Evaluation of Cellulolytic Endo-1,4-β-D-Glucanase Activity in the Digestive Fluid of Adult Phytophagous Beetle *Hoplasoma unicolor*

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DOI: https://doi.org/10.21315/tlsr2021.32.3.4

Highlights

- Presence of cellulolytic endoglucanase activity was detected in digestive fluid of the beetle *Hoplasoma unicolor*.

- Quantitative estimation showed a relatively high cellulase activity with specific activity measured as 0.69 (±0.01) units per mg of protein.

- Zymogram analysis revealed the presence of two active cellulolytic proteins in the gut extract.
Evaluation of Cellulolytic Endo-1,4-β-D-Glucanase Activity in the Digestive Fluid of Adult Phytophagous Beetle *Hoplasoma unicolor*

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Publication date: 30 September 2021

To cite this article: Mohammad Mosleh Uddin, Suzana Afrin Lima, Tanim Jabid Hossain, Newton Kar, Yeasmin Zahan and Babamale AbdulKareem Olarewaju. (2021). Evaluation of cellulolytic endo-1,4-β-D-glucanase activity in the digestive fluid of adult phytophagous beetle *Hoplasoma unicolor*. Tropical Life Sciences Research 32(3): 53–68. https://doi.org/10.21315/tlsr2021.32.3.4

Abstract: Insects of the taxonomic order Coleoptera are recognised for considerable cellulolytic activity in their digestive fluid. The cellulolytic activity of the gut fluid in *Hoplasoma unicolor*, a member of Coleoptera, however, remains unexplored. In this study, we, for the first time, report the qualitative and quantitative analysis of cellulolytic activity in the digestive fluid of this insect. The cellulolytic endo-1,4-β-D-glucanase activity was confirmed in the supernatant of the insect’s digestive fluid by agar plate assay using carboxymethyl cellulose as the substrate. To determine the optimum pH, enzyme activity was further assessed in an acidic pH range of 5 to 6, and the highest activity was observed at pH 5.3. For quantitative analysis, endoglucanase activity was measured using 3,5-dinitrosalicylic acid method which revealed that the specific activity of the gut sample was 0.69 (±0.01) units per mg of protein. For further characterisation of the cellulases in the sample, SDS-PAGE and zymogram analysis were carried out. Two active cellulolytic bands were detected on the zymogram suggesting the presence of two distinct endoglucanases which completely disappeared upon heating the sample at 55°C. Our study, therefore, highlights prospect of the gut fluid of *H. unicolor* as an important source of cellulase enzymes that merits further investigations into their extensive characterisation for potential industrial applications.

Keywords: Cellulolytic Activity, Endo-1,4-β-D-Glucanase, Beetle, Insect Gut Fluid, Insect Cellulase, Carboxymethyl Cellulose, Zymography

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INTRODUCTION

Cellulases are hydrolytic enzymes that catalyse the cleavage of 1,4-β-glycosidic bonds between glucose residues present in cellulose, the most abundant biopolymer produced on earth and the major constituent of agricultural and industrial wastes (Bayer et al. 1998; Dashtban et al. 2010). Consequently, substantial efforts are being made during the past few decades into prospecting for novel cellulolytic enzymes and elucidation of their catalytic properties so that the enzymes could be employed in many important bioprocesses, most particularly in: (1) the production of cost-effective and sustainable biofuels from the cellulosic and lignocellulosic biomass, (2) efficient and ecofriendly management of waste disposal and (3) industrial processes specially the textile, paper, food and detergent industries (Bayer et al. 2007; Jayasekara & Ratnayake 2019; Phitsuwan et al. 2013). The cellulase enzymes refer to three distinct types of cellulolytic hydrolases, e.g., endo-1,4-β-D-glucanase (endoglucanase; EC 3.2.1.4), exo-1,4-β-D-glucanase (cellobiohydrolase; EC 3.2.1.91 and 3.2.1.176), and β-glucosidase (EC 3.2.1.21) (Okano et al. 2014). The endoglucanase catalyses hydrolysis of soluble and insoluble β-(1,4)-glucan substrates, either directly on the polymer and/or shorter (poly)-oligosaccharides, both internally as well as from reducing and nonreducing ends in a nonprocessive or processive manner to produce oligosaccharides of various lengths (Girfoglio et al. 2012; Hobdey et al. 2015). Exoglucanases progressively hydrolyse cellulose at the reducing and non-reducing ends to release cellobiose moieties. Whereas β-glucosidases catalyse the final step: hydrolysis of the products generated by both endo- and exo-glucanases, i.e., soluble cellulodextrins and cellobioses to produce glucose. Although complete degradation of cellulose into glucose requires synergistic action among the three hydrolases, endoglucanase is believed to be the most important of the three (Annamalai et al. 2016). These cellulolytic hydrolases are commonly referred to as carboxymethylcellulase or CMCase, since carboxymethylcellulose (CMC) is the substrate most widely used for the determination of functional cellulase activity in experimental procedures (Ali et al. 1995).

