



Optimal sugar release from macroalgal feedstock with dilute acid pretreatment and enzymatic hydrolysis

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Abstract

Production of bioethanol from macroalgal feedstock involves pretreatment and hydrolysis of biomolecules for release of sugar followed by the fermentation of the sugar. The most commonly used pretreatment method for macroalgal biomass is dilute acid hydrolysis using mineral acids H_2SO_4 and HCl . The optimization of dilute acid hydrolysis of *Ulva lactuca* yielded 214.67 mg g^{-1} using $0.5 \text{ N } H_2SO_4$, 5% w/w substrate concentration at $121 \text{ }^\circ\text{C}$ for 45 min, and hydrolysis of *Enteromorpha intestinalis* yielded 239.94 mg g^{-1} of reducing sugar using $0.7 \text{ N } H_2SO_4$, 5% w/w substrate concentration at $120 \text{ }^\circ\text{C}$ for 45 min. Crude enzyme extracted from marine bacteria *Vibrio parahaemolyticus* and purified by the two-step purification produced 61.82% yield with 2.97-fold purification. Enzymatic hydrolysis of pretreated macroalgal biomass produced onefold higher reducing sugar than acid hydrolysis for *Ulva lactuca* ($261.76 \pm 0.9 \text{ mg g}^{-1}$) and *Enteromorpha intestinalis* ($289.89 \pm 2.4 \text{ mg g}^{-1}$).

Keywords Pretreatment · Dilute acid · Enzyme · Optimization · Macroalgae · Reducing sugar

1 Introduction

The exploitation of finite fossil fuel resources has given rise to increased price fluctuations and elevated greenhouse gas emissions, contributing mainly to global warming. These drawbacks have escalated the need for alternative, renewable, sustainable, and economically viable energy resources such as carbohydrate-rich biomass to produce bioethanol. Bioethanol production is obtained from the carbohydrate fraction of the biomass, which is extracted and fermented [1–4]. Biomass conversion involves the separation of carbohydrate fraction to simple sugar through pretreatment methods, a vital step in biofuel production [5]. Several attempts have been made towards pretreatment of macroalgae or

seaweeds [6–11], including chemical, physical, or biological or combination of these techniques. Pretreatment is carried out to enhance the surface area of the feedstock for the release of the constituent fermentable (reducing) sugars, which depends on the characteristics of chosen feedstock, catalysts, operation parameters, and strength [8, 12, 13]. Macroalgal biomasses are composed of a wide range of polysaccharides such as cellulose, ulvan, laminarin, Floridan starch, etc. These polysaccharides are broken down into monosaccharides which serve as raw materials for bioethanol production. Constituents of these monosaccharides vary in the macroalgal biomass, as summarized in Table 1.

Macroalgae with higher moisture (80–85%) content and devoid of lignin polymer is well suited for microbial conversion than combustion or thermochemical conversion [14, 33, 34]. Also, the absence of lignin avoids the necessity of employing harsh pretreatment processes required in lignocellulosic biomass [14, 33]. Pretreatment of biomass is carried out for (i) size reduction and (ii) alter or remove structural and compositional impediments before enzymatic hydrolysis. Pretreatments are required to be cost-effective with minimal inhibitor formation while releasing a higher quantum of sugar.

Physicochemical pretreatment involves liquid catalysts with higher process conditions to treat the feedstock.

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Table 1 Reducing sugar composition of macroalgae

Sources	Glu	Xyl	Man	Gal	Ara
<i>Ulva</i> sp.	8.2	4.5	0.29	1	0.08
<i>Enteromorpha</i> sp.	26.3	3.5		6	
<i>Gracilaria</i> sp.	24	0.3	0.07	42.8	0
<i>Eucheuma</i> sp.	0.78			22.39	
<i>Porphyridium</i> sp.	16.9	4.7	0.1	5.3	0
<i>Padina</i> sp.	1.4	1.3	0	0.75	0.04
<i>Sargassum</i> sp.	22.5	0.5	4	4	
<i>Palmaria</i> sp.	3.8	31.1	0	5.5	0
<i>Laminaria</i> sp.	18.5	0.24	0.33	0.49	0.05
<i>Undaria</i> sp.	0.11	0.13	0.1	0.58	0.05

Glu, glucose; Xyl, xylose; Gal, galactose; Man, mannose; Ara, arabinose

Source [14–32]:

Pretreatment using chemicals such as acid, alkaline, and ammonia fiber expansion as well as soaking in aqueous ammonia and inorganic salts has been tried and is economical [8, 9, 35]. *Ulva lactuca* feedstock was subjected to four different pretreatments: ethanol organ solvent, alkaline, liquid hot water, and ionic liquid treatments. Organosolvent and liquid hot water treatment produced the highest sugar recovery of 808 mg g⁻¹ dry weight (DW) and 629 mg g⁻¹ DW, respectively [35]. In the hot water pretreatment, holes on algal feedstock surface (observed under scanning electron microscopy) indicated crystallinity (index of 97.5%) and cracks, which has enhanced enzyme digestibility of the feedstock. *Gelidium amansii* pretreated with 0.05–0.2 N Ca (OH)₂ at 121 °C for 15 min resulted in gel formation. Hence, alkaline pretreatment is not opted for pre-processing of macroalgal feedstock, especially red and brown macroalgae containing hydrocolloids such as agar, carrageenan, and algin [36]. The most commonly used chemical pretreatment method employs mineral acids such as H₂SO₄ and HCl at milder concentrations of 0.3–0.9 N [9, 10]. Various reaction parameters such as reaction time, acid concentration,

and substrate concentration are involved for efficient sugar release from the macroalgal feedstock. Pretreatment with dilute H₂SO₄ at different concentrations (~0.5–1%) and moderate temperature (~140–190 °C) [37] has been used widely for macroalgal cell wall depolymerization. Energy consumption in acid pretreatment is comparatively low compared to other pretreatments as it requires lower temperature as well as lesser incubation time (Table 2). Sulfuric acid reduces the production of inhibitors and improves the solubilization of seaweed polysaccharides [38]. The US National Renewable Energy Laboratory study reveals that the use of dilute acid (0.5–1%; 160–180 °C for 10 min) pretreatment aided in the release of different simple sugars (xylose, arabinose, galactose, glucose) [39, 40].

