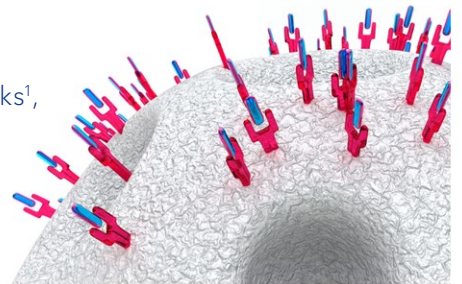


Purification of *p*-coumaric acid esterase by immobilized metal ion affinity chromatography

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SUMMARY

The *p*-coumaric acid esterase (*p*-CAE) has a wide range of applications and is therefore an enzyme of interest. *p*-CAE was produced by *Pichia pastoris* and purified by immobilized metal ion affinity chromatography (IMAC). Different IMAC systems were tested and compared. 35.4% of the total protein was identified as active *p*-CAE. A total activity loss of 21.4% was recorded. The *p*-CAE could be successfully purified.

INTRODUCTION

The structure of the plant cell wall consists of covalent cross-linked polysaccharides which provide stability [1]. Hydroxycinnamic acids are not freely present in plants but can be cleaved by substrate-specific enzymes from the complex structure. One common hydroxycinnamic acid is *p*-coumaric acid (*p*-CA) [2]. The *p*-coumaric acid esterase (*p*-CAE) is known for its very high *p*-CA activity. It hydrolyzes the ester linkages between the polysaccharides and various hydroxycinnamic acids [3]. The *p*-CAE is particularly

attractive for the industry due to its broad substrate specificity [4]. The *p*-CAE can be used in many ways in the food and pharmaceutical industries, as well as in the area of renewable resources, e. g. bioethanol production. A 6xHis-Tag was attached to the *p*-CAE which was recombinantly expressed in *Pichia pastoris* and purified by immobilized metal ion affinity chromatography (IMAC) promising better handling and a higher yield compared to the classical production.

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RESULTS

p-CAE was previously purified by IMAC with a three-step elution [4], resulting in our case in a broad elution. Thus, four different IMAC systems were further tested. Two different materials Ni-IDA and Ni-NTA and sodium phosphate buffer pH 8 and pH 6 were used. For qualitative evaluation, **Fig. 1** shows the chromatograms of the elution peaks and corresponding SDS-PAGE bands. The SDS-PAGE image shows the eluate and the enzyme treated with Endo H. All impurities were separated qualitatively equal. The recovery rate of the activity and the protein content for each system are shown in **Fig. 2**. The largest proportion of the total activity can be found in the eluate. The total protein concentration used is entirely recovered

in the pH 6 systems. Due to an additional buffering in the conditions favored for activity measurement, namely pH 8, a loss of the total protein concentration was detected. The four systems are similar in terms of recovery rates. For this reason, the system with the largest elution peak (IDA system pH 8) is selected as the optimal system. The results of this purification are summarized in **Tab. 1** and the chromatogram and SDS-PAGE of all fractions are shown in **Fig. 3**.

Tab. 1 Purification results of p-CAE by IMAC IDA pH 8 system

Protein content (%)	Total activity (%)
35.4 ± 0.1	78.6 ± 1.0

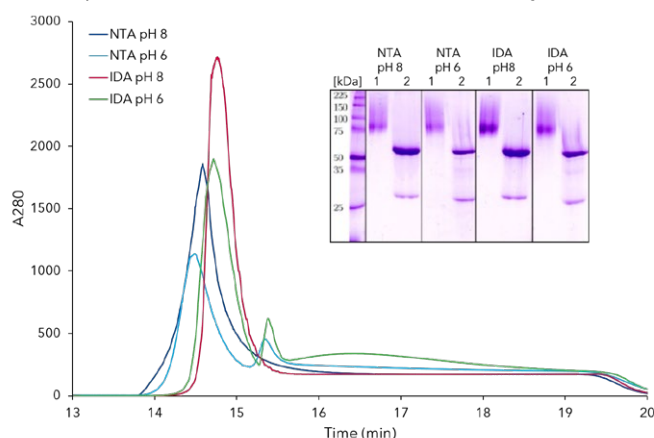


Fig. 1 Elution chromatograms at 280 nm of four IMAC systems and SDS-PAGE of eluate (1) and eluate treated with Endo H (2)

