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1 ***Giardia duodenalis* arginine deiminase modulates the phenotype**  
2 **and cytokine secretion of human dendritic cells by depletion of**  
3 **arginine and formation of ammonia<sup>1</sup>**

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13 Running title: *Giardia ADI enzyme modulates human DC response*

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## 26 **Abstract**

27

28 Depleting arginine is a recognized strategy of pathogens to evade immune effector  
29 mechanisms. Depletion depends on microbial enzymes such as arginases which are  
30 considered as virulence factors. The effect is mostly interpreted as being a consequence of  
31 successful competition with host enzymes for the substrate. However, both arginases and  
32 arginine deiminases (ADI) have been associated with pathogen virulence. Both deplete  
33 arginine but their reaction products differ. An ADI has been implicated in the virulence of  
34 *Giardia duodenalis*, an intestinal parasite that infects humans and animals causing significant  
35 morbidity. Dendritic cells (DC) play a critical role in host defense, also in a murine *G.*  
36 *duodenalis* infection model. Functional properties of these innate immune cells depend on the  
37 milieu in which they are activated. Here, the dependency of the response of these cells on  
38 arginine was studied using *Giardia* ADI and LPS-stimulated human monocyte-derived DC.  
39 Arginine depletion by ADI significantly increased TNF- $\alpha$  and decreased IL-10 and IL-12p40  
40 secretion. It also reduced up-regulation of surface CD83 and CD86 molecules that are  
41 involved in cell-cell interactions. Arginine depletion also reduced phosphorylation of S6K in  
42 DC suggesting the involvement of the mTOR signaling pathway. The changes were due to  
43 arginine depletion and reaction product formation, in particular ammonium ions. Comparison  
44 of  $\text{NH}_4^+$  and urea revealed distinct immunomodulatory activities for these products of  
45 deiminases and arginases, respectively. The data suggest that a better understanding of the  
46 role of arginine-depleting pathogen enzymes for immune evasion will have to take enzyme  
47 class and reaction products into consideration.

## 48 **Introduction**

49 Many pathogens are thought to compete with the host for arginine as part of their virulence  
50 mechanisms. This is best known for pathogens expressing arginases that compete for arginine  
51 with host nitric oxide synthases and thereby are considered to prevent anti-microbial nitric  
52 oxide (NO) formation (1, 2). However, also other arginine-metabolizing enzymes have been  
53 implicated in microbial virulence, in particular arginine deiminases (ADI). The latter enzymes  
54 are thought to be relevant in streptococcal infections (3, 4) and infections with the non-  
55 invasive, gastrointestinal protozoan parasite *Giardia duodenalis* (5, 6), a medically significant  
56 cause of diarrheal syndromes and malabsorption (7, 8).

57 Arginine is not only necessary for generation of NO but it plays also other important roles  
58 during immune response. Lack of arginine was shown to inhibit T-cell function (9), and  
59 arginine levels affect signaling via the mammalian target of rapamycin (mTOR) pathway as  
60 reported for other cells (10, 11). The mTOR pathway, in turn, was shown to contribute to the  
61 regulation of co-stimulatory surface marker levels on dendritic cells (DC; (1, 12, 13)). These  
62 cells play a crucial role during induction of an adaptive immune response by secreting  
63 cytokines and by cell-contact dependent interactions with T- and other immune cells.  
64 Although DC are important for adaptive immunity against microbial infections, the effect of  
65 pathogen-mediated arginine-depletion on their function is not known.

66 Arginine-dependent virulence mechanisms of pathogens rely on enzymes such as arginases or  
67 deiminases that deplete arginine but also generate reaction products at the same time. Of note,  
68 these products differ between these classes of enzymes that generate ornithine and urea or  
69 citrulline and  $\text{NH}_4^+$ , respectively. Commonly, changes of immune cell responses due to  
70 different arginine levels have been studied by comparing responses in the presence or absence  
71 of arginine. However, this does not reflect the situation when arginine is depleted by an  
72 enzymatic reaction as it can be the case during infections. Yet, the combined effect of arginine

73 depletion by an enzymatic reaction and the ensuing product formation on immune cells has  
74 largely been ignored.

75 Referring to *G. duodenalis* as a relevant model, we studied here the immunomodulatory  
76 effects of arginine-depletion by exposing human monocyte-derived DC to recombinant  
77 *G. duodenalis* ADI during DC activation with LPS. The effect of this treatment on IL-10,  
78 IL-12p40, and TNF- $\alpha$  secretion as well as cell surface expression of CD83 and CD86 was  
79 monitored. We show that both, arginine-depletion and  $\text{NH}_4^+$  formation by the active parasite  
80 enzyme are immunomodulatory on monocyte-derived dendritic cells (moDC) causing an  
81 increase in TNF- $\alpha$ , as opposed to a decrease in IL-10 and IL12p40 production and a reduction  
82 of surface located CD86 and CD83. In particular the latter effect correlated with an inhibition  
83 of the mTOR pathway since phosphorylation of the mTOR S6K target protein was decreased.  
84 We furthermore show that  $\text{NH}_4^+$  but not urea exacerbated the inhibition of IL-10 production  
85 and surface marker up-regulation compared with arginine depletion alone, suggesting a  
86 difference between the immunomodulatory activities of the products of arginases and  
87 deiminases.

88

## 89 **Material and methods**

90

### 91 **Cell culture**

92

93 DC were cultured in RPMI 1640 (Sigma, Germany) supplemented with 10% FCS (Biochrom,  
94 Germany) or arginine-free RPMI (PanBiotech, Germany) supplemented with 10% dialyzed  
95 FCS (Biochrom). Both media contained 10 mM Hepes (Biochrom), penicillin (100 U/mL) -  
96 streptomycin (100 µg/mL) (PAA, Germany) and 50 µM 2-mercaptoethanol (Roth, Germany).

97

98 *G. duodenalis* trophozoites, strain WB-C6 (ATCC 50803), were propagated in TYI-S-33  
99 medium as previously described (14). Genomic DNA was isolated with the DNeasy Blood &  
100 Tissue kit (Qiagen, Germany).