Reducing sugar (RS) released using H₂SO₄ from various macroalgal species such as *Gracilaria verrucosa* (430 mg g⁻¹ RS, 1.5% H₂SO₄) [46]; *Kappaphycus alvarezii* (300 mg g⁻¹ RS, 0.9 N H₂SO₄) [47]; *Gracilaria verrucosa* (7 g L⁻¹ RS, 373 mM H₂SO₄) [48]; *Laminaria japonica* (29.09% RS, 0.06% H₂SO₄) [49]; *Kappaphycus alvarezii* (81.62 g L⁻¹ RS, 1% v/v H₂SO₄) [15]; *Gelidium amansii* (33.7% RS, 3% H₂SO₄) [10]; *Gracilaria verrucosa* (7.47 g L⁻¹ RS, 0.1 N H₂SO₄) [50]; and *Kappaphycus alvarezii* (30.5 g L⁻¹ RS, 0.2 M H₂SO₄) has been reported [16]. However, drawback of dilute acid (higher concentration > 0.9 N) pretreatment is the generation of a higher concentration of 5-hydroxymethyl furfural (HMF) and levulinic acid (LA) (with the degradation of hexose sugars and furfurals from pentose sugar degradation), which acts as inhibitors for microorganisms during the fermentation process by reducing enzymatic and biological activities, breaking down the DNA and inhibiting protein and RNA synthesis [51]. In order to overcome this, enzyme saccharification or biological pretreatment using either cellulase enzyme (of commercial-grade) or enzymes isolated from fungi or bacteria has been tried.

The most common enzymes employed for seaweed hydrolysis in earlier studies are commercial enzymes such as Cellulase, Celluclast 1.5 L, Viscozyme L, Novozyme

Table 2 Various parameters involved in pretreatment process

Pretreatment process	Temperature, °C	Pressure, atm absolute	Reactions times, min
Alkaline hydrolysis	208		2 h
Acid hydrolysis	121	3–15	45
Ionic liquids (ILs)	80–115		60
Organosolvent process	50–200		1–2 h
Steam explosion	160–260	6.8–47.6	24 h
Ammonia fiber explosion (AFEX)	60–100	15–20	<5
Liquid hot water	170–230	49.34	1 h

[41–45]

188, Termamyl 120 L, β -glucosidase, Multifect, Meicelase, and Amyloglucosidase operated at pH 4.5–5.5 and temperature 35–55 °C; incubation time varies based on the algal feedstock [8, 15, 51–59]. The current study focuses on the evaluation of sugar release from *Enteromorpha intestinalis* and *Ulva lactuca* using dilute acid hydrolysis and enzyme (extracted from *Vibrio parahaemolyticus*) saccharification of the dilute acid pretreated biomass.

2 Materials and methods

2.1 Sample collection

E. intestinalis and *U. lactuca* seaweed samples were collected from Aghanashini estuary, Kumta taluk, Uttara Kananda district, Karnataka. Samples were thoroughly cleaned of epiphytes, then shade dried and powdered to obtain uniform size, and stored in polythene sachets for further analysis.

2.2 Dilute acid hydrolysis

Pretreatment of macroalgal feedstock was carried out at different acid concentrations, substrate concentrations, reaction time, and temperatures. Initially, acid hydrolysis was carried out using H_2SO_4 and HCl with concentrations ranging from 0.05, 0.1, 0.3, 0.5, 0.7, 0.9 to 1 N and keeping other parameters constant. The concentration of acid required for optimal sugar production was assessed, and further optimization was carried out. The reaction temperature for pretreatment was carried out with 2% w/v substrate at 30, 60, 90, and 120 °C for 45 min. Reaction time pretreatment with 2% w/v substrate concentration was carried out for 30, 60, 90, 105, and 120 min. Pretreatment for algal biomass at different substrate concentrations of 1%, 2%, 3%, 5%, 7%, and 9% w/v was carried out at 120 °C for 45 min. After hydrolysis, residues were separated by filtration, and total sugar and reducing sugar were determined by phenol sulfuric acid [60] and dinitrosalicylic acid (DNS) [61] methods, respectively. DNS method is widely used to determine reducing sugar content in fields of food, bioprocess, medicine, etc. [62, 63]. Neutralization was carried out for acid hydrolysate using Na_2CO_3 , NaOH, activated charcoal, and $Ca(OH)_2$ [64, 65]. The significance of the above factors in influencing sugar release was determined using ANOVA. Thin-layer chromatography analysis of algal hydrolysate is obtained following optimized acid hydrolysis on silica gel plates using mobile phase butanol/ethanol/water (3:2:1 v/v/v). Later, these plates were dried at room temperature and dipped in $AgNO_3$ solution for 1 min, and when dried, the plates were sprayed with ethanolic sodium hydroxide solution until dark brown spots appeared [66].

The efficiency of acid hydrolysis pretreatment (%) is calculated using Equation 1:

$$E_p(\%) = \frac{(\Delta S)}{TS} \times 100 \quad (1)$$

where:

- E_p is the efficiency of acid hydrolysis pretreatment (%).
- ΔS is the disaccharide increase (mg) during acid hydrolysis pretreatment.
- TS is the total sugar (mg).

2.3 Response surface method

Response surface method (RSM) was used to evaluate the relationship between independent variables, reaction temperature (°C, X_1), reaction time (min, X_2), and substrate concentration (% w/v, X_3), and dependent variable, reducing sugar ($mg\ g^{-1}$, Y). The experimental data was analyzed and the probable relationship follows the second-order polynomial Eq. 2:

$$Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_3 X_3 + \beta_{11} X_1^2 + \beta_{22} X_2^2 + \beta_{33} X_3^2 \quad (2)$$

where Y is a response variable and X_1 , X_2 , and X_3 are independent variables, β_0 is the offset term; β_1 , β_2 , and β_3 are the linear coefficients as per least squares method; and β_{11} , β_{22} , and β_{33} are the first, second, and third linear coefficients, respectively [48]. The Student's t test was performed for the determination of the statistical significance of the regression coefficient [55, 67].

2.4 Crude enzyme production from *Vibrio parahaemolyticus*

Crude enzyme is extracted from *Vibrio parahaemolyticus* [68] using CMC as a sole source of carbon for screening cellulose-degrading bacteria. Endoglucanase was determined using the carboxymethyl cellulase method (CMCase) [69]; the endoglucanase enzyme cleaves the intermolecular β -1–4-glycosidic bonds present in cellulose. Cellulase was purified by centrifuging bacterial culture at 12,000 rpm for 15 min at 4 °C and supernatant was collected. Proteins were precipitated to 80% saturation with $(NH_4)_2SO_4$ at 4 °C and pelletized through centrifugation. Pellet was dissolved in Tris–HCl (pH 7) and purified using ion exchange chromatography, wherein the sample was applied to Superdex 200 column equilibrated with Tris–HCl. Fractions were collected, and fractions with the highest enzyme activity were pooled and considered for other characterization. Enzymatic activity refers to the amount of enzyme that releases 1 μ mol of reducing sugar per minute.

2.5 Enzyme characterization

The effect of pH on the enzyme activities was estimated using a buffer of different concentrations: 50 mM citrate buffer (pH 3–4), citrate phosphate buffer (pH 5–6), Tris–HCl (pH 7), and potassium phosphate buffer (pH 8). The effect of temperature on enzyme activity was determined by incubating the enzyme assay mixture of optimum pH at different temperatures ranging from 25 to 60 °C. The effect of salinity on enzyme assay mixture and salinity (NaCl) ranging from 4 to 20% was determined. Samples were incubated for 1 h with CMC as a sole source of carbon.

