

Feb 13, 2024

Version 1

ssDNA2.0: Ligation mix I V.1

DOI

dx.doi.org/10.17504/protocols.io.6qpvr3j6bvmk/v1

Matthias Meyer¹, Sarah Nagel¹, Anna Schmidt¹

¹Max Planck Institute for Evolutionary Anthropology

MPI EVA Ancient DNA C...



Matthias Meyer

Max Planck Institute for Evolutionary Anthropology

Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN  ACCESS



DOI: <https://dx.doi.org/10.17504/protocols.io.6qpvr3j6bvmk/v1>

Document Citation: Matthias Meyer, Sarah Nagel, Anna Schmidt 2024. ssDNA2.0: Ligation mix I. **protocols.io**
<https://dx.doi.org/10.17504/protocols.io.6qpvr3j6bvmk/v1>

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

Created: February 13, 2024



Last Modified: April 10, 2024

Document Integer ID: 95158

Keywords: stranded dna library preparation, dna library preparation, stranded dna library, dna libraries for the sequencing, stranded dna, ligation mix, first adapter ligation, dna from ancient biological remain, degraded dna, dna, protocol for the preparation, sequencing, ancient biological remain

Funders Acknowledgements:

Max Planck Society

Abstract

Protocol for the preparation of Ligation mix I for the first adapter ligation in automated single-stranded DNA library preparation using the ssDNA2.0 method (Gansauge et al. 2020).

References

Gansauge, M.-T., Aximu-Petri, A., Nagel, S., & Meyer, M. (2020). Manual and automated preparation of single-stranded DNA libraries for the sequencing of DNA from ancient biological remains and other sources of highly degraded DNA. *Nature Protocols*, 15, 2279-2300.

Troubleshooting

Note

The volume of Ligation mix I suffices for one 96-well library preparation plate (96 + 20 reactions to account for dead volumes and loss of reagent). It is advisable to prepare 10-20 mixes at once.

Materials

Reagent/consumable	Supplier	Catalogue number	Decontamination *
Reagents			
Water, HPLC-grade	Sigma Aldrich/Merck	1153332500	UV
50% PEG-8000 (w/v)	Jena Bioscience	CSS-256	UV
100 mM ATP	Thermo Fisher Scientific	R0441	-
Consumables			
5 ml screw cap tubes (rack 2d Lp W/barcode)	VWR	NUNC374320-BR	-

* Decontamination of reagents and consumables should be performed as detailed in the documents below.

Equipment

- Label printer (e.g. Brady M611, cat. no. M611-EU-LABS) and tube labels (e.g. Labels for TLS2200/TLS PC Link/Polyester, cat. no. PTL-82-499)

Protocol

1. Prepare a 5 ml tube with Ligation mix I by adding the following reagents. Pre-mix reagents by slowly inverting the tube at least two times and spin down (full mixing is performed later, after enzyme addition).

Reagent	Volume (μl)	Final concentration in reaction
Water	116	
50% PEG-8000 (w/v)	3712	20%

	Reagent	Volume (μl)	Final concentration in reaction
	100 mM ATP	46.4	0.5 mM
	<i>sum</i>	<i>3874.4</i>	

Note

[Labeling]

Prepare tube labels using Brady printer including name of the mix, date (dd.mm.yyyy) and the name of the person who prepared the Ligation mix I.

2. Freeze at -20 °C until used.

Note

[Documentation]

Note the lot/batch numbers of the reagents used for master mix preparation in Labfolder (orange fields).

Appendix

Document

NAME

UV decontamination of reagents/buffers

CREATED BY

Elena Essel

Preview

