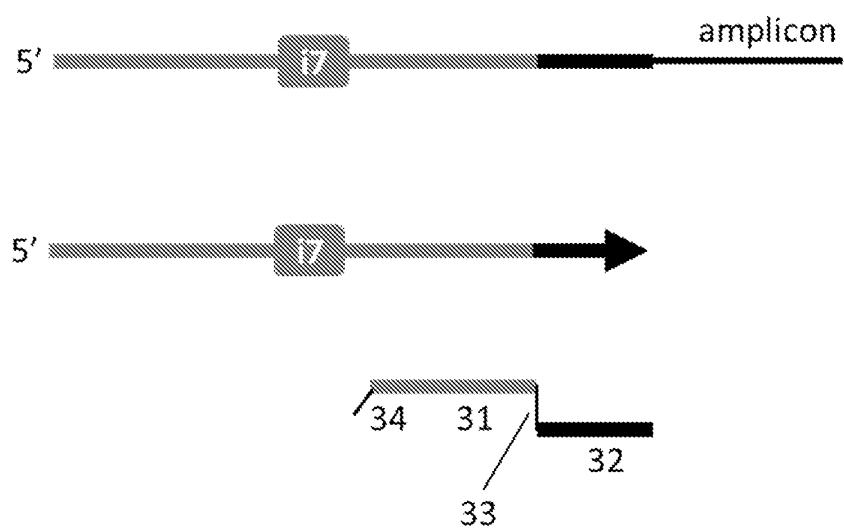
Fig. 3L

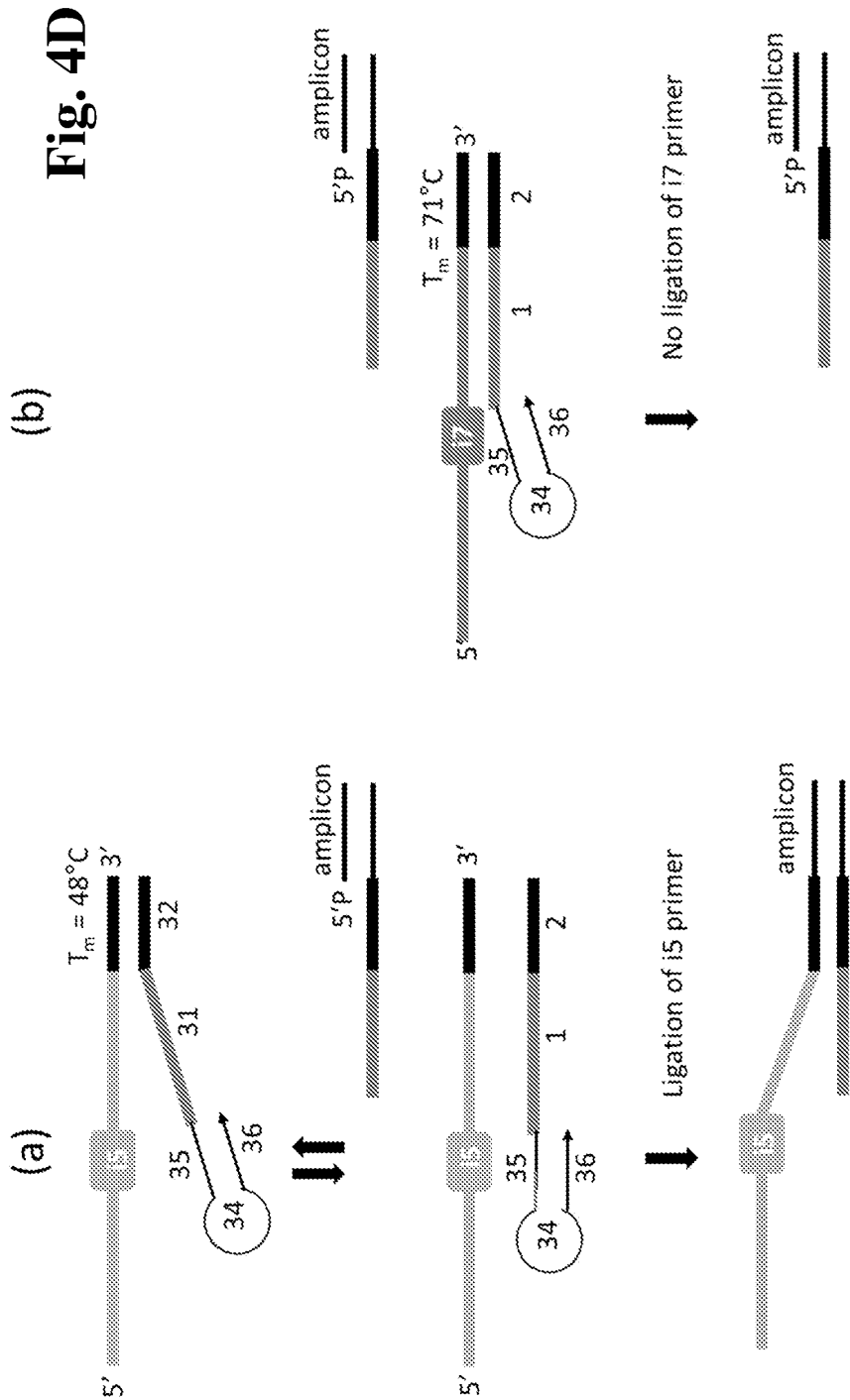


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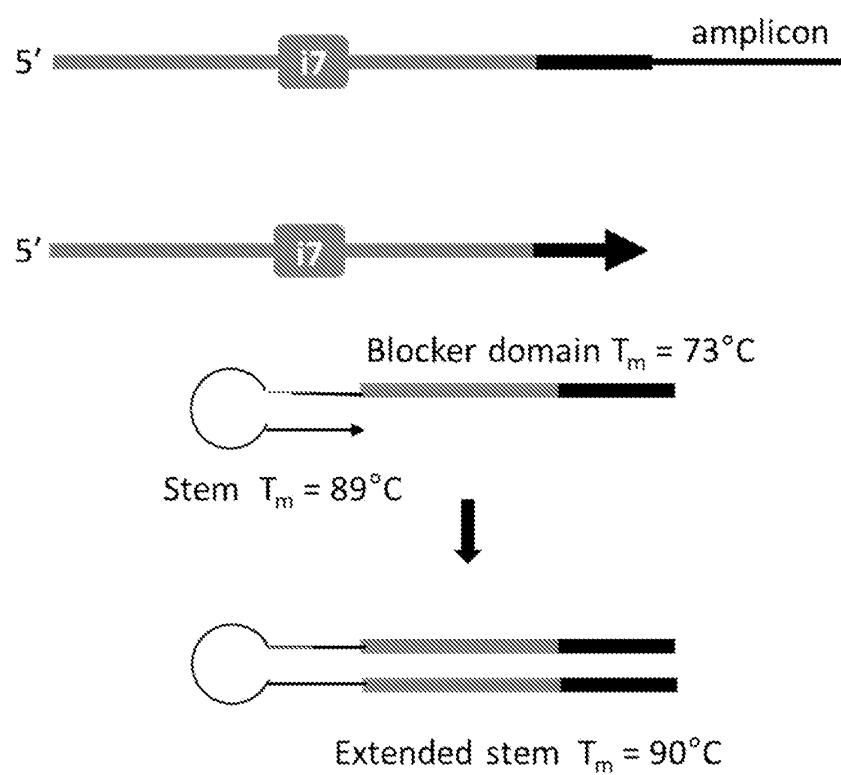
Fig. 4A