2.6 Efficient sugar release from macroalgal biomass through pretreatment: dilute acid hydrolysis and enzyme saccharification

Macroalgal feedstock *Enteromorpha intestinalis* (EI) and *Ulva lactuca* (UL) samples were subjected to dilute acid hydrolysis using an optimized acid concentration of 0.7 N and 0.5 N H₂SO₄ at optimal temperature and time of 121 °C and 45 min. Dilute acid pretreated macroalgal biomass EI and UL was subjected to enzymatic hydrolysis at 55 °C pH 6 for 36 h. The reducing sugar released from the above processes was recovered through a centrifuge and was estimated using DNS method [61].

2.7 SEM analysis

Macroalgal biomass surface morphology (untreated, acid-treated, and enzyme-treated biomass) was qualitatively analyzed using SEM (JEOL-IT 300). Macroalgal samples were placed on an aluminum specimen mount using conductive carbon tape. Sputter gold coating was performed to prevent charging. Samples were then examined in SEM under vacuum condition at accelerating voltage of 10 kV.

3 Results and discussion

Total sugar and reducing sugar (after dilute acid hydrolysis) were estimated for both *E. intestinalis* and *U. lactuca*. The optimization variables considered are acid concentration,

reaction temperature, substrate concentration, and reaction time.

3.1 Effect of dilute acid concentration

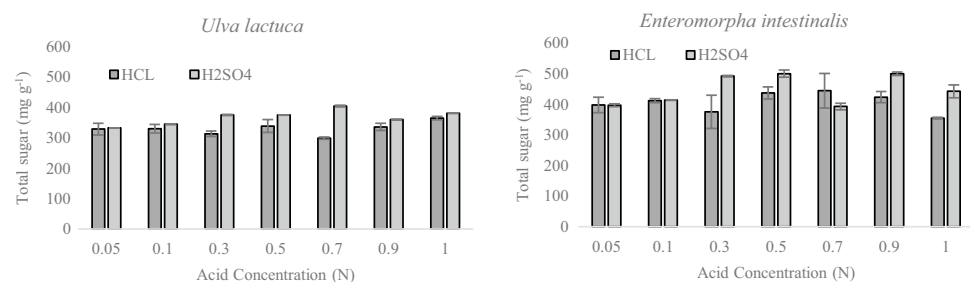
The effect of dilute acid concentration on *E. intestinalis* and *U. lactuca* was determined using acid concentrations of 0.01, 0.05, 0.3, 0.5, 0.7, and 1 N, for 1% substrate concentration at 121 °C for 1 h. Higher total sugar for *E. intestinalis* and *U. lactuca* biomass was obtained for pretreatment using H₂SO₄ (Fig. 1). The sugar content gradually decreased with increase in acid concentrations as sugar decomposition varies based on concentrations and different acid catalytic activities [70]. Acid-catalyzed glucose decomposition is more dependent on the concentration of hydrogen ions at a particular temperature than on hydrogen ion sources [71].

Reducing sugar estimation was carried out for *E. intestinalis* and *U. lactuca* using HCl and H₂SO₄ (Fig. 2). Reducing sugar increased gradually with the increase in H₂SO₄ concentration for *E. intestinalis*, whereas for *U. lactuca*, reducing sugar increased up to 0.5 N H₂SO₄ and then decreased drastically. Acid hydrolysis efficiency was calculated for both the acid, and it was found that 0.7 N H₂SO₄ with the conversion efficiency of 80.18% was suitable for *E. intestinalis* and 0.5 N H₂SO₄ achieved a conversion efficiency of 60.07% for *U. lactuca*. These concentrations were kept constant for further optimization study. H₂SO₄ hydrolysis exhibited better reducing sugars compared to HCl and was considered for optimization. Hydrolysis using different acid concentrations released different concentrations of reducing sugars from the macroalgal biomass depending on their structure and biochemical composition, demonstrating that a customized approach is needed for hydrolysis.

3.2 Effect of reaction temperature on acid hydrolysis

Reducing sugar at different temperatures of 30, 60, 90, and 120 °C was recorded (Fig. 3). The highest reducing sugar of 549.45 mg g⁻¹ and 528.46 mg g⁻¹ was obtained for *E. intestinalis* and *U. lactuca* at 120 °C with 90.9% and 97.7% sugar conversion, respectively. Pretreatment of terrestrial biomass involves higher temperature (165–210 °C); this is

Fig. 1 Total sugar released using different concentrations of acid for *U. lactuca* and *E. intestinalis* ($p < 0.05$)



attributed to their rigid structures [59], whereas macroalgal biomass requires milder temperatures. Studies involving red algae *Kappaphycus alvarezii* pretreated using 1% H₂SO₄ at 120 °C for 60 min obtained 81 g L⁻¹ of reducing sugar [15], whereas earlier study [16] of similar pretreatment conditions at 130 °C obtained 22.4 g L⁻¹ of reducing sugar. Reducing sugar of 65 mg g⁻¹ was obtained after pretreatment of *U. pinnatifida* at 120 °C for 24 h [17]. Pretreatment of *Saccharina japonica* using 40 mM H₂SO₄ at 121 °C for 60 min yielded 20.6 g L⁻¹ of reducing sugar [57]. An earlier study [59] of similar pretreatment conditions using 1 mM H₂SO₄ for 120 min achieved 34 g L⁻¹ reducing sugar, indicating that the concentration of acid in hydrolysis plays a critical role in incubation temperature. Higher pretreatment conditions lead to degradation of sugars to hydroxymethyl furfural, which inhibits yeast growth by reducing the biological enzymatic activities, causing DNA and cell wall damage, inhibition of RNA, and protein synthesis [72].

3.3 Effect of reaction time on hydrolysis

To investigate the release of total sugar and reducing sugar, *E. intestinalis* and *U. lactuca* were hydrolyzed by 0.7 N and 0.5 N H₂SO₄, respectively, at 121 °C and different reaction times varying from 15, 30, 60, 90 to 120 min (Fig. 4). Maximum total sugar of 399 mg g⁻¹ was obtained at 105 min for *E. intestinalis*; maximum reducing sugar (121 mg g⁻¹) was recorded at 45 min with a conversion efficiency of 42.1%. Maximum total and reducing sugars were produced for *U. lactuca* at 45 min and were seen decreasing with the increase in incubation time. *G. verrucosa* was subjected to pretreatment using 0.1 N H₂SO₄, and maximum total sugar (12.06 g L⁻¹) and reducing sugar (6.99 g L⁻¹) were obtained at 15 min incubation time. A shorter incubation time is required for red algae as a major fraction of sugar (i.e., Floridean starch) is composed in the cytoplasm of the red algae [73], which gets released easily. Hence, the longer incubation time was not considered as it leads to increase

Fig. 2 Reducing sugar release using different concentrations of acid for *U. lactuca* and *E. intestinalis* ($p < 0.05$)

