Inositol Phosphate Phosphatases of Microbiological Origin: the Inositol Pentaphosphate Products of Aspergillus ficuum Phytases

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The fungus Aspergillus ficuum NRRL ³¹³⁵ is known to produce an extracellular nonspecific orthophosphoric monoester phosphohydrolase (EC 3.1.3.2) with a pH optimum of 2.0, as well as an extracellular $m\nu$ o-inositol hexaphosphate phosphohydrolase (EC 3.1.3.8; phytase) with pH optima of 2.0 and 5.5. Both these enzymes are also known to hydrolyze myo-inositol hexaphosphate. The pentaphosphates liberated in the first step of this hydrolysis have been isolated and identified by ion-exchange chromatography and optical rotation. The nonspecific orthophosphoric monoester phosphohydrolase produces a single pentaphosphate, D-myo-inositol-1,2,4,5,6-pentaphosphate, whereas the phytase, at both pH 2.0 and 5.5, produces ^a mixture of two pentaphosphates. The major component of this mixture is $D\text{-}myo$ -inositol-1,2,4,5,6-pentaphosphate and the other is D-myo-inositol-1,2,3,4,5-pentaphosphate. Thus the pathways of dephosphorylation of $m\gamma$ ^o-inositol hexaphosphate by these two enzymes differ from that of wheat-bran phytase which forms $L-my_0$ -inositol-1,2,3,4,5-pentaphosphate.

Phytases (EC 3.1.3.8) isolated from wheatbran (12) and a soil bacterium (Pseudomonas sp.) (2) have been shown to dephosphorylate myo-inositol hexaphosphate in a stepwise manner. The predominant *myo*-inositol pentaphosphates formed by these enzymes are, respectively, L-myo-inositol-1,2,3,4,5-pentaphosphate and $p\text{-}myo\text{-}inositol-1,2,4,5,6\text{-}pentaphos$ phate. No similar study of phytases of fungal origin has been published although Theodorou (11), by using electrophoresis, has shown that the phytase of a mycorrhizal fungus, Rhizopogan luteolus, produces a mixture of at least two pentaphosphates. The major component of this mixture corresponds to the major pentaphosphate liberated by Neurospora crassa phytase (7) and by Pseudomonas phytase (2), whereas the minor component corresponds to the pentaphosphate liberated by wheat-bran phytase (12).

Shieh, Wodzinski, and Warl (10) recently described the isolation and properties of two fractions with phytase activity from culture filtrates of the fungus Aspergillus ficuum NRRL 3135. One of these fractions had a single pH optimum at 2.0, whereas the other had pH optima at 2.0 and at 5.5.

The work reported in this paper examines the pentaphosphates produced by both the extracellular phytases from A. ficuum NRRL 3135 to establish whether the pathways of dephosphorylation of m y₀-inositol hexaphosphate by these enzymes resemble that of wheat-bran phytase (12) or Pseudomonas phytase (2).

MATERIALS AND METHODS

All chemicals were reagent grade unless otherwise specified.

Culture of organism. A. ficuum 3135 was maintained on agar slopes consisting of Czapek Dox medium (Difco) in which the inorganic orthophosphate had been replaced by dodecasodium myo-inositol hexaphosphate (1.75 g/liter; Sigma). Cultures of the organism were prepared by using the low-phosphate medium (2 mg of P per 100 ml) and technique of Shieh and Ware (9).

Enzyme activity. Phytase activity was assayed at 40 C by determining the rate of release of inorganic orthophosphate (5). Glycine-hydrochloric acid was used as buffer at pH 2.0, and succinic acid-sodium hydroxide was used as buffer at pH 5.5. The assay solutions were 0.02 M with respect to buffer.

Protein determinations. Protein was determined by the method of Lowry et al. (8) with bovine serum albumin as a standard.

Polysaccharide determinations. Polysaccharide was determined by the procedure of Dubois et al. (3) with sucrose as a standard.

