Contents lists available at ScienceDirect





journal homepage: www.elsevier.com/locate/yabbi

# Catalytic activity of human indoleamine 2,3-dioxygenase (*h*IDO1) at low oxygen



CrossMark

### Avodele O. Kolawole<sup>1</sup>, Brian P. Hixon<sup>2</sup>, Laura S. Dameron, Ian M. Chrisman, Valeriy V. Smirnov<sup>\*</sup>

Department of Chemistry and Biochemistry, Center for Biomolecular Structure and Dynamics, University of Montana, Missoula, MT 59812, United States

#### ARTICLE INFO

Article history: Received 25 November 2014 and in revised form 12 February 2015 Available online 21 February 2015

Keywords: Indoleamine dioxygenase Oxygen electrode Enzyme kinetics Substrate analog Inhibition Global fit

#### ABSTRACT

A cytokine-inducible extrahepatic human indoleamine 2,3-dioxygenase (*h*IDO1) catalyzes the first step of the kynurenine pathway. Immunosuppressive activity of *h*IDO1 in tumor cells weakens host T-cell immunity, contributing to the progression of cancer. Here we report on enzyme kinetics and catalytic mechanism of *h*IDO1, studied at varied levels of dioxygen (O<sub>2</sub>) and L-tryptophan (L-Trp). Using a cyto-chrome  $b_5$ -based activating system, we measured the initial rates of O<sub>2</sub> decay with a Clark-type oxygen electrode at physiologically-relevant levels of both substrates. Kinetics was also studied in the presence of two substrate analogs: 1-methyl-L-tryptophan and norharmane. Quantitative analysis supports a steady-state rather than a rapid equilibrium kinetic mechanism, where the rates of individual pathways, leading to a ternary complex, are significantly different, and the overall rate of catalysis depends on contributions of both routes. One path, where O<sub>2</sub> binds to ferrous *h*IDO1 first, is faster than the second route, which starts with the binding of L-Trp. However, L-Trp complexation with free ferrous *h*IDO1 is more rapid than that of O<sub>2</sub>. As the level of L-Trp increases, the slower route becomes a significant contributor to the overall rate, resulting in observed substrate inhibition.

© 2015 Elsevier Inc. All rights reserved.

#### Introduction

In the human body, L-tryptophan (L-Trp) is one of the nine essential dietary amino acids [1]. It functions as a building block of proteins and as a precursor of niacin, an intermediate in the biosynthesis of NAD<sup>+</sup> [2], and serotonin, a mood-modulating neurotransmitter and physiological regulator [3]. Up to 90% of dietary L-Trp is catabolized via the kynurenine pathway [4,5]. The kynurenine pathway starts with a dioxygenation reaction of L-Trp that is catalyzed by indoleamine 2,3-dioxygenase (IDO) and tryptophan 2,3-dioxygenase (TDO) [6,7]. Both IDO and TDO contain a type-b heme and use dioxygen  $(O_2)$  to open the five-membered ring of L-Trp to form N-formyl-L-kynurenine (NFK), as shown in Scheme 1 [6]. Excessive exhaustion of L-Trp via the kynurenine pathway significantly hinders T-cell proliferation, differentiation, effector function, and viability, resulting in a suppressed immune response [8]. Catabolites of L-Trp are capable of promoting immunosuppression and tumor tolerance during cancer [9], formation of cataracts [10,11], HIV-related neurological damage, and ischemic brain injury [12].

IDO is induced extrahepatically throughout the body [6]. The strongest inducer of IDO is a proinflammatory cytokine interferon- $\gamma$ 

\* Corresponding author.

(IFN- $\gamma$ ) [13]. IDO-initiated L-Trp degradation accelerates during conditions that cause cellular activation of the immune response such as malignancy, inflammation, autoimmune disorder, and pregnancy [12]. The induction of IDO by IFN- $\gamma$  is greatly diminished in a hypoxic environment, where the O<sub>2</sub> concentration is between 5% and 10% of air-saturation level [14–16]. During hypoxia, cells stimulated with IFN- $\gamma$  demonstrate reduced levels of IDO compared to normoxic conditions. IDO antimicrobial and immunoregulatory functions are also significantly impaired [14–16].

Even under normoxic conditions, the physiological levels of  $O_2$  are quite low (between 14% and 24% of air-saturation level or between ~35 µM and ~65 µM  $O_2$ ) [17]. The steady-state kinetic data for IDO are known primarily in solutions saturated with air [18–28]. Here, we report on enzyme kinetic studies of human IDO isoform-1 (*h*IDO1)<sup>3</sup> (EC 1.13.11.52) at physiologically-relevant levels of  $O_2$  as a function of both substrates –  $O_2$  and L-Trp. This

E-mail address: valeriy.smirnov@umontana.edu (V.V. Smirnov).

<sup>&</sup>lt;sup>1</sup> Present address: Department of Biochemistry, Federal University of Technology, P.M.B. 704, Akure 340001, Nigeria.