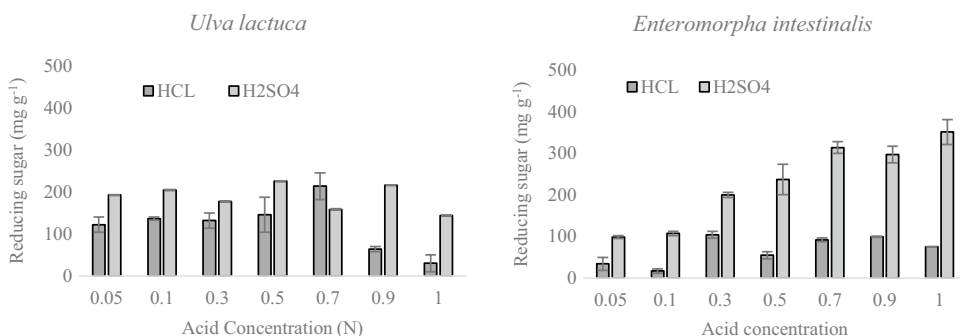


Fig. 3 Effect of reaction temperature on acid hydrolysis of *U. lactuca* and *E. intestinalis* ($p < 0.05$)

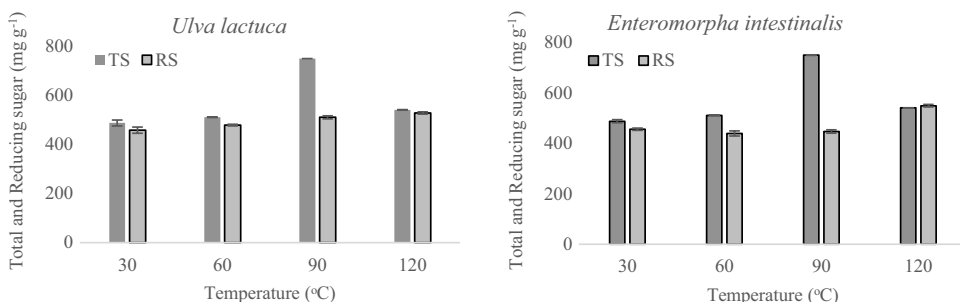
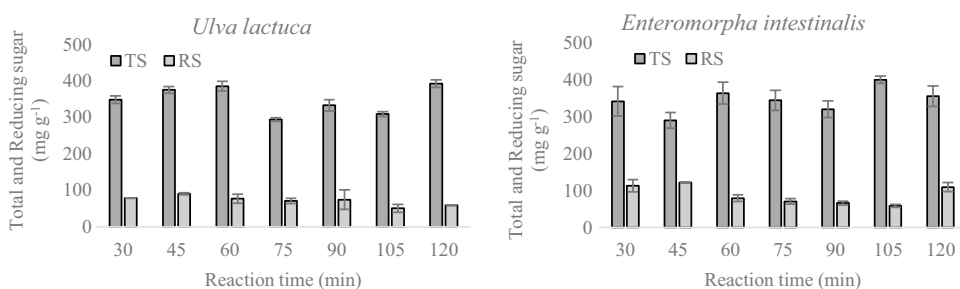


Fig. 4 Effect of incubation time on total and reducing sugar release during acid hydrolysis from *U. lactuca* and *E. intestinalis* ($p < 0.05$)



energy and cost, as well as accelerates the degradation of sugars to 5-HMF, levulinic acid, and formic acid [8, 10, 52, 65]. Therefore, 45 min was considered as the optimum incubation time for further studies.

3.4 Effect of substrate concentration on acid hydrolysis

The effect of varying substrate concentrations (1–9% w/v) on acid hydrolysis of *E. intestinalis* and *U. lactuca* was investigated at 121 °C for 45 min (Fig. 5). Total and reducing sugar concentration decreased with an increase in substrate concentrations. Similar results were observed for *E. intestinalis*, and the total reducing sugar decreased with an increase in solid to liquid ratio [8]. In a conventional simultaneous saccharification and fermentation process, substrate concentration of 10% (w/v) is considered optimal due to high viscosity and difficulty in handling the slurry [70]. Hydrolysis of *Kappaphycus alvarezii* required a 10% substrate concentration [16]. The highest sugar conversion rate was achieved at 2% (w/v) of *G. verrucosa* [50]. Higher efficiency of 85.43% and 62.97% was obtained for 5% (w/v) of *E. intestinalis* and *U. lactuca* substrate. Therefore, 5% (w/v) was considered as optimum substrate concentration.

3.5 Optimized sugar from *E. intestinalis* and *U. lactuca*

Acid hydrolysis of *E. intestinalis* and *U. lactuca* feedstock was carried out at an optimum acid concentration of 0.7 N and 0.5 N H₂SO₄, respectively at 121 °C for 45 min 5% (w/v) substrate concentration (Table 3). The highest reducing sugar of 206.82 ± 14.96 mg g⁻¹ was recorded from *U. fasciata* using sodium acetate (pH 4.8) buffer pretreatment process at 120 °C for 60 min. *Undaria pinnatifida* was

pretreated at a higher concentration of acid, 5% H₂SO₄ at 120 °C for 24 h to obtain 65 mg glucose/g biomass [17]. *U. pertusa* was subjected to a high thermal liquefaction process (HTLP), with a process condition of 400 °C at 40 mPa and obtained 352 mg g⁻¹ of reducing sugar. HTLP pretreatment loosens the complex structure and increases the porosity of the cell membranes allowing the entry of the solvent for further degradation [74]. Reducing sugar concentration of 145 ± 2.1 mg g⁻¹ was obtained from pretreatment of *Saccharina japonica* (10% w/v) at 121 °C for 60 min using 40 mM H₂SO₄ [57]. Pretreatment of red seaweed *Gracilaria* sp. was carried out using 0.1 N H₂SO₄ at 121 °C for 1 h at 20% w/v biomass loading and obtained 277 mg g⁻¹ of reducing sugar [75]. Inhibitors from acid hydrolysate hydroxymethyl furfural (HMF) and levulinic acid (LA) were detoxified using activated charcoal [65], which removed 70.37% HMF and 38.8% LA; similarly, Na₂CO₃ detoxified the 56.1% from *U. lactuca* and 23.3% from *E. intestinalis* [64] indicating that hydrolysis using dilute acid concentration resulted in a lower concentration of inhibitors.

TLC analysis showed glucose and xylose in the both acid hydrolysate of *E. intestinalis* and *U. lactuca* (Fig. 6).

