

Microplate Assay to Study Carboxypeptidase A Inhibition in Andean Potatoes

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[Abstract] Metalloproteinases (MCP) are zinc-dependent exopeptidases that catalyze the hydrolysis of C-terminal amide bonds in proteins and peptides. They are involved in a wide range of physiological processes and have recently emerged as relevant drug targets in biomedicine (Arolas *et al.*, 2007). In this context, the study and discovery of new MCP inhibitors from plants constitute a valuable approach for the development of new therapeutic strategies. Herein we describe a simple and accessible microplate method for the study of the specific and dose-response carboxypeptidase A inhibitory activities present in Andean potato tubers. Our protocol combines an extraction method optimized for small protein inhibitors in plant tissues, with the measurement of enzyme kinetics using a microplate reader. These instruments are capable of reading small sample volumes, for many samples in a very short time-frame, therefore reducing the time and costs of high-throughput screening experiments. Although this protocol describes the study of Andean potatoes, our approach is also applicable to the analysis other plant samples.

Keywords: Metalloproteinase, Carboxypeptidase A, Inhibitor, Inhibitory activity, Microplate assay, Potatoes

[Background] In higher plants, small proteinaceous protease inhibitors are wound-induced molecules produced as a part of its defense system against insect attack (Graham *et al.*, 1981; Villanueva *et al.*, 1998). Among the studied inhibitors, only two are specific for MCP, *i.e.*, the potato carboxypeptidase inhibitor (PCI) and its close homolog found in tomato plants (TCI). Over the last few decades, the presence of MCP inhibitors in Solanaceae has been extensively reported, revealing potato (*Solanum tuberosum*) as one of the most important sources of MCP inhibitors (Hass *et al.*, 1979; Obregon *et al.*, 2012; Lufrano *et al.*, 2015). In humans, MCP action is exquisitely regulated and dysregulation of its function might lead to disease or even to cell death (Arolas *et al.*, 2007). In fact, MCP have been associated with human pathologies such as acute pancreatitis (Appelros *et al.*, 1998), diabetes (Cool *et al.*, 1997), several types of cancer (Ross *et al.*, 2009; Sun *et al.*, 2016; Abdelmagid *et al.*, 2008; Tsakiris

et al., 2008), fibrinolysis (Valnickova *et al.*, 2007), inflammation (Deiteren *et al.*, 2009) or neurodegeneration (Rogowski *et al.*, 2010). In this context, there is an interest in the discovery of new MCP inhibitors, and thus we focus our studies in potatoes that are native from the Andean region of South America. In this region, thousands of different potato varieties coexist, constituting a natural reservoir for the discovery of novel MCP inhibitors (Figure 1).

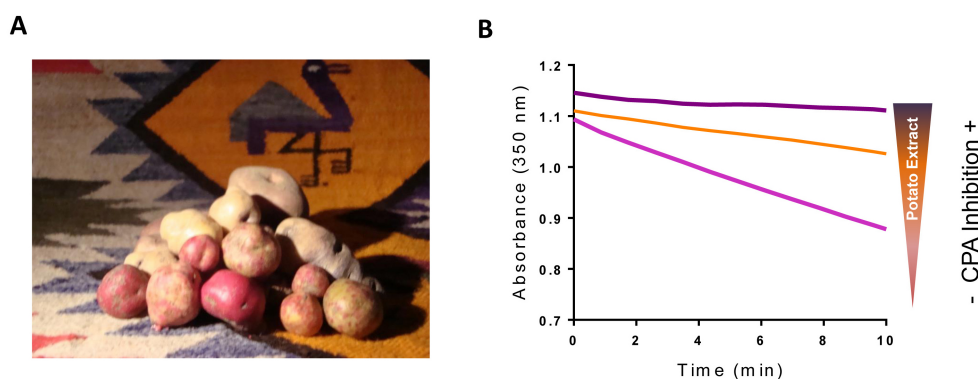


Figure 1. Andean potatoes and potato extract CPA inhibitory activity. A. Picture displaying the large number of potato varieties found within the Andean region. Currently in this region coexist thousands of Andean varieties of *Solanum tuberosum* (Machida-Hirano, 2015; Clausen *et al.*, 2010). B. Effects of potato extracts on bCPA activity. The activity of bovine CPA (bCPA) was measured using the substrate N-(4-methoxyphenylazoformyl)-Phe-OH determining the decrease in absorbance at 340 nm in function of the time. Due to its high content in MCP inhibitors, the addition of potato extract to the reaction decreases the rate of substrate hydrolysis in a dose-response manner.