The endoglucanase enzymes characterised so far are mostly those of microbial and plant origin. This enzyme activity has long been believed to be limited to only bacteria, fungi and plants. While activity was also detected in the digestive fluid of lower animals such as the insects, it was totally attributed to the microbial symbionts in the insect gut until 1998 when Watanabe et al. first described the identification of an endo-1,4-β-D-glucanase gene in the termite Reticulitermes speratu. Today, insects from at least 20 families such as Acrididae, Buprestidae, Amphisbatidae, Tenthredinidae, etc. are known to produce their own cellulolytic enzymes (Su et al. 2013). Insects, therefore, are regarded as very promising candidates to search for novel cellulases. The highly adapted phytophagous insects appear to be the most important species in this regard since they feed on very fibrous and lignocellulose-rich plant tissues (Oppert et al. 2010).
In the present study we sought to evaluate cellulolytic activity in gut fluid of the beetle *Hoplasoma unicolor*, a phytophagous insect of the taxonomic order Coleoptera. Previously, a number of beetles have been shown to possess considerable amount of cellulolytic activity in their gut or head fluid (Table 1). Su *et al.* (2013), for example, investigated cellulolytic activities in the gut fluids of 54 insect species from seven orders and detected highest CMCase activities in the insects of Coleoptera and Orthoptera. Other groups also reported high endoglucanase activity in the gut fluids of several other beetle species (Table 1). Beetles, therefore, are considered to be attractive candidates to prospect for novel cellulolytic enzymes with remarkable catalytic potential.

We herein report the determination of endo-1,4-β-D-glucanase activity in the digestive fluid of adult *H. unicolor*, a beetle that feeds on foliage of herbs (Mathew *et al.* 2005). This beetle was found abundantly feeding on its host plant, *Clerodendrum villosum*, a tomentose shrub (Hazmi *et al.* 2019). To our knowledge, cellulolytic activity in the gut fluid of this beetle was not reported previously. Therefore, qualitative and quantitative measurement of the enzyme activity was carried out on CMC using the partially purified gut fluid prepared from this insect. SDS-PAGE and zymogram analyses were also performed to further understand the multiplicity of the active cellulolytic proteins present in the enzyme extract.

**MATERIALS AND METHODS**

**Insect Collection**

This study investigated cellulolytic activity in the gut fluid of the beetle *H. unicolor*. Several beetles are already known to possess notable amount of cellulase enzymes in their gut fluid. Being a phytophagous beetle, *H. unicolor* is also speculated to produce considerable amount of cellulase enzymes in its gut which still remains unstudied. Hence in the present study, twenty adult beetles, ~1.2 cm in length, were collected from the host plant *C. villosum* near the campus of Mawlana Bhashani Science and Technology University, Bangladesh in the month of September 2019. The insects were actively feeding on or in close proximity to the host plant at the time of collection (Fig. 1a). Insects were kept on the leaves during transportation to the laboratory and placed in ice for 10 min before dissection.

**Dissection and Preparation of Digestive Extract for Enzyme Assays**

Dissections of the insects were performed on ice. Insect guts were removed and collected in microcentrifuge tube emerged in ice. The dissected guts were combined, cut into small pieces, homogenised by vortexing and centrifuged at 10,000 rpm for 10 min at 4°C. The 130 µL of supernatant was collected and transferred to a fresh microcentrifuge tube and stored at −20°C until it is ready to use. This gut sample has been designated hereafter as gut extract or enzyme extract.
| Family          | Name of species     | Life stages | Body segment | Substrate | Specific activity (U/mg) | References          |
|-----------------|---------------------|-------------|--------------|-----------|-------------------------|---------------------|
| Cerambycidae    | *Morimus funereus*  | Larva       | G            | CMC       | 1.20                    | Dojnov et al. 2013  |
|                 |                     | Larva<sub>1</sub> | G            | CMC       | 2.40                    |                     |
|                 |                     | Larva<sub>2</sub> | G            | CMC       | 2.80                    |                     |
|                 |                     | Larva<sub>3</sub> | G            | CMC       | 2.80                    |                     |
|                 | *Anoplophora glabripennis* | Adult<sup>d</sup><sub>4,m</sub> | G            | CMC       | 15.49<sup>u</sup>       | (Li et al. 2008)   |
|                 |                     | Adult<sup>d</sup><sub>4,f</sub> | G            | CMC       | 9.61<sup>u</sup>        |                     |
|                 |                     | Adult<sup>d</sup><sub>5,m</sub> | G            | CMC       | 13.06<sup>u</sup>       |                     |
|                 |                     | Adult<sup>d</sup><sub>5,f</sub> | G            | CMC       | 10.34<sup>u</sup>       |                     |
|                 |                     | Adult<sup>d</sup><sub>6,m</sub> | G            | CMC       | 13.54<sup>u</sup>       |                     |
|                 |                     | Adult<sup>d</sup><sub>6,f</sub> | G            | CMC       | 8.02<sup>u</sup>        |                     |
|                 |                     | Adult<sup>d</sup><sub>7,m</sub> | G            | CMC       | 17.12<sup>u</sup>       |                     |
|                 |                     | Adult<sup>d</sup><sub>7,f</sub> | G            | CMC       | 10.76<sup>u</sup>       |                     |
|                 | *Cerambycid sp.*    | Adult       | G            | CMC       | 0.29                    | (Shi et al. 2011)  |
|                 |                     | Larva       | G            | CMC       | 0.17                    | (Oppert et al. 2010) |
|                 |                     |              |              | MCC       | 0.03                    |                     |
|                 | *Elaphidion mucronatum* | Larva       | G            | CMC       | 0.42                    |                     |
|                 |                     |              |              | MCC       | 0.025                   |                     |
|                 | *(Say)*              | Larva       | G            | CMC       | 0.35                    |                     |
|                 |                     |              |              | MCC       | 0.08                    |                     |
|                 | *Neoclytus a. acuminatus* | Larva       | G            | CMC       | 0.42                    |                     |
|                 | *(Fabricius)*        |              |              | MCC       | 0.025                   |                     |