3.6 Assessing the optimal pretreatment conditions through RSM (Response Surface Method)

RSM involved assessing the optimal pretreatment conditions (Table 4) for maximum reducing sugar yield from *E. intestinalis* and *U. lactuca*. The possible combinations of independent variables were chosen through stepwise regression, and the probable relationship with the yield of sugar (Y) is expressed in Eqs. 3 for *E. intestinalis* and 4 for *U. lactuca*, respectively ($p < 0.05$). Response surface curves were generated using 14 data points of each variable as depicted in Fig. 7 for *E. intestinalis* and *U. lactuca* reducing sugar

Fig. 5 Effect of substrate concentration of acid hydrolysis of *U. lactuca* and *E. intestinalis*

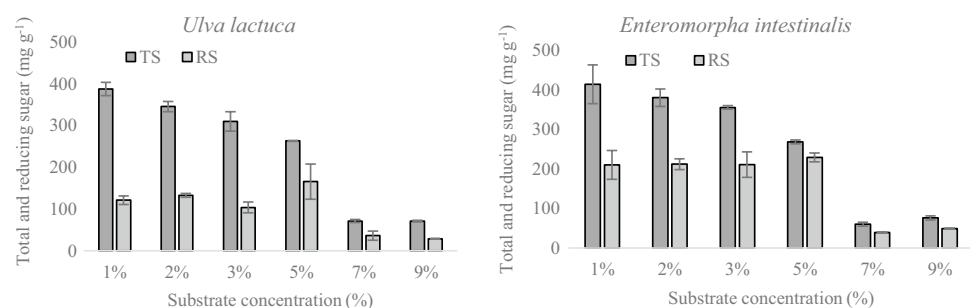
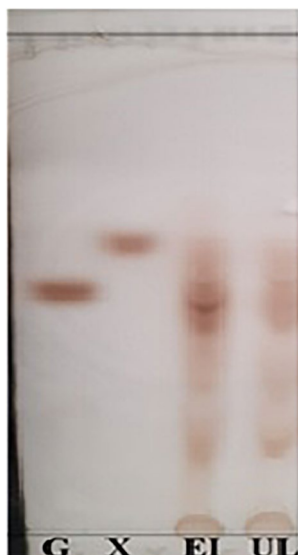


Table 3 Optimized sugar release using dilute acid pretreatment of *E. intestinalis* and *U. lactuca*

Biomass	Dilute acid pretreatment	Reducing sugar (mg g ⁻¹)
<i>Enteromorpha intestinalis</i>	5% w/w, 0.7 N H ₂ SO ₄ at 121 °C for 45 min	239.94 ± 1.36
<i>Ulva lactuca</i>	5% w/v, 0.5 N H ₂ SO ₄ at 121 °C for 45 min	214.67 ± 0.97

Fig. 6 TLC analysis of hydrolysate obtained after optimized acid hydrolysis of algal biomass; *E. intestinalis* (EI) and *U. lactuca* (UL) with glucose (G) and xylose (X) as standards



yield at different reaction temperatures, time, and substrate concentrations, which aided in arriving at the optimum level of each variable for maximum response. An increase in substrate concentrations led to a decline in reducing sugar release, which could be due to sorption loss.

$$Ye = 584.9 + 4.8X_1 - 4.7X_2 - 5.7X_3 - 0.05X_1^2 - 0.017X_2^2 - 1.45X_3^2 \tag{3}$$

$$Yu = 293.2 + 11.6X_1 - 3.2X_2 + 17.9X_3 - 0.09X_1^2 - 0.01X_2^2 - 3.8X_3^2 \tag{4}$$

The effect of reaction temperature and incubation time on hydrolysis of *E. intestinalis* and *U. lactuca* when substrate concentration was kept constant as shown in Fig. 7

and reducing sugar yield decreased with an increase in incubation temperature and incubation time. Higher reducing sugar yield was recorded at lower temperatures (30–60 °C) and incubation time (30–90 min). In order to obtain reducing sugar yield between 400 and 600 mg/g, the optimum reaction temperature of 75 °C, reaction time 75 min, and substrate concentration of 5% w/v were recorded from the RSM 3D plot. However, in this study, dilute acid hydrolysis of algal biomass for efficient reducing sugar yield between 200 and 240 mg/g was achieved at temperature 121 °C and time 45 min at 5% w/v substrate concentration. Higher glucose yield for *Sargassum* spp. is achieved at optimized acid concentration of 3.75 and 4.5% (w/v) substrate concentration and optimum temperature 115 °C for 86–90 min [55]. It is seen that the pretreatment temperature and incubation time obtained in this study to treat *E. intestinalis* and *U. lactuca* were milder than the terrestrial biomass. The presence of cellulose, hemicellulose, and lignin imparts the rigidity to the terrestrial biomass. It hence requires a temperature between 165 and 210 °C at a high concentration of acids for a longer incubation time (4 weeks) [55, 57].

It is seen that the pretreatment temperature and incubation time obtained to treat *E. intestinalis* and *U. lactuca* were milder than the terrestrial biomass requiring temperature between 165 and 210 °C at a high concentration of acids for longer incubation time (4 weeks). A harsh pretreatment condition was required due to the rigidity of the biomass with cellulose, hemicellulose, and lignin [55, 59]. Estimated effects, standard errors (SE), Student’s *t* test, and significance value for the model representing

Table 4 Experimental design displaying reducing sugar yield for individual runs of the response surface methodology design for *E. intestinalis* and *U. lactuca*

Design points	Reaction temperature (°C)	Reaction time (min)	Substrate concentration (%w/v)	Reducing sugar (mg g ⁻¹)	
				<i>E. intestinalis</i>	<i>U. lactuca</i>
	X ₁	X ₂	X ₃	Ye	Yu
1	30	45	2	477.5	473.3
2	90	45	2	447.1	513.3
3	120	45	2	549.4	628.1
4	120	30	2	124.1	78.8
5	120	105	2	58.5	50.7
6	120	120	2	109.1	133.5
7	60	45	2	439.6	480.0
8	120	60	2	79.1	77.3
9	120	45	5	103.1	166.1
10	120	45	7	157.7	116.2
11	120	45	1	210.8	121.6
12	120	45	3	238.1	104.2
13	120	45	9	49.28	29.02
14	120	75	2	70.53	70.9