<sup>&</sup>lt;sup>2</sup> Present address: National Foundation for Fertility Research.

<sup>&</sup>lt;sup>3</sup> Abbreviations used: hIDO1, human indoleamine 2,3-dioxygenase isoform-1; TDO, tryptophan 2,3-dioxygenase; ι-Trp, ι-tryptophan; 1-Me-ι-Trp, 1-methyl-ι-tryptophan; NFK, N-formyl-ι-kynurenine; O<sub>2</sub>, dioxygen; O<sub>2</sub><sup>--</sup>, superoxide anion; H<sub>2</sub>O<sub>2</sub>, hydrogen peroxide; NHM, norharmane (β-carboline); IFN-γ, interferon-γ; NAD<sup>+</sup>, nicotinamide adenine dinucleotide; β-NADH, β-nicotinamide adenine dinucleotide hydride; Cu,Zn-SOD, copper-zinc superoxide dismutase; DNase I, deoxyribonuclease I; IPTG, isopropyl β-D-1-thiogalactopyranoside; PMSF, phenylmethylsulfonyl fluoride; EDTA, ethylenediaminetetraacetic acid; MOPS, 3-(N-morpholino)propanesulfonic acid; SDS-PAGE, sodium dodecyl sulfate polyacrylamide gel electrophoresis; LRF MALDI-TOF, linear-reflectron matrix-assisted laser desorption/ionization time-of-flight mass spectrometer.



Scheme 1. Reaction catalyzed by IDO.

information is vital for understanding enzyme-substrate interactions in *h*IDO1 and for designing inhibitors with enhanced therapeutic responses [29–31]. Using a new assay methodology, we quantify the effects of O<sub>2</sub> concentration on the initial rates of *h*IDO1 catalysis. We analyze the data within a mechanistic model that allows for initial complexation of either O<sub>2</sub> or L-Trp with a free ferrous form of *h*IDO1 – <sup>Fe(II)</sup>*h*IDO1 [6,23]. This model also considers that both resulting substrate-bound species – <sup>Fe(III)</sup>*h*IDO1·0<sup>-</sup><sub>2</sub> and <sup>Fe(II)</sup>*h*IDO1·L-Trp [32], lead to a ternary complex – *h*IDO1·O<sub>2</sub>-L-Trp.

The ternary complex has been previously characterized in human and rabbit IDO by the stepwise mixing of substrates with the ferrous enzyme either manually, at a low temperature, or on a stopped-flow, at room temperature [24-28,33,34]. Here we analyze the kinetics of the formation of the ternary complex. We observe that under steady-state conditions, the ternary complex forms via two separate pathways. Even though the O2-first/L-Trpsecond addition route is faster than the L-Trp-first/O<sub>2</sub>-second path [23,35], the organic substrate binds to the free ferrous *h*IDO1 at a higher rate than O<sub>2</sub> [35]. At low physiologically-relevant concentrations of both substrates, the slower L-Trp-initiated pathway is a significant contributor to the overall catalytic rate, resulting in pronounced substrate inhibition of catalysis. Such kinetic control of hIDO1 activity could be operational in vivo where, depending on tissue oxygenation [17], O<sub>2</sub> supply may be limited relative to L-Trp level [36,37].

#### Materials and methods

#### Reagents

L-tryptophan (Cat. # T0254), β-NADH (Cat. # N1161), Cu,Zn-SOD (Cat. # S8160), catalase (Cat. # C100), lysozyme (Cat. # L6876), DNase I (Cat. # D5025), PMSF (Cat. # P7626), Trizma (Cat. # T1503), MOPS (Cat. # M3183), imidazole (Cat. # 56750), EDTA (Cat. # ED4SS), kanamycin sulfate (Cat. # K1377), 1-methyl-L-tryptophan (Cat. # 447439), norharmane (Cat. # N6252) and norharmane hydrochloride (Cat. # N6377) were from Sigma–Aldrich; agar (Cat. # N833) and yeast extract (Cat. # J850) were from Amresco; tryptone (Cat. # 95039) was from Fluka; sodium chloride (Cat. # SX0425), monobasic sodium phosphate (Cat. # SX0710), dibasic sodium phosphate (Cat. # SX0715) were from EMD; δ-aminolevulinic acid hydrochloride (Cat. # 01433) and IPTG (Cat. # 00194) were from Chem-Impex International, Inc. in Wood Dale, IL; N<sub>2</sub> (pre-purified grade) and O<sub>2</sub> (USP grade) were from Norco, Inc. in Boise, ID. The pETevIDO plasmid for hIDO1 expression and the plasmids for cytochrome  $b_5$  and cytochrome  $b_5$  reductase were kindly provided by Prof. A. Grant Mauk (University of British Columbia).

