

Dec 12, 2022

Chloroform-methanol protein precipitation from microalgae and Pierce BCA assay

DOI

dx.doi.org/10.17504/protocols.io.yxmvm2e25g3p/v1

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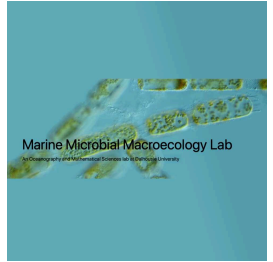
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Protocol Citation: Ying-Yu Hu, Christopher Lord, Zoe V Finkel 2022. Chloroform-methanol protein precipitation from microalgae and Pierce BCA assay. **protocols.io** <https://dx.doi.org/10.17504/protocols.io.yxmvm2e25g3p/v1>

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Protocol status: Working

We use this protocol and it's working

Created: November 28, 2022

Last Modified: December 12, 2022

Protocol Integer ID: 73284

Keywords: Protein , Microalgae, Chloroform-methanol precipitation, Pierce BCA assay, methanol protein precipitation from microalgae, methanol protein precipitation, bca assay, pierce bca protein, extracted protein, methanol, chloroform, detergent in crude protein, crude protein, protein, chlorophyll, microalgae, phospholipid

Funders Acknowledgements:

Simons Collaboration on Computational Biogeochemical Modeling of Marine Ecosystems

Grant ID: 549937


Ocean Processes and Ecology

Grant ID: 723789


Abstract

Chlorophyll, phospholipids, sucrose, glycerol and some detergent in crude protein extracted from microalgae can interfere the Pierce BCA protein assay. In order to remove these interference, bead miller extracted protein is precipitated by chloroform-methanol prior to BCA assay. The resulting precipitation is dissolved into Sarcosine-Tris solution. Low limit of detection is about 5 ug/mL.


Protocol materials

 Tris(hydroxymethyl)aminomethane hydrochloride 1M pH 8.0 RNase free **Fisher Scientific Catalog #AAJ60080AK**

 N-lauroylsarcosine sodium salt solution (30%) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #61747**

 Methanol **Merck MilliporeSigma (Sigma-Aldrich) Catalog #34860**

 Chloroform (HPLC grade) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #439142-4L**

 Thermo Scientific™ Pierce™ Bovine Serum Albumin Standard 2 mg/mL (50 mL) **Thermo Scientific Catalog #Thermo Scientific™ 0023210**

 Pierce BCA Protein Assay Kit **Thermo Fisher Scientific Catalog #23225**

Troubleshooting



Safety warnings

! Use fume-hood when handling methanol and chloroform.

All waste containing methanol and chloroform shall be collected in waste container for halogenated organic solvents.



Reagent preparation

1 Tris buffer [M] 5 Mass Percent (pH 8.0)

Add  500 μ L [M] 1 Mass Percent  8.0 Tris into 100 mL MilliQ water



Tris(hydroxymethyl)aminomethane hydrochloride 1M pH 8.0 RNase free **Fisher Scientific Catalog #AAJ60080AK**

2 20% Sarcosine

Dilute 2 part 30% N-lauroylsarcosine sodium salt with 1 part [M] 5 Mass Percent (pH 8.0) Tris buffer



N-lauroylsarcosine sodium salt solution (30%) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #61747**

Protein precipitation

1h 12m

3 Thaw protein extract

4 Turn on refrigerate centrifuge

Equipment

CENTRIFUGE 5430 R


NAME

Eppendorf

BRAND

MP2231000510

SKU

5 Turn on incubator/shaker, preheat to  37 °C

**Equipment****SHAKING INCUBATOR**

NAME

71L

TYPE


Corning® LSE™

BRAND

6753

SKU

6 Prepare ice-bath

7 Well mix the extract and then transfer  100 µL of extract to 2 mL microtube (Abdos tubes give better precipitation results), in replicate.

Equipment**Micro Centrifuge Tubes**

NAME

Abdos

BRAND

P10203

SKU


Note


If extract has debris, spin down debris by




13300 rpm, Room temperature, 00:05:00 and transfer only clear supernatant.

Debris can cause overestimation of protein content.

