Dec 11, 2020

O Amplification and Pooling

In 5 collections

DOI

dx.doi.org/10.17504/protocols.io.beqkjduw

Franziska Aron¹, Guido Brandt²

¹Friedrich-Schiller Universität Jena; ²Max Planck Institute for the Science of Human History

WarinnerGroup MPI EVA Archaeogenetics



Franziska Aron

Friedrich-Schiller Universität Jena, Max Planck Institute fo...





DOI: <u>dx.doi.org/10.17504/protocols.io.beqkjduw</u>

Protocol Citation: Franziska Aron, Guido Brandt 2020. Amplification and Pooling . **protocols.io** <u>https://dx.doi.org/10.17504/protocols.io.beqkjduw</u>

License: This is an open access protocol distributed under the terms of the <u>Creative Commons Attribution License</u>, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working We use this protocol and it's working

Created: April 07, 2020

Last Modified: December 11, 2020

Protocol Integer ID: 35308

Keywords: DNA library, NGS, dual-index, ancient DNA, sequencing, nonUDG, double-stranded, DNA, genomic DNA, genomics, palaeogenetics, archaeogenetics, paleogenetics, archeogenetics, aDNA, Illumina, library preparation, nucleic acids, Amplification, PCR, Index Amplification



Abstract

This protocol describes the amplification procedure of dual-indexed double-stranded DNA libraries, for shotgun Illumina sequencing. It is typically used for libraries indexed using the following protocol: (<u>https://dx.doi.org/10.17504/protocols.io.bakticwn</u>)

Image Attribution

Franziska Aron

Guidelines

Working in an Molecular Biology Laboratory

This protocol can place in a typical DNA-based molecular biology lab. Please keep in mind the safety guidelines of your specific country and institution.

Recommendations include wearing of:

- lab coats
- closed shoes and trousers
- safety glasses
- nitril or latex gloves

Materials

MATERIALS

8 0.2 ml PCR Tube strips Eppendorf Catalog #0030124359

X DNA LoBind Tube 1.5ml Eppendorf Catalog #022431021

2 ml LoBind Tubes Eppendorf Catalog #0030108078

X Eppendorf Tubes[®] 5.0 mL with snap cap Eppendorf Catalog #30119460

X dNTP Mix (25 mM each) Thermo Fisher Scientific Catalog #R1121

Sodium Acetate buffer solution 3M pH 52 for molecular biology **Merck MilliporeSigma (Sigma-Aldrich) Catalog** #S7899-500ML

X Tween 20 Merck MilliporeSigma (Sigma-Aldrich) Catalog #P9416-50ML

Water HPLC Plus Merck MilliporeSigma (Sigma-Aldrich) Catalog #34877-2.5L-M

X D1000 Ladder Agilent Technologies Catalog #5067-5586

X D1000 ScreenTape Agilent Technologies Catalog #5067-5582

X D1000 Reagents Agilent Technologies Catalog #5067-5583

X Herculase II Fusion DNA Polymerase Agilent Technologies Catalog #600679

X High Sensitivity D1000 Ladder Agilent Technologies Catalog #5067-5587

🔀 High Sensitivity D1000 Reagents Agilent Technologies Catalog #5067-5585

🔀 High Sensitivity D1000 ScreenTape Agilent Technologies Catalog #5067-5584

X MinElute PCR Purification Kit Qiagen Catalog #28004

Primers

	Oligo_ID	Sequence (5'-3')	Coce ntrati on
ſ	IS5	AATGATACGGCGACCACCGA	10 µM
ſ	IS6	CAAGCAGAAGACGGCATACGA	10 µM

Lab equipment

PCR Thermocycler (e.g. Eppendorf Thermomaster Nexus) Centrifuge 1.5/2.0 ml (e.g. Eppendorf 5424) Rotor 1.5/2.0ml (e.g. Eppendorf F-45-24-11) Mini table centrifuge TapeStation (e.g. Agilent Technologies, 4200 Tapestation System, SKU: G2991AA) Vortex mixer (e.g. Scientific Industries Vortex-Genie® 2)

Safety warnings

Reagents

Sodium Acetate

- H139: Causes serious eye irritation



Ethanol

- H225 Highly flammable liquid and vapour.
- H319 Causes serious eye irritation.