101

102

### 103 **Generation of recombinant proteins**

104

105 The ADI-coding sequence (*Gd-adi*) was amplified from genomic DNA of *G. duodenalis*  
106 strain WB-C6 by PCR using *Pfu* polymerase (Fermentas, Germany) and specific primers (5'-  
107 aatgactgactctccaaggataaaga-3' and 5'- tccctcacttgatatcgacgcagatgtca-3'). The resulting 1.8 kb  
108 fragment was cloned into the expression vector pASG-IBA35 (StarGate cloning, IBA GmbH,  
109 Germany) according to the manufacturer's manual to produce a recombinant protein with an  
110 N-terminal His<sub>6</sub>-tag. After transformation into *E. coli* DH5αZ1 cells (15), the His<sub>6</sub>-tagged ADI  
111 (pASG-IBA35\_ADI) was purified from cultures grown in LB medium supplemented with  
112 100 µg/mL ampicillin (Roth) and 50 µg/mL spectinomycin (Sigma) overnight at 37°C  
113 without induction by anhydrotetracycline. Cells were harvested by centrifugation (8200 x g,  
114 10 min, 4°C) and then washed in ice-cold PBS. The pellet was resuspended in 25 mM Hepes

115 (Fluka, Germany), 150 mM NaCl (Merck, Germany), 5 mM imidazole (Merck) plus EDTA-  
116 free protease inhibitor cocktail (Roche, Germany), pH 7.5 (Buffer A) and disrupted using a  
117 high pressure homogenizer (EmulsiFlex, Avestin, Germany). After centrifugation (15000 x g,  
118 30 min, 4°C), the supernatant was passed through a 0.45 µm pore size filter (Sartorius,  
119 Germany) and loaded onto a HisTrap<sup>TM</sup>FF column (GE Healthcare, Germany) pre-  
120 equilibrated with Buffer A on an ÄktaFPLC system (GE Healthcare). Finally, protein was  
121 eluted in 25 mM Hepes, 150 mM NaCl, and 200 mM imidazole, pH 7.5 (Buffer B). Finally,  
122 imidazole was removed by desalting using a PD-10 column (GE Healthcare) and residual LPS  
123 by an EndoTrap® kit (Hyglos, Germany). Recombinant protein in 1x PBS was finally  
124 concentrated using a Vivaspin concentrator (5 kDa PES membrane, Sartorius). To obtain a  
125 catalytically inactive arginine deiminase mutant (ADI<sub>C424A</sub>), cysteine424 was changed to  
126 alanine using the QuikChange® mutagenesis kit (Stratagene). Site-directed mutagenesis PCR  
127 was performed using the primers 5'-gtacggctctctgcacgccgcatctcaggttgg-3' and 5'-  
128 aacaacctgagatgcggcgtgcagagagccgtac-3' and, the vector pASG-IBA35\_ADI as template.  
129 Recombinant ADI<sub>C424A</sub> was expressed and purified as described for the wild-type enzyme.  
130 Purity of both recombinant proteins was controlled by SDS-PAGE and verified using Western  
131 blot. Protein concentration was determined using a BCA protein assay kit (Thermo Scientific  
132 Fisher, Germany). Enzymatic activity of recombinant ADI was measured by colorimetric  
133 determination of citrulline formation as described (16).

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**141 Production of polyclonal antibody**

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143 An alpaca was immunized four times (day 0, 21, 35, 49; Preclinics, Potsdam, Germany) with  
144 300 µg of purified recombinant ADI resuspended in 100 µL PBS plus adjuvant (complete  
145 Freund's adjuvant for the 1<sup>st</sup> injection and incomplete Freund's adjuvant for subsequent  
146 booster injections). Pre-immune serum was collected prior to the first immunization. After the  
147 last booster injection, the antiserum was obtained.

148 For Western blot analysis the alpaca antiserum was diluted 1:2500 and the secondary  
149 horseradish-peroxidase-conjugated goat anti-llama IgG antibody (Bethyl Laboratories, USA)  
150 1:20000, both in 5% non-fat dried milk (Roth), 1xPBS, 0.05% Tween20 (Roth).

151

152

**153 Generation of human moDC**

154

155 Peripheral blood mononuclear cells (PBMC) were isolated from buffy coats of healthy  
156 volunteers (German Red Cross, Berlin, Germany) by density gradient centrifugation (Ficoll-  
157 Paque<sup>TM</sup> Plus, GE Healthcare). Monocytes were separated by magnetic cell sorting using  
158 CD14 MicroBeads (Miltenyi Biotec, Germany). Typically, cells were thereby enriched to  
159  $\geq 90\%$  CD14<sup>+</sup> as determined by flow cytometry. To generate moDC,  $3.0 \times 10^6$  monocytes were  
160 seeded into 6-well tissue culture plates. DC growth medium additionally contained 1000U/mL  
161 human rGM-CSF (Bayer, USA) and 10 ng/mL human rIL-4 (R&D Systems, Germany). After  
162 6 days, immature DC were harvested and the cell population was characterized by analyzing  
163 an aliquot by flow cytometry. Staining of surface markers revealed low levels of CD14 and  
164 CD86 and high levels of HLA-DR, whereas CD25 and CD83 were not expressed.

165

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## 167 **DC stimulation and analysis of cytokine production and cell surface markers**

168

169  $5 \times 10^5$  immature DC/mL were seeded into a 12-well tissue culture plate and cultured at 37°C  
170 either in DC growth medium or in arginine-free DC growth medium supplemented or not with  
171 2 mM of arginine (Merck), or citrulline (Sigma), and/or ammonium chloride (Sigma),  
172 respectively. Cells were exposed to different concentrations of recombinant ADI or ADI<sub>C424A</sub>  
173 and 1 µg LPS/mL (*Escherichia coli* 026:B6 from Sigma) was added at the same time to  
174 trigger DC maturation. Cell-culture supernatants were collected after 24 h and were frozen at -  
175 80°C until determination of TNF-α, IL-10, IL-12p70 (all from eBioscience, Germany), IL-23  
176 (U-CyTech, Netherlands), and IL-12p40 (Biolegend, Germany) by ELISA following  
177 manufacturers' protocols. DC were stained with anti-human FITC-CD14 (Immunotools),  
178 PerCP-CD86 (Abcam), PE-HLA-DR (Immunotools), AlexaFluor488-CD83 (Biolegend), or  
179 Dyomics647-CD25 (Immunotools) conjugated mAb or the relevant isotype controls  
180 (Immunotools). Data were acquired on a LSR II flow cytometer (Becton Dickinson,  
181 Germany) and analyzed using FlowJo software (Tree Star, Inc.). Dead cells were excluded by  
182 gating according to FSC/SSC characteristics.

