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Enzyme kinetics of hevamine, a chitinase from the rubber tree
Hevea brasiliensis

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Abstract The enzyme kinetics of hevamine, a chitinase from the rubber tree Hevea brasiliensis, were studied in detail with a new enzyme assay. In this assay, the enzyme reaction products were derivatized by reductive coupling to a chromophore. Products were separated by HPLC and the amount of product was calculated by peak integration. Penta-N-acetylglucosamine (penta-nag) and hexa-N-acetylglucosamine (hexa-nag) were used as substrates. Hexa-nag was more efficiently converted than penta-nag, which is an indication that hevamine has at least six sugar binding sites in the active site. Tetra-N-acetylglucosamine (tetra-nag) and allosamidin were tested as inhibitors. Allosamidin was found to be a competitive inhibitor with a $K_i$ of 3.1 μM. Under the conditions tested, tetra-nag did not inhibit hevamine. © 2000 Federation of European Biochemical Societies. Published by Elsevier Science B.V. All rights reserved.

Key words: Chitinase; Competitive inhibition; Chitin oligomer; Hevea brasiliensis

1. Introduction

Chitin, a $\beta$ (1,4)-linked $N$-acetylglucosamine (GlcNAc) polymer, is a major component of the exoskeleton of fungi and invertebrates. These organisms produce chitinases to remodel their exoskeleton during growth and cell division [1]. Other organisms make chitinases as well. For instance plants use chitinases as a defense against pathogenic fungi [2]. Hevamine, a chitinase from the rubber tree Hevea brasiliensis belongs to the family 18 glycosyl hydrolases. This class of chitinases has been found in a wide range of eukaryotes and prokaryotes [3,4]. Primary [5] and tertiary structures of hevamine [6] have been determined. The enzyme cleaves not only chitin, but also the sugar moiety of peptidoglycan between the C-1 of a $N$-acetylglucosamine and the C-4 of a $N$-acetylmuramate residue [7] and was the first enzyme found with this cleavage specificity.

The enzyme uses a retaining catalytic mechanism, which is also the mode of action of hen egg white lysozyme (HEWL), but there is no sequence homology between HEWL and hevamine. Structural data also indicate that the cleavage mechanisms of HEWL and hevamine are completely different [8,9]. In contrast to HEWL, which uses two carboxylate residues in its reaction mechanism, hevamine has only one carboxylate in its active site. Instead of an oxocarbenium ion, which is formed during the HEWL reaction, structural and theoretical data suggest that in hevamine the reaction proceeds with an achimeric assistance of the neighboring $N$-acetyl group [10,11] and that an oxazolinium ion intermediate is formed. Support for this mechanism has been obtained from X-ray studies of the binding mode of allosamidin in the active site of hevamine. Allosamidin is thought to be a transition state analog (Fig. 1).

Although many chitinases have been isolated, detailed kinetic analyses of these proteins have been scarce. The assays for activity measurements mostly use colloidal chitin or chitin derivatives. Because of the heterogeneous nature of these substrates, it is not possible to determine $k_{cat}$ and $K_M$ values. An alternative assay uses short chitin oligomers coupled to a 4-methylumbelliferyl group and measures the fluorescence of the released umbelliferyl group, which is a very sensitive method. A disadvantage is that with this substrate only the cleavage of the umbelliferyl group is measured and not cleavage reactions that occur between two sugar residues. Sometimes cleavage does not occur between the umbelliferone group and the oligomer, but between two sugar residues in the oligomer. In such a case, there is no fluorescence and no activity will be measured [12].

In this paper, we describe the production of short $N$-acetylglucosamine oligomers and the kinetics of hevamine with these substrates by a new assay, which does not suffer from the disadvantages of the existing assays. After incubation with enzyme, the products are coupled to an aromatic chromophore so that sensitive detection is possible in the near UV. Coupling to the chromophore also enhances the separation of the products by high performance liquid chromatography (HPLC). With this assay, the inhibition kinetics of hevamine with allosamidin and chitotetraose were studied in detail.

