Identification and Localization of Gold Nanoparticles in Potassium Ion Pores: Implications for K<sub>ir</sub> Blockade

Chur Chin · Yu Shin Park

ABSTRACT

Introduction: In our previous study, we found that negatively charged gold nanoparticles with spermidine have the potential of blocking inwardly rectifying potassium channels (K<sub>ir</sub>), both at the cellular and the tissue level.

Methods: For the purpose of the present study, we purified the cytoplasmic domain of the K<sub>ir</sub> 3.1 receptor from Escherichia coli. Using single particles with surface coating by transmission electron microscopy, we identified the gold nanoparticles at the cytoplasmic side of the human K<sub>ir</sub> channel.

Results: Energy-dispersive X-ray spectroscopy showed the presence of the gold deposits in the cytoplasmic domain of the K<sub>ir</sub> receptor.

Conclusion: In conclusion, we could identify undecagold in the ion pore of the K<sub>ir</sub>3.1 channel in order to clarify its direct blocking effect in the K<sub>ir</sub> ion pore by undecagold.

Keywords: CryoEM; Cytoplasmic domain; K<sub>ir</sub>3.1; STEM

INTRODUCTION

Inwardly rectifying potassium channels (K<sub>ir</sub>) allow greater potassium ion (K<sup>+</sup>) flow into the cell than in the outward direction [1]. Nanoparticle penetration into the cells is an important process in drug delivery [2]. The penetration efficiency of nanoparticles may reach approximately 75%, especially for those with anisotropic shapes or asymmetric surface decoration [3]. Nontoxic vitamin complexes such choline and spermidine have been used as composites containing nanoparticles [4]. The identification of new types of molecules targeting ion channels is of significant interest in biological research. Therefore, the aim of our
work was to explore the mechanistic evidence of using novel materials such as undecagold, as ion channel blockers.

Gold nanoparticles can block the function of a channel by fitting into its pore and preventing further conformational change [5]. Kir channels are characterized by an extensive pore encompassing continuous transmembrane and cytosolic portions [6]. The key elements in G-loop gating transition involve the movement of the N-terminus and C-linker that removed constraining intermolecular interactions. Negatively charged gold nanoparticles (0.8 nm) with spermidine are able to enter the cell and block the human Kir channel at its intracellular face [7]. The G loops are located towards the cytoplasmic site, playing a role in G-loop disrupted gating or inward rectification. The G loop is the narrowest portion of the cytoplasmic domain in the Kir3.1 ion pore. This ion pore which has a diameter of approximately 8 nm has never been permeated by negatively charged undeca-gold [8]. The cytoplasmic ion-permeation pathways are occluded by four cytoplasmic loops that form a girdle around the central pore (G-loop). Kir channels, similar to other K+ channels, have a tetrameric architecture, but, unlike other channels, have a long ion permeation pathway that consists of both a transmembrane and a cytosolic portion. The gating of Kir channels has been shown to involve the binding of divalent cations and polyamines to its cytoplasmic domain [9]. G-protein-gated K+ channels (Kir3.1–Kir3.4) control electrical excitability, regulating the heart rate by opening a G-loop gate in the cytoplasmic domain [10]. G loop crystal structures revealed two possible conformations of the cytoplasmic domains, one with a constricted cytosolic gate and another with a dilated G-loop gate. In cryo-electron microscopy (EM) images, undecagold-spermidine complex and negative charged gold nanoparticles (0.8 nm) bind ionically to two NH2+ and one NH+ portions of the spermidine molecules. Spermidine (1.2 nm) is a suitable candidate for introducing gold nanoparticles into the cytoplasmic vestibule (3 nm diameter) of the receptors. Spermidine passes through the pore domain of the Kir channel with the incapacity for blocking other channels.

Kir3.1 normally functions as a heteromultimer with other members of the Kir3 family, whereas the chimera are homomultimer. The chimera used in this study might lack proper coupling between the cytoplasmic and transmembrane pores. However, single channel activity has yet to be demonstrated for the prokaryotic Kir channels. We have purified the cytoplasmic domain of the Kir3.1 receptor from E. coli (Figs. 1, 2, 3.).

Fig. 1 Control (Kir receptor cytoplasmic domain): a purified protein of the cytoplasmic domain of the Kir channel have collapsed lumens. Kir inward-rectifier potassium ion channel
MATERIALS AND METHODS

