Human plasma inositol hexakisphosphate (InsP⁶) phosphatase identified as the Multiple Inositol Polyphosphate Phosphatase 1 (MINPP1)

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Abstract

Inositol hexakisphosphate (InsP₆), also known as phytic acid, is a potent chelator of bivalent cations. Intracellular InsP₆ molecules are associated with magnesium. Calcium is the prevalent bivalent cation outside the cell and its association with Ins P_6 could lead to the formation of insoluble complexes. To avoid the formation of dangerous Ins P_6 /Calcium precipitates in the bloodstream, mammals must possess a robust $InsP_6$ phosphatase in their plasma. Here we identify the Multiple Inositol Polyphosphate Phosphatase 1 $(MINPP1)$ as the InsP₆ phosphatase present in human plasma.

Figure 1. Identification of Minpp1 as the plasma InsP⁶ phosphatase preventing InsP⁶ precipitation.:

A) The indicated amount of $InsP_6$ was added to 1 ml of Ringer's solution (147 mM NaCl, 4 mM KCl, 3 mM CaCl₂) Krebs-Ringer's solution (120 mM NaCl, 5 mM KCl, 2 mM CaCl2, 1 mM MgCl2, 25 mM NaHCO3) and control salt solution NaCl KCl (147 mM NaCl, 4 mM KCl) and incubated at 37^oC for 30 minutes with rotation. InsP₆-Ca precipitated were observed by phase contrast microscopy. B) 5 nmol of InsP₆ was added to 1 ml of human plasma and incubated at 37°C for the indicated time before acidification and extraction of inositol phosphates with the TiO₂ procedure (Wilson & Saiardi, 2018). The extracts were loaded on 33% polyacrylamide gel and stained with toluidine blue (Lonetti et al., 2011). To orientate the gel polyP100 (50 nmol [Pi]) was loaded on the left and $InsP_6$ (2 nmol) was used as migration standard. OrangeG (OG) dye was used to monitor gel electrophoresis. C) Albumin and immunoglobulin-depleted male and female plasma (50 mg) were used to perform Western blot with anti Minpp1 antibody. HEK293T and isogenic minpp1-/- extracts (10 mg) (Ucuncu et al., 2020) were used as positive and negative controls respectively. Shown are the representative results of at least three biological repeats.

Description

Inositol phosphates (InsPs) represent a diverse and important class of intracellular signalling molecules (for review see (Kim et al., 2024)). The fully phosphorylated inositol ring of inositol hexakisphosphate (Ins P_6) represents the most abundant Ins Ps present in mammalian cells, with intracellular concentrations ranging from 20 to 100 µM (Qiu et al., 2020; Shears, 2001). InsP⁶ not only acts as a structural component of proteins such as the HIV capsid (Mallery et al., 2019) and the RNA-editing deaminase ADAR2 (Macbeth et al., 2005), but also regulates signalling pathways such as necroptosis by activating mixed lineage kinase domain-like [\(MLKL](https://www.alliancegenome.org/gene/HGNC:26617)) protein (Dovey et al., 2018) and protein phosphorylation by activating casein kinase 2 (CK2) (Solyakov et al., 2004). Additionally, InsP₆ is the main precursor of inositol pyrophosphates such as InsP₇ and InsP₈, which are important signalling molecules themselves (Nguyen Trung, et al., 2022; Wilson et al., 2013). The charged nature of InsPs prevents their diffusion across the plasma membrane, and therefore, InsP's metabolic and signalling networks are exclusively intracellular with many InsPs-kinases localized within the cytosol or into the nucleus (Otto et al., 2007; Shears, 2004). We could envisage, however, that cell death mechanisms and subsequent cell lysis could lead to the release of intracellular InsPs to our bloodstream. What is the fate of this theoretical extracellular pool of InsPs? Here we focus our attention on InsP₆.