Here, we describe a simple protocol to determine the specific and dose-response carboxypeptidase A inhibitory activity present in Andean potatoes and in other biological extracts using microplates. The major advantage of this protocol over other available approaches (Yanes *et al.*, 2007) is that the use of microplates allows multiple enzymatic measurements to be done in a single experiment, therefore reducing the time and costs.

Materials and Reagents

1. 50 ml tubes
2. 96-well microplates, clear flat bottom (Corning, catalog number: 3364)
3. Syringe filters, 0.45 μ m pore size (EMD Millipore, catalog number: SLHP033RS)
4. Potato tubers
5. General laboratory materials and instrumentation (e.g., micropipettes, microtubes, tips)
6. Bradford assay kit, e.g., Coomassie Plus Assay Kit (Thermo Fisher Scientific, Thermo Scientific™, catalog number: 23236)
7. Bovine serum albumin (BSA)

8. N-(4-methoxyphenylazoformyl)-Phe-OH·potassium salt (Bachem, catalog number: M-2245)
9. Trizma® base (Sigma-Aldrich, catalog number: T1503)
10. Sodium chloride (NaCl) (Sigma-Aldrich, catalog number: S7653)
11. Hydrochloric acid (HCl) (Sigma-Aldrich, catalog number: 258148)
12. Dimethyl sulfoxide (DMSO) (Sigma-Aldrich, catalog number: D4540)
13. Bovine carboxypeptidase A (bCPA) (Sigma-Aldrich, catalog number: C9268)
14. Carboxypeptidase A reaction buffer/Extraction buffer (see Recipes)
15. 2 mg/ml bCPA stock solution (see Recipes)
16. 10x bCPA working solution (see Recipes)
17. 1,000x substrate stock solution (see Recipes)
18. 10x substrate working solution (see Recipes)

Equipment

1. Laboratory blender or equivalent (Oster, catalog number: 004093-008-NP0)
2. Refrigerated centrifuge (suitable for volumes of 50 ml) (Beckman Coulter, model: Avanti J-26 XPI)
3. UV-Vis microplate spectrophotometer system capable of operating at 340 and 595 nm (*e.g.*, PerkinElmer, model: Victor X 2030-0050 or other equivalent spectrophotometer)
4. pH meter (HACH LANGE SPAIN, Crison, model: GLP 21)
5. 37 °C oven (*e.g.*, Thermo Fisher Scientific, Thermo Scientific™, model: Heratherm Compact Microbiological Incubator)
6. Multichannel pipette (*e.g.*, Technology Networks, model: CappAero Multichannel Pippete 25-200 µl)

Software

1. GraphPad Prism 5 software (GraphPad Software, Ing USA)

Procedure

A. Preparation of Andean potato extracts and protein quantification

1. Wash two mature potato tubers (stage five, according to Johnson, 2008) with distilled water. Peel, weigh (in our case 11.50 g) and dice them into small pieces of about 2 x 2 cm in size (see Figures 2A and 2B).