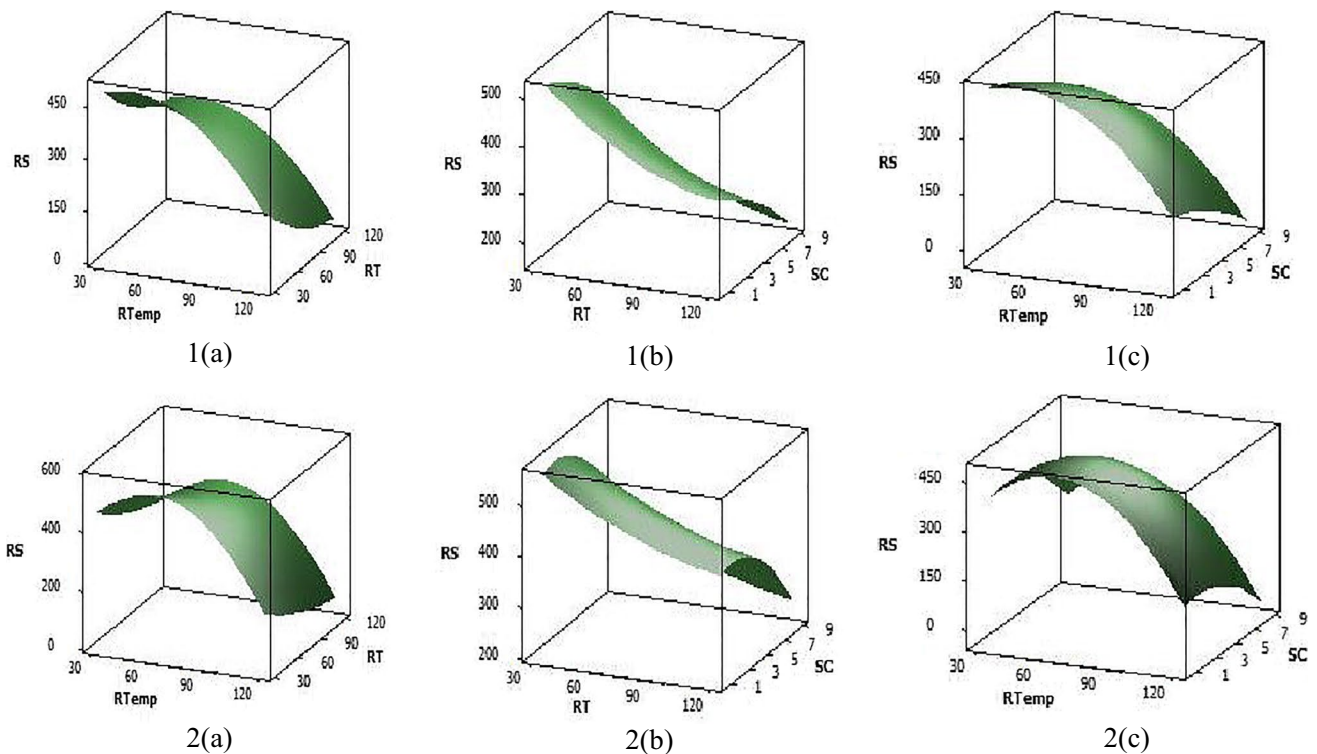


Fig. 7 Response surface plots of reducing sugar yield for *E. intestinalis* (1a–b) and *U. lactuca* (2a–b) after dilute acid pretreatment at different reaction temperature, time, and substrate concentrations. **1a**

and **2a** Reducing sugar yield at substrate concentration = 5% w/v; **1b** and **2b** reducing sugar yield at temperature = 75 °C; **1c** and **2c** reducing sugar yield at reaction time = 75 min ($p < 0.05$)

reducing sugar yield from *U. lactuca* and *E. intestinalis* are represented in Table 5.

3.7 Enzyme purification and characterization

The purification of cellulase enzyme is summarized in Table 6, which is a two-step purification and includes ammonium sulfate precipitation and size exclusion chromatography. The purified enzyme exhibited 7.24 U mg^{-1} of specific activity, and 61.82% yield was obtained with 2.97-fold purification. Purification was further confirmed by observing a single protein band on SDS-PAGE (Fig. 8) with an estimated molecular mass of 29 kDa. Similarly, molecular mass was obtained for cellulase extracted from *Salinivibrio* sp. NTU-05 exhibiting 32.4 U mg^{-1} specific activity and 18.9% recovery with 29.5-fold purification [76]. Extraction of Endo- β -1, 4-glucanase Cel5A from *Vibrio* sp. exhibited a molecular mass of 50 kDa, indicating functional cellulase gene in *Vibrio* genus (Gao et al., 2010).

The enzyme exhibited the highest activity at pH 6, and the activity profile showed that the enzyme was active over a wide range of pH, retaining 90% of its activity (Fig. 9). Similar characteristic pH tolerance over a wide range has been studied earlier for *Paenibacillus* sp. pH 7 [77]; *Marinobacter* sp. MS1032 [78]; *Vibrio* sp. G21 pH 6.5–7.5 [79]; *Bacillus*

sp. H1666 pH 3–9 [80]; *Bacillus* sp. [81]; *Stachybotrys atra* BP-A [82], *Bacillus flexus* pH 8–12 [83]; and *Salinivibrio* sp. pH 6.5–8.5 [76].

Table 5 Estimated effects, standard errors (SE), Student's t test, and significance value for the model representing reducing sugar yield from *U. lactuca* and *E. intestinalis*

<i>U. lactuca</i>	Coefficient	SE	t stat	p value
Intercept	293.26	200.95	4.01	0.002
X_1	11.61	1.79	-2.11	0.061*
X_2	-3.25	1.96	-0.98	0.34
X_3	17.92	1.25	-1.05	0.31
X_1^2	-0.09	0.02	-0.01	0.04
X_2^2	-0.01	0.001	-0.001	0.25
X_3^2	-3.8	1.2	-0.01	0.36
<i>E. intestinalis</i>	Coefficient	SE	t stat	p value
Intercept	584.94	155.11	5.01	0.0005
X_1	4.8	1.38	2.29	0.04*
X_2	-4.74	1.51	-1.6	0.13
X_3	-5.78	2.21	-1.4	0.18
X_1^2	-0.05	0.01	-0.001	0.03
X_2^2	-0.017	0.001	-0.001	0.21
X_3^2	-1.45	1.05	-0.13	0.30

* $p < 0.05$

Table 6 Purification steps of cellulase enzyme isolated from *Vibrio parahaemolyticus*

Purification steps	Protein (mg ml ⁻¹)	Total activity IU ml ⁻¹	Specific activity U mg ⁻¹	Purification fold	Yield %
Crude	38.75	94.44	2.44	1.00	100.00
80% ammonium precipitation	13.37	88.00	6.58	2.70	93.18
Superdex 200 column fractions	8.07	58.38	7.24	2.97	61.82

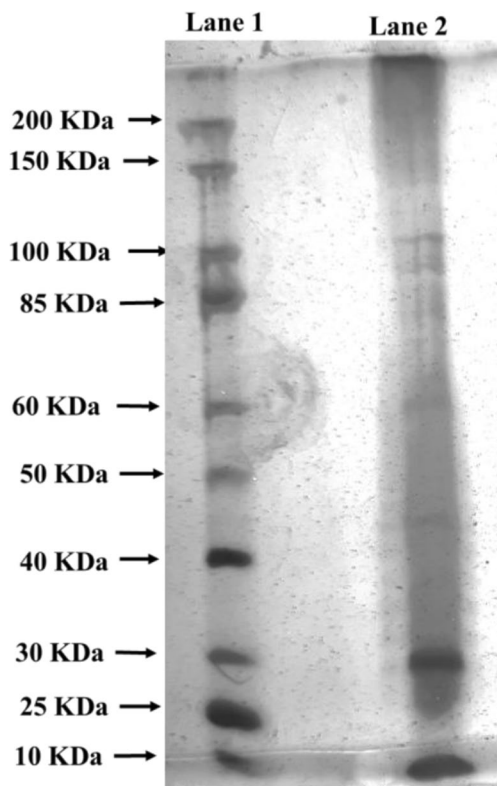


Fig. 8 SDS-PAGE of purified cellulase from *Vibrio parahaemolyticus*. Lane 1, protein ladder; lane 2, cellulase enzyme in 10% SDS-PAGE

The highest activity of the enzyme was recorded at 50 °C with higher stability between 40 and 55 °C (Fig. 10). Enzyme activity decreased due to the fluidity of protein conformation with an increase in temperature above 55 °C. It was seen that around 60% of the activity remained at 60 °C. The optimum temperature for cellulase-producing bacteria was 40–60 °C [76–78, 80–83].