(Continued on next page)
| Family          | Name of species                      | Life stages | Body segment | Substrate* | Specific activity (U/mg) | References                  |
|-----------------|--------------------------------------|-------------|--------------|------------|--------------------------|-----------------------------|
| Tenebrionidae   | *Megopis sinica*                      | Larva       | G            | CMC, FP    | 0.53, 0.21               | (Su et al. 2013)            |
|                 | *Trichoferus campestris*              | Larva       | G            | CMC, FP    | 0.22, 0.03               |                             |
|                 | *Batocera horsfieldi*                 | Larva       | G            | CMC, FP    | 0.21, 0.008              |                             |
|                 | *Massicus raddei*                     | Larva       | G            | CMC, FP    | 0.27, 0.01               |                             |
|                 | *Anoplophora glabripennis*            | Larva       | G            | CMC        | 0.41                     | (Geib et al. 2010)          |
| Tenebrio molitor| *Tribolium castaneum*                | Adult       | G            | CMC        | 0.02                     | (Oppert et al. 2010)        |
|                 |                                      | H           | CMC          | 0.02       |                          |                             |
|                 |                                      |             | MCC          | 0.01       |                          |                             |
| Tenebrio molitor| *Tenebrio molitor*                   | Larva       | G            | CMC        | 0.02                     |                             |
|                 |                                      | H           | CMC          | 0.03       |                          |                             |
|                 |                                      | G           | MCC          | 0.05       |                          |                             |
|                 |                                      | H           | MCC          | 0.01       |                          |                             |
| Tribolium castaneum | *Tribolium castaneum*           | Adult       | Whole insect | CMC        | –                        | (Rehman et al. 2009)        |
| Tribolium castaneum | *Tribolium castaneum*               | Larva       | –            | CMC        | 12.9c                    | (Willis et al. 2011)        |
| Tenebrio obscurus | *Tenebrio obscurus*                  | Larva       | G            | CMC, FP    | 0.07, 0.08               | (Su et al. 2013)            |
| Zophobas morio  | *Zophobas morio*                     | Larva       | G            | CMC        | 0.222                    | (Szentner et al. 2019)      |
|                 |                                      | Adult       | G            | CMC        | 0.517                    |                             |
### Table 1 (Continued)

| Family          | Name of species          | Life stages | Body segment | Substrate* | Specific activity* (U/mg) | References            |
|-----------------|--------------------------|-------------|--------------|------------|---------------------------|-----------------------|
| Chrysomelidae   | *Leptinotarsa decemlineata* | –           | G            | CMC        | 0.28                      | (Oppert et al. 2010)  |
|                 |                          |             | H            | CMC        | 0.18                      |                       |
|                 | Podontia quatuordecimpunctata | Larva   | G            | CMC        | 1.73                      | (Uddin et al. 2012)  |
|                 | *Aulacophora foveicollis* | Larva       | G            | CMC        | –                         | (Sami & Shakoori 2008) |
|                 | *Xanthogaleruca luteola*  | Larva       | G            | CMC        | –                         | (Vatanparast et al. 2012) |
|                 | *Gastrophysa viridula*    | Larva       | G            | CMC        | –                         | (Busch et al. 2018)  |
|                 |                          | Adult       | G            | CMC        | –                         |                       |
|                 | *Hoplasoma unicolor*      | Adult       | G            | CMC        | 0.69                      | Present study         |
| Curculionidae   | *Rhynchophorus ferrugineus* | Larva     | G            | CMC        | –                         | (Vatanparast et al. 2014) |
|                 | *Graphognathus leucoloma* | Larva       | G            | CMC        | 0.27                      | (Oppert et al. 2010)  |
| Scarabaeidae    | *Phyllophaga* sp          | Larva       | G            | CMC        | 0.01                      | (Oppert et al. 2010)  |
|                 |                          |             | H            | CMC        | 0.01                      |                       |
|                 | *Popilia japonica*        | –           | G            | CMC        | 0.03                      |                       |
|                 |                          |             |              | MCC        | 0.02                      |                       |
| Lyctidae        | *Lyctus [prob. planicollis Lec.]* | Adult | G            | CMC        | 0.20                      | (Oppert et al. 2010)  |
|                 |                          |             |              | MCC        | 0.06                      |                       |