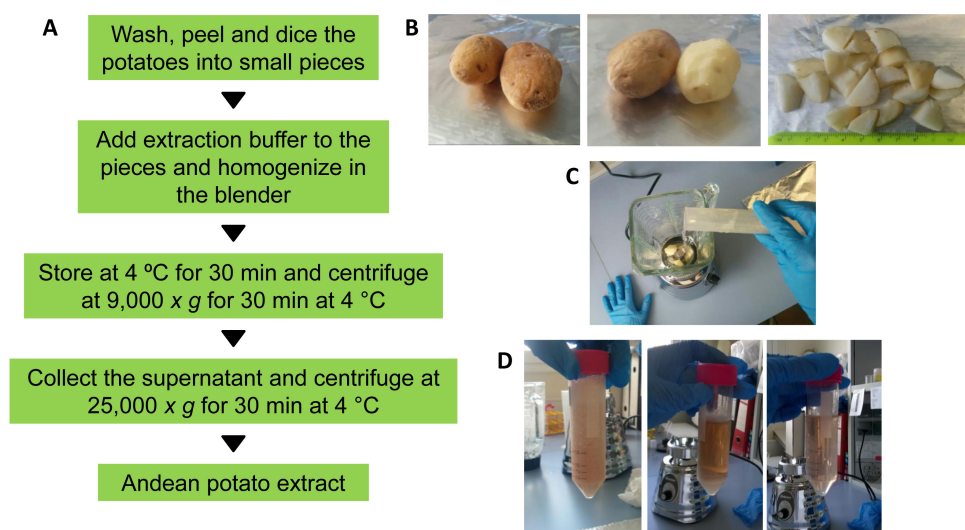


Figure 2. Preparation of potato extracts. A. Workflow followed to obtain the crude potato extract enriched in MCP inhibitors. B. Potato peeling and dicing. C. Mixing of the diced potatoes with the extraction buffer into the blender. D. The appearance of the potato extract before and after two centrifugation steps (from left to right).

- Mix the potato pieces with three volumes of ice-cold extraction buffer (in our case 34.50 ml; a mass/volume ratio of 1:3) into the cooled blender and homogenize at gentle speed for 5 min (Figures 2A and 2C).

Note: The homogenization should be performed in several sessions of 15-30 sec each, with 60 sec intervals between sessions to prevent excessive heating of the sample.

- After homogenization, transfer the potato homogenate to 50 ml tubes and centrifuge the sample at 9,000 x g for 30 min at 4 °C.
- Transfer the supernatant to a clean 50 ml centrifuge tube and spin at 25,000 x g for 30 min at 4 °C (Figures 2A and 2D). After the second centrifugation step, collect the resultant supernatant and filter through a 0.45 µm syringe filter to eliminate protein aggregates. The resultant potato extract can be stored at -20 °C until analyzed.
- Determine the protein concentration of the samples using the Coomassie Plus Assay Kit, according to the manufacturer's instructions. In brief, prepare a final volume of 500 µl of each of the six standard solutions containing 0, 2, 5, 10, 15 and 20 µg/ml of BSA and appropriate dilutions of the sample/s. Transfer 150 µl of each standard and potato extract samples into different wells of a 96-well microplate. BSA standards and potato extract samples should be assayed in triplicate. Then, add 150 µl of the Coomassie Plus reagent (see Materials and Reagents section) to each well and mix using the micropipette, by pipetting up and down carefully. After 5 min incubation at room temperature, measure the absorbance at 595 nm using a UV-Vis microplate spectrophotometer. Typically, we obtained 1-2 mg/ml of protein in the extracts.

B. Enzymatic assays

1. Determination of bCPA specific inhibitory activity

- a. Prepare triplicates of the reaction mixtures (Table 1) in a microplate without adding the substrate. Calculate the volume of potato extract to be added, in order to obtain 20% to 80% of bCPA inhibition. To obtain these inhibition levels, we typically add around 5-30 µg/ml of final protein concentration to the assay.

Table 1. Reaction mixture

	Control reaction	Reaction + potato extract
bCPA reaction buffer (µl)	200	200 - X
Enzyme working solution (µl)	25	25
Potato extract (µl)	–	X
Substrate (µl)	25	25
Final reaction volume (µl)	250	250

Where X is the volume (in µl) of potato extract to be assayed. The substrate should be added immediately before plate reading (see step B1c below).