![Fig. 1. Structure of allosamidin.](image-url)
2. Materials and methods

2.1. Preparation of chitin oligomers

Chitin oligomers were prepared according to the method of Aiba et al. [13] by partial acetylation of chitosan, followed by cleavage with a chitinase and complete acetylation of the product. 1 g of chitosan was dissolved in 20 ml of 3% acetic acid. After one night of solubilization, 80 ml of methanol was slowly added under stirring until a homogeneous solution was obtained. Subsequently 100 µl of acetic acid anhydride was added to 100 ml reaction mixture in 5 min period under vigorous mixing conditions. Mixing continued for at least 1 h. Methanol and acetic acid were removed by evaporation in a film evaporator. The partially acetylated chitosan was dissolved in 20 ml 0.2 M ammonium acetate, pH 5.4. 1.5 U chitinase from Streptomyces griseus (Sigma) was added and the solution was incubated for 1 week at 37°C. After 1 week, another 1.5 U of chitinase was added and incubation was continued for another week. After drying, the sample was fully acetylated as described above, by using 1.5 ml instead of 100 µl acetic acid anhydride. Again the sample was dried and dissolved in 20 ml of water. 100-150 mg of oligomers were loaded on a Biogel P4 extra fine grade (Bio-Rad) gel filtration column of 98×3.2 cm. Several peaks were observed in duplicate or triplicate per substrate concentration. Approximately 1 pmol of hevamine was added to 1.5-3 ml reaction mixture in Greiner tubes. After 30 min, the reaction was stopped, by cooling the samples in liquid nitrogen. After freeze-drying, the samples were derivatized by reductive coupling of p-aminoethylbenzamido with sodium cyanoborohydride. The water phase was extracted with chloroform to remove unreacted ABEE. The water phase was added. The reagent solution was prepared by dissolving 165 mg penta-acetylglucosamine (pentamer) and 35 mg of sodium cyanoborohydride in 20 ml glacial acetic acid and 350 µl of methanol. The tubes were capped and heated at 80°C for 30 min.

After drying, the samples were dissolved in 350 µl of water containing approximately 0.50 µM of internal standard and extracted with chloroform to remove unreacted ABEE. The water phase was analyzed by HPLC. With penta-N-acetylglucosamine (pentamer) as a substrate, di-N-acetylgulosamine was used as an internal standard. When hevamine cleaves oligomers that are larger than four residues, pentamer is cleaved into a tetramer and a monomer, hexa-nag is cleaved into a heptamer and a dimer [10]. In Fig. 3, a chromatogram is shown of the digestion of pentamers by hevamine. Dimer is added as an internal standard. The substrate, product and internal standard are baseline separated, meaning that the product peak can be measured accurately by peak integration. The retention time of derivatized oligosaccharrides decreases with increasing degree of polymerization. Tetra-N-acetylglucosamine (tetra-nag)-ABEE and penta-nag-ABEE hardly differ in retention time. Therefore the small tetra-nag-ABEE peak is completely overlapped by penta-nag-ABEE and cannot be measured.

Hevamine shows Michaelis–Menten behavior with penta-nag and hexa-nag as substrates (Fig. 4). The $k_{cat}$ and $K_M$ of hevamine with penta-nag and hexa-nag are given in Table 1. Hexa-nag was converted more efficiently than penta-nag, as hevamine has a higher $k_{cat}$ and a lower $K_M$ value. From modelling studies, it was already proposed that hevamine can bind six sugar residues in the active site. The kinetic data show that this indeed is the case. Adding 3.13 µM of allosamidin to the incubation mix gave a $K_M$ of (27.6 ± 2.7) µM and a $k_{cat}$ of (0.385 ± 0.016) s⁻¹ (Fig. 4).

There was no significant change in $k_{cat}$, but a significant change in $K_M$. The $K_i$ value was 3.1 µM. Adding 15 µM of hevamine decreases with increasing molecular weight. These results are consistent with the results of Aiba [13]. This procedure yielded oligomers with a purity of more than 80%. Further purification of the oligomers by reversed phase HPLC yielded sufficient pure products (>95%) for enzyme kinetic measurements.

3. Results

In Fig. 2, the result of the separation of N-acetylgulosamine oligomers, prepared as described, is shown. The amount of hevamine is less than 10% of the total amount of oligomers. The amount of shorter oligomers increases with decreasing molecular weight. These results are consistent with the results of Aiba [13]. This procedure yielded oligomers
tetra-nag to the incubation mixture did not inhibit the enzyme reaction, showing that tetra-nag, although it is binding in the active site, is not a high affinity inhibitor.