#### Enzyme preparation

hIDO1 (EC 1.13.11.52) was overexpressed in One Shot BL21 Star (DE3) chemically competent *Escherichia coli* cells (Invitrogen)

using a published procedure [38]. Greater than 95% homogeneity of *h*IDO1 was confirmed by the SDS–PAGE after the second Ni-affinity column and dialysis into 50 mM Tris-HCl pH 8.0 buffer, containing 100 mM NaCl and 4 mM EDTA. The purified enzyme exhibited an [M+H]<sup>+</sup> peak at m/z = 45,586 on a Microflex LRF MALDI-TOF mass spectrometer (Bruker), which is in good agreement with the expected value of m/z = 45,643. Cytochrome  $b_5$  and cytochrome  $b_5$  reductase, overexpressed in One Shot BL21 Star (DE3) cells, were purified as described elsewhere [39,40]. The homogeneity of these proteins was confirmed by 10% SDS–PAGE analysis. Protein concentrations were determined spectrophotometrically. All enzyme stocks were stored at -75 °C.

## Measurement of initial rates of hIDO1 catalysis by monitoring $O_2$ consumption

Initial rates of hIDO1 catalysis of L-Trp dioxygenation were measured at 25.0 °C using a Clark-type oxygen electrode consisting of a digital ammeter (Biological Oxygen Monitor, model 5300A, Yellow Springs Instruments (YSI)) and a polarographic oxygen probe (5331A, YSI). Highly reproducible O<sub>2</sub> depletion traces (Fig. S1) [41] were recorded in a reaction chamber thermostated on a modified bath assembly (5301B, YSI) interfaced to a recirculating water bath (DC10, Thermo Scientific; K20, Haake). A VWR  $10'' \times 10''$  professional stirrer was used to maintain the stirring rate at 700 rpm. O<sub>2</sub> and N<sub>2</sub> gases were metered using a Riteflow flow meter (PMR1-010976, Bel-Art Scienceware). This experimental setup allowed for precise control of temperature, stirring rate, and oxygen level. The oxygen probe was calibrated daily to the dissolved oxygen in air-saturated water (resistivity  $\ge 18.2 \text{ M}\Omega \times \text{cm}$ ) at normal pressure. Due to the elevation of Missoula, the 100% airsaturation level of pure water at 25.0 °C corresponds to  $[O_2] = 230 \,\mu\text{M}$ , which is lower than  $[O_2] = 258 \,\mu\text{M}$  at sea level [42]. Solubility of O<sub>2</sub> was calculated using atmospheric pressure measured with a mercury manometer interfaced to a high-vacuum line (Chemglass). Atmospheric pressure at sea level was taken as 760 mmHg [42].

A typical reaction mixture (3.000 mL) contained 20 mM MOPS buffer at pH 7.0, 150 μM β-NADH, 1.0 μM cytochrome b<sub>5</sub>, 140 nM cytochrome b<sub>5</sub> reductase, 54 nM Cu,Zn-SOD, 12 nM catalase, and varying concentrations of L-Trp [43]. The solution was equilibrated against air or a mixture of pure O<sub>2</sub> and N<sub>2</sub>, saturated with water vapor at atmospheric pressure by passing through a fritted gas washing bottle (Chemglass, Cat. # CG-1114-13). During inhibition studies, the reaction mixture was also supplemented with 3-15 µL of inhibitor stock solution to the final inhibitor concentration of 1-5 µM. Inhibitor stocks (100 mL of 1.00 mM solution of either 1-methyl-L-tryptophan or norharmane hydrochloride) were prepared in 100 mM potassium phosphate buffer at pH 7.0. The reaction was initiated by injecting a 5- $\mu$ L aliquot of *h*IDO1 stock (~120  $\mu$ M) into a reaction chamber to an expected final concentration of ~200 nM, using a gas-tight syringe (Hamilton). The exact concentrations of hIDO1 stock solutions were measured spectrophotometrically on an AVIV Model 14 Spectrophotometer (Aviv Biomedical), or a NanoDrop 2000 (Thermo Scientific) using molar absorptivity  $\varepsilon_{\lambda=404\text{nm}}$  = 172,000 M<sup>-1</sup> cm<sup>-1</sup> [44]. Initial rates of O<sub>2</sub> consumption were measured in the range of the steepest  $[O_2]$  decline (12 s long), normally 2–4 s after injecting the ferric *h*IDO1. All measurements were performed in triplicate or greater. Readouts of the oxygen probe were transmitted to a PC workstation with a 1-Hz sampling rate. The slopes of the oxygen consumption traces were determined in Excel 2010 (Microsoft) and their absolute values were expressed in  $\mu$ M/s. These initial velocities of O<sub>2</sub> consumption were converted to specific activity of *h*IDO1 by dividing by the exact final micromolar enzyme concentration –  $[hIDO1]_T$ . All nonlinear Download English Version:

https://daneshyari.com/en/article/1925053

Download Persian Version:

https://daneshyari.com/article/1925053

Daneshyari.com