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Fig. 4C



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Fig. 4E

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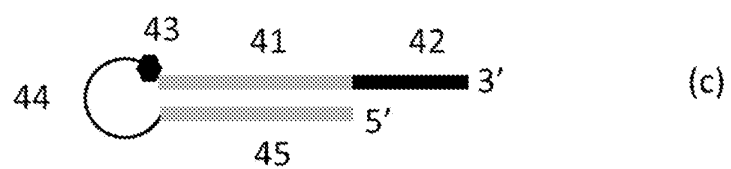
Fig. 5A

Fig. 5B

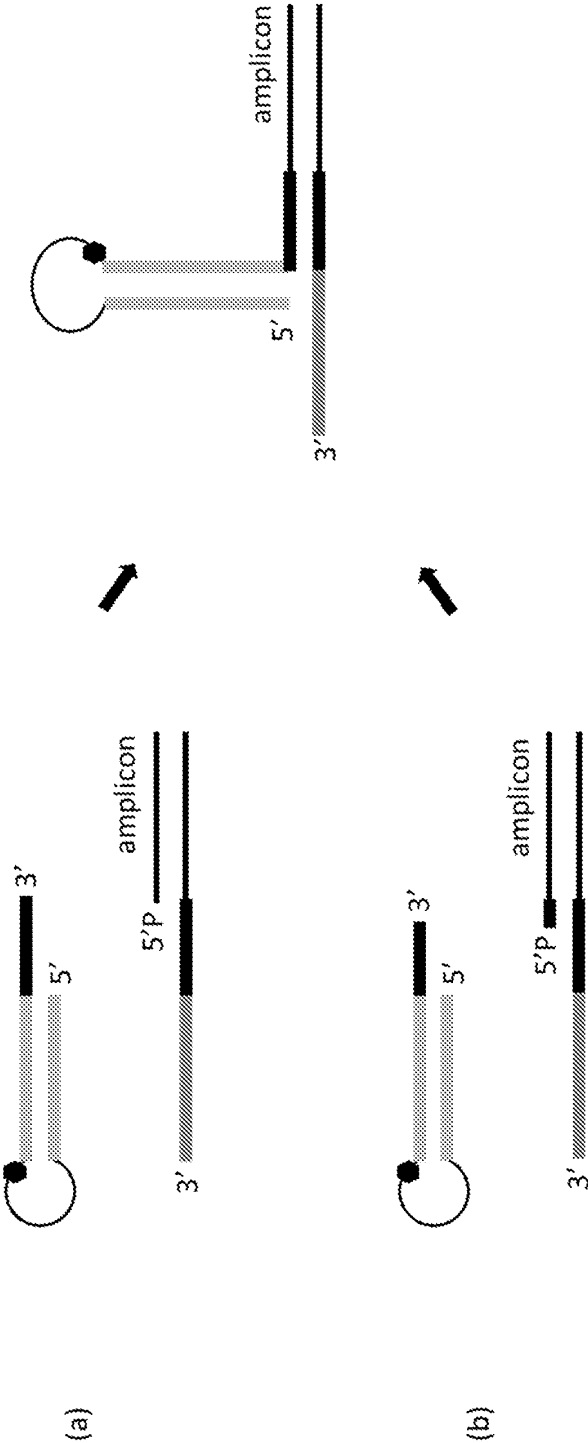


Fig. 5C

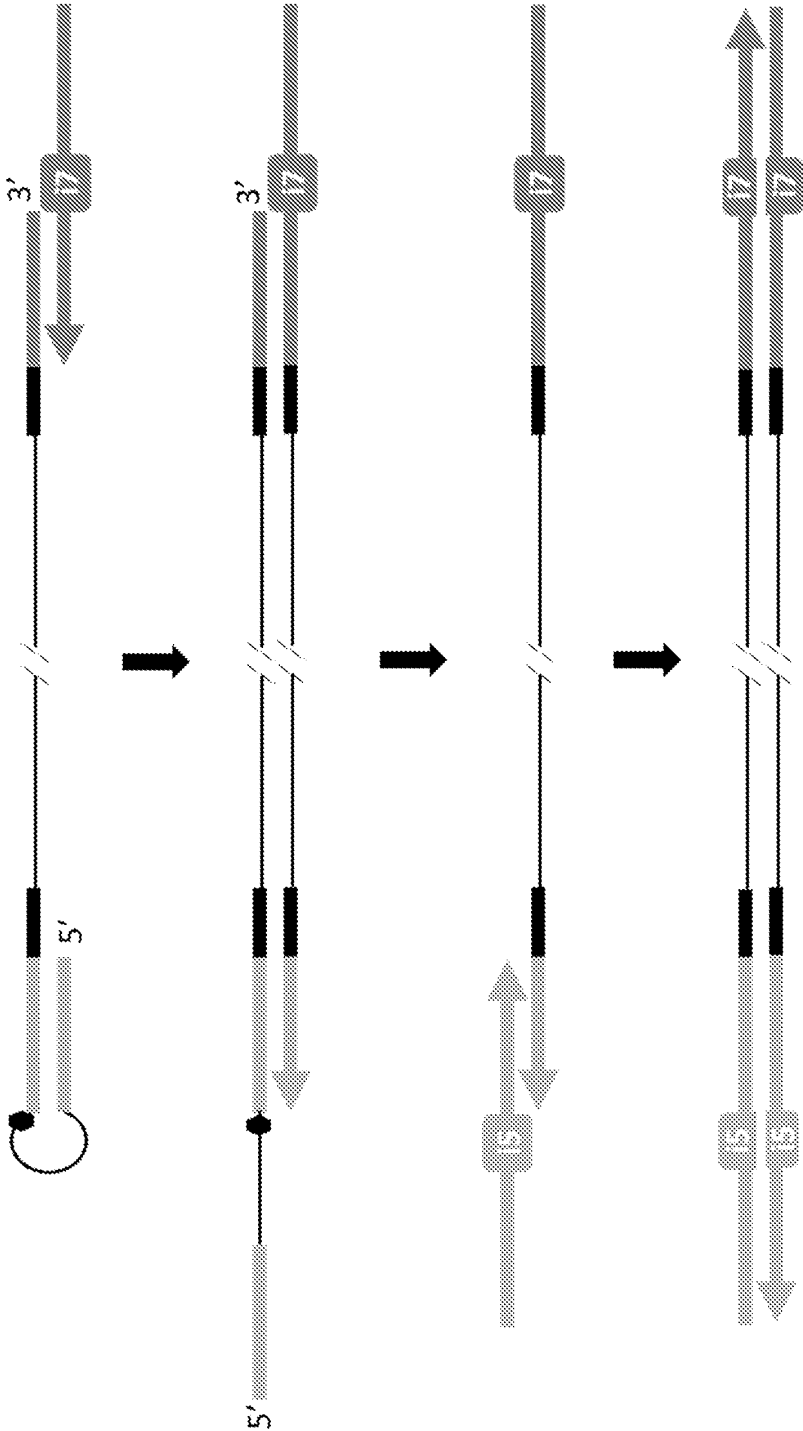


Fig. 6A

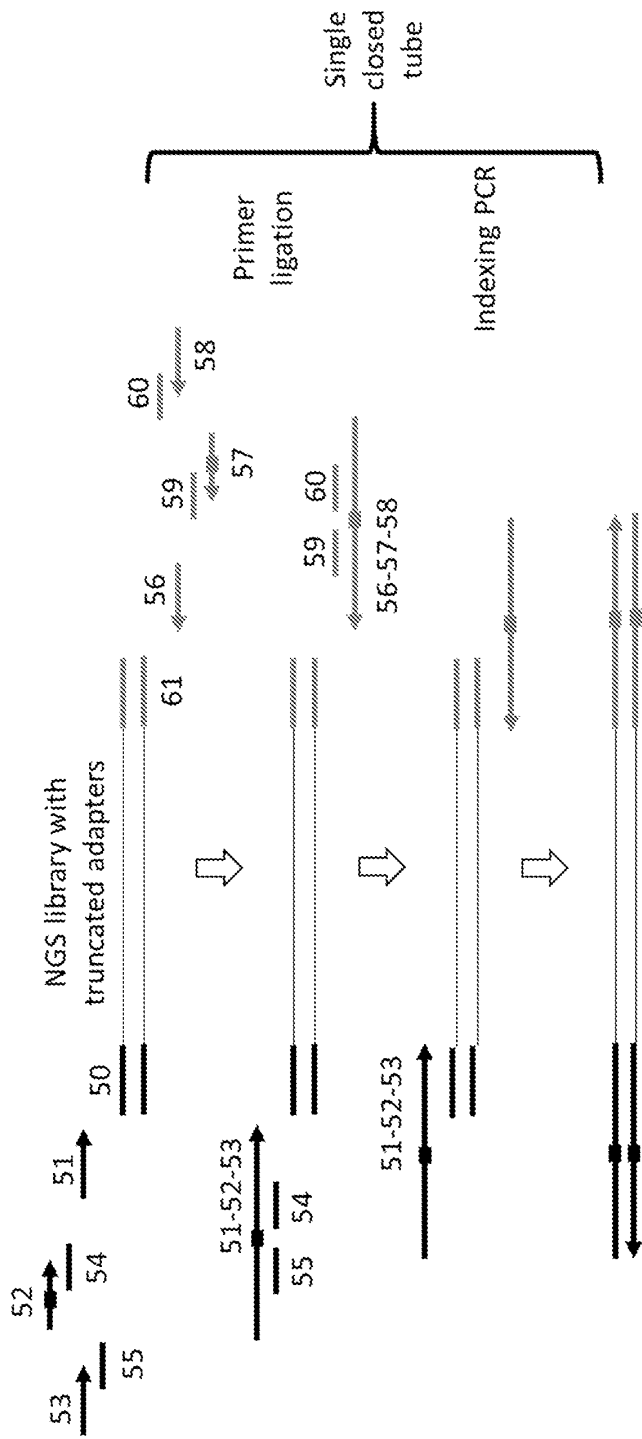