183

## 184 **Western blot analysis**

185

186 One million immature DC/mL were seeded into a 12-well tissue culture plate in arginine-free  
187 DC medium containing rGM-CSF and rIL-4, further supplemented or not with 2 mM of  
188 arginine, citrulline and/or ammonium chloride. After 30 min of resting, cells were treated with  
189 2 µM rapamycin (Sigma), recombinant ADI or ADI<sub>C424A</sub> and further cultured at 37°C for  
190 90 min. The immature cells were then exposed to 1 µg/mL LPS for 30 min. Cells were  
191 harvested, centrifuged at 300 x g at 4°C for 10 min, and supernatants were then collected for  
192 citrulline detection. Cell pellets were washed twice in ice-cold PBS before resuspension in

193 lysis buffer [10 mM Tris-HCl (Roth) pH 7.2, 150 mM NaCl (Merck), 1% Triton X-100  
194 (Merck), 1% sodium deoxycholate (Sigma), EDTA-free protease inhibitor cocktail (Roche)].  
195 To remove cell debris, samples were centrifuged for 5 min at 14000 x *g* at 4°C. Protein  
196 concentration of the supernatant was determined by using BCA protein assay (Thermo  
197 Scientific Fisher, Germany) according to the manufacturers' protocol. A total of 50 µg protein  
198 was separated by SDS-PAGE and transferred to nitrocellulose membranes (Biorad,  
199 Germany). Membranes were blocked for 1 h in 5% non-fat dried milk (Roth), 1xTBS, 0.1%  
200 Tween20 (Roth) and incubated over night at 4°C with primary antibodies (Cell signaling,  
201 Germany) against p70-S6K (#9202), phospho-p70-S6K (#9205), E4-BP1 (#9452), phospho-  
202 E4-BP1 (#9455) and β-actin (#4967) diluted 1:1000 in 5% BSA (Roth), 1xTBS, 0.1%  
203 Tween20. Bound antibodies were detected with horseradish peroxidase-conjugated goat anti-  
204 rabbit IgG (Jackson ImmunoResearch, USA) diluted 1:2000 in 5% BSA, 1xTBS, 0.1%  
205 Tween20 and visualized on X-ray films (GE Healthcare) using ECL Plus Western Blotting  
206 Detection System (GE Healthcare). Quantification of Western blot signals was performed  
207 with ImageJ 1.42q software (NIH, USA).

208

209

## 210 **Descriptive statistics**

211

212 Data are given as mean ± SD. Statistical significance was assessed by paired t-test (two-  
213 tailed). All analyses were performed using GraphPad Prism 5 software (GraphPad Software,  
214 Inc., USA) at the level of  $p < 0.05$ .

## 215 **Results**

216

217 *Defining experimental conditions for arginine-depletion by recombinant, enzymatically active*218 *G. duodenalis* ADI during DC activation in vitro

219

220 The average daily intake of arginine by humans has been estimated to be 5 g of arginine,

221 corresponding to 28 mmol, (17). It has been reported that an infected person sheds up to  $10^9$ 222 *G. duodenalis* organisms per day (18). We first aimed to estimate the effect of ADI activity on223 the amount of arginine that may be available to *G. duodenalis* during an infection. Thus, we

224 determined the ADI activity per million trophozoites to be equivalent to approx. 0.02 U by

225 preparing lysates from different isolates and determining their ADI activity (data not shown).

226 Therefore, at least 20 units of enzyme may be produced per day of infection. One unit

227 corresponds to 1  $\mu$ mol arginine metabolized per minute, hence  $>28$  mmol could potentially be228 turned over in one day. Thus, the infection with *Gd* has the potential to substantially deplete

229 arginine in the gastrointestinal tract.

230 We cloned the *adi* gene present in strain WB-C6 and produced the respective recombinant231 enzyme as well as an enzymatically inactive mutant ADI<sub>C424A</sub> (19), in which Cys at residue232 424 was replaced by Ala, as hexahistidine-tagged proteins in *E. coli*. Analysis by SDS-PAGE

233 and Western blot of the two affinity-purified proteins demonstrated that the recombinant

234 proteins were highly pure and behaved similar to the endogenous ADI parasite protein of

235 66 kDa (Fig. 1). That was also described by others, but unlike we could not confirm the

236 previously described higher molecular weight of 85 kDa for native *Gd* ADI (20). The purified

237 wild type protein had an ADI activity of 6.8 U/mg.

238 To achieve arginine-depletion by recombinant ADI in a standard 24 h DC stimulation assay

239 enabling to study the consequences of arginine-depletion on DC, we calculated that 4  $\mu$ g of

240 active recombinant ADI will convert all free arginine in 43 min (culture volumes of 1 ml

241 RPMI containing 1.15 mM arginine), 10-fold less enzyme will achieve this in approx. 7 h and  
242 100-fold less enzyme would need nearly 72 h assuming a constant activity over time. Thus,  
243 we choose these amounts of the active enzyme to study dose dependence in the following DC  
244 activation experiments and used the inactive mutant to control for possible effects unrelated to  
245 enzyme activity.

246

247 *Enzymatic arginine depletion by ADI modifies pro- and anti-inflammatory cytokine secretion*  
248 *of LPS-activated moDC*

249

250 In the intestine, DC project cellular extensions between epithelial cells (IEC) into the lumen  
251 of the gut to sample antigens (21, 22). Through these projections they are thought to recognize  
252 microbe-associated molecular patterns by pattern recognition receptors and become  
253 stimulated to mature and migrate to peripheral lymph nodes where they initiate antigen-  
254 specific immunity (23, 24). Maturing DC up-regulate cell surface markers (e.g. MHC, CD83,  
255 CD86) and release cytokines (e.g. IL-12, IL-10, TNF- $\alpha$ ) to enable communication with and  
256 activation of T- and other immune cells (25). To model such conditions and investigate  
257 potential immunomodulatory effects of arginine limitation on DC function, we used human  
258 moDC that can be readily prepared in the necessary quantities, stimulated them with LPS, and  
259 exposed the cells to ADI. Citrulline concentrations in the culture supernatants were  
260 determined at the end of the 24 h assay period and confirmed the calculated effects of the ADI  
261 dilutions (see above and Fig. 2A) on arginine turn-over. As expected, addition of ADI<sub>C424A</sub>  
262 did not result in citrulline formation at all tested concentrations (Fig. 2A and data not shown).  
263 Cytokine concentrations in these supernatants were then determined by ELISA (Fig. 2B).  
264 LPS-activation of cells stimulated roughly a 100-fold increase in IL-12p40, TNF- $\alpha$  and IL-10  
265 in arginine-replete medium compared to non-activated controls (data not shown). In contrast,  
266 LPS-stimulated moDC exposed to 4  $\mu$ g/mL of ADI produced significantly less IL-10 and IL-

267 12p40. These values were 45% ( $P < 0.001$ ,  $n = 7$ ) and 17% ( $P < 0.01$ ,  $n = 6$ ) lower compared  
268 to control cells exposed to mutant ADI<sub>C424A</sub>, respectively. In contrast, TNF- $\alpha$  secretion by  
269 ADI-treated and LPS-activated moDC was increased and values were on average 74% ( $P <$   
270 0.01,  $n = 6$ ) higher than values of LPS-stimulated, mutant enzyme exposed control cells.  
271 Both, the decreased IL-10 and IL-12p40 and the increased TNF- $\alpha$  production, depended on  
272 ADI activity, indicating that the kinetics of arginine-depletion did matter. These data show  
273 that arginine depletion by ADI modulates cytokine secretion of LPS-activated moDC and,  
274 importantly, that the effects differ depending on the cytokine analyzed.

