CLAIMS

WHATS IS CLAIMED IS:

1. A method for preparing a polynucleotide library, the method comprising:
    a. in a first tailing reaction, adding a first tail to each of a plurality of target polynucleotides by template-independent polymerization, wherein the first tailing reaction comprises a first adapter comprising an overhang that hybridizes to the first tail;
    b. in a first ligation reaction, ligating a strand of the first adapter to the first tail;
    c. amplifying target polynucleotides comprising the strand of the first adapter by extending a first primer hybridized to the strand of the first adapter;
    d. in a second tailing reaction, adding a second tail to each of a plurality of the amplified target polynucleotides by template-independent polymerization, wherein the second tailing reaction comprises a second adapter comprising an overhang that hybridizes to the second tail; and
    e. in a second ligation reaction, ligating a strand of the second adapter to the second tail.
2. The method of claim 1, wherein the method comprises one or more of: (a) fragmenting polynucleotides to produce the target polynucleotides; (b) dephosphorylation of one or both ends of the target polynucleotides; and (c) denaturing double-stranded polynucleotides to single-stranded polynucleotides to produce the target polynucleotides.
3. The method of any one of the preceding claims, wherein the plurality of target polynucleotides comprises single-stranded DNA.
4. The method of any one of the preceding claims, wherein the target polynucleotides are treated prior to step (b) of claim 2 to differentially modify methylated cytosines or unmethylated cytosines.
5. The method of claim 4, wherein the differentially modifying comprises treating the target polynucleotides with bisulfite.

~~6. The method of any one of the preceding claims, further comprising amplifying target polynucleotides comprising the strand of the second adapter by extending a second primer hybridized to the strand of the second adapter.~~

~~7. The method of claim 6, further comprising an amplification reaction with a third primer and a fourth primer, wherein (i) the third primer hybridizes to a complement of at least a portion of the first primer, and (ii) the fourth primer hybridizes to a complement of at least a portion of the second primer.~~

~~8. The method of claim 7, wherein the third primer, the fourth primer, or both comprise an index sequence that identifies a sample source of the target polynucleotides.~~

~~9. The method of claim 7 or 8, further comprising sequencing amplification products of the amplification comprising the third and fourth primer.~~

~~10. The method of claim 9, further comprising grouping sequencing reads according to the index sequence.~~

~~11. The method of claim 9 or 10, wherein sequencing comprises detecting a sequence variant or a difference in nucleotide methylation, relative to a reference sequence.~~

~~12.~~ 1. A method for preparing a polynucleotide library, the method comprising:

a. in a first tailing reaction, adding a first tail to each of a plurality of target polynucleotides by template-independent polymerization, wherein the plurality of target polynucleotides comprises single-stranded polynucleotides, wherein the first tailing reaction further comprises hybridizing to the first tail an overhang within one strand of a first double-stranded adapter ~~comprising an overhang that hybridizes to the first tail~~;

b. in a first ligation reaction, ligating ~~a~~the other strand of the first adapter to the first tail;

c. amplifying target polynucleotides comprising the strand of the first adapter by extending a first primer hybridized to the strand of the first adapter; and

d. in a second ligation reaction, ligating a strand of a second adapter to the amplified target polynucleotides at the end opposite to the first primer.

~~13.~~ 2. The method of claim ~~12~~1, wherein the second adaptor is double-stranded, wherein the method further ~~ligation reaction~~ comprises:~~,~~

a. in a second tailing reaction, adding a second tail to each of a plurality of the amplified target polynucleotides by template-independent polymerization, wherein the second tailing reaction further comprises hybridizing to the second tail an overhang within one strand of the second adaptor; and

b. in the second tailing reaction, ligating the other strand of the second adaptor to the second tail.