Fig. 6B

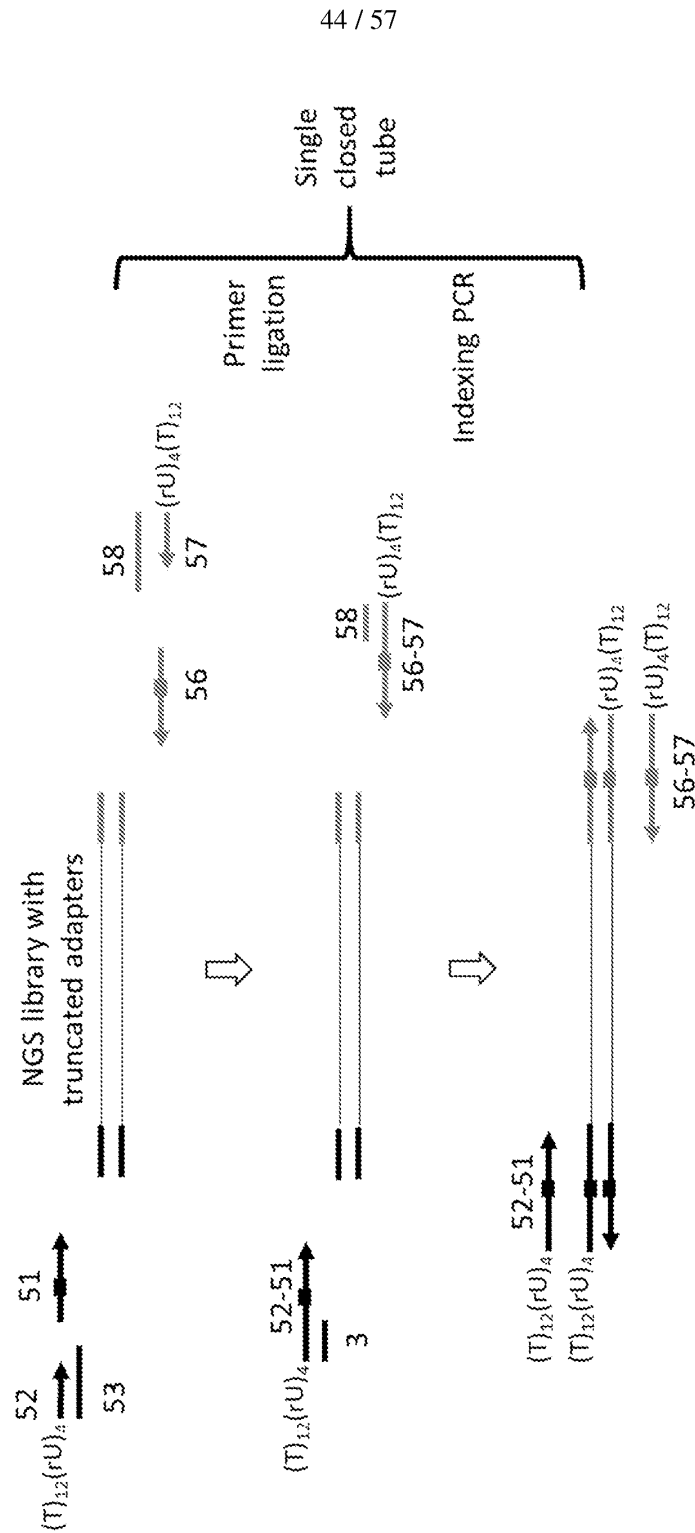


Fig. 6C

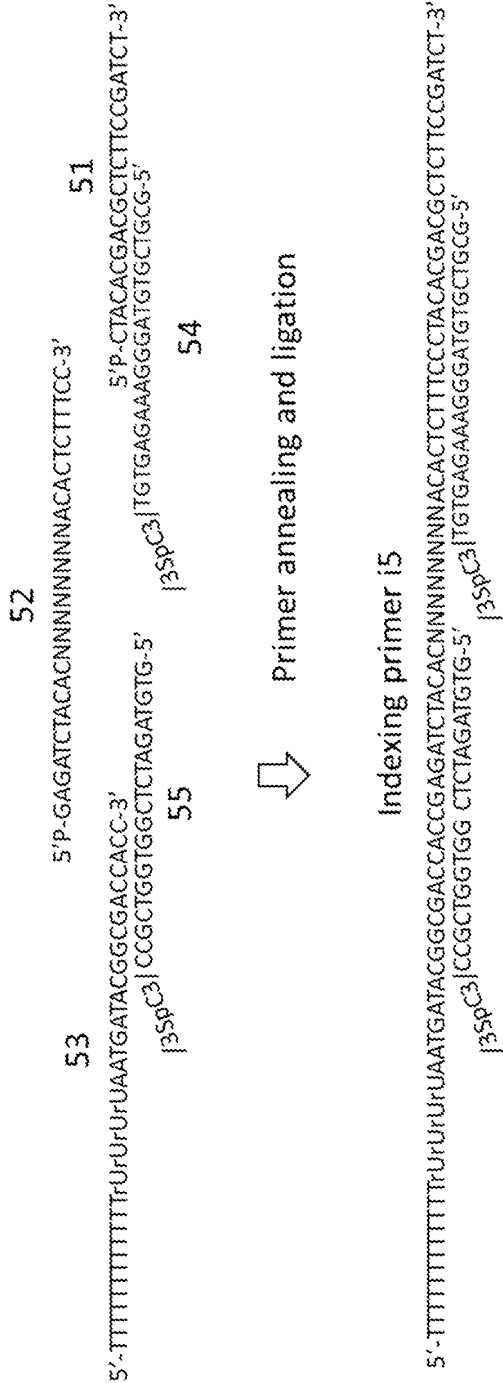
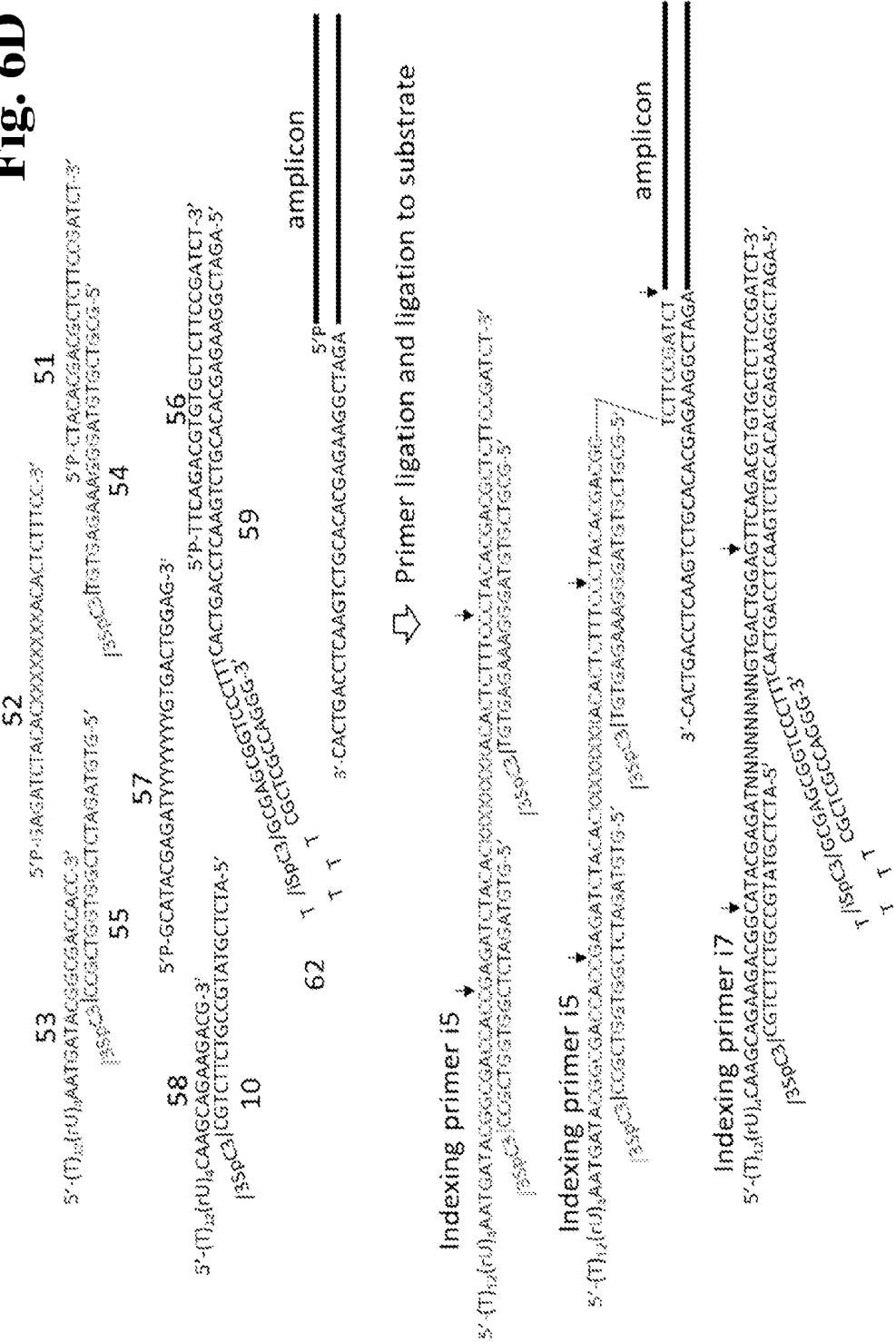
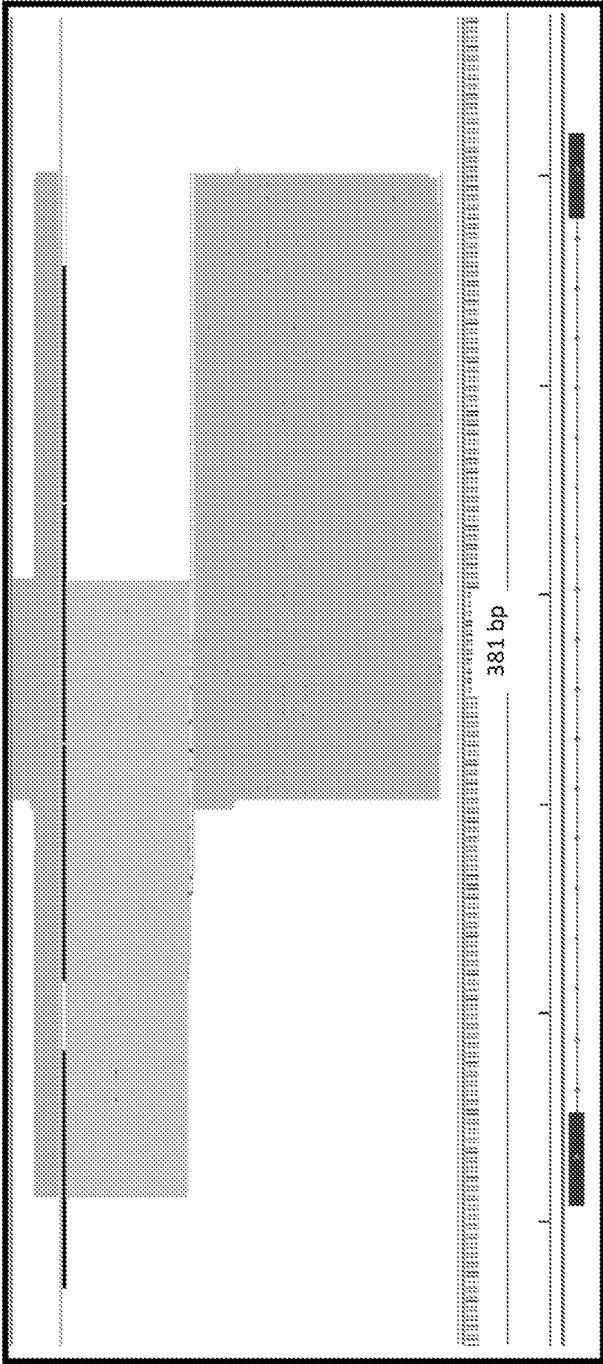