Inositol hexakisphosphate possesses a unique charge density, with its twelve negative charges constrained around the carbon backbone of inositol. The biophysical properties of $InsP₆$ and its association with bivalent cations have been intensively studied (Hager et al., 2016; Kurz et al., 2023). Seminal thermal analysis and solubility measurements studies have determined that in the cytosolic environment, in which magnesium is the prevalent bivalent cation (Romani, 2011), Ins P_6 could exist as a soluble penta-magnesium complex up to a concentration of 49 μ M (Torres et al., 2005; Veiga et al., 2006). To verify InsP₆ solubility in conditions mimicking the salt composition of the human plasma, we took advantage of two intravenous fluids used to treat dehydration: the glucose-depleted Ringer's solution $(147 \text{ mM NaCl}, 4 \text{ mM KCl}, 3 \text{ mM CaCl}_2)$ and the glucosedepleted Krebs-Ringer's solution (120 mM NaCl, 5 mM KCl, 2 mM CaCl₂, 1 mM MgCl₂, 25 mM NaHCO₃) and as a control the same mix of sodium and potassium salts (147 mM NaCl, 4 mM KCl) omitting bivalent cations. Using phase contrast microscopy, we observed the formation of insoluble precipitates when physiological levels of InsP $_6$ (10 μ M and 100 μ M) were added to Ringer's or Krebs-Ringer's solution, in a calcium-dependent manner (Figure 1A). Hence, these precipitates must be $InsP₆/Calcium complexes. This qualitative visual result is consistent with the quantitative measurements reporting the$ insolubility of Ins P_6 in solutions containing calcium (Veiga et al., 2006), which is the most prevalent bivalent cation present in plasma (Allgrove, 2009). The prompt precipitation of InsP₆/Calcium complexes in plasma-mimicking solutions suggests that mammals must have evolved mechanisms to cleanse ${\rm InsP_6}$ from plasma, to avoid the formation of harmful precipitates in their circulatory system. Indeed, we previously demonstrated the presence of a robust $InsP_6$ dephosphorylation activity in mammalian plasma (Irvine et al., 2015; Wilson et al., 2015). Here we repeated the Ins P_6 dephosphorylation assay using human plasma samples from both male and female donors. The incubation of InsP₆ (5 μ M) in plasma from both genders at 37^oC resulted in InsP₆ dephosphorylation over time (Figure 1B), thus confirming previous findings (Irvine et al., 2015; Wilson et al., 2015).

The human genome carries a single gene encoding a phosphatase active towards InsP_6 , namely Multiple Inositol Polyphosphate Phosphatase 1 [\(MINPP1\)](https://www.alliancegenome.org/gene/HGNC:7102). This enzyme belongs to a conserved family of histidine acid phosphatases (IPR016274), commonly referred to as phytases because acting on phytic acid another name for $InsP_6$. The thoughtful characterization of [MINPP1](https://www.alliancegenome.org/gene/HGNC:7102) enzymatic activities *in vitro* and *in vivo* revealed that this enzyme dephosphorylates InsP⁶ primarily to Ins(1,2)P₂ (Nguyen Trung, Kieninger, et al., 2022). By regulating InsP₆ metabolism, Minnp1 controls numerous cellular processes; additionally, recent Mendelian genetic studies have revealed the importance of [MINPP1](https://www.alliancegenome.org/gene/HGNC:7102) in the pathophysiology of a specific form of [pontocerebellar](https://www.alliancegenome.org/disease/DOID:0060264) hypoplasia (PCH) (Appelhof et al., 2021; Ucuncu et al., 2020), a severe

neurodegenerative disorder. To verify whether $MINPP1$ is present in human plasma to account for the observed InsP₆ phosphatase activity (Figure 1B), we utilized an immunochemical assay. To apply this approach to plasma, it is crucial to remove the albumin and immunoglobulins (mainly IgG), which represent about 60% and 20% of the proteome in the plasma, respectively and could interfere with immunoblotting. Using a commercially available albumin and IgG depletion kit we enriched plasma proteome. The western blot performed on the eluate enriched for non-albumin and non-IgG proteins employing anti $MINPP1$ antibody demonstrates the presence of this critical InsP₆ phosphatase in human plasma (Figure 1C). Our direct analysis confirms a mass spectrometry (MS) based study identifying Minpp1 in human plasma (Farrah et al., 2011). Different sample preparation approaches, MS techniques, and algorithms used to extract MS data could lead to the identification of different sets of proteins from human plasma. In fact, a recent meta-analysis aimed at generating a reference set of plasma proteome to be used for targeted MS does not include **[MINPP1](https://www.alliancegenome.org/gene/HGNC:7102)** (Kliuchnikova et al., 2023). Nevertheless, [MINPP1](https://www.alliancegenome.org/gene/HGNC:7102) is one of the 4072 plasma proteins listed by the Human Protein Atlas [\(https://www.proteinatlas.org](about:blank)), here Minpp1 presence is recorded as non-validated by blood-based immunoassay. Our study unequivocally provides this important evidence.

Our confirmation of [MINPP1](https://www.alliancegenome.org/gene/HGNC:7102) in human plasma should put to rest the debate on the presence of InsP₆ in plasma (Irvine, 2014; Irvine et al., 2015) that few authors have been able to detect using obviously unreliable analytical methods. The presence of Minpp1 in plasma deemed it unlikely for plasma to contain InsP_6 . Our finding also prompts a revaluation of the literature that suggest a direct health-beneficial role of orally administrated Ins P_6 . Any positive dietary benefit of Ins P_6 is likely to have derived from its catabolism to propionate by gut bacteria (De Vos et al., 2024) or from its dephosphorylation to inositol in the gut, which is subsequently absorbed by the intestine.