275

276 *Enzymatic arginine depletion by ADI reduces upregulation of surface CD83 and CD86 levels*  
277 *of LPS-activated moDC*

278

279 To investigate the influence of arginine depletion by ADI on the phenotype of maturing DC,  
280 moDC were stimulated with LPS and treated with ADI or its mutant ADI<sub>C424A</sub> as described  
281 above and analyzed for selected surface marker proteins levels by flow cytometry. Activation  
282 of immature moDC with LPS induced up-regulation of CD83 and CD86 (data not shown) as  
283 expected. In contrast, LPS-stimulated moDC treated with arginine-depleting levels of ADI  
284 expressed significantly less CD83 and CD86 than control cells. Surface CD83 was on average  
285 22% ( $P < 0.05$ ,  $n = 3$ ) and that of CD86 15% ( $P < 0.01$ ,  $n = 4$ ) lower than on LPS-treated,  
286 mutant enzyme-exposed control cells. The effects were again ADI dose dependent (Fig. 3B).  
287 The reduced up-regulation of surface CD83 and CD86 unlikely reflects a general effect on  
288 surface protein levels since HLA-DR abundance on LPS-activated moDC was not affected by  
289 ADI (data not shown).

290

291

292 *ADI immunomodulatory effects on LPS-activated moDC result from both arginine depletion*  
293 *and product formation*

294

295 We asked next whether the modulatory effects of ADI on moDC functional phenotype  
296 resulted from arginine depletion and/or the formation of ADI reaction products citrulline  
297 and/or  $\text{NH}_4^+$ . To investigate possible cause effect relationships, moDC were generated as  
298 before, harvested and then seeded into arginine-free culture medium that was only  
299 subsequently supplemented with arginine, citrulline, and/or ammonium chloride, respectively.  
300 Cells were then treated as before with 4  $\mu\text{g}$  of ADI or  $\text{ADI}_{\text{C424A}}$  and were activated with LPS.  
301 After 24 h of incubation, supernatants were collected and citrulline content and cytokine  
302 concentration were determined. Furthermore, moDC were harvested and cell surface markers  
303 were analyzed.

304 IL-10 and TNF- $\alpha$  secretion of cells grown in arginine-replete media were modulated as  
305 described above, i.e. ADI-treated LPS-activated moDC produced less IL-10 and more TNF- $\alpha$   
306 than control cells (Fig. 4). MoDC stimulated in the absence of arginine showed a small  
307 decrease in IL-10 production but levels remained significantly higher than those of cells  
308 treated with ADI in the presence of arginine where citrulline and  $\text{NH}_4^+$  could be formed  
309 (Fig. 4). The addition of the ADI products, in particular of  $\text{NH}_4^+$  in the form of  $\text{NH}_4\text{Cl}$ ,  
310 reduced IL-10 levels to those observed in the presence of arginine and ADI. Similarly, CD83  
311 and CD86 surface levels were also reduced, although this did not reach statistical significance  
312 for the latter marker (data not shown). In contrast, increase in TNF- $\alpha$  production was mainly  
313 driven by arginine depletion (Fig. 4) and not affected by  $\text{NH}_4^+$ . The increase due to arginine  
314 depletion was reduced by citrulline, indicating that citrulline could substitute for arginine in  
315 this respect (Fig. 4). Supplementing DC with  $\text{NH}_4^+$  during stimulation in the presence of  
316 arginine but with no ADI present also reduced IL-10 but had no effect on TNF- $\alpha$  production  
317 (Fig. 5).

318 Thus, the immunomodulation of moDC by ADI resulted from a combination of distinct  
319 effects of arginine depletion and/or citrulline and  $\text{NH}_4^+$  product formation.

320

321  *$\text{NH}_4^+$  and urea, the reaction products of ADI and arginases, differ in their effect on cytokine*  
322 *secretion and surface marker profile of LPS-stimulated moDC*

323

324 As mentioned before, many pathogens are thought to evade NO-mediated immune clearance  
325 through arginine depletion by arginases, which will result in the formation of ornithine and  
326 urea (2). Immunomodulatory effects may thus be different from conditions where deiminases  
327 are relevant. We were therefore interested in comparing the immunomodulatory effects of  
328  $\text{NH}_4^+$  and urea. LPS-stimulated cells were incubated in medium devoid of arginine but  
329 supplemented with ammonium chloride or urea for 24 h and then supernatants were collected  
330 for the detection of IL-10 and TNF- $\alpha$  by ELISA (Fig. 6A) and CD83 and CD86 proteins by  
331 flow cytometry (Fig. 6B). Notably, cells stimulated in medium supplemented with urea  
332 produced significantly more IL-10 and displayed significantly higher levels of CD83 and  
333 CD86 surface proteins than cells stimulated in the presence of  $\text{NH}_4^+$ . In contrast and  
334 corroborating our result that TNF- $\alpha$  production was not affected by  $\text{NH}_4^+$  (Fig. 4), no  
335 differences were noted between cells treated with  $\text{NH}_4^+$  or urea with respect to TNF- $\alpha$  release.