Fig. 6D





65% fungal coverage
145 to 695bp

18S	ITS1	5.8S	ITS2	28S
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Fig. 7

Fig. 8

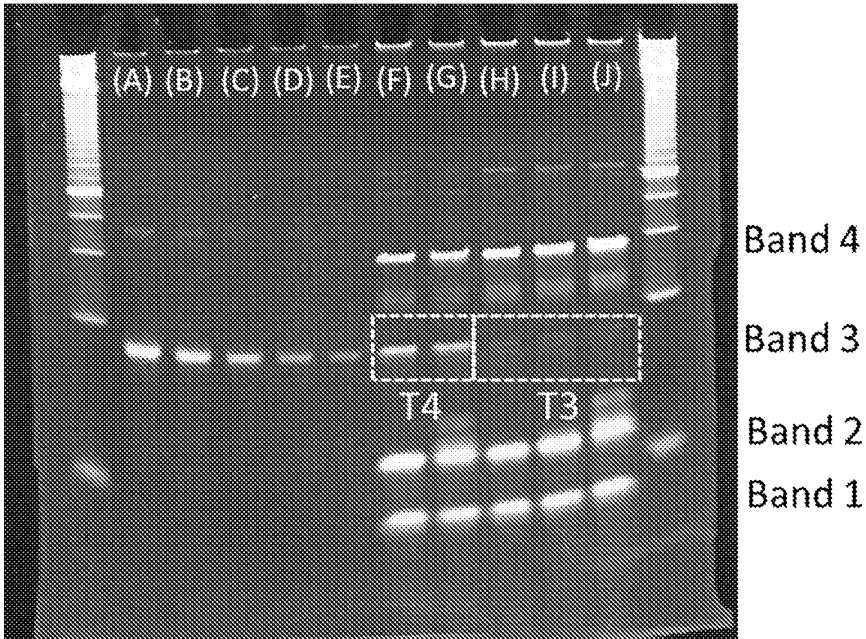
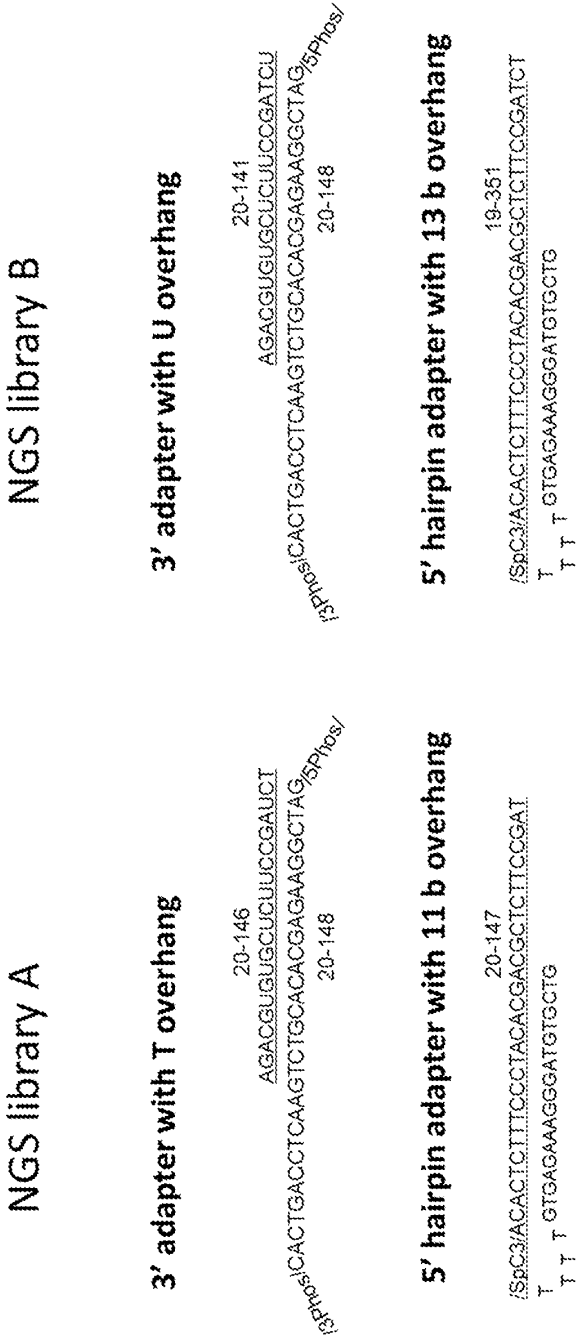