(Continued on next page)
| Family       | Name of species | Life stages | Body segment | Substrate | Specific activity (U/mg) | References                      |
|--------------|-----------------|-------------|--------------|-----------|--------------------------|---------------------------------|
| Scolytinae   | *Scolytus* [prob. *rugulosus* (Müller)] | Larva       | G            | CMC       | 0.55                     | (Oppert et al. 2010)            |
|              |                 |             |              | MCC       | 0.03                     |                                 |
| Buprestidae  | *Chrysobothris* sp. | Larva       | G            | CMC       | 0.29                     |                                 |
| Lucanidae    | *Dorcus titanus* | Larva       | G            | CMC       | 0.05                     | (Su et al. 2013)                |
|              |                 |             |              | FP        | 0.03                     |                                 |
| Anobiidae    | *Lasioderma serricorne* | Larva       | G            | CMC       | 0.03                     | (Su et al. 2013)                |
|              | *Leptinotarsa decemlineata* | Larva       | G            | CMC       | 0.015                    |                                 |
|              | *Lasioderma serricorne* | Larva       | G            | CMC       |                          | (Minoo et al. 2012)            |
| Elateridae   | *Pyrearinus termitilluminans* | Larva       | G            | CMC       | 0.014                    | (Colepicolo-Neto et al. 1986)   |

*Note: d1 to d7 = Fed with experimental diets; m = Male, f = Female; s = CMC: CarboxyMethyl Cellulose, MCC: Micro Crystalline Cellulose, FP: Filter Paper; a = In most papers the specific activity was not exactly mentioned and the info was rather extracted from the figures; u = µmol (glucose)·g⁻¹ (fresh weight)·h⁻¹; c = Cellulase gene was cloned and expressed.*
Qualitative Endoglucanase Assay

Cellulolytic activity of the gut extract was determined essentially as described previously (Hossain et al. 2020). Briefly, 100 µL of the gut extract was placed inside wells in agar media (pH 5, 5.3 and 6) supplemented with 1% (w/v) CMC (Sigma-Aldrich) in 0.1 M sodium acetate buffer. A control plate was also included where gut extract was replaced with the same volume of buffer only. After overnight incubation at 37°C, the plates were stained with 0.1% Congo red solution for 5–10 min and destained with 0.1M NaCl for 10–15 min. Formation of clear zone was considered positive result for endoglucanase activity.

Protein Estimation

Amount of proteins in the gut extract was estimated by Lowry protein assay method using bovine serum albumin (BSA) as standard (Lowry et al. 1951).

Electrophoresis

The enzyme extract was examined by sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) following the method as described by Laemmli (1970). Briefly, aliquots containing samples equivalent to 0.8 µg and 4 µg of protein were applied to pre-casted gel and electrophoresis was conducted with running buffer (25 mM Tris, 192 mM glycine with 0.1% SDS) at constant voltage of 110 V. Proteins were stained with Coomassie Brilliant Blue R-250.

Zymogram Analysis

We carried out zymogram analysis to detect the specific proteins having cellulolytic activity in the gut extract as described in a previous work (Schwarz et al. 1987) with small modifications as in Uddin et al. (2012). Gel was prepared including 0.1% CMC before polymerisation of resolving gel. Gut sample (0.8 µg or 4 µg of protein) containing 1× loading buffer (50 mM Tris HCl pH 6.8, 1.5% SDS, 0.02% bromophenol blue, 10% glycerol and 2% β-mercaptoethanol) was loaded onto the gel, with or without a pre-heat treatment at 55°C for 30 min. Electrophoresis was conducted at 4°C at 110 V. After electrophoresis, gel was washed five times (5–6 min each) in 50 mL of washing buffer (0.1 M sodium succinate pH 5.3) at room temperature followed by a final wash for 30 min with the same buffer. Gel was stained with 0.1% Congo red solution for 10–15 min and destained using 1 M NaCl until clear depolymerisation bands became visible.

Quantitative Endoglucanase Assay

Cellulase activity was quantified using a modified 3,5-dinitrosalicylic acid (DNSA) assay (Miller 1959). Enzyme extract (20 µg) was mixed with 235 µL CMC (1% CMC sodium salt in 0.1 M Na-acetate buffer pH 5.3). After incubation at 37°C
for 45 min in water bath, 450 µL of DNS was added to stop the reaction followed by heating in boiling water bath for 10 min and then 40% of 230 µL of Rochelle salt was added. Absorbance was measured at 540 nm by UV spectrophotometer against a reaction blank in which DNS solution was added before the enzyme extract. A standard curve of absorbance against glucose (50–400 µg) was constructed to calculate the amount of reducing sugar released (glucose equivalents) during the assay. One unit of cellulolytic activity was defined as the amount of enzyme required to produce 1 μmol of reducing sugar (glucose equivalents) per min at 37°C at pH 5.3. Specific activity is described as units per mg of protein. The experiment was done in triplicate and the mean value is presented.