~~14. The method of claim 13, wherein the second tailing reaction comprises a second adapter comprising an overhang that hybridizes to the second tail.~~

~~15. The method of claim 14, wherein, in the second ligation reaction, ligating a strand of the second adapter to the second tail.~~

~~16.~~ 3. The method of claim ~~12~~1, wherein the second adaptor is double-stranded, wherein (i) the second ligation reaction does not comprise a second tailing reaction; and (ii) the second ligation reaction comprises ~~a second adapter comprising an overhang that hybridizes~~ hybridizing to the amplified target polynucleotides an overhang within one strand of the second adaptor and directly ligating the other strand of the second adaptor to the amplified target polynucleotides.

~~17.~~ 4. The method of any one of claims ~~12-16~~1-3, wherein the method comprises one or more of: (a) fragmenting polynucleotides to produce the target polynucleotides; (b) dephosphorylation of one or both ends of the target polynucleotides; and (c) denaturing double-stranded polynucleotides to single-stranded polynucleotides to produce the target polynucleotides.

~~18. The method of any one of claims 12-17, wherein the plurality of target polynucleotides comprises single-stranded DNA.~~

~~19.~~ 5. The method of any one of claims ~~12-18~~1-4, wherein the target polynucleotides are treated prior to step (b) of claim ~~17~~4 to differentially modify methylated cytosines or unmethylated cytosines.

~~20.~~ 6. The method of claim ~~19~~5, wherein the differentially modifying comprises treating the target polynucleotides with bisulfite.

~~21.~~ 7. The method of any one of claims ~~12-20~~1-6, further comprising amplifying target polynucleotides comprising the strand of the second adapter by extending a second primer hybridized to the strand of the second adapter.

~~22.~~ 8. The method of claim ~~21~~7, further comprising an amplification reaction with a third primer and a fourth primer, wherein (i) the third primer hybridizes to a complement of at least a portion of the first primer, and (ii) the fourth primer hybridizes to a complement of at least a portion of the second primer.

~~23.~~ 9. The method of claim ~~22~~8, wherein the third primer, the fourth primer, or both comprise an index sequence that identifies a sample source of the target polynucleotides.

~~24.~~ 10. The method of claim ~~22 or 23~~8 or 9, further comprising sequencing amplification products of the amplification comprising the third and fourth primer.

~~25.~~ 11. The method of claim ~~24~~10, further comprising grouping sequencing reads according to the index sequence.

12. The method of claim 10 or 11, wherein sequencing comprises detecting a sequence variant or a difference in nucleotide methylation, relative to a reference sequence.

~~26.~~ 13. A kit for preparing a polynucleotide library, the kit comprising:

a. a template-independent polymerase;

b. a first pool of nucleotides that can be polymerized by the template-independent polymerase;

c. a second pool of nucleotides that can be polymerized by the template-independent polymerase;

d. a first adapter comprising an overhang that is hybridizable to tails formed by polymerizing the first pool of polynucleotides for producing amplified target polynucleotides; and

e. a second adapter comprising an overhang that is hybridizable to the amplified target polynucleotides.

~~27.~~ 14. The kit of claim ~~26~~13, further comprising a first primer that is hybridizable to a strand of the first adapter under conditions for a primer extension reaction.

~~28.~~ 15. The kit of claim ~~26 or 27~~13 or 14, further comprising a second primer that is hybridizable to a strand of the second adapter under conditions for a primer extension reaction.

~~29.~~ 16. The kit of claim ~~28~~15, wherein the sequence of the first primer that is hybridizable to the strand of the first adapter is different from the sequence of the second primer that is hybridizable to the second adapter.

~~30.~~ 17. The kit of claim ~~28 or 29~~15 or 16, further comprising a third primer and a fourth primer, wherein (i) the third primer is hybridizable to a complement of at least a portion of the first primer under conditions for a primer extension reaction, and (ii) the fourth primer is hybridizable to a complement of at least a portion of the second primer under conditions for a primer extension reaction.

18. The kit of any one of claim 13 to 17, wherein the overhang of the second adaptor comprises random bases.