[MINPP1](https://www.alliancegenome.org/gene/HGNC:7102) is predominantly localized inside the endoplasmic reticulum (ER) since it possesses an ER retention signal (KDEL) (Kilaparty et al., 2014; Yu et al., 2023). Secretory vesicles may emerge from ER-Golgi vesicular trafficking pathways which ultimately facilitate the release of **[MINPP1](https://www.alliancegenome.org/gene/HGNC:7102)** into the plasma. Since secretion of KDEL protein has been reported (Palazzo et al., 2022), further studies aimed at characterizing the secretory mechanisms of [MINPP1](https://www.alliancegenome.org/gene/HGNC:7102) should be performed to fully appreciate the physiological functions of this important InsP_6 phosphatase outside the cell.

The demonstration of extracellular [MINPP1](https://www.alliancegenome.org/gene/HGNC:7102) opens new perspectives to interpret the role of this InsP₆ phosphatase might play in disease conditions (Appelhof et al., 2021; Ucuncu et al., 2020). In the absence of **[MINPP1](https://www.alliancegenome.org/gene/HGNC:7102)**, neural cell death could be associated with a release of InsP₆, leading to InsP₆/Calcium precipitates in the extracellular space, potentially contributing to pathogenicity. Pathology mechanism could result from a "vicious circle" combining both intra and extracellular deleterious consequences on neural cell differentiation and survival during brain development. Our result raises the possibility of [MINPP1](https://www.alliancegenome.org/gene/HGNC:7102)'s presence in the cerebrospinal fluid (CSF). This is a tantalising prospect that warrant the urgent need for further studies in the physiology of **MINPP1**.

Methods

InsP₆ solubility study.

To study the solubility of InsP₆, we used three different solutions: glucose-depleted Ringer's solution (147 mM NaCl, 4 mM KCl, 3 mM CaCl₂), glucose-depleted Krebs-Ringer's solution (120 mM NaCl, 5 mM KCl, 2 mM CaCl₂, 1 mM MgCl₂, 25 mM NaHCO₃), and a control solution identical to the Ringer's solution but without calcium (147 mM NaCl, 4 mM KCl). InsP₆ was added to each solution at final concentrations of 10 μ M and 100 μ M. The mixtures were incubated with rotation at 37°C for 30 minutes. Following incubation, samples were examined under a phase contrast microscope (Olympus CX41) to detect the presence of insoluble $\mathrm{InsP}_6\text{-}\mathrm{Ca}$ precipitates.

$InsP₆$ dephosphorylation assay.

Male and Female human plasma were bought from TCS Biosciences (Cat: PR200-F-100-H2). Each 1 mL plasma sample was $supplemented$ with a HEPES-MgCl₂ solution to achieve final concentrations of 2 mM HEPES and 1 mM MgCl₂. InsP₆ (5 µM) (Calbiochem, Sigma-Aldrich, Cat: 407125) was added, and reactions were incubated at 37°C for 20, 60, and 120 minutes. Reactions were stopped by adding 20 µL of a stop solution (100 mM EDTA; 100 mM EGTA). Following incubation, inositol phosphates were purified using the TiO2 method as previously described (Wilson & Saiardi, 2018) and analysed by polyacrylamide gel electrophoresis PAGE followed by toluidine blue staining (Losito et al., 2009).

Albumin/IgG depletion.

Albumin and IgG were depleted from plasma using the ProteoExtract® Albumin-IgG Removal Kit MAXI (Calbiochem, Sigma-Aldrich, Cat: 122643). Plasma samples were diluted 1:10 in 10X Binding Buffer, and columns were equilibrated with

1X Binding Buffer. The diluted samples were passed through the column, albumin and IgG depleted eluate was collected by washing the column with 2M salt solution as for manufacturer's instructions.

Western Blot Assay.

Plasma samples depleted of IgG and albumin were concentrated and desalted using Amicon® Ultra-0.5 Centrifugal Filter Devices (Millipore, Cat.: UFC501096) according to the manufacturer's protocol. The eluted and concentrated fractions were quantified using the DC™ Protein Assay (Bio-Rad, Reagent A, Cat.: #5000113; Reagent B, Cat.: #5000114; Reagent S, Cat.: #500-0115). Plasma proteins (50 µg) were resolved by electrophoresis using 4-12% Bis-Tris polyacrylamide gels (NuPAGE™, Invitrogen, REF. NP0321BOX) and transferred to nitrocellulose membranes (GE Healthcare Life Sciences Whatman™, Cat.: 10401396). Membranes were blocked in 5% non-fat milk in TBS-T (0.1%) and incubated overnight at 4°C with [MINPP1](https://www.alliancegenome.org/gene/HGNC:7102) primary antibody (Santa Cruz, Cat: SC-514214). After three washes in TBS-T (0.1%), membranes were incubated with a secondary anti-mouse IgG1 antibody (Invitrogen, Cat: PA-74421) for 1 hour at room temperature. Detection was performed using the Clarity™ Western ECL substrate (Bio-Rad, Cat.: #170-5060) and images were acquired with the Alliance Q9 imaging system.

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