336

337 *Arginine turn-over by ADI results in decreased phosphorylation of the mTOR-signaling*  
338 *pathway target S6K in LPS-activated moDC*

339

340 Previous studies suggested an inhibitory effect of branched-chain amino acids on the  
341 mTOR/S6K-signaling pathway, resulting in impaired maturation of moDC and particularly  
342 affecting CD83 expression (26). mTOR is a serine/threonine kinase that is present in two  
343 distinct protein complexes, mTORC1 and mTORC2. Activation of mTORC1 leads to

344 phosphorylation of the protein S6 kinase (S6K) and the eukaryotic initiation factor 4E (eIF-  
345 4E) binding protein 1 (4E-BP1) that both are involved in the regulation of protein translation  
346 (27). Arginine levels have been shown to affect mTOR signaling in T-cells (10, 11). Since  
347 ADI by arginine depletion had immunomodulatory effects on moDC, we hypothesized that  
348 this may also involve the mTOR pathway in moDC. MoDC were seeded in arginine-free  
349 medium, then supplemented or not with arginine. The cells were treated with recombinant  
350 ADI, catalytically inactive ADI<sub>C424A</sub> or, as a positive control, with the mTOR inhibitor  
351 rapamycin and were activated with LPS. Cells were incubated for 24 h before determining  
352 CD83 surface levels (Fig. 7A). In the presence of arginine CD83 levels were lower on cells  
353 treated with ADI than on cells exposed to the mutant enzyme, as shown before. MoDC  
354 stimulated in the absence of arginine or in the presence of arginine and rapamycin also  
355 showed reduced levels of CD83 protein (Fig. 7A).

356 To investigate whether the reduced CD83 surface levels caused by ADI activity correlated  
357 with mTOR signaling, treated cells were harvested 30 min after LPS stimulation to assess  
358 mTOR-dependent phosphorylation events. Cells were lysed and equal amounts of total  
359 protein were separated by SDS-PAGE. The abundance of phosphorylated protein of the  
360 mTOR target S6K was then determined by Western blot analysis and results were quantified.  
361 Suppression of S6K phosphorylation in moDC correlated with reduced CD83 surface protein  
362 levels and this depended on arginine levels (Fig. 7B). Control cells treated with rapamycin  
363 showed the expected suppression of S6K phosphorylation. Phosphorylation of 4E-BP1,  
364 however, did not depend on arginine availability (data not shown).

365 These data suggest that arginine levels, similar to what has been shown for branched-chained  
366 amino acids (26), affect mTOR activity in moDC but also indicate a difference between  
367 amino acids.



## 368 **Discussion**

369

370 We show here that arginine depletion by *G. duodenalis* ADI modulates surface markers and  
371 cytokine response of in vitro activated human moDC. The immunomodulation depended on  
372 both, depletion of arginine and the ADI reaction products, in particular  $\text{NH}_4^+$ . By  
373 investigating the consequences of arginine depletion we found that the mTOR pathway is  
374 implicated in the molecular signaling and leads to the modulation of DC responses. These  
375 findings add further evidence to the hypothesis suggested by others (28) that ADI is a  
376 molecularly defined virulence and pathogenicity factor of *G. duodenalis*. Arginine-depleting  
377 enzymes of pathogens have been implicated in immune evasion mechanisms because of  
378 competing for the same substrate as host NO synthases. Little attention has been paid to date  
379 to effects beyond this inhibition of microbial effector function. Our data implicate that  
380 arginine-metabolizing enzymes of pathogens are more widely involved in immunomodulation  
381 and suggest distinct roles for the reaction products of different enzyme classes in this process.  
382 Currently, the relevance of our in vitro findings for the understanding of the pathogenesis of  
383 giardiasis must remain speculative and further studies are required. Nonetheless, we would  
384 like to discuss their potential implications: Firstly and as mentioned before, DC in the  
385 intestine project extensions between epithelial cells into the lumen of the gut to sample  
386 antigens (21, 22). Although only two studies with murine cells have been published to date  
387 that investigated the interaction of Giardia parasites with DC (29, 30), they provided evidence  
388 for the importance of DC in controlling Giardia infection (30). The ability to modulate DC  
389 function could thus provide a selective advantage to the parasite and our calculations suggest  
390 that arginine depletion may indeed occur in situ.

391 Secondly, arginine-dependent response modulation would be consistent with observations on  
392 the pathophysiology of human giardiasis. Atrophy of villi has been detected microscopically  
393 in intestinal biopsies of chronically infected patients, and symptomatic disease has been

394 correlated with a dysfunction of the epithelial barrier (31) but the process leading to this is not  
395 understood. These sequelae can, however, also be observed when intestinal biopsies are  
396 treated with TNF- $\alpha$  (32). TNF- $\alpha$  may thus have pathogenic properties in this context. Studies  
397 in mice showed that peak level Giardia parasite loads were around 10-fold higher in animals  
398 devoid of TNF- $\alpha$  (33) hence it was proposed that it has a protective function in giardiasis.  
399 However, in the same study transepithelial resistance was reduced to the same extent despite a  
400 much lower parasite burden in TNF- $\alpha$  responsive mice. An explanation consistent with the  
401 results from human biopsies exposed to TNF- $\alpha$  (31) could be that reduced epithelial integrity  
402 during giardiasis is due to parasite factors and TNF- $\alpha$ , the latter exerting a dual (protective  
403 and pathogenic) role.

404 Thirdly, children suffering from symptomatic giardiasis were shown to have increased  
405 mucosal levels of pro-inflammatory cytokines including TNF- $\alpha$ , which decreased after anti-  
406 parasitic treatment and resolution of symptoms while local levels of IL-10 increased after  
407 treatment (34). This is in agreement with our in vitro findings and may indicate that parasite  
408 ADI mediated arginine depletion impaired IL-10 and enhanced TNF- $\alpha$  secretion by mucosal  
409 DC in situ in these children. The relative abundance of IL-10 and TNF- $\alpha$  is recognized as a  
410 critical parameter in intestinal diseases in mice (35) and in humans (36, 37).

411 Fourthly, although *G. duodenalis* ADI is found intracellularly as part of the arginine  
412 dehydrolase (ADH) pathway and this pathway is thought to exploit arginine as an energy  
413 source to produce ATP in these amitochondrial organisms (38), the protein is also released by  
414 the parasite. Microarray analysis of the transcriptional response to host cell contact had  
415 revealed an up-regulation of *ADI* mRNA (28), and ADI as well as ornithine carbamoyl  
416 transferase (OCT, the next enzyme in the ADH pathway) were released by the parasite in  
417 contact with intestinal cells in vitro and in vivo ((39) and S.B. unpublished data). The facts  
418 that ADI and OCT were detected in these assays and are also immunodominant antigens  
419 during infection (40, 41) indicate that significant amounts of the enzymes are extracellular

420 and therefore free citrulline may even be further metabolized to ornithine. It is tempting to  
421 speculate that this may exacerbate TNF- $\alpha$  production because the negative feedback of  
422 citrulline on this parameter would be reduced. Thus, release of ADH compounds could have  
423 evolved in part due to selective pressure by the host's immune response.