RESULTS AND DISCUSSION

Endo-1,4-β-D-Glucanase Activity of the Gut Extract

This work aimed at assessing the endo-1,4-β-D-glucanase activity in digestive fluid of the phytophagous beetle *H. unicolor*. To this end, we collected digestive fluid from 20 adult insects and obtained its protein fraction by centrifugation. This partially purified gut extract was examined for the presence of endoglucanase activity in an agar plate assay using CMC as the substrate. A distinct zone of clear halo was formed in the agar media when spotted with the gut extract that clearly indicated depolymerisation of CMC in the media confirming presence of cellulolytic proteins in the sample [Figs. 1(a) and 1(b)]. With several beetles from the order Coleoptera are already known to have cellulolytic activity in its digestive fluid (Table 1), our study, therefore, makes a new inclusion, *H. unicolor*, to that list. Although most of the previous studies didn’t confirm whether the respective enzymes are encoded within the insect genome or of microbial origin (Geib *et al.* 2010; Oppert *et al.* 2010; Rehman *et al.* 2009; Sami & Shakoori 2008; Su *et al.* 2013; Uddin *et al.* 2012). In our study as well, whether the endoglucanase activity is endogenous or, secreted from microbial symbionts, or, contributed by both, needs to be investigated by more extensive analysis in the future. In the digestion of recalcitrant plant matters, enzymes from both the insects and the symbionts are required for complete breakdown of the polymers, e.g., cellulose into simple sugars (Shelomi *et al.* 2020). Availability of the genome sequence of the beetle might also provide us gaining more insights into its endogenous cellulases.

In previous works, the endoglucanase activity assay of the gut fluid of Coleoptera insects was mostly carried out at pH ~5 to 6 (Geib *et al.* 2010; Oppert *et al.* 2010; Su *et al.* 2013; Vatanparast *et al.* 2014; Willis *et al.* 2010). Whereas in a study with *Podontia quatuordecimpunctata*, an insect from the same family as *H. unicolor*, the enzyme assay was also performed at pH 5.3 (Uddin *et al.* 2012). Therefore, to determine the optimum pH for endoglucanase activity of the enzyme extract, we initially decided to perform the agar plate assay at three different pH ranging from 5 to 6. Area of clear halos formed on the agar plates, which indicates the enzyme intensity (Hossain *et al.* 2020), was visually inspected. We found that
the endoglucanase activity of the gut fluid was highest at pH 5.3, whereas the activity decreased below and above this pH (i.e., pH 5 and 6) with the smallest zone of clearance formed at pH 6. Since lower enzyme activity was found on either side of pH 5.3, therefore, the endoglucanase activity was not examined at other pH ranges. Hence, it appears from the above findings that the cellulolytic enzymes present in the gut sample exert maximum activity at acidic pH. This finding is consistent with those reported in a number of previous research. For example, the endoglucanases from gut fluid of a number of beetles and termites as well as bacteria and fungi showed highest activity at slightly acidic pH mostly in the range of pH 5 to 6 (Busch et al. 2018; Hatefi et al. 2017; Tokuda et al. 1997; Zhang et al. 2011). Insect cellulases with optimum activity at alkaline pH were also reported though (Sami & Shakoori 2008; Willis et al. 2011).

![Figure 1](image1.png)

**Figure 1**: Endoglucanase activity of the digestive fluid from *H. unicolor* adults. (a) Beetles on its host plant at the time of collection; (b) and (c) Screening endo-1,4-β-D-glucanase activity in the gut extract prepared from digestive fluid of the beetle. Wells were made in agar media containing CMC and loaded with either buffer solution (b) or the gut extract (c). Clear zone was only produced by the gut extract indicating the presence of endoglucanase activity.
SDS-PAGE and Zymogram Analysis

SDS-PAGE analysis of the digestive extract was carried out to examine the purity and number of proteins present in the partially purified extract and to reveal their molecular weight (MW). The gut extract was used at two different concentrations, 0.8 µg or 4 µg of protein, in the analysis. Five distinct protein bands were detected at both concentrations with the MW ranging from ~25 to 65 kDa (Fig. 2), suggesting that the MW of the cellulolytic proteins in the sample lies within this range. Detection of only five protein bands also indicates that the gut sample could be fairly purified.

![SDS-PAGE Image]

**Figure 2:** Separation of proteins of the gut extract by SDS-PAGE. Samples containing 0.8 µg or 4 µg of proteins were loaded onto the gel and, after electrophoresis, stained with Coomassie Brilliant Blue. Each of the protein bands detected is indicated by an arrow-head.

To find out which of the five proteins were the active endoglucanase enzymes in the digestive extract, we performed a zymogram analysis using CMC as the substrate. Two prominent bands of clearance due to the degradation of CMC were observed that were in close proximity to each other which suggested the presence of at least two endoglucanase enzymes in the gut sample (Fig. 3). When the sample was pre-heated at 55°C for 30 min before loading onto the gel, no hydrolysis band was detected indicating complete loss of enzyme activity due to heat inactivation; although a number of insect cellulases were previously demonstrated to be enzymatically active at similar temperatures (Sami & Shakoori 2008; Tokuda *et al.* 1997; Watanabe *et al.* 1997). Together, the results of SDS-PAGE and zymogram analysis suggested that only two of the five proteins detected in the gut sample possessed endoglucanase activity in the experimental conditions used. Detection of multiple cellulase enzymes in insect gut sample is not uncommon. Similar to our findings, two cellulolytic protein bands were reported in other insects such as *Tribolium castaneum, Syrbula admirabilis* and *Tenebrio molitor* (Oppert *et al.* 2010; Rehman *et al.* 2009). Some studies also described the detection of more than two cellulolytic proteins in zymography (Su *et al.* 2013; Uddin *et al.* 2012; Willis *et al.* 2010 ).
The two active enzymes detected in the zymography were very close to each other having similar molecular masses. Of the five proteins detected in SDS-PAGE, on the other hand, only the two at bottom of the gel were closely placed. Taken together, we assume that the proteins which were found in close vicinity in SDS-PAGE analysis with MW of ~25 and ~30 kDa, might be the endoglucanases. The MW of these two enzymes is similar to those reported in various other insects as well as in other organisms. For example, *Rhagium inquisitor*, *Dissosteira carolina*, *S. admirabilis*, *T. molitor* and others were documented to have cellulases in their digestive fluids with the MW lying in the range of 20 to 35 kDa (Oppert et al. 2010; Rehman et al. 2009; Willis et al. 2010).