Ion-exchange dextran chromatography of culture filtrate. Cell-free culture filtrate (50 ml) was buffer exchanged, at room temperature, into citrate buffer $(0.05 \text{ M}; pH 3.5)$ by using a column of Sephadex (2.5 by 40 cm; G25 Pharmacia). The bufferexchanged, active material (70 ml) was chromatographed at room temperature by using a column of SE-Sephadex (2.5 by ³⁴ cm; C50 Pharmacia). A linear pH gradient in citrate buffer (0.05 M) was used for elution. The mixing chamber contained 200 ml of pH 3.5 buffer, whereas the reservoir contained ²⁰⁰ ml of pH 5.5 buffer. The fraction volume was approximately 7.0 ml.

Enzyme hydrolysis of myo-inositol hexaphosphate. Dodecasodium myo-inositol hexaphosphate (50 mg; Sigma) was dissolved in 3.0 ml of the appropriate 0.05 M buffer, glycine-hydrochloric acid at pH 2.0 or succinic acid-sodium hydroxide at pH 5.5. The pH was adjusted to that of the buffer, and sufficient further buffer was added to bring the final incubation mixture volume to 10.0 ml. An appropriate volume of enzyme solution was added to initiate hydrolysis, and the mixture was incubated at 40 C until between 5.8 and 9.1% of the total ester-phosphate had been released as inorganic orthophosphate. The reaction was terminated by the addition of sufficient 1 M NaOH to raise the pH to about 10. The resulting solution was suitable for immediate ion-exchange chromatography. In preparations of pentaphosphates for optical rotation determinations, the quantity of dodecasodium myo-inositol hexaphosphate was increased to 5.0 g, and the incubation mixture volume was increased to 100 ml. The products of the large-scale preparations were isolated as their barium salts which were dissolved in 0.48 N HCI for the ion-exchange isolation of the pentaphosphates or in 0.28 N HCl for the isolation of the tetraphosphates.

Ion-exchange chromatography of myo-inositol penta- and tetraphosphates. The enzymic hydrolysates from the small-scale preparations were chromatographed by using method B described by Cosgrove (1). The fraction volume was 10.0 ml. The barium salts of the products of the large-scale preparations were chromatographed under similar conditions, but the column dimensions were increased to 2.5 by 80 cm and the fraction volume was increased to 25.0 ml. The resolved penta- and tetraphosphates were isolated as their barium salts.

Optical properties. Optical rotatory dispersion curves and optical rotations at 300 nm were determined with a Perkin-Elmer P22 spectropolarimeter. Rotations were determined by using the free acids of the tetraphosphate and of the pentaphosphates. The free acids were formed by treating the barium salts with a fivefold excess of Dowex AG-50W $(x8; H⁺)$ form; 100-200 mesh) for 4 hr at room temperature. Previous experience has shown that this technique vields a sample free from Ba^{2+} ions, by the sulfate test.

RESULTS

Ion-exchange dextran chromatography. The phytase activity of the culture filtrate was resolved into two major and possibly three minor peaks (Fig. 1). Peak A had ^a single activity maximum at pH 2.0, whereas peak E had activity maxima at both pH 2.0 and 5.5. Peak E was largely free of polysaccharide and inactive protein. The contents of tubes ¹ through 10 were combined, as were the contents of tubes 44 through 57. These two solutions were used as enzyme stock solutions for the subsequent experiments.

Ion-exchange chromatography. The phytase from peak A produced ^a single pentaphosphate peak (Fig. 2) and a single tetraphosphate peak (Fig. 3). The pentaphosphate corresponds to a 1,2,4,5,6-pentaphosphate (1), whereas the tetraphosphate corresponds to a 1,2,5,6-tetraphosphate (1). The phytase from peak E produced a mixture of two pentaphosphates at both pH 2.0 and 5.5 (Fig. 4 and 5). The major pentaphosphate component corresponds to a 1,2,4,5,6-pentaphosphate (1), and the minor pentaphosphate corresponds to a 1,2,3,4,5 pentaphosphate (1). The major tetraphosphate (Fig. 6) component corresponds to mvo -tetra II (1), a 1,2,5,6-tetraphosphate, and the minor component corresponds to myo-tetra ^I (1).

Optical properties. The optical rotatory dispersion curves of the pentaphosphates were plain and negative in the range 300 to 600 nm. The absolute magnitude of the rotation increased towards the shorter wavelength. The molecular rotations at ³⁰⁰ nm are recorded in Table 1.