424 To our knowledge this is the first report showing an immunomodulatory effect of arginine  
425 depletion and  $\text{NH}_4^+$  formation on the response of DC other than on NO formation. Precedence  
426 for a clinically relevant role of amino acids in modulating immune responses exists (42, 43).  
427 For DC it has been shown that branched-chain amino acids affect maturation, and it has been  
428 proposed that this modulation is a consequence of inhibition of the mTOR pathway (26). Our  
429 results are consistent with this hypothesis but extend the concept to arginine. Of note,  $\text{NH}_4^+$   
430 has recently been reported to modulate mTOR activity in yeast cells (44). This suggests that  
431 reaction products formed by arginine-depleting enzymes could further affect mTOR signaling.  
432 However, additional signaling pathways are likely to be involved in mediating the distinct  
433 effects of arginine depletion alone and  $\text{NH}_4^+$  formation. The latter has recently also been  
434 invoked in the T cell inhibition mediated by *Salmonella* L-asparaginase II (45). Further  
435 studies are required to understand this comprehensively.

436 In summary, we describe immunomodulatory effects of arginine depletion on human DC.  
437 Using ADI from *G. duodenalis* we show that this immunomodulation depends on arginine  
438 depletion and the products formed by the enzyme. This reveals novel facets of DC response  
439 modulation by arginine-metabolizing enzymes and may have implications for the general  
440 understanding of arginine-dependent virulence mechanisms of pathogens.

441

442 **Acknowledgment**

443

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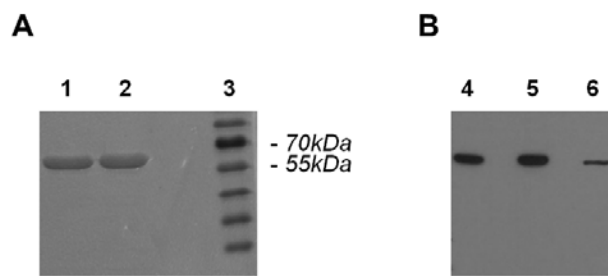
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580 **Figure Legends**

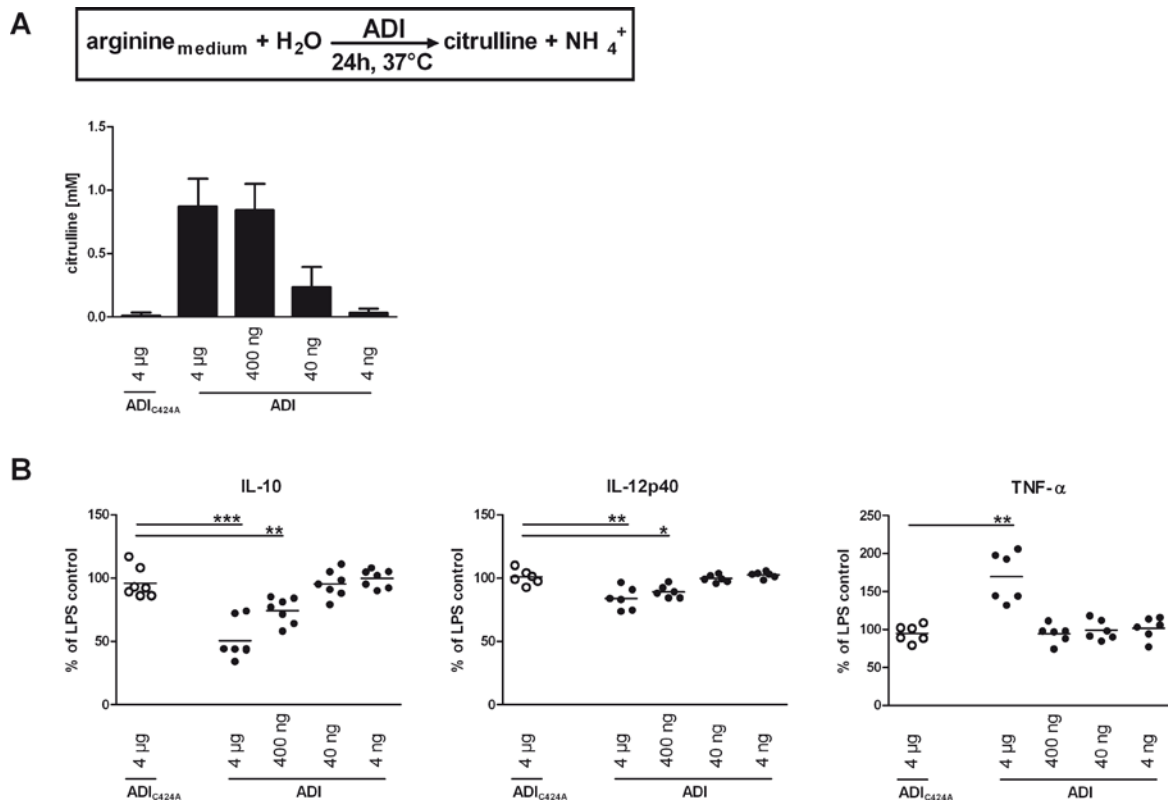
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582 **Figure 1. SDS-PAGE and Western blot analysis of purified recombinant *G. duodenalis***  
583 **arginine deiminase**

584 A, Coomassie-stained SDS-PAGE loaded with affinity-purified recombinant catalytically  
585 inactive ADI<sub>C424A</sub> (lane 1), recombinant enzymatically active ADI (lane 2) and molecular  
586 mass standard (lane 3). B, Antigenic identification of 0.5 µg affinity-purified recombinant  
587 ADI (lane 4), 0.5 µg catalytically inactive ADI<sub>C424A</sub> (lane 5) and native ADI in 2.9 µg of  
588 *G. duodenalis* (strain WB-C6) lysate (lane 6) by Western blot analysis using a polyclonal  
589 alpaca antiserum raised against ADI. Pre-immune serum used as control did not react with  
590 any *G. duodenalis* protein (data not shown).