- b. Cover the microplate with a lid and incubate at 37 °C for 15 min.
- c. Add 25 µl of substrate working solution to each well, mix carefully and thoroughly, by pipetting up and down carefully with a multichannel micropipette (A graphical demonstration of steps B1a, B1b and B1c is shown in Video 1).
Note: the homogenization should be performed in no longer than 60 sec, to avoid significant consumption of the substrate before absorbance monitoring.
- d. Perform absorbance measurements at 340 nm every 30 sec for 10 min.
- e. One unit of inhibitory activity is defined as the amount of inhibitor able to reduce one unit of bCPA activity, which in turn corresponds to the amount of enzyme that hydrolyzes 1.0 µmol of N-(4-methoxyphenylazoformyl)-Phe-OH per min at 25 °C. Consequently, Equation 1 can be used to calculate the Specific Inhibitory Activity (SIA) found in potato extracts.

Equation 1:

$$IA (U/ml) = \left(\left(\frac{\Delta Abs(340nm)}{\Delta t(min)} \right)_{Control} - \left(\frac{\Delta Abs(340nm)}{\Delta t(min)} \right)_{Potato\ ext} \right) \frac{1}{\xi} \frac{V_{total}}{V_{Potato\ ext}} \frac{D}{L}$$

Where,

IA is the Inhibitory Activity in U/ml,

ΔAbs/Δt is the absorbance variation per unit of time (in min) during the reaction in absence (control), and in presence of the potato extract (Potato ext), respectively,

ξ is the extinction coefficient for the substrate N-(4-methoxyphenylazoformyl)-Phe-OH (ξ 350 nm = 19 [µmoles/ml]⁻¹ cm⁻¹),

V_{total} is the assay volume and V_{Potato ext} is the volume of extract added to the reaction,

D and L are the dilution factor for the extract and the path length (in cm), respectively. Typically, the path length in a microplate for a volume of 250 µl is 0.7 cm. However, for different reaction volumes, or for a more accurate calculation, check your 96-well microplate manufacturer instructions.

Then, calculate directly the SIA by dividing the resultant IA value by the protein concentration of the sample in mg/ml (see Equation 2 and examples in Table 2).

Equation 2:

$$SIA = \frac{IA}{[Protein]_{Potato Ext}}$$

2. Determination of the IC50: Dose–Response curve assay (see Figure 3)

- a. Prepare the same control reaction as described in step B1a and prepare at least 12 additional reaction mixtures with different final concentrations of the potato extract in the assay, ranging from 0 to 300 µg/ml, (or even with higher concentration to obtain the complete inhibition of bCPA activity).

Note: We typically assay a total of 15 different extract final concentrations, containing 0, 1, 2, 3, 5, 10, 15, 20, 30, 75, 100, 150, 200, 250 and 300 µg/ml of protein.

Fit the results obtained to the following Equation 3 and determine the IC50 value.

Equation 3:

$$Y = \frac{100}{1 + 10^{(X - \log IC_{50})}}$$

Where,

X is the log-transformed protein concentration assayed,

Y is the normalized bCPA activity (relative to the control condition and expressed as a percentage of the maximal activity).

Y values can be calculated for each extract concentration, using the following Equation 4.

Equation 4:

$$Y = \frac{\left(\frac{\Delta Abs_{340nm}}{\Delta t(\min)}\right)_{Potato ext}}{\left(\frac{\Delta Abs_{340nm}}{\Delta t(\min)}\right)_{Control}} \times 100$$

The IC50 value is the extract concentration necessary to reach a 50% of bCPA inhibition (Copeland, 2005). See representative examples of dose-response bCPA inhibitory plots in Figure 3 and the corresponding IC50 values in Table 2.