4. Discussion

The kinetic data indicate that the active site of hevamine contains at least six sugar binding sites. This was shown with enzyme kinetic experiments with short chitin oligomers. The results showed that hevamine has a higher \( k_{\text{cat}} \) and a lower \( K_M \) with hexa-nag compared to penta-nag as substrate, meaning that binding of a sixth residue enhances catalysis. This was already proposed for hevamine from modelling studies. Soaking of penta-nag or hexa-nag in the active site of hevamine was unsuccessful, because these substrates were degraded in the hevamine crystals (unpublished results). It was not possible to accurately determine the \( K_M \) and \( k_{\text{cat}} \) values of hevamine for hexa-nag, because the method used is not sensitive enough to measure product peaks accurately below concentrations of 0.2 \( \mu \)M. These kinetic studies also show that binding of the sixth sugar residue in the active site enhances the enzyme reaction. The additional binding energy of hexa-nag compared to penta-nag can be calculated with the following formula [16]:

\[
\frac{(k_{\text{cat}}/K_M)A}{(k_{\text{cat}}/K_M)B} = \exp\left(\frac{-\Delta\Delta G_b}{RT}\right)
\]

The additional binding energy \( \Delta\Delta G_b \) of the sixth residue is approximately \(-6\) kJ/mol.

Of the two compounds tested, only allosamidin gave a clear competitive inhibition. In the literature, there are many studies of inhibition of chitinases by allosamidin and allosamidin analogues [17]. In most cases reported, allosamidin gave a competitive inhibition [18–21]. The \( K_i \) values found were approximately one order of magnitude lower than the value found for hevamine, meaning that hevamine has a relatively low binding affinity for allosamidin, compared to other family 18 chitinase. Only a cytosolic chitinase from Neurospora crassa gave non-competitive inhibition [22]. Milewski et al. [21] studied inhibition of Candida albicans chitinase, using N-umbelliferone chitooligosaccharides as substrates, and found that the mode of inhibition also depends on the buffer system used. In PIPES, MES and phosphate buffers, allosamidin was a competitive inhibitor. In citrate containing buffers, the mode of inhibition was mixed or even non-competitive. Our buffer system also contained citrate but we did not observe such a buffer effect.

These experiments also showed, that tetra-nag is not a high affinity inhibitor. Our results indicate that its \( K_i \) value is much higher than 20 \( \mu \)M, which contrasts with the \( K_i \) value of 20 \( \mu \)M of tetra-nag bound to HEWL [23]. This was somewhat surprising, because soaking studies showed that tetra-nag binds in positions \(-4 \) to \(-1 \) in the active site [6]. Allosamidin binds in the active site in positions \(-3 \) to \(-1 \) [10]. This mode of binding combined with the low \( K_i \) value of allosamidin is an indication that the oxazoline ring is essential for the binding in the active site of hevamine.

This new enzyme assay is very useful for accurate determination of \( K_M \) and \( k_{\text{cat}} \) values and will be used in the future to characterize the kinetic constants of hevamine mutants.

References


### Table 1

Kinetic parameters, with their standard deviations, of hevamine with penta- and hexa-nag as substrates

<table>
<thead>
<tr>
<th>Substrate</th>
<th>( K_M ) (( \mu )M)</th>
<th>( k_{\text{cat}} ) (s(^{-1}))</th>
<th>( K_M/k_{\text{cat}} )</th>
</tr>
</thead>
<tbody>
<tr>
<td>Penta-nag</td>
<td>(13.8 ± 0.7)</td>
<td>(0.355 ± 0.010)</td>
<td>(2.57 ± 0.21)( \times )10(^8) M(^{-1}) s(^{-1})</td>
</tr>
<tr>
<td>Hexa-nag</td>
<td>(3.2 ± 0.8)</td>
<td>(1.0 ± 0.06)</td>
<td>(3.1 ± 1.0)( \times )10(^3) M(^{-1}) s(^{-1})</td>
</tr>
</tbody>
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