Fig. 9

Blockers

19-356	CAAGCAGAAAGACGGGCATACGAGATCACCTATGCCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT TTCACTGACCCUCAAGUCTGCACATGAGAAGGCUAGA
19-357	CAAGCAGAAAGACGGGCATACGAGATCACCTATGCCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT TTCACTGACCCUCAAGUCTGTACATGAGAAGGCUAGA
19-354	CAAGCAGAAAGACGGGCATACGAGATCACCTATGCCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT TTCACTGACCCUCAAGUCTGCACACGAGAAGGCUAGA
19-353	CAAGCAGAAAGACGGGCATACGAGATCACCTATGCCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT TTCACTGACCCUCAAGUCTGCACACGAGAAGGCTAGA
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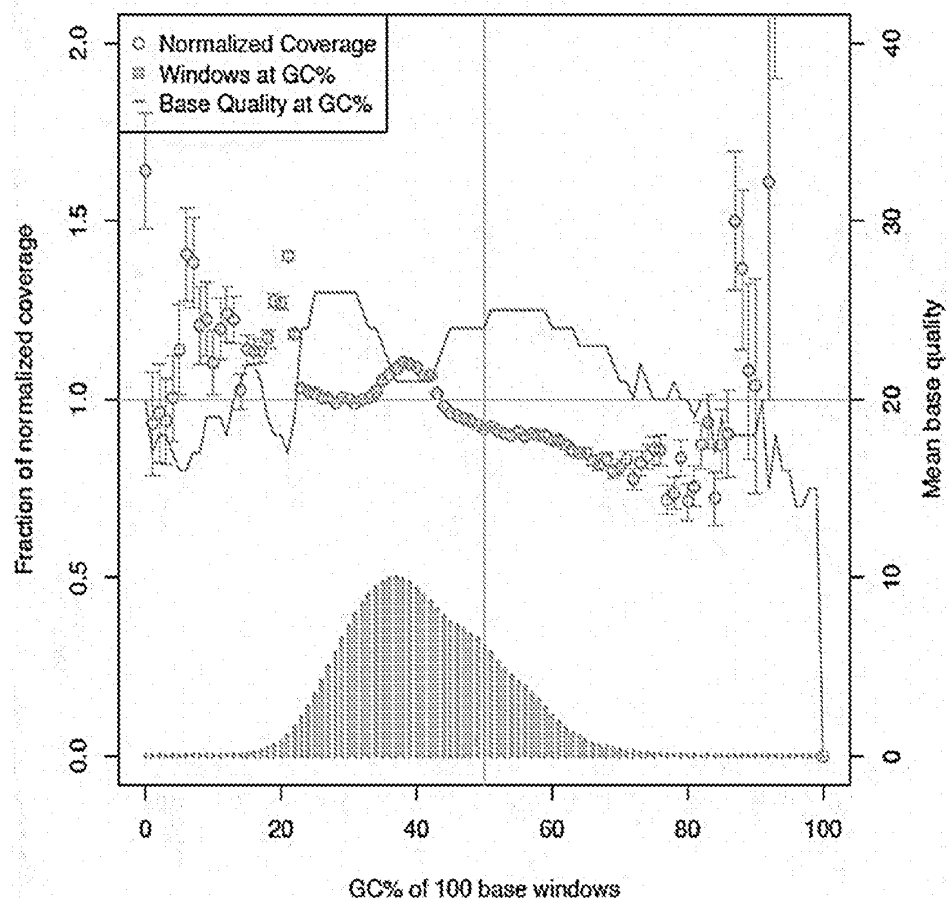
Fig. 10A



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Fig. 10B

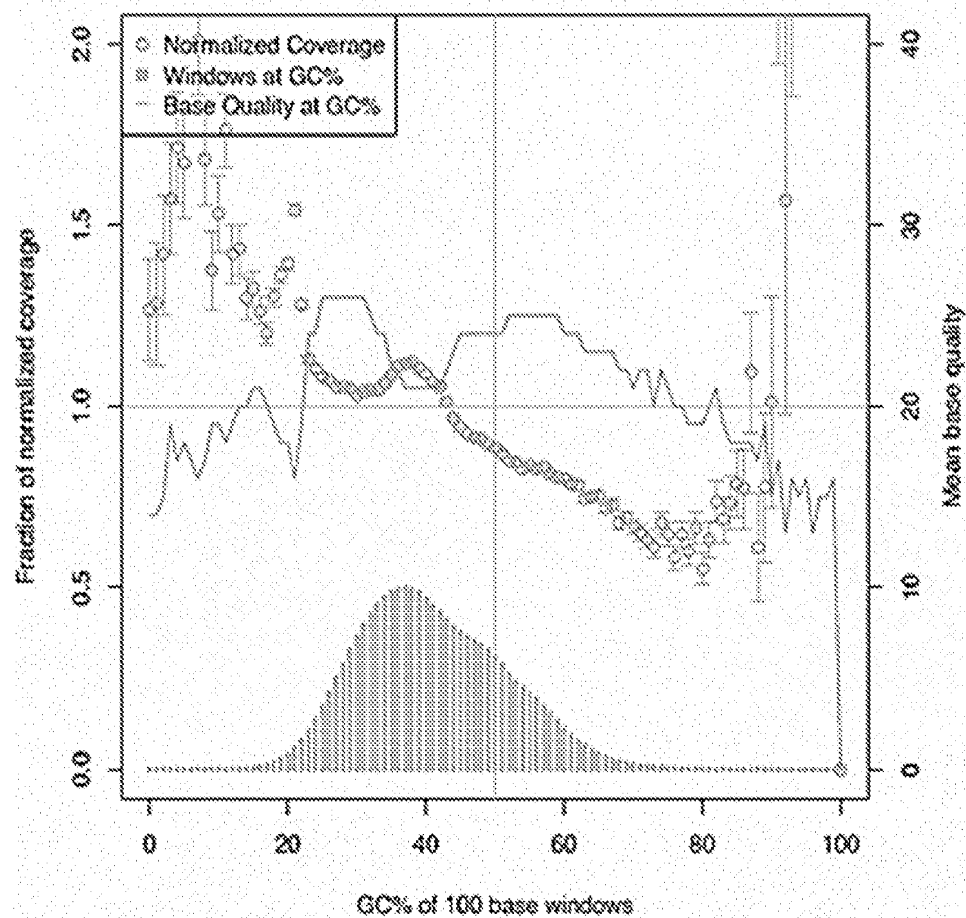
ed-Adapter-P5tr-Hairpin-No-CT-50ng-2_S2.bwa.sort.markdup.bam.Accel
Total clusters: 670,886, Aligned reads: 1,337,512