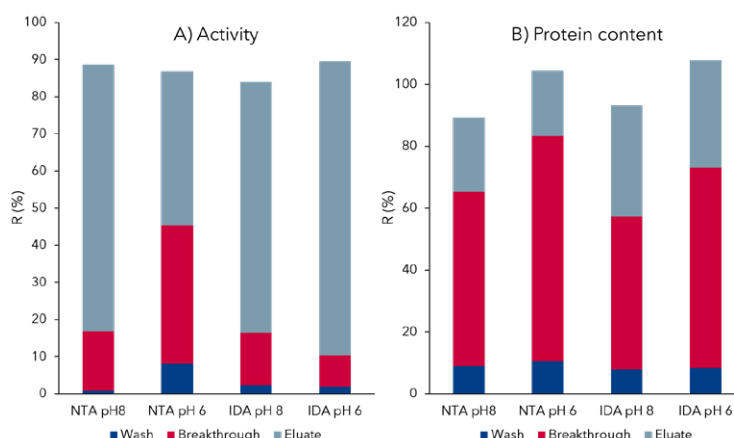


Fig. 2 Validation of four IMAC systems by recovery rate for enzyme activity and protein content

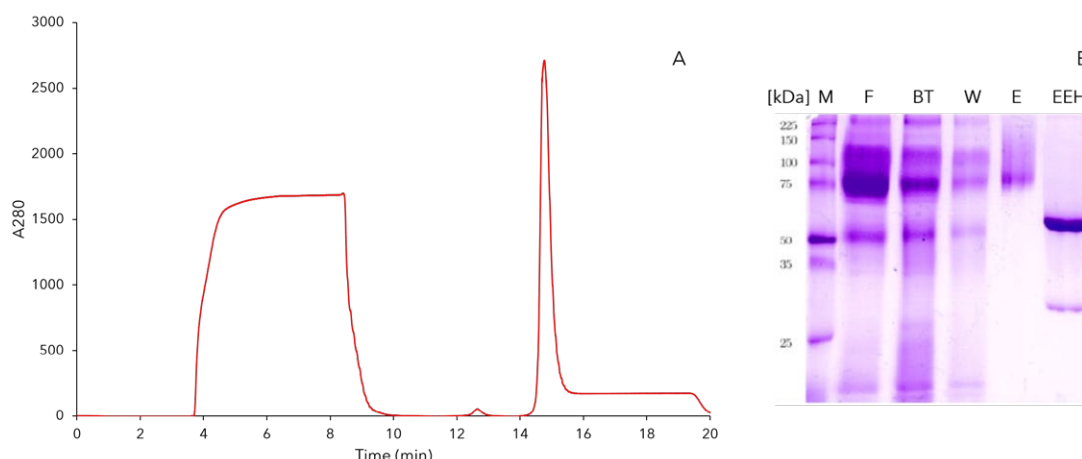
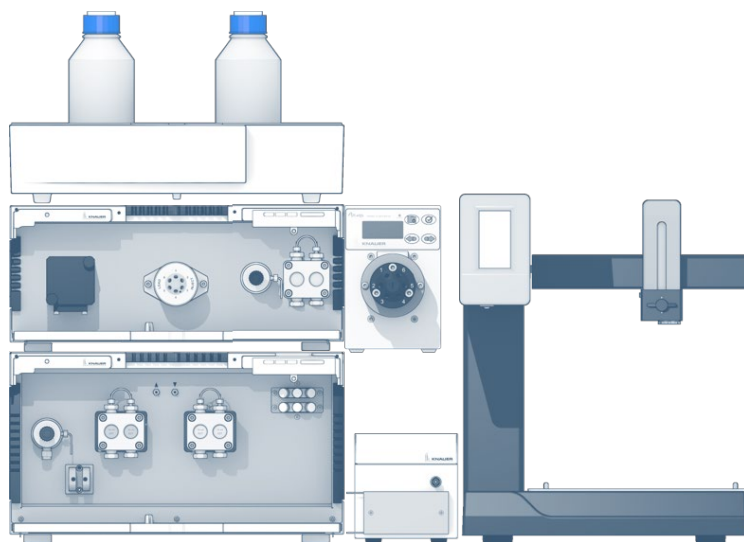