Guanidinium hydrochloride (GuHCI) (in PB buffer of Qiagen MinElute kit)

- H302 Harmful if swallowed.
- H332 Harmful if inhaled.
- H315 Causes skin irritation.
- H319 Causes serious eye irritation.



Kits

Check manufacturer's safety information for the TapeStation Kits used in this protocol.

Check manufacturer's safety information for the MinElute PCR Purification kit used in this protocol.

- Note that PBI must be stored at room temperature in the dark. PBI is light sensitive.

Before start

Planning

This protocol takes 1 day.

Check all waste disposal guidance for all reagents in this protocol against your corresponding laboratory regulations.

Preparation of buffers (Qiagen MinElute kit):

- Add ethanol to PE wash buffer acccording to manufacturer's instructions.
- Add $\underline{\square}$ 200 µL pH-Indicator and $\underline{\square}$ 300 µL Sodium Acetate to $\underline{\square}$ 48.5 mL of PB binding buffer. This solution is referred to as **PBI** throughout the protocol. Must be stored at room temperature in the dark. PBI is light sensitive.
- Add Tween-20 to EB elution buffer to a final concentration of 0.05% Tween-20 in EB. This solution is referred to as EBT throughout the protocol.

Equipment

Make sure all necessary equipment is available (see Materials).

Abbreviations

EBT = modified EB-Buffer (MinElute Kit), see Preparation of buffers HPLC = High Performance Liquid Chromatography (-Grade Water) PBI = modified PB-Buffer (MinElute Kit), see Preparation of buffers PE = PE-Buffer from Qiagen MinElute Kit

Samples

This protocol is designed for the amplification of indexed libraries as prepared by the protocol described in (<u>https://dx.doi.org/10.17504/protocols.io.bakticwn</u>). The indexing protocol generates $450 \,\mu$ of indexed library, of which $420 \,\mu$ will be used for this protocol. Ensure sufficient indexed library is available before starting this protocol.

Calculations

```
1 Prepare amplification assay [ \_ 100 \muL per reaction ]
```

Based on the quantification results of the indexed libaries (<u>https://dx.doi.org/10.17504/protocols.io.bakticwn</u>) calculate the number of PCR cycles (amplification factor) needed to reach 10^13 copies of DNA per indexed Library.

Note

Formula in Excel to get the Cycles needed

```
=LOG((1*10^13/Copies per rxn),2)
(log base 2)
```

Example: The following calculation is for $45 \text{ }\mu\text{L}$ per reaction, with two indexed library

samples (A and B) having different concentrations of DNA copies.

Optonial Changes: 1. If the Calculation shows up less then 3 Cycles, you also have the Option to add less then 5 μ l.

2. Instead of 4 reactions of 5 μ l each you can also split in 8 reactions

of 2 µl each

Instead of 4 × 5 μ l reactions you can also split in 8 × 2 μ l reactions

Samp le Nam e	Copi es per µl	µl per rxn	Copi es per rxn	Cycle s need ed	Real Cycle s	Ampl ificati on Facto r	Outp ut per rxn [Copi es]
A	7.32E +10	5	3.66E +11	4.772 998	5	32	1.17E+ 13
В	5.79E +06	5	2.32E +07	18.72 0182 51	19	5242 88	1.21E +13

Note

Do not calculate the amount of cycles for a higher amount of copies than 1.4 *10^13 to avoid heteroduplexes.

. .

Preparation

2 Prepare cleaned workspace with all necessary reagents and equipment.

Note		
Label all 🖉 0.2 mL PCR	strips for the PCR reactions.	

PCR

3

Set up four amplification reactions of $4 \text{ IOO } \mu \text{L}$ each per library

Reagent	Stock concentration	Final concentration	1x Volu me [μl]
Herculase II Reaction buffer	5x	1x	20
IS5 primer	10 μΜ	0.4 μΜ	4
IS6 primer	10 μΜ	0.4 μΜ	4
dNTP's	25 mM	0.25 mM	1
Herculase II Fusion	1U	0.01 U	1
DNA			5
HPLC-Water			65
Total			100

- 3.1 Vortex master mix before adding the enzyme. After adding the enzyme, mix by pipetting or inverting the tube.
- 3.2 Pipette Δ 95 µL mastermix and Δ 5 µL indexed library into each tube (use Δ 0.2 mL PCR strips).