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Fig. 10C

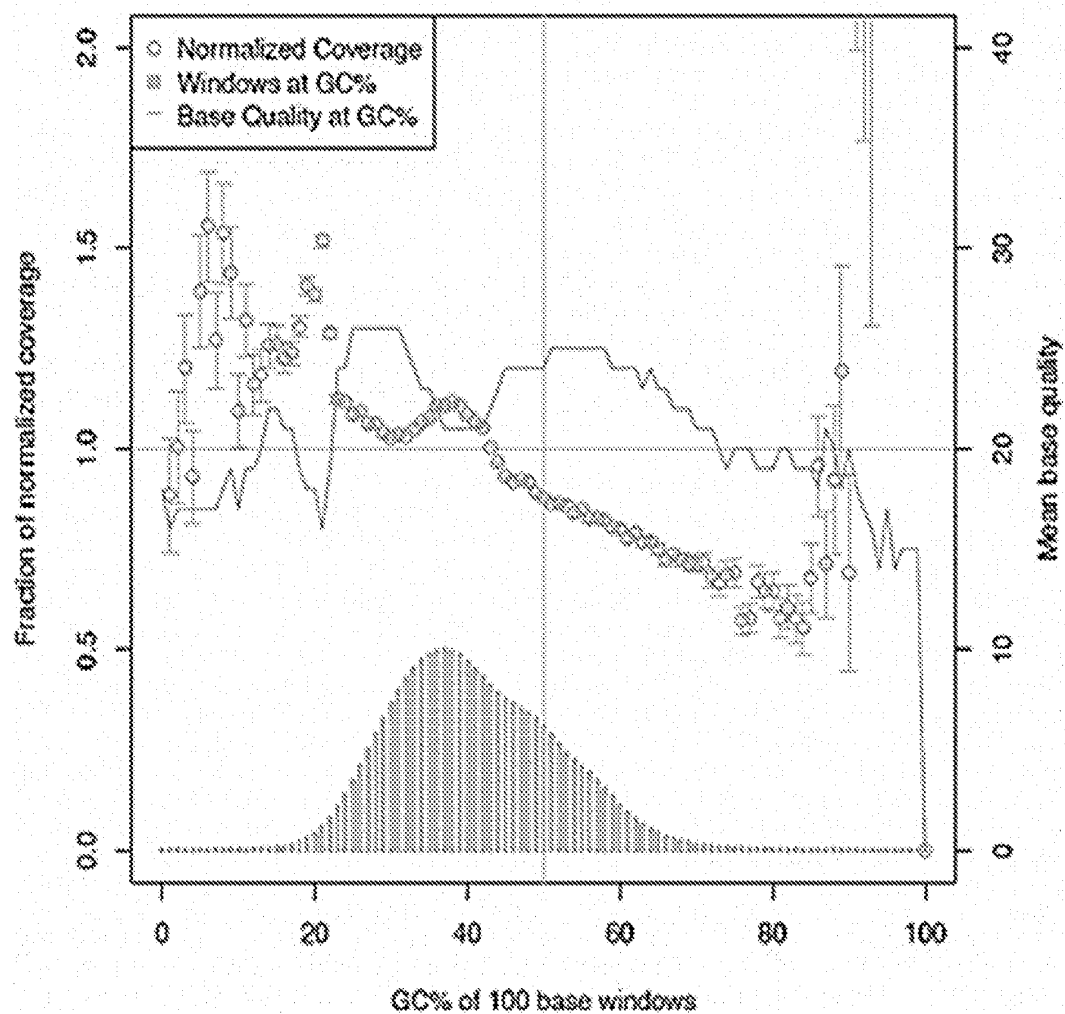
ad-Adapter-P5tr-Hairpin-No-CT-250ng-1_S5.bwa.sort.markdup.bam.Accel
Total clusters: 689,094, Aligned reads: 1,372,223



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Fig. 10D

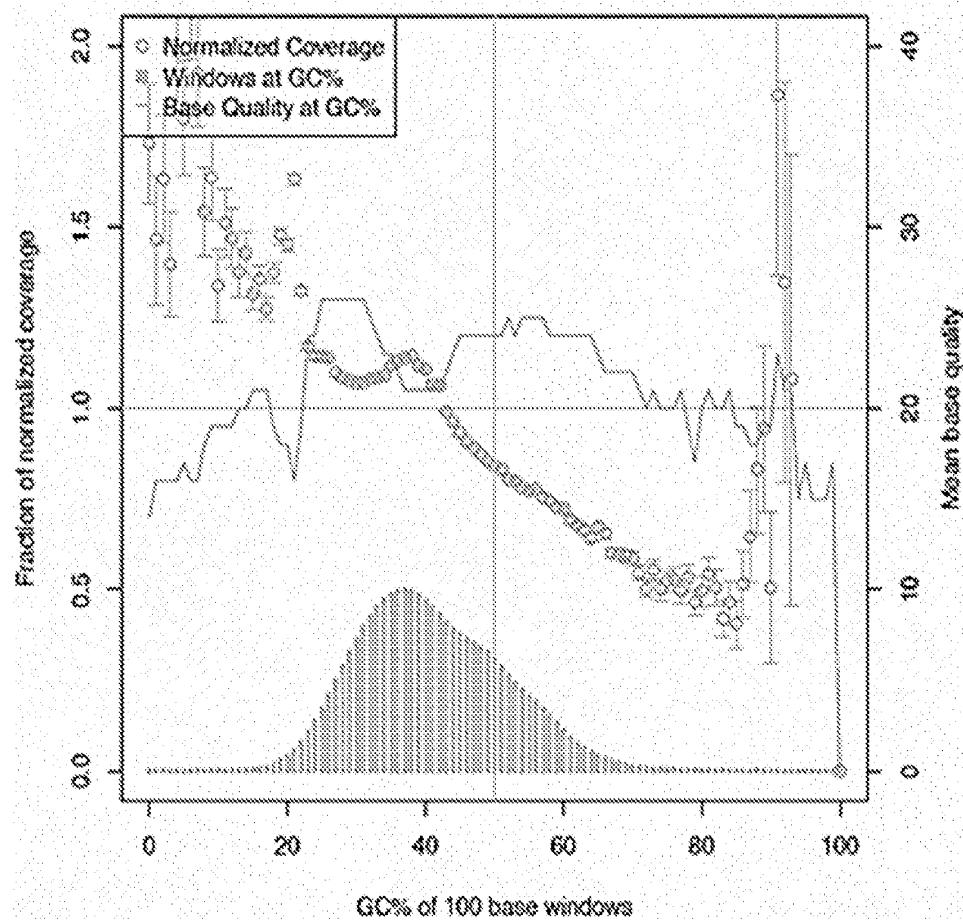
J-Adapter-P5tr-Hairpin-19-351-50ng-1_S7.bwa.sort.markdup.bam.Accel
Total clusters: 670,370, Aligned reads: 1,336,590