8 In the fume hood, add  400 μ L methanol

 Methanol Merck MilliporeSigma (Sigma-Aldrich) Catalog #34860

9 Gently vortex for  00:00:30 by using a tube insert

30s

Equipment

VWR ANALOG VORTEX MIXER

NAME

VWR


BRAND

10153-838


SKU

With tube insert


SPECIFICATIONS


10 In the fume hood, add  100 μ L chloroform

 Chloroform (HPLC grade) Merck MilliporeSigma (Sigma-Aldrich) Catalog #439142-4L

11 Gently vortex for  00:00:30 by using a tube insert

30s


12 In the fume hood, add  300 μ L MilliQ

13 Gently vortex for  00:00:30 by using a tube insert


30s

14 Incubate  On ice for  00:30:00

30m


15  20000 rcf, 4°C, 00:10:00

10m

16 In the fume hood, remove upper phase by leaving about  250 μ L liquid

**Note**


Do not disturb the interphase

17 In the fume hood, add  300 μ L methanol

18 Gently mix the liquid until bottom layer disappear and the solution is homogenous.

Note

The formation of small pellet might be observed, but might be invisible due to low protein mass.



19  20000 rcf, 4°C, 00:10:00

10m

20 In the fume hood, remove all solvent.



Note

Watch the pipet closely. Do not remove pellets with the solvent.

21 If pellet tends to be aspirated with solvent, add another  300 μ L methanol, gently vortex, and  20000 rcf, 4°C, 00:10:00

10m

22 In the fume hood, remove most solvent by using 1000 μ L pipet tip, and then remove the rest by using 100 μ L pipet tip. Do not remove pellet with solvent.

23 Dry pellet in vacuum desiccator for at least  00:30:00 at  Room temperature

30m



Note

Any methanol and chloroform residue can affect the re-dissolving of pellet in BCA assay. However, do not dry protein pellet for too long, otherwise it might be difficult to re-dissolve.



BCA assay

30m

24 Add  5 μL 20% sarcosine and  95 μL [M] 5 Mass Percent (pH 8.0) Tris buffer to dry protein pellet, incubate at  37 $^{\circ}\text{C}$ for 15 to 30 min.


25 Use tube insert, vortex all tubes for 15 to 30 min until pellet is completely re-dissolved.

26 BSA standard solutions



Thermo Scientific™ Pierce™ Bovine Serum Albumin Standard 2 mg/mL (50 mL) **Thermo Scientific Catalog #**Thermo Scientific™ 0023210

	Standard	20% sarcosine (uL)	5 mM Tris (uL)	2 mg/mL BSA (uL)	Final Conc. (mg/mL)
	SD1	5	95	0	0
	SD2	25	470	5	0.02
	SD3	25	463	12	0.048
	SD4	25	450	25	0.1
	SD5	25	425	50	0.2
	SD6	25	375	100	0.4
	SD7	25	275	200	0.8
	SD8	25	225	250	1

27 Vortex and then use reverse pipetting: transfer  100 μL standard solutions into the corresponding tubes, except for SD1 (it has already been 100 uL).

28 Use the following formula to determine the total volume of working reagent (WR) required. Consider leaving several mL of extra volume:


$$(\# \text{ standards} + \# \text{ samples}) \times \left(\text{img alt="pipette tip icon" data-bbox="378 751 398 771"} 800 \mu\text{L} \right) = \text{total volume WR required}$$

29 Prepare WR by mixing 50 parts of BCA reagent A with 1 part of BCA Reagent B in a 50 mL falcon tube



Pierce BCA Protein Assay Kit **Thermo Fisher Scientific Catalog #**23225



- 30 Use one tip and reverse pipetting: Add  800 μL WR into each tube, make sure that the tip doesn't have contact with the solution, so that samples are not cross-contaminated.

Note

Since BCA assay is sensitive to duration, although reagent is aqueous, it is more efficient to use reverse pipetting and quickly dispense reagent into all tubes, therefore the duration difference amongst standards and samples can be minimized.



- 31 Vortex each tube, shake and incubate at  37 °C for  00:30:00

30m

- 32 Remove samples from the incubator.