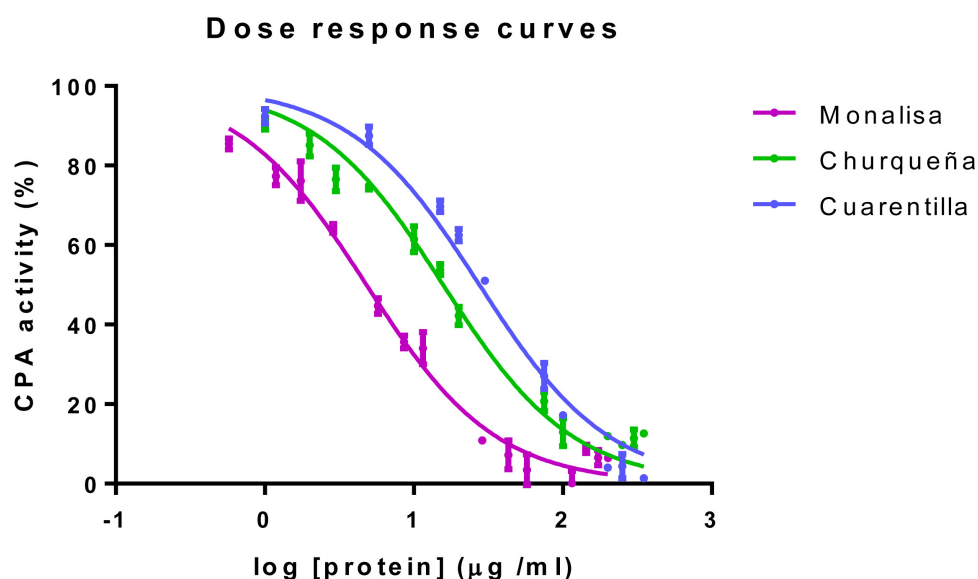


Figure 3. Examples of dose-response inhibitory plots. Representative examples of dose-response inhibitory curves determined for three different varieties of potatoes; *Solanum tuberosum* subsp. *tuberosum* var. monalisa (Monalisa, magenta solid line), *Solanum tuberosum* subsp. *andigenum* var. churqueña (Churqueña, green solid line), *Solanum tuberosum* subsp. *andigenum* var. cuarentilla; (Cuarentilla, blue solid line).

Table 2. Summary of the inhibitory activities found in different varieties of potatoes

Potato specie	Protein concentration (mg/ml)	Inhibitory activity (mU/ml)	Specific inhibitory activity (mU/mg)	IC50 (µg/ml)
A	1.56 ± 0.032	3.59 ± 0.06	2.30 ± 0.021	4.81 ± 0.3
B	2.71 ± 0.033	2.07 ± 0.11	0.76 ± 0.012	15.73 ± 0.9
C	0.79 ± 0.014	0.56 ± 0.06	0.71 ± 0.018	27.54 ± 1.8

A: *Solanum tuberosum* subsp. *tuberosum* var. monalisa; B: *Solanum tuberosum* subsp. *andigenum* var. churqueña; C: *Solanum tuberosum* subsp. *andigenum* var. cuarentilla

The two protocols for the determination of SIA and IC50 are summarized in the scheme of Figure 4 and Video 1.