**Figure 3:** Zymogram analysis for detection of the active endoglucanases in the digestive fluid of adult *H. unicolor*. Proteins in the sample (~800 ng) were separated by electrophoresis on gel containing CMC, and the active endoglucanases were subsequently visualised by staining with Congo red. Clear bands in the gel indicate cellulolytic activity due to the degradation of CMC. Arrow-heads indicate the position of the active enzymes. Lanes 1 and 3: 4 µg sample pre-heated at 55°C. Lanes 2 and 4: 4 µg sample without pre-heat treatment. Lanes 5 and 7: 0.8 µg sample pre-heated at 55°C. Lanes 6 and 8: 0.8 µg sample without pre-heat treatment.

**Quantitation of Endo-1,4-β-D-Glucanase Activity**

Finally, to determine specific activity, we carried out a quantitative analysis of endoglucanases present in the enzyme extract towards CMC. The specific enzyme activity was measured as 0.69 (± 0.01) U/mg of protein which seems quite high as compared to those reported in the insects of the same order (Table 1) and in other insects as well (Oppert et al. 2010). The endoglucanase activity in the members of Coleoptera has been reported to vary over a very wide range, from as little as 0.01 to as much as 2.80 U/mg (Table 1). The highest activity...
was found in the beetle *Morimus funereus* (Dojnov et al. 2013). Direct comparison of the activities among various endoglucanases is not, however, fully coherent due to the different experimental conditions used in the studies.

**CONCLUSION**

Findings of this paper represent the first description and preliminary characterisation of the cellulolytic activity in the digestive fluid of *H. unicolor*. The endo-1,4-β-D-glucanases of the gut fluid showed prominent efficiency in digestion of cellulose with specific activity higher than those of most other insects. Further research for purification and complete characterisation of the endoglucanases are, therefore, needed to understand their origin and catalytic efficacy which can provide foundation for development of an efficient and inexpensive system for biofuel production and waste management based on the hydrolytic enzymes of insect digestive fluid.

**ACKNOWLEDGEMENTS**

We are grateful to Dr. Dwaipayan Sikdar, Department of Biochemistry and Molecular Biology, University of Chittagong (BMB, CU) for his help and advice. We thank Sumaiya Islam Chowdhury, Biochemistry and Pathogenesis of Microbes Research Group, BMB, CU, and all laboratory members of Department of Biochemistry and Molecular Biology, Mawlana Bhashani Science and Technology University for their help.

**REFERENCES**

Ali S, Hall J, Soole K L, Fontes C M G A, Hazlewood G P, Hirst B H and Gilbert H J. (1995). Targeted expression of microbial cellulases in transgenic animals. In S B Petersen, B Svensson and S Pedersen (eds.). *Progress in biotechnology*. Amsterdam: Elsevier, 279–293. https://doi.org/10.1016/S0921-0423(06)80111-5

Annamalai N, Rajeswari M V and Balasubramanian T. (2016). Endo-1,4-β-glucanases: Role, Applications and Recent Developments. In V K Gupta (ed.). *Microbial enzymes in bioconversions of biomass*. Cham: Springer International Publishing, 37–45. https://doi.org/10.1007/978-3-319-43679-1_3

Bayer E A, Chanzy H, Lamed R and Shoham Y. (1998). Cellulose, cellulases and cellulosomes. *Current Opinion in Structural Biology* 8(5): 548–557. https://doi.org/10.1016/S0959-440X(98)80143-7

Bayer E A, Lamed R and Himmel M E. (2007). The potential of cellulases and cellulosomes for cellulosic waste management. *Current Opinion in Biotechnology* 18(3): 237–245. https://doi.org/10.1016/j.copbio.2007.04.004
Busch A, Kunert G, Wielsch N and Pauchet Y. (2018). Cellulose degradation in Gastrophysa viridula (Coleoptera: Chrysomelidae): Functional characterization of two CAZymes belonging to glycoside hydrolase family 45 reveals a novel enzymatic activity. *Insect Molecular Biology* 27(5): 633–650. https://doi.org/10.1111/imb.12500

Colepicolo-Neto P, Bechara E J H, Ferreira C and Terra W R. (1986). Evolutionary considerations of the spatial organization of digestion in the luminescent predaceous larvae of *Pyrearinus termitilluminans* (Coleoptera: Elateridae). *Insect Biochemistry* 16(5): 811–817. https://doi.org/10.1016/0020-1790(86)90118-6

Dashtban M, Maki M, Leung K T, Mao C and Qin W. (2010). Cellulase activities in biomass conversion: Measurement methods and comparison. *Critical Reviews in Biotechnology* 30(4): 302–309. https://doi.org/10.3109/07388551.2010.490938