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Fig. 10E

-Adapter-P5tr-Hairpin-19-351-250ng-1_S11.bwa.sort.markdup.bam.Acce
Total clusters: 687,575, Aligned reads: 1,368,142



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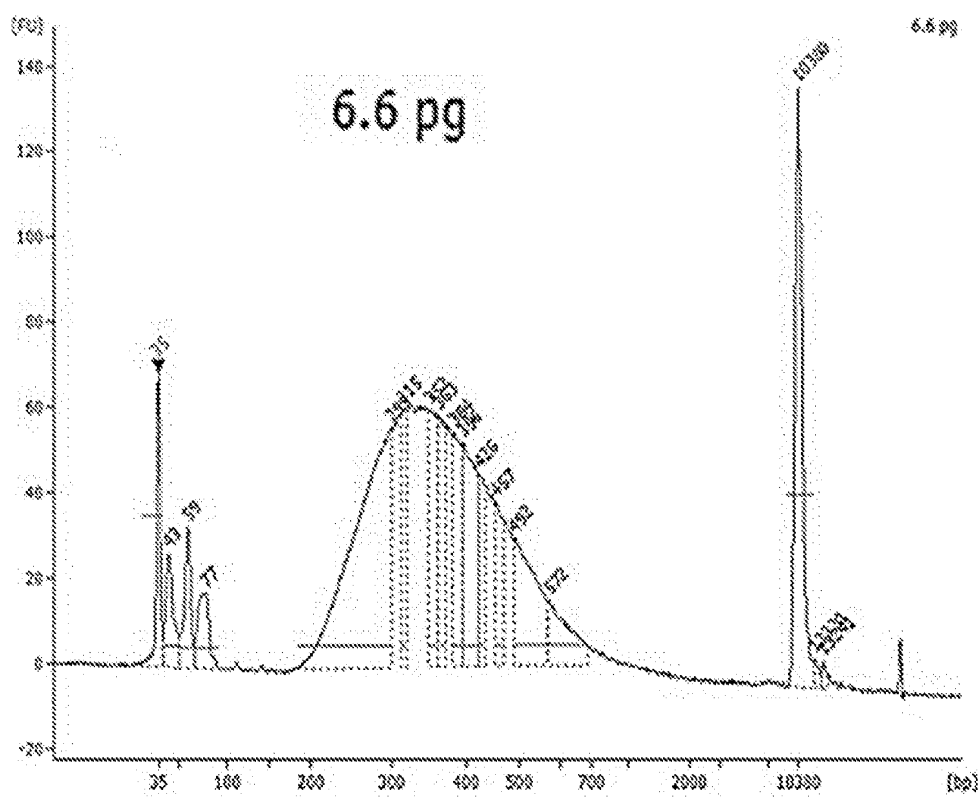
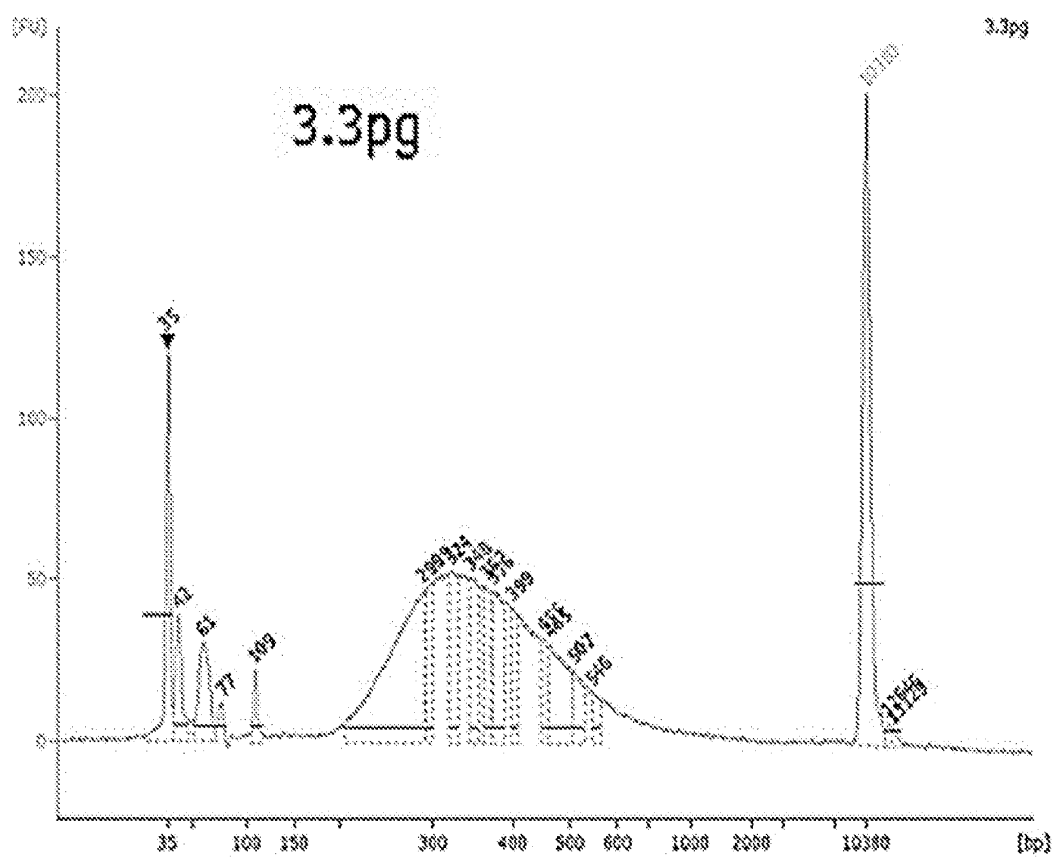
Fig. 11A

Fig. 11B



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Fig. 11C