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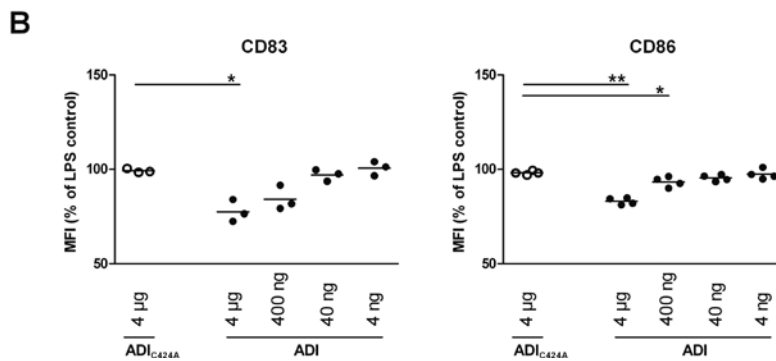
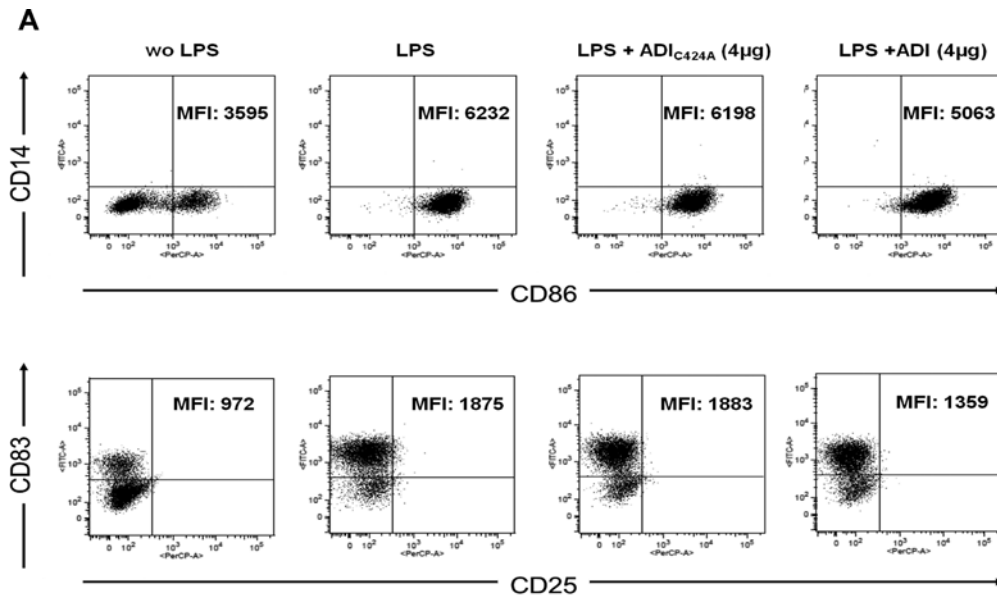


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594 **Figure 2. Enzymatic arginine depletion by ADI modulates cytokine secretion of LPS-**  
 595 **activated human moDC**

596 Immature moDC ( $5 \times 10^5$  per sample) were exposed to the indicated amounts of recombinant  
 597 ADI or, as control, corresponding levels of ADI<sub>C424A</sub> (only highest amount shown) and  
 598 1 µg/mL LPS was added. Citrulline content was determined as a measure of the cumulated  
 599 ADI activity and arginine depletion over the 24 h assay time (A). DC cytokine secretion into  
 600 supernatants was assessed by ELISA and is expressed as percentage of the amount secreted  
 601 by LPS stimulated cells not exposed to any ADI protein (B). Bars in (A) represent the mean  $\pm$   
 602 SD from experiments with DC prepared from seven different donors. Symbols in (B)  
 603 represent values from individual donors with means indicated by horizontal lines. Differences  
 604 between the amount of cytokines secreted by cells exposed to mutant (control) or active ADI  
 605 were analyzed by paired t test (two-tailed); \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

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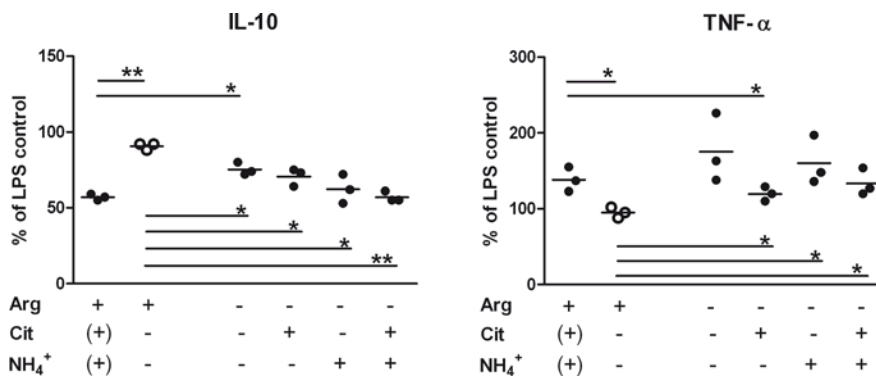
608 **Figure 3. Enzymatic arginine depletion by ADI reduces CD83 and CD86 surface**  
 609 **markers induction by LPS activation on moDC**

610 Cells were treated as described in Fig. 1 and all cells, except the non-stimulated control cells,  
 611 were treated with 1 μg/mL LPS. After incubation for 24 h at 37°C, moDC were harvested,  
 612 stained with cell surface marker-specific antibodies and analyzed by flow cytometry.

613 (A) Representative dot plots for CD14/CD83 (top panel) and CD25/CD86 (bottom panel)  
 614 expression with the respective mean fluorescence intensity (MFI) values for CD86 and CD83

615 are shown. (B) Relative MFI in % of control LPS-stimulated cells for CD83 (n=3) and CD86  
 616 (n=4) with moDC from different donors are shown. Horizontal lines correspond to mean

617 values and differences between the respective MFI on cells exposed to mutant (control) or  
 618 active ADI were analyzed by paired t test (two-tailed); \* $p < 0.05$ , \*\* $p < 0.01$ .



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621 **Figure 4. Depletion of arginine and formation of ADI reaction products citrulline and**  
 622 **ammonium ions modulate moDC response to LPS activation**

623 Immature moDC after harvest were seeded into arginine-free growth medium. The medium  
 624 was then supplemented as indicated with arginine, citrulline and/or ammonium chloride  
 625 (2mM each; brackets indicate addition of ADI to arginine-replete medium to reflect formed,  
 626 not supplemented products). Cells were activated with LPS and treated with ADI (●) or  
 627 mutant ADI<sub>C424A</sub> (○) for control. Values correspond to % of the respective parameter  
 628 determined for cells stimulated with LPS only in arginine-replete medium. After 24 h,  
 629 supernatants were collected and cytokine concentrations determined by ELISA. Horizontal  
 630 lines correspond to mean values and differences between the respective parameter determined  
 631 with cells exposed to mutant (control) or active ADI were analyzed by paired t test (two-  
 632 tailed); \* $p < 0.05$ , \*\* $p < 0.01$ .