DISCUSSION

By the use of ion-exchange chromatography (1), it is now possible to decide whether a $m\gamma$ inositol pentaphosphate is a 1,3,4,5,6-, a 1,- 2,4,5,6-, a 1,2,3,4,6-, or a 1,2,3,4,5-pentaphosphate. The myo-inositol pentaphosphates isolated in this study were identified by using a two-stage ion-exchange chromatographic procedure standardized against myo-inositol pentaphosphates isolated and identified in a previous investigation (1). The assignment of structures to these standard myo-inositol pentaphosphates is supported by a separate investigation (6) which involved phosphorus: myo inositol ratios, paper chromatography, and electrophoresis.

The assignment of sign of optical rotation to

FIG. 1. Ion-exchange dextran chromatography of A. ficuum NRRL 3135 culture filtrate. Symbols: \Box , phytase activity at pH 2.0; O, phytase activity at pH 5.5; \bullet , protein; \blacksquare , polysaccharide.

FIG. 2. Ion-exchange chromatography of the hydrolysate $(pH 2.0)$ of myo-inositol hexaphosphate by the phytase of peak A. Elution was by 0.48 N HCl, and fraction volume was 10 ml. Abbreviations: Pi, inorganic phosphate; DP, myo-inositol diphosphate; TrP, myo-inositol triphosphate; TP, myo-inositol tetraphosphate.

FIG. 3. Ion-exchange chromatography of the hydrolysate $(pH 2.0)$ of myo-inositol hexaphosphate by the phytase of peak A. Elution was by 0.28 N HCl, and fraction volume was 10 ml.

FIG. 4. Ion-exchange chromatography of the hydrolysate (pH 2.0) of myo-inositol hexaphosphate by the phytase of peak E. For conditions see Fig. 2.

the enantiomers of the myo -inositol-1,2,3,4,5pentaphosphates rests on the work of Tomlinson and Ballou on wheat-bran phytase (12). They deduced that the major $m\gamma_0$ -inositol pentaphosphate produced by wheat-bran phytase is $L-my$ o-inositol-1,2,3,4,5-pentaphosphate because its further enzymic dephosphorylation yielded $p\text{-}m\gamma o\text{-}inositol-1,2,6\text{-}triphosphate$ and also myo-inositol-1,2,3-triphosphate. Johnson and Tate (7) confirmed this conclusion and found that L -myo-inositol-1,2,3,4,5-pentaphosphate has a positive optical rotation at pH_1 and 589 nm. The assignment of sign of optical rotation to the enantiomers of the m y₀-inositol- $1,2,4,5,6$ -pentaphosphates rests on the work of Johnson and Tate on the phytase from a strain

FIG. 5. Ion-exchange chromatography of the hydrolysate (pH 5.5) of myo-inositol hexaphosphate by the phytase of peak E. For conditions see Fig. 2.

FIG. 6. Ion-exchange chromatography of the hydrolysate (pH 5.5) of myo-inositol hexaphosphate by the phytase of peak E. For conditions see Fig. 3.

of Neurospora crassa (7). The myo-inositol-1,2,4,5,6-pentaphosphate produced by this enzyme was identified as D-myo-inositol-1,- 2,4,5,6-pentaphosphate because its further enzymic dephosphorylation yielded D-myoinositol-1,2-diphosphate. D-myo-inositol-1,-2,4,5,6-pentaphosphate was found to have a negative optical rotation at pH ¹ and ⁵⁸⁹ nm. The sign of optical rotation was confirmed by Cosgrove (2) with $D-my_0$ -inositol-1,2,4,5,6pentaphosphate produced by a phytase from a bacterium of the genus Pseudomonas.

