GeneAll[®] Application Note

Total RNA extraction from *Ulva linza Linnaeus* samples using Ribospin™ Plant kit

Experimental Conditions

Materials Required

- Ribospin[™] Plant kit (307-150, 50 preps)
- Tissuelyser II (85300, Q) (or another bead beating device)
- Liquid nitrogen (LN₂)
- Absolute ethanol (C₂H₆O, CAS No. : 64-17-5, ≥99.0%)
- 70% ethanol (C₂H₆O, CAS No. : 64-17-5, 70.0%)
- 1.5 ml or 2.0 ml microcentrifuge tube
- Vortex mixer
- Centrifuge (Max. speed 14,000 rpm)
- Pipette & sterile pipette tips
- Suitable protector (ex. lab coat, disposable gloves, goggles, etc.)
- Ice (for incubation)

Sample Information

• Sample type :



Ulva linza Linnaeus

- Sampling : After collecting fresh *Ulva linza Linnaeus* samples, put them in a bag and seal it.
- How to store : Store in -70°C deep freezer
- Homogenizing : Tissuelyser II (85300, Supplier : Q)
- Conditions :
- Sample amount : 25, 50 and 100 mg
- Elution volume : 50 μl

Protocol

Before experiment

- Before using for the first time, add absolute ethanol (ACS grade or better) into Buffer RBW and RNW as indicated on the bottle.
- Prepare DNase I reaction mixture just before step 8.
 Prepare aliquot DNase I and thaw on ice.
 - Mix 2 µl DNase I with 70 µl Buffer DRB.

Sample preparation

- 1. After measuring 25, 50 and 100 mg of each sample, put it into a 1.5 ml microcentrifuge tube. Then, it is rapidly frozen using LN_2 .
- 2. Grind the sample using TissueLyser II (30 Hz, 10 sec).
- 3. The next step is according to the protocol of Ribospin[™] Plant.

Ribospin[™] Plant brief protocol

- * For more details, refer to the handbook of Ribpsin[™] Plant
 - 1. Transfer the powder into a 1.5 ml microcentrifuge tube.
 - 2. Add 350 μl of Buffer RPL and incubate the mixture for 3 min at room temperature.
 - 3. Transfer the lysate to EzPure[™] Filter and centrifuge at ≥10,000 x g for 30 sec.
 - 4. Transfer the supernatant into a 1.5 ml microcentrifuge tube.
 - 5. Add 1 volume of 70% ethanol to the supernatant and mix well.
 - 6. Apply the mixture into Column Type W and centrifuge at \geq 10,000 x g for 30 sec.
 - 7. Add 500 μl of Buffer RBW to Column Type W and centrifuge at \geq 10,000 x g for 30 sec.
 - 8. Apply the DNase I mixture into Column Type W.
- 9. Incubate the mixture for 10 min at room temperature.
- 10. Add 500 μ l of Buffer RBW to Column Type W and incubate for 2 min and centrifuge at \geq 10,000 x g for 30 sec.
- 11. Add 500 μ l of Buffer RNW to Column Type W and centrifuge at \geq 10,000 x g for 30 sec (twice).
- 12. Centrifuge at \geq 10,000 x g for an additional 1 min.
- 13. Add 50 μI of Nuclease-free water to the center of the membrane.
- 14. Centrifuge at $\geq 10,000 \times g$ for 1 min.

Result

#	Sample	Conc. (ng/µl)	A260/280	A260/230	Yield (µg)
1	Ulva linza Linnaeus (25 mg)	100.2	2.19	2.33	5.0
2		111.3	2.24	2.45	5.6
3	Ulva linza Linnaeus (50 mg)	187.7	2.21	2.24	9.4
4		183.6	2.25	2.42	9.2
5	Ulva linza Linnaeus (100 mg)	256.6	2.24	2.44	12.8
6		334.2	2.25	2.46	16.7

Table 1. The concentration, yield and and purity of RNA extracted from 25~100 mg of Ulva linza Linnaeus samples.

% Remark

Absorbance measurement instrument : NanoDrop[™] 2000/2000 c (ND-2000, Supplier : T)

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Figure 1. The result of electrophoresis of total RNA from Ulva linza Linnaeus sample. Lanes $1 \sim 2 : 2 \mu$ l of eluate from 25 mg of Ulva linza Linnaeus Lanes $3 \sim 4 : 2 \mu$ l of eluate from 50 mg of Ulva linza Linnaeus Lanes $5 \sim 6 : 2 \mu$ l of eluate from 100 mg of Ulva linza Linnaeus Lane M : GENESTA^M 1 kb DNA ladder with 5X loading dye (GA-100, GeneAll[®], 1 μ l loading)

Lane M : GENESIA^{....} 1 KO DNA ladder with 5X loading dye (GA-100, GeneAli[®], 1 µ loading) ※ Remark

Electrophoresis conditions : 1.0% agarose gel (150 V, 17 min, 2 µl loading)



Figure 2. The result of electrophoresis of PCR products after RT-PCR. Lanes 1~2 : 2 µ of eluate from 25 mg of Ulva linza Linnaeus Lanes 3~4 : 2 µ of eluate from 50 mg of Ulva linza Linnaeus Lanes 5~6 : 2 µ of eluate from 100 mg of Ulva linza Linnaeus Lane M : GENESTA[™] 100 bp DNA ladder with 5X loading dye (GA-010, GeneAll®, 1 µl loading) Lanes P1~P2 : Positive control (total RNA template from pine leaf)

% Remark

Primer: NAd5 gene primer (gene of mitochondrial plant cell) Electrophoresis condition : 2.0% agarose gel (150 V, 20 min, 3 μl loading)

Lane N : Negative control (nuclease-free water)

Conclusion

The *Ulva Linnaeus* is well known for the wide distribution throughout the world and the research about its genomic information is widely conducted owe to its major contribution to biofouling.

The isolation of high-quality RNA from *Ulva Linnaeus* is the initial but the most import step for the successful research but there has always been problematic due to a high content a polyphenolic compounds and polysaccharides in the sample.

This study presents simple and effective method for the high-quality RNA isolation from *Ulva Linnaeus* in different starting amounts using Ribospin[™] Plant Kit.

The RNA obtained from *Ulva Linnaeus* was found to be suitable for many downstream applications such as RT-PCR, PCR, qPCR, and etc.