Dojnov B, Pavlović R, Božić N, Margetić A, Nenadović V, Ivanović J and Vujčić Z. (2013). Expression and distribution of cellulase, amylase and peptidase isoforms along the midgut of *Mormus funereus* L. (Coleoptera: Cerambycidae) larvae is dependent on nutrient substrate composition. *Comparative Biochemistry and Physiology Part B: Biochemistry and Molecular Biology* 164(4): 259–267. https://doi.org/10.1016/j.cbpb.2013.02.001

Geib S M, Tien M and Hoover K. (2010). Identification of proteins involved in lignocellulose degradation using in gel zymogram analysis combined with mass spectrscopy-based peptide analysis of gut proteins from larval Asian longhorned beetles, *Anoplophora glabripennis*. *Insect Science* 17(3): 253–264. https://doi.org/10.1111/j.1744-7917.2010.01323.x

Girfoglio M, Rossi M and Cannio R. (2012). Cellulose degradation by *Sulfolobus solfataricus* requires a cell-anchored endo-β-1-4-glucanase. *Journal of Bacteriology* 194(18): 5091–5100. https://doi.org/10.1128/JB.00672-12

Hatefi A, Makhdoumi A, Asoodeh A and Mirshamsi O. (2017). Characterization of a bi-functional cellulase produced by a gut bacterial resident of Rosaceae branch borer beetle, *Osphranteria coerulescens* (Coleoptera: Cerambycidae). *International Journal of Biological Macromolecules* 103: 158–164. https://doi.org/10.1016/j.ijbiomac.2017.05.042

Hazmi I R, Sulaiman A, Azhari M L H and Samsudin S F. (2019). A checklist of beetles in Pulau Pangkor, Perak, Malaysia. *The Malaysian Forester* 82(1): 237–246.

Hobdey S E, Donohoe BS, Brunecky R, Himmel M E and Bomble Y J. (2015). Chapter 7: New insights into microbial strategies for biomass conversion. In M E Himmel (ed.). *Direct microbial conversion of biomass to advanced biofuels*. Amsterdam: Elsevier, 111–127. https://doi.org/10.1016/B978-0-444-59592-8.00007-5

Hossain T J, Chowdhury S I, Mozumder H A, Chowdhury M N A, Ali F, Rahman N and Dey S. (2020). Hydrolytic exoenzymes produced by bacteria isolated and identified from the gastrointestinal tract of bombay duck. *Frontiers in Microbiology* 11: 2097. https://doi.org/10.3389/fmicb.2020.02097

Jayasekara S and Ratnayake R. (2019). Microbial cellulases: An overview and applications. In A R Pascual and M E E Martin (eds.). *Cellulose*. IntechOpen [Online]. https://doi.org/10.5772/intechopen.84531

Laemmli U K. (1970). Cleavage of structural proteins during the assembly of the head of bacteriophage T4. *Nature* 227(5259): 680–685. https://doi.org/10.1038/227680a0

Li X, Yan X, Luo Y, Tian G and Sun H. (2008). Cellulase in *Anoplophora glabripennis* adults emerging from different host tree species. *Forestry Studies in China* 10(1): 27–31. https://doi.org/10.1007/s11632-008-0004-z
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Lowry O H, Rosebrough N J, Farr A L and Randall R J. (1951). Protein measurement with the Folin phenol reagent. The Journal of Biological Chemistry 193(1): 265–275. https://doi.org/10.1016/S0021-9258(19)52451-6

Mathew G, Shamsudeen R and Chandran R. (2005). Fauna of protected areas - 23: Insect fauna of Peechi-Vazhani Wildlife Sanctuary, Kerala, India. Zoo’s Print Journal 20(8): 1955–1960. https://doi.org/10.11609/JOTT.ZPJ.1280.1955-60

Miller G L. (1959). Use of dinitrosalicylic acid reagent for determination of reducing sugar. Analytical Chemistry 31(3): 426–428. https://doi.org/10.1021/ac60147a030

Minoo S S, Vahid H N and Mohammad V. (2012). Cellulase activity in the larval digestive tract of the Colorado potato beetle, Leptinotarsa decemlineata (Coleoptera: Chrysomelidae) and the cigarette beetle, Lasioderma serricorne(Coleoptera: Anobiidae). Journal of Crop Protection 1(3): 201–210.

Okano H, Ozaki M, Kanaya E, Kim J-J, Angkawidjaja C, Koga Y and Kanaya S. (2014). Structure and stability of metagenome-derived glycoside hydrolase family 12 cellulase (LC-CelA) a homolog of Cel12A from Rhodothermus marinus. FEBS Open Bio 4: 936–946. https://doi.org/10.1016/j.fob.2014.10.013

Oppert C, Klingeman W E, Willis J D, Oppert B and Jurat-Fuentes J L. (2010). Prospecting for cellulolytic activity in insect digestive fluids. Comparative Biochemistry and Physiology. Part B, Biochemistry & Molecular Biology 155(2): 145–154. https://doi.org/10.1016/j.cbpb.2009.10.014

Phitsuwan P, Laohakunjit N, Kerchoechuen O, Kyu K L and Ratanakhanokchhai K. (2013). Present and potential applications of cellulases in agriculture, biotechnology, and bioenergy. Folia Microbiologica 58(2): 163–176. https://doi.org/10.1007/s12223-012-0184-8

Rehman F U, Aslam M, Tariq M I, Shaheen A, Sami A J, Naveed N H and Batool A I. (2009). Isolation of cellulolytic activity from Tribolium castaneum (red flour beetle). African Journal of Biotechnology 8(23): 6710–6715. https://doi.org/10.4314/ajb.v8i23.66387

Sami A J and Shakoori A. (2008). Biochemical characterization of endo-1, 4-β-D-glucanase activity of a green insect pest Aulacophora foveicollis (Lucas). Life Science Journal 5(2): 30–36.