Enzyme activity declined with an increase in salt concentration; at NaCl concentration > 10%, the enzyme retained 5% of its activity after 24 h (Fig. 11). Enzyme activity was above 20% up to 10% NaCl concentration. It is seen that NaCl concentration induces the activity of endo-β-1, 4-glucanase Cel5A from *Vibrio* sp. G21 and EgI-AG from alkaliophilic *Bacillus agaradhaerens* [79, 84]. Enzyme displayed activity in a broad range of 0–10% NaCl concentration with optimum NaCl concentration of

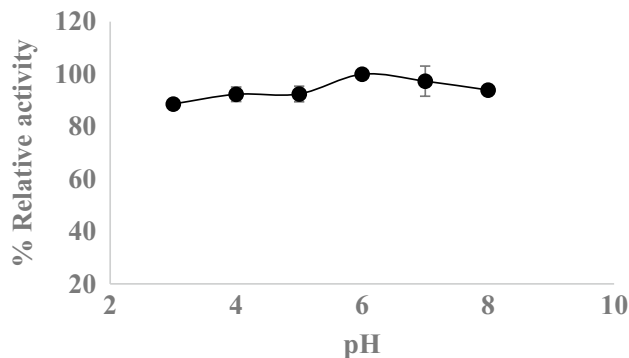


Fig. 9 Effect of different pH on enzyme activity

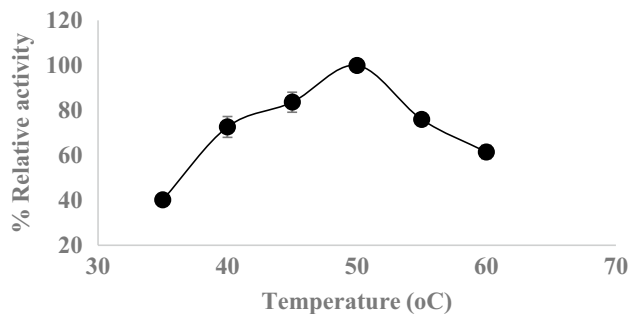


Fig. 10 Effect of different temperatures on enzyme activity

3%. Enzyme activity was stable only up to 10% NaCl concentration despite the enzyme being extracted from marine bacteria *V. parahaemolyticus*.

3.8 Dilute acid pretreatment and enzymatic hydrolysis

Acid hydrolysis of *E. intestinalis* and *U. lactuca* biomass was carried out at an optimum acid concentration of 0.7 N and 0.5 N H₂SO₄, respectively, at a temperature of 121 °C for 45-min incubation time and 5% (w/v) substrate concentration (Table 7).

In similar studies, the highest reducing sugar of 206.82 ± 14.96 mg g⁻¹ was recorded from *U. fasciata* using sodium acetate (pH 4.8) buffer pretreatment at 120 °C for 60 min. *Undaria pinnatifida* was pretreated at a higher concentration of acid, 5% H₂SO₄ at 120 °C for 24 h to obtain

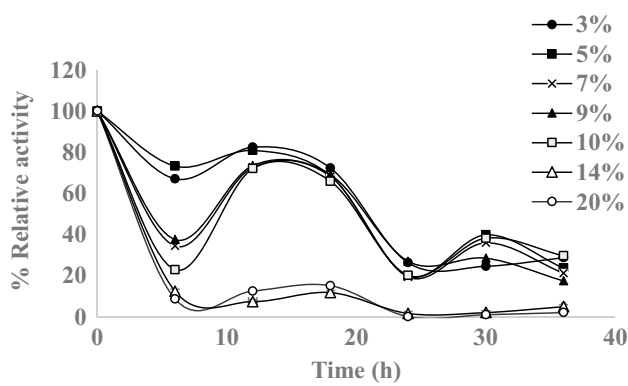


Fig. 11 Effect of NaCl concentration on enzyme activity

65 mg glucose g^{-1} biomass [17]. *U. pertusa* was subjected to a high thermal liquefaction process (HTLP), with a process condition of 400 °C at 40 mPa, and obtained 352 mg g^{-1} of reducing sugar. HTLP pretreatment loosens the complex structure and increases the porosity of the cell membranes allowing the entry of the solvent for further degradation [74]. Reducing sugar concentration of 145 ± 2.1 mg g^{-1} was obtained from pretreatment of *Saccharina japonica* (10% w/v) at 121 °C for 60 min using 40 mM H_2SO_4 [57]. Pretreatment of red seaweed *Gracilaria* sp. using 0.1 N H_2SO_4 at 121 °C for 1 h at 20% w/v biomass loading yielded 277 mg g^{-1} of reducing sugar [75].

Enzyme hydrolysis is affected by various factors, such as temperature, pH, and concentration (enzyme/substrate). Increasing enzyme concentration will speed up the reaction, as long as there is substrate availability; however, if once all of the substrates are bound, the reaction will cease to speed up. On the other hand, increasing substrate concentration also increases the rate of reaction to a certain extent. But once all enzymes are bound, any increase in substrate will have no effect on the reaction rate due to saturation of available enzymes. Trivedi et al. (2015) isolated cellulase enzyme from *Cladosporium sphaerospermum* and subjected *Ulva lactuca*, green seaweed to enzymatic hydrolysis and obtained 112 mg/g of reducing sugar. However, in this study,

the highest reducing sugar of 107.6 mg/g was obtained from *U. lactuca*, whereas 135.9 mg/g reduced sugar from *E. intestinalis* indicating enzyme ability to hydrolyze the macroalgal polysaccharide.

Acid pretreated macroalgal biomass (*E. intestinalis* and *U. lactuca*) was subjected to enzymatic hydrolysis using purified enzyme and incubated for 24 h and observed two-fold increase in reducing sugar in both biomass and 1.2-fold increase from dilute acid pretreatment, compared to crude enzymatic hydrolysis. Enzymatic hydrolysis of *U. lactuca* using purified enzyme extracted from *Bacillus* sp. H1666 yielded 450 mg g^{-1} of reducing sugar, indicating the potential applicability of the enzyme for algal biomass saccharification [80]. Enzymes secreted from the cell are generally found along with other proteins, lipids, polysaccharides, and nucleic acids. The measurement of enzyme purity is defined as the relation of the activity of the enzyme to the total protein present (i.e., the specific activity). Enzyme purification is carried out in order to remove the contaminants and increase the specific activity [85]. In this study, purified enzyme yielded higher reducing sugar due to increased specific activity (Table 6).