- 33 Load samples into microplate in duplicate:

Note

1. Reverse pipetting: aspire  200 μL sample from the middle of solution
2. Tip gently touches the side of the well, avoid bending. Dispense 200 μL into the microplate
3. Dispose the tip
4. Use a new tip, reverse pipet another  200 μL as replicate
5. Tip gently touches the side of the well, avoid bending. Dispense 200 μL into the microplate

Equipment**96-Well Microplates**

NAME

Polystyrene, Clear,

TYPE

Greiner Bio-One

BRAND

82050-760

SKU



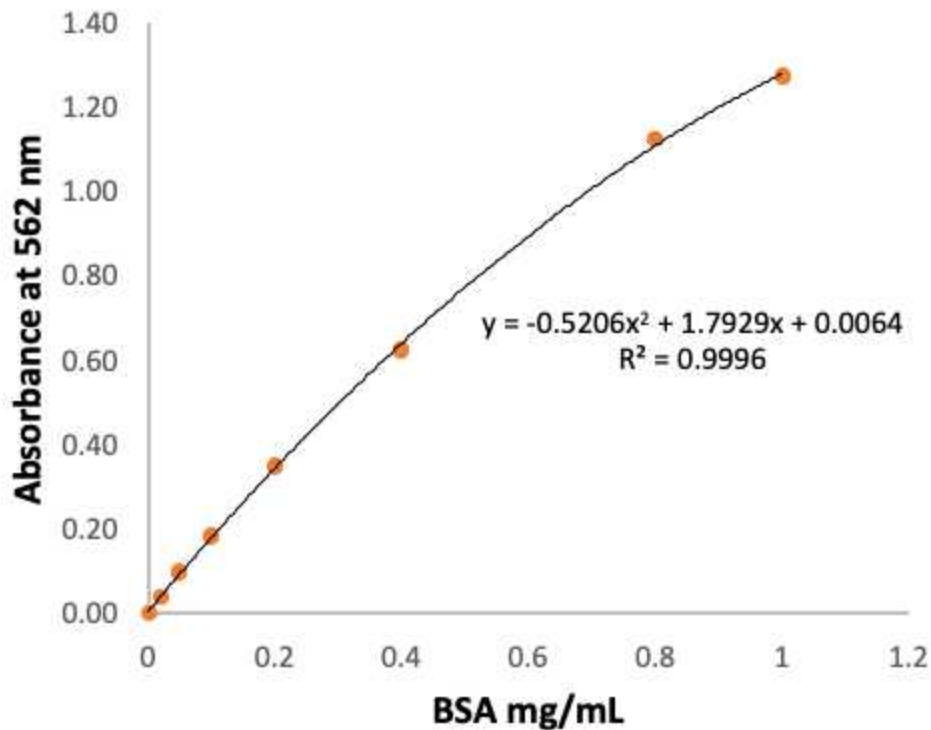
- 34 Shake for 5 s at 600 rpm in a continuous and high force mode
Read endpoint 562 nm with a measurement time 100 ms

Equipment

Varioskan LUX Multimode Microplate Reader	NAME
Thermo Fisher	BRAND
VL0L00D0	SKU

Calculation

- 35 Subtract the average 562 nm absorbance measurement of the blank standard replicates from the 562 nm measurements of all other individual **standard**.
- 36 Subtract the average 562 nm absorbance measurement of the blank sample (filter) replicates from the 562 nm measurements of all other individual **sample**.
- 37 Prepare a standard curve by plotting the average Blank-corrected 562 nm measurement for each BSA standard versus its concentration in mg/ml. The standard curve is quadratic.



- 38 For the calculation convenience, plot BSA concentration (Conc) versus Corrected absorbance (Abs) to obtain a standard curve as following:
$$\text{Conc_mg/mL} = a \times \text{Abs}^2 + b \times \text{Abs} + c$$

Use the corrected measured absorbance of samples (Abs) to calculate the total protein concentration (Conc_mg/mL) from each sample.

- 39
$$\text{Protein_mg/filter} = \text{Conc_mg/mL} \times \text{PEB_mL}$$

Where PEB is the volume of protein extraction buffer used to extract protein from microalgae sample.