Schwarz W H, Bronnenmeier K, Gräbnitz F and Staudenbauer W L. (1987). Activity staining of cellulases in polyacrylamide gels containing mixed linkage beta-glucans. Analytical Biochemistry 164(1): 72–77. https://doi.org/10.1016/0003-2697(87)90369-1

Shelomi M, Wipfler B, Zhou X and Pauchet Y. (2020). Multifunctional cellulase enzymes are ancestral in Polyneoptera. Insect Molecular Biology 29(1): 124–135. https://doi.org/10.1111/imb.12614

Shi W, Ding S-Y and Yuan J S. (2011). Comparison of insect gut cellulase and xylanase activity across different insect species with distinct food sources. BioEnergy Research 4(1): 1–10. https://doi.org/10.1007/s12155-010-9096-0

Su L-J, Zhang H-F, Yin X-M, Chen M, Wang F-Q, Xie H, Zhang G-Z and Song A-D. (2013). Evaluation of cellulolytic activity in insect digestive fluids. Genetics and Molecular Research 12(3): 2432–2441. https://doi.org/10.4238/2013.January.4.11

Szentner K, Waśkiewicz A, Kaźmierczak S, Wojciechowicz T, Goliński P, Lewandowska E and Wasielewski O. (2019). Enzymatic hydrolysis of cellulose using extracts from insects. Carbohydrate Research 485: 107811. https://doi.org/10.1016/j.carres.2019.107811
Tokuda G, Watanabe H, Matsumoto T and Noda H. (1997). Cellulose digestion in the wood-eating higher termite, *Nasutitermes takasagoensis* (Shiraki): Distribution of cellulases and properties of endo-β-1,4-glucanase. *Zoological Science* 14(1): 83–93. https://doi.org/10.2108/zsj.14.83

Uddin M M, Chowdhury M M H, Mojumder S and Dwipayan S. (2012). Multiple endo-beta-1,4-glucanases present in the gut fluid of a defoliating beetle, *Podontia quatuordecimpunctata* (Coleoptera: Chrysomelidae). *Pakistan Journal of Biological Sciences* 15(7): 333–340. https://doi.org/10.3923/pjbs.2012.333.340

Vatanparast M, Hosseininaveh V, Ghadamyari M and Sajjadi S M. (2012). Pectinase and cellulase activity in the digestive system of the elm leaf beetle. *Xanthogaleruca luteola* Muller (Coleoptera: Chrysomelidae). *Journal of Asia-Pacific Entomology* 15(4): 555–561. https://doi.org/10.3923/pjbs.2012.05.017

Vatanparast M, Hosseininaveh V, Ghadamyari M and Sajjadi S M. (2014). Plant cell wall degrading enzymes, pectinase and cellulase, in the digestive system of the red palm weevil, *Rhynchophorus ferrugineus* (Coleoptera: Curculionidae). *Plant Protection Science* 50(4): 190–198. https://doi.org/10.17221/43/2013-PPS

Watanabe H, Nakamura M, Tokuda G, Yamaoka I, Scrivener A M and Noda H. (1997). Site of secretion and properties of endogenous endo-β-1,4-glucanase components from *Reticulitermes speratus* (Kolbe), a Japanese subterranean termite. *Insect Biochemistry and Molecular Biology* 27(4): 305–313. https://doi.org/10.1016/S0965-1748(97)00003-9

Watanabe H, Noda H, Tokuda G and Lo N. (1998). A cellulase gene of termite origin. *Nature* 394(6691): 330–331. https://doi.org/10.1038/28527

Willis J D, Klingenman W E, Oppert C, Oppert B and Jurat-Fuentes J L. (2010). Characterization of cellulolytic activity from digestive fluids of *Dissosteira carolina* (Orthoptera: Acrididae). *Comparative Biochemistry and Physiology Part B: Biochemistry and Molecular Biology* 157(3): 267–272. https://doi.org/10.1016/j.cbpb.2010.06.012

Willis J D, Oppert B, Oppert C, Klingenman W E and Jurat-Fuentes J L. (2011). Identification, cloning, and expression of a GHF9 cellulase from *Tribolium castaneum* (Coleoptera: Tenebrionidae). *Journal of Insect Physiology* 57(2): 300–306. https://doi.org/10.1016/j.jinsphys.2010.11.019

Zhang D, Lax A R, Bland J M and Allen A B. (2011). Characterization of a new endogenous endo-β-1,4-glucanase of *Formosan subterranean* termite (*Coptotermes formosanus*). *Insect Biochemistry and Molecular Biology* 41(4): 211–218. https://doi.org/10.1016/j.ibmb.2010.12.006