The 0.8 nm negatively charged gold nanoparticles were purchased from Nanoprobes, Inc (New York, NY, USA). The ionic bond was induced by vortexing two samples with different concentrations of positively charged spermidine (i.e., 10 mM and 100 nM), 1 mg of meso-2.3-dimercaptosuccinic acid (DMSA) powder, and 50 nM of negatively charged gold nanoparticles for approximately 15 min followed by mixing with a pipette for 15 min. The negatively charged CoO– particles were individually coated with a thin film of three nanoparticle molecules with DMSA, each bound to one NH₂ and two NH₁ spermidine molecules (Supplementary video clip). The spermidine hydroxide solution and the DMSA were obtained from Sigma Aldrich (St. Louis, MO, USA). Undecagold has a core of 11 gold atoms only 0.8 nm in diameter and is ideal for ultra-high-resolution EM work such as cryo-EM, scanning transmission EM. In this study, we present the structure, obtained by single particle analysis with cryoEM of the cytoplasmic Kᵢᵣ domain containing undeca-gold inside the complex (Supplementary Figs. 1, 2). Undeca-gold binding occurs at the outer end of the G-loop [11]. Purified protein solutions were diluted to 10 μg mL⁻¹ in a buffer solution containing 20 Mm Tris–HCl (pH 8.0), 100 mM NaCl, and the corresponding detergent. A diluted protein solution of the cytoplasmic domain of the Kᵢᵣ channel was added at a volume ratio of 1:10 to reach a concentration of

Fig. 2 Two dimensional TEM images show the tetramer of Kᵢᵣ3.1 cytoplasmic domain with collapsed lumen of the ion pore. TEM transmission electron microscopy, Kᵢᵣ inward-rectifier potassium ion channel

Fig. 3 Mass spectra of the tetramer of Kᵢᵣ3.1 cytoplasmic domain from the MALDI set-up shows the peptide fragments of the protein. Kᵢᵣ inward-rectifier potassium ion channel, MALDI matrix-assisted laser desorption/ionization

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Protein complexes were adsorbed for 1 min onto glow-discharged, 400 mesh carbon-coated grids. The grids were washed with four drops of distilled H₂O. Micrographs were taken at a low dose on a JEM-2100 operated at 120 kV (JEOL Inc, Peabody, MA 01960, USA). Tilt pairs were recorded at 0° and 45° or 0° and 60°, at a nominal magnification of 69000× on a Gatan Ultrascan 4 k CCD (Gatan Inc., Pleasanton, CA 94588, USA) [12].

RESULTS AND DISCUSSION

Spermidine–0.8 nm gold nanoparticles complexes, which is not determined by a transmission EM (TEM) of JEM-2100 (Figs. 4, 5, 6) with energy dispersive X-ray spectroscopy (EDS), surface-coated with DMSA (Figs. 7, 8), have been successfully used to produce a high signal in TEM images in the ion pore or surrounding tetramer components.

Dimercaptosuccinic acid can form chelates with gold nanoparticles via the two sulfur atoms. DMSA is water soluble with extracellular distribution and cannot penetrate the lipid bilayer, but is able enter to the cytoplasmic domain of Kᵢᵣ ion pore accompanied by spermidine molecules, already coated by the gold nanoparticles [13]. Moreover, in atomic force microscope observations, the point less than 0.8 nm cannot be counted as the gold nanoparticles because of the substrate roughness originating from the natural convex point. Evidence of the existence of gold nanoparticles in the ion pore interaction has been supported by high density surface-coated with DMSA in TEM images [14].

Energy dispersive X-ray spectroscopy was also used to confirm the presence of the gold deposits in the cytoplasmic domain of the Kᵢᵣ receptor (Figs. 9, 10, 11). X-ray photoelectron spectroscopy measurements were performed using an ultrahigh vacuum chamber with a base pressure below 5–10⁻⁹ Torr. XPS data was collected using an X-ray photoelectron spectrometer PHI 5400 (Physical Electronics) with a nominal energy resolution of 0.7 eV. Spectra were acquired using a photon beam of 1486.6 eV, selected from a conventional (nonmonochromatic) Al/Mg dual-anode X-ray source. Data was collected at room temperature under vacuum, and samples were analyzed as soon as they were prepared, without any extra cleaning procedure. The EDS spectrum of the receptors containing the gold nanoparticles indicates the presence of C and O in the receptor molecules. The small gold peak evidenced in the EDS spectrum confirms that a validated amount of the 0.8 nm gold nanoparticle is present in the ion pore of the Kᵢᵣ channels [15].

Gold nanoparticles (0.8 nm) belonging to a new class of ion channel inhibitors directly
block the human K_\text{ir} channel I(KAch) when accompanied by the antagonist spermidine. Blocking of the I(KAch) channel by the gold nanoparticle-spermidine complex has been shown to occur in atrial cardiomyocytes. A reduced K^+ current delays the repolarization of the complex in patients with atrial fibrillation (AF) [16].

This complex is a new class of three anti-arrhythmic agents that may potentially treat AF [17]. The complex is designed to be a selective I(KAch) blocker in contrast to multiple class 1 and 3 anti-arrhythmic agents with less proarrhythmic potential [18].
A limitation of this study was that the authors could not identify the 3-D images of the gold nanoparticles in the ion pore because of technical limitations.

CONCLUSION

In conclusion, we could identify undeca-gold in the ion pore of the Kir3.1 channel in order to...
clarify its direct blocking effect in the $K_{ir}$ ion pore by undecagold.

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Compliance with ethics guidelines. This article does not contain any new studies with human or animal subjects performed by any of the authors.

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