Fig. 3 Optimal IMAC system with IDA and pH 8 for purification of p-CAE; A) Chromatogram 280nm, B) SDS page (M - marker, F - filtrate, BT - breakthrough, W - wash, E - eluate, EEH - eluate treated with Endo H)

MATERIALS AND METHOD

The bioinert AZURA® system consisted of a 50 mL high pressure gradient pump, a single wavelength detector, an injection valve, a 50 mL sample pump and a fraction collector. Two different 1mL IMAC columns, IDA and NTA, columns were used for the purification after immobilization with nickel ions. In addition, a comparison of two different buffers (sodium phosphate buffer pH 6 and pH 8) was made. All measurements were performed in duplicate. Bradford Assay was used to determine the total protein concentration. The activity determination of the p-CAE was carried out using an optimized enzyme assay in combination with a reverse phase HPLC method [4].



CONCLUSION

For the purification of p-CAE, two column materials and two pH values were tested. Comparing the Ni-IDA with the Ni-NTA column, the absorption using IDA is higher and delayed. IDA can adsorb larger amounts of protein than NTA [5]. The IDA systems are in this case beneficial. Evaluation of two different pHs shows that, the target protein elutes as one sharp peak from the column when adjusting pH to 8. At pH 6, a second smaller elution peak can be seen, which was identified also as pure target protein. IMAC should be carried out at neutral to slightly basic pH values such as pH 8, to make sure the histidyl residues of imidazole are not protonated during adsorption [6]. The slightly acidic pH system and the NTA column material were not optimal for p-CAE purification. These results led to the final procedure where p-CAE was purified with Ni-IDA column at pH 8.

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ADDITIONAL MATERIALS AND METHODS

Tab. A2 Method parameters

Column temperature	RT
Injection volume	5 mL
Injection mode	Sample loop
Detection wavelength	UV 280 nm
Data rate	2 Hz

Tab. A3 Pump parameters (program 1)

Eluent A	20mM Sodium phosphate buffer pH6 + 5mM imidazole + 300 mM NaCl	
Eluent B	20mM Sodium phosphate buffer pH6 + 250mM imidazole + 300 mM NaCl	
Flow rate	1 mL/min	
Pump program [Vol in mL]	[%] A	[%] B
0	100	0
3	100	0
13	100	0
13.1	0	100
17	0	100
17.1	100	0
20	100	0

Tab. A4 Pump parameters (program 2)

Eluent A	50 mM Sodium phosphate buffer pH8 + 5 mM imidazole + 300 mM NaCl	
Eluent B	50 mM Sodium phosphate buffer pH8 + 250mM imidazole + 300 mM NaCl	
Flow rate	1 mL/min	
Pump program [Vol in mL]	[%] A	[%] B
0	100	0
3	100	0
13	100	0
13.1	0	100
17	0	100
17.1	100	0
20	100	0

Tab. A5 System configuration

Instrument	Description	Article No.
Pump	AZURA P 6.1L High Pressure Pump with 50 mL pump head, Ceramic, without Degasser	APH68FB
Assistant	AZURA ASM 2.1L Left: UVD2.1S single wavelength detector Middle: 6Port2Pos, 1/16", PEEK Right: P4.1L pump with 10 ml pump head	AYCAECBK
Conductivity monitor	AZURA CM 2.1S with flow cell - up to 100 ml/min	ADG30GD
Fraction collector	Foxy R1	A59100
Column switching valve	Bioinert Multifunction Selection Valve	AWB00FC
Column	NTA HisTrapTMFF Crude 1ml IDA HiTrapTM 1ml	
Software	PurityChrom5 Basic	A2650

RELATED KNAUER APPLICATIONS

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VBS0064 - Comparison of IgG purification by two different protein A media

VBS0065 - Separation of two model proteins with ion exchange chromatography

VBS0067 - Automated two-step purification of 6xHis-tagged GFP with AZURA Bio purification system

VBS0068 - Fast and robust purification of antibodies from human serum with a new monolithic protein A column

VBS0069 - Purification of Sulfhydryl Oxidase

VBS0070 - Ion Exchange Chromatography with AZURA® Bio purification system

VBS0071 - Comparison of two column sets for antibody purification in an automated two step purification process

VBS0072 - Separation of proteins with cation exchange chromatography on Sepapure SP and CM

VBS0073 - Separation of proteins with anion exchange chromatography on Sepapure Q and DEAE

VBS0074 - Comparison of ion exchange columns