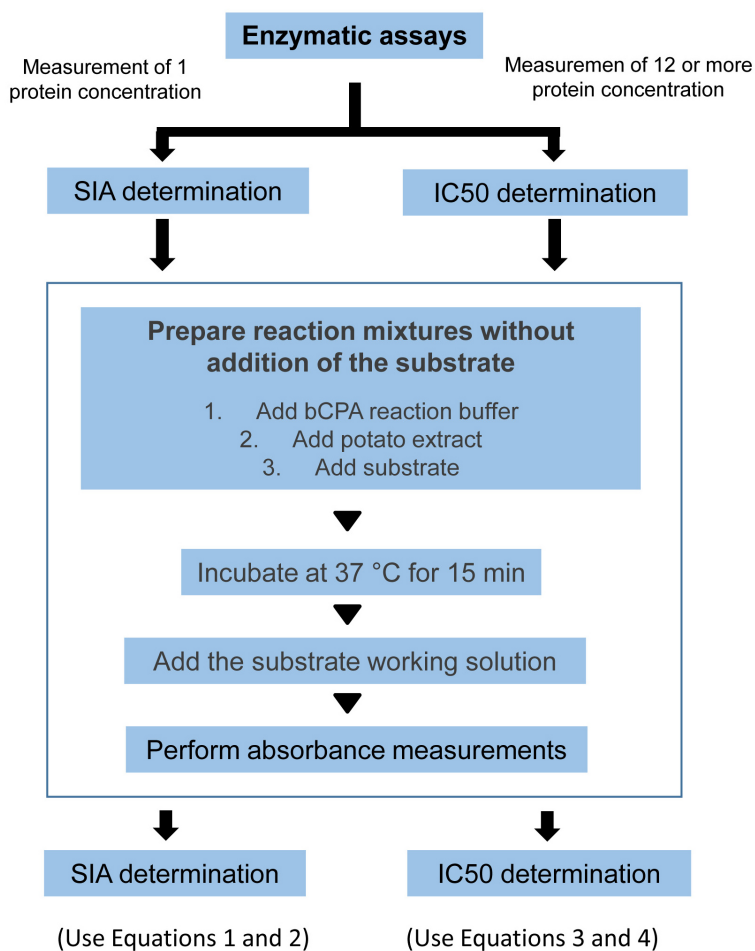


Figure 4. Enzymatic assays. General workflow for the measurement of inhibitory activity and determination of the SIA and IC50.

Video 1. Video demonstration of the protocol to perform the enzymatic measurements



Data analysis

Data fittings and IC50 determination were performed using GraphPad Prism 5 software (GraphPad Software, Ing USA) (Motulsky *et. al.*, 2007). Only fittings with a $R^2 > 0.97$ were considered.

Notes

1. Freshly collected samples display higher inhibitory activities; however, frozen samples can also be used.
2. The protocol described here is suitable for any scale of sample preparation, therefore volumes and reagent quantities should be scaled proportionally.
3. Here we used a laboratory blender that allows a complete sample blending and homogenization. Alternatively, other appropriate homogenization systems can be used.
4. The soluble fraction resultant from the step A4 is a crude extract rich in MCP inhibitors. Other proteinaceous and chemical compounds soluble in the homogenization buffer can be also present in the sample. For a higher MCP inhibitor enrichment (*e.g.*, for K_i determination), further purification procedures could be addressed (Pearce and Ryan, 1983).
5. We used a commercial Bradford assay kit; however, non-commercial Bradford reagents or other different quantification assays can be used (Bradford, 1976; He, 2011a and 2011b).
6. The microplate reactions can alternatively be performed in conventional spectrophotometer cuvettes by scaling up the reagents proportionately.
7. Mix the enzyme suspension thoroughly to ensure complete mixing. It is strongly recommended to follow manufacturer's instructions for an accurate enzyme preparation.
8. The substrate concentration used is such that the enzyme is at the maximum velocity. The enzyme concentration might be adjusted to consume less than 5-10% of substrate.

Recipes

1. Carboxypeptidase A reaction buffer/Extraction buffer (1 L, 20 mM Tris-HCl, 500 mM NaCl, pH 7.5)
 - 2.42 g of Trizma base
 - 29.22 g of NaCl
 - MilliQ water up to 900 ml
 - Adjust solution to pH 7.5 by addition of 5 N HCl
 - Adjust the final volume with MilliQ water to 1,000 ml
 - Store at 4 °C up to two weeks
2. 2 mg/ml bCPA stock solution
 - Prepare 1 ml of a 2 mg/ml bCPA solution (~57 μ M) in reaction buffer
 - Note: We typically dilute 100 μ l of commercial bCPA with 900 μ l of carboxypeptidase A reaction*

buffer. Note that different batches of bCPA may have different enzyme concentrations, therefore the dilutions should be adjusted accordingly.

Divide into 20 µl aliquots

For short-term storage, store at 4 °C; for long-term storage, store at -20 °C

3. 10x bCPA working solution

Prepare 10 ml of a 50 nM enzyme solution by diluting 8.77 µl of bCPA stock solution in 9.991 ml of carboxypeptidase A reaction buffer

For short-term storage, store at 4 °C; for long-term storage, store at -20 °C

4. 1,000x substrate stock solution (100 mM N-[4-methoxyphenylazoformyl]-Phe-OH)

Dissolve 100 mg of N-(4-methoxyphenylazoformyl)-Phe-OH·potassium salt (Mock *et al.*, 1996) in 2.74 ml of DMSO

Divide into 100 µl aliquots and store at -20 °C

5. 10x substrate working solution, 1 mM (N-[4-methoxyphenylazoformyl]-Phe-OH)

Dilute 100 µl of the substrate stock solution to a final volume of 10 ml with carboxypeptidase A reaction buffer

Store at -20 °C

Acknowledgments

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