Without a knowledge of the molecular rotations of the pure enantiomers it is not possible to know whether the molecular rotations recorded in Table ¹ are due to mixtures of enan-

TABLE 1. Molecular rotations of the free acids of myo-inositol tetra- and pentaphosphates^a

Compound	Enzyme source	Concn (g/100) ml)	$[M]_{\rm iso}^{\rm ss}$
m yo-Inositol-1,2,5,6-te- traphosphate	Peak A, pH 2.0	0.383	-233°
m yo-Inositol-1,2,4,5,6- pentaphosphate	Peak A, pH 2.0	0.449	$-115°$
m yo-Inositol-1,2,4,5,6- pentaphosphate	Peak E, $pH 2.0$	0.390	$-147°$
myo -Inositol-1,2,4,5,6- pentaphosphate	Peak E, pH 5.5	0.400	-160°
m yo-Inositol-1,2,3,4,5- pentaphosphate	Peak E, pH 5.5	0.341	37°

^a Molecular rotations were determined at ³⁰⁰ nm and ²⁵ C by using aqueous solutions of the free acids. The pathlength was ¹ dm. The concentrations of the samples were calculated from determinations of total phosphate. Determinations were made at 300 nm because at the conventional 589 nm the measured rotations were negative but of the order of 10-' degrees. The use of ³⁰⁰ nm increased the measured rotation by a factor of up to 22 over that at 589 nm and left no doubt as to the sign of rotation.

tiomers or not. It is therefore only possible to identify the major component of a possible mixture of enantiomers. The sign of rotation (Table 1) and the chromatographic behavior (Fig. 2) of the pentaphosphate produced by peak A phytase identify it as D-myo-inositol-1,2,4,5,6-pentaphosphate (1), and, similarly, the tetraphosphate as D-myo-inositol-1,2,5,6 tetraphosphate (1). The signs of rotation (Table 1) and the chromatographic behavior (Fig. 4 and 5) of the pentaphosphates produced by peak E phytase at both pH 2.0 and 5.5 identify the major pentaphosphate component as D-myo-inositol-1,2,4,5,6-pentaphosphate (1) and the minor pentaphosphate component as $D-my_0$ -inositol-1,2,3,4,5-pentaphosphate (1). The structures of $p\text{-}myo\text{-}inositol-1,2,4,5,6-pen$ taphosphate and D-myo-inositol-1,2,3,4,5-pentaphosphate are illustrated in Fig. 7.

The extracellular phytase of A. ficuum NRRL ³¹³⁵ produces the same pentaphosphate as the major pentaphosphate component produced by Pseudomonas phytase (2). The pathway of hydrolysis of myo-inositol hexaphosphate by these enzymes from A. ficuum NRRL 3135 therefore resembles that of Pseudomonas phytase (2) and not that of wheatbran phytase (12).

The phytase activity of peak A is more stereospecific in its action than either Pseudomonas phytase or the phytase activity of peak E, since it produces a single pentaphosphate, whereas the other enzymes produce two. This conclusion is supported by the fact that peak A phytase also produces ^a single tetraphos-

FIG. 7. a, The structure of D-myo-inositol-l,- 2,4,5,6-pentaphosphate; b, the structure of D-myoinositol-1,2,3,4,5-pentaphosphate.

pentaphosphate

phate. Such stereospecificity was unexpected since Shieh, Wodzinski, and Ware (10) had reported that the nonspecific orthophosphoric monoester phosphohydrolase had a much wider substrate specificity than the pH 5.5 phytase.

The phytase activity of peak E closely parallels that of Pseudomonas phytase, since both produce the same two pentaphosphates. Previous work on Pseudomonas phytase (4) had shown that the minor pentaphosphate component produced by that enzyme was either Lmyo-inositol-1,2,3,4,5-pentaphosphate, its enantiomer or a mixture. Not enough of this minor component was available for determination of an optical rotation, but it is probable on the basis of the work reported in this paper that the minor pentaphosphate component of Pseudomonas phytase is also D-myo-inositol-1,2,3,4,5-pentaphosphate.

The action of the phytase of peak A at pH 2.0 and of the phytase of peak E at both pH 2.0 and 5.5 is consistent with the model of the active center of a bacterial (Pseudomonas sp.) phytase proposed by Irving and Cosgrove (4). In all cases, the phosphate group which is hydrolyzed to yield the pentaphosphate is a member of a vicinal *trans*-equatorial pair with the same stereochemistry as $(+)$ -trans-cyclohexane-1, 2-diol-diphosphate.

The major pentaphosphate component produced by wheat-bran phytase is L-myo-inositol-1,2,3,4,5-pentaphosphate (7). It is therefore now possible, using both the A. ficuum NRRL 3135 enzymes and the phytase from wheatbran, to prepare three of the four possible optically active forms of myo-inositol pentaphosphate.

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