X

Ø

Safety information

Keep the remaining library at $[-20 \circ C]$ until further use.

4 Amplify in a thermocycler with the following program:

Tempreature	Time	
95°C	2 min	Inital denat uratio n
95°C	30 sec	Cycle
60°C	30 sec	(see
72°C	30 sec	1)
72°C	5 min	Final elong ation
Finally ho	ld the reactions at 10 °C.	

Note

Adjust the number of cycles according to the amplification factor as calculated in step 1.

During this incubation take **MinElute columns** out of the fridge so they warm up to **room temperature** before use in the next step.

This is an ideal point to prepare downstream steps, including labelling of final elution tubes, MinElute Columns etc.

MinElute Purification

5	5 Purify with MinElute kit with the following modifications to the manufacturer's protocol:				
	Use one column for all four reactions [= 400μ L PCR product] of a sample.				
5.1	Add $\boxed{4}$ 2400 µL PB or PBI* buffer to a $\boxed{4}$ 5 mL tube for each sample (this is				

 \triangleq 600 µL buffer for each PCR reaction). Add all 4 PCR reactions per sample to the

X

Z

same tube with PB buffer and vortex briefly.

	Safety information
	After the PCR product is mixed with the PBI, the PBI should keep its yellow colour. If it turns purple the pH is too high and the efficency of the MinElute columns is not guaranteed.
5.2	Load $\boxed{1}$ 700 µL of the mixture onto one MinElute column, incubate for $\textcircled{0}$ 00:02:00, spin $\textcircled{1}$ 15800 x g, 00:01:00, and discard flow-through.
	Note
	Pour off the liquid into a waste tube, and pat the rim of the collection tube dry on a paper tissue or towel. Use just one spot on the paper tissue per sample. Be careful not to touch the rim of the tube on the waste container. After you are finished with all samples, discard the paper and wipe clean the surface underneath with water and soap.
53	Descet les dis substitutes complete mistrue une mus through the column
5.5	ED go to step #5.2
5.4	Add \blacksquare 700 µL PE (wash) buffer, spin $\textcircled{15800 \times g, 00:01:00}$, and discard flow-through.
5.5	Dry spin 😯 15800 x g, 00:01:00 ,
5.6	Put column into new 📕 1.5 mL LoBind tubes.
5.7	Add $4 50 \mu$ EBT buffer to the center of the filter, incubate for $2000000000000000000000000000000000000$

Note

Carefully pipette EBT directly onto the center of the membrane without touching the membrane.

Measurement and Dilution

6 Dilute amplified index library 1:10 with HPLC- water and check for fragment size, concentration, and heteroduplexes. (for example with the D1000 Kit's Tape, Reagent and Buffer - following the manufacturer's protocol on the TapeStation)

Note

if you see heteroduplexes you need to perform a reconditoning PCR.

Reconditoning PCR: one cycle PCR using 100 ng library template in a 100 μ l Herculase PCR reaction (same set up as in 3) and amplified with 1 cycles of 95°C for 2 min, 58°C for 2 min, and 72°C for 5 min. Purify with MinElute kit following the instructions from Step 5, but elute in 20 μ l EBT.

7 Dilute each amplified indexed library to <u>A 10 nM</u> with EBT buffer or HPLC-water for shotgun sequencing. Then pool the <u>A 10 nM</u> amplified indexed libraries in equimolar amounts (take the same volume for each sample).

Note

The final concentration of a pool of several amplfied indexed libraries should be $_$ 10 nM $_{-}$

8 Check the <u>I 10 nM</u> library or the <u>I 10 nM</u> library pool for the correct concentration, (for example with the HighSensitivity D1000 Kit's Tape, Reagent and Buffer following the manufacturer's protocol on the TapeStation.)