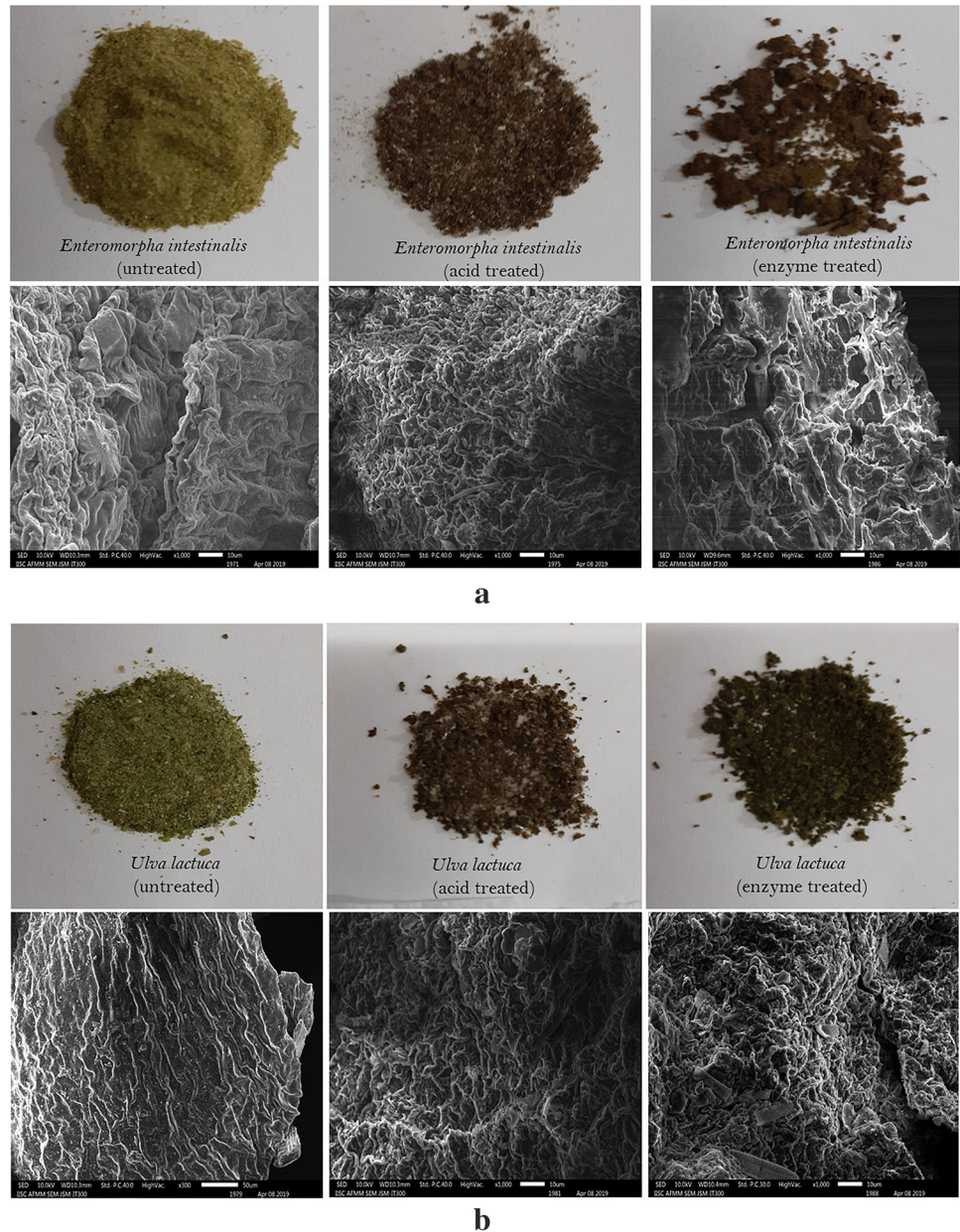
3.9 Scanning electron microscopy

Scanning electron microscopic (SEM) analysis of macroalgal biomass revealed ultrastructural changes in the biomass during dilute acid pretreatment. Figures 12a and 12b depict the untreated surface of *E. intestinalis* and *U. lactuca*; raw or untreated biomass had continuous, even, and smooth surfaces, whereas biomass after dilute acid pretreatment had loosened the rugged surface, which increased the surface area, exposing more internal cellulose for enzymatic hydrolysis. The roughness of the seaweed surface after dilute acid hydrolysis makes it more liable for enzymatic hydrolysis. The presence of strong hydrogen bonding of cellulose and Van der Waal forces of glucose molecules imparts the crystalline structure to biomass [52, 86, 87]. Scanning electron microscope (SEM) images revealed cracks and holes on the pretreated algal

Table 7 Optimized sugar release using dilute acid pretreatment of *E. intestinalis* and *U. lactuca*

Biomass	Dilute acid pretreatment	Reducing sugar (mg)	Acid Pretreated biomass	Enzymatic hydrolysis	Reducing sugar (mg)	
					Crude enzyme	Purified enzyme
<i>E. intestinalis</i> (500 mg)	5% w/w, 0.7 N H_2SO_4 at 121 °C for 45 min	239.94 ± 1.3	<i>E. intestinalis</i> (320 mg)	pH 6 at 50 °C for 24 h	135.93 ± 11.48	289.89 ± 2.4
<i>U. lactuca</i> (500 mg)	5% w/v, 0.5 N H_2SO_4 at 121 °C for 45 min	214.67 ± 0.9	<i>U. lactuca</i> (320 mg)	pH 6 at 50 °C for 24 h	107.68 ± 9.55	261.76 ± 0.9

Fig. 12 **a** Scanning electron micrograph of *E. intestinalis* depicting ultrastructural changes in the feedstock — untreated sample compared with the acid and enzyme treated. **b** Ultrastructural changes evident in the scanning electron micrograph of *U. lactuca* macroalgal biomass — untreated, acid, and enzyme treated



surface. *Gelidium amansii* treated at 121 °C were observed under SEM; electron micrographs revealed fibers exposed in autoclaved samples allowing enzymes to easily degrade the cells [52].

4 Conclusions

The present study demonstrates the potential of *E. intestinalis* and *U. lactuca* as marine source for production of sugar for biofuel production. Feedstock *E. intestinalis* and *U. lactuca* were subjected to dilute acid pretreatment and

yielded $239.94 \pm 1.3 \text{ mg g}^{-1}$ and $214.67 \pm 0.9 \text{ mg g}^{-1}$ of reducing sugar. An enzyme extracted from marine bacteria *Vibrio parahaemolyticus* hydrolyzed the algal biomass efficiently, releasing onefold higher reducing sugar than dilute acid pretreatment. Enzymatic hydrolysis of pretreated macroalgal biomass produced onefold higher reducing sugar than the dilute acid pretreatment. Dilute acid pretreatment prior to enzymatic hydrolysis improves algal biomass saccharification and releases higher reducing sugar, increasing bioethanol yield. However, tailor-made approaches need to be employed depending on algal species, as some sugars are easily hydrolyzed by acid and few by a direct enzyme.

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