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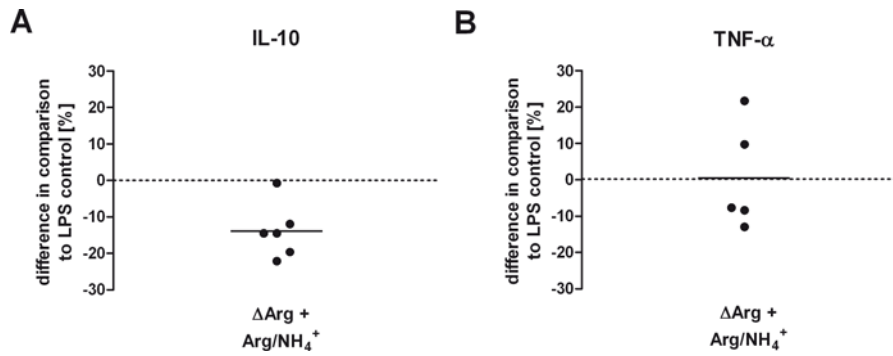
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642 **Figure 5. Addition of NH<sub>4</sub><sup>+</sup> in the presence of arginine reduces IL-10 secretion of moDC**  
 643 **after LPS activation**

644 Immature moDC were prepared and stimulated as described (Fig. 4) and medium was  
 645 supplemented with arginine or arginine and ammonium chloride. After 24 h, supernatants were  
 646 taken and cytokine concentrations determined by ELISA. Each dot represents an independent  
 647 experiment with moDC prepared from an individual donor and values are expressed as %  
 648 difference to cytokine amounts produced by arginine supplemented, LPS-stimulated cells.  
 649 Significance was tested against the null hypothesis that there is no differences between cells  
 650 stimulated in the presence of arginine or arginine plus ammonium ion using paired t test (two-  
 651 tailed); P values were  $p < 0.01$  for IL-10 levels and non-significant for TNF- $\alpha$ .

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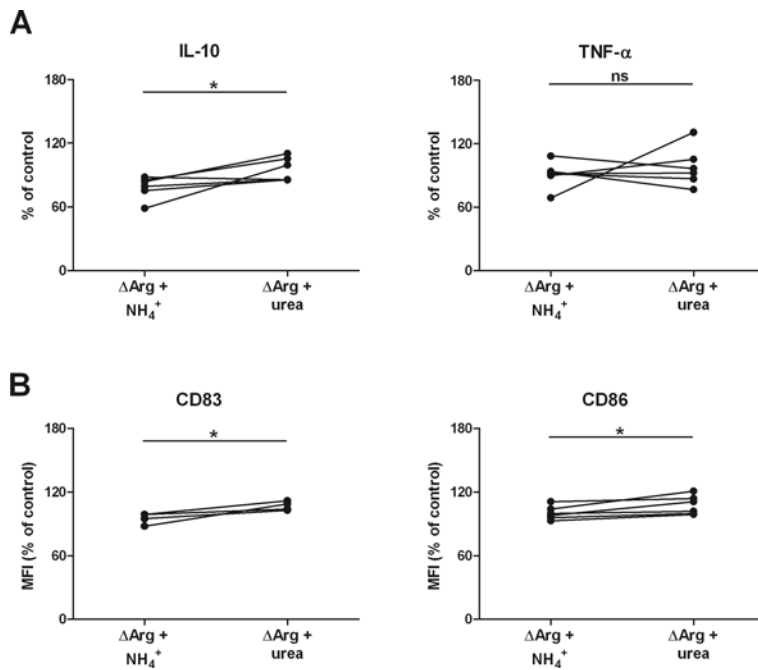
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660 **Figure 6. Immunomodulation of LPS-activated moDC under arginine starvation is**  
 661 **different between  $\text{NH}_4^+$  and urea.**

662 MoDC were prepared as described in Fig. 4 and stimulated in arginine-free medium  
 663 supplemented with either 2 mM of ammonium chloride or urea. After 24 h, supernatants were  
 664 collected and cytokine concentrations determined by ELISA (A). The moDC were harvested  
 665 and surface marker proteins were analyzed by flow cytometry (B). Dots represent values from  
 666 experiments with cells from six (A) or four (B) individual donors and are expressed as % of  
 667 control values obtained with LPS-activated cells with no supplement (paired t test (two-  
 668 tailed); \* $p < 0.05$ ).

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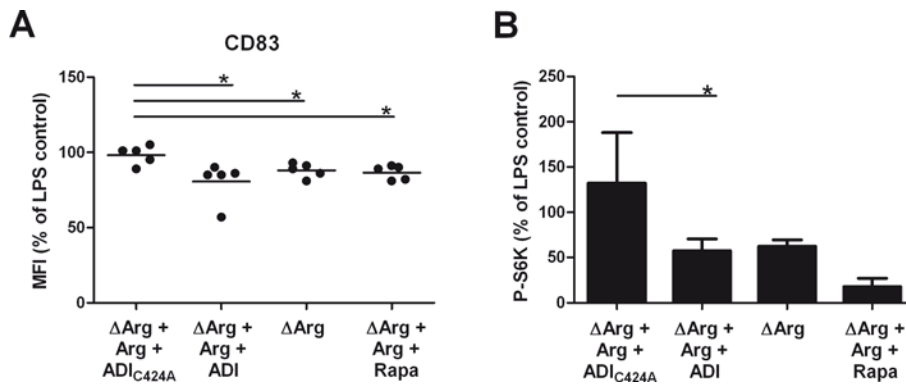
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677 **Figure 7. ADI-mediated arginine depletion decreases mTOR-signaling in LPS-**  
 678 **stimulated moDC**

679  $10^6$  immature moDC were seeded in arginine-free growth medium into each well of a 12-well  
 680 tissue culture plate. Arginine-free medium ( $\Delta$ Arg) was supplemented or not with 2 mM  
 681 arginine and cells were treated or not with 4  $\mu$ g of ADI, 4  $\mu$ g of ADI<sub>C424A</sub>, or 2  $\mu$ M  
 682 rapamycin. For activation 1  $\mu$ g/mL LPS was added after 2 h to each sample. (A) MoDC CD83  
 683 surface marker levels 24 h after stimulation compared to stimulated control cells are shown.  
 684 Bars represent the mean  $\pm$  SD of independent experiments with five different donors (paired  
 685 t test (two-tailed), \* $p$  < 0.05 compared with the respective control). (B) In parallel, 30 min  
 686 after LPS stimulation, cells were harvested, washed and lysed. 50  $\mu$ g of cell extracts were  
 687 separated by SDS-PAGE and p70-S6K, phospho-p70-S6K and  $\beta$ -actin were detected by  
 688 immunoblotting and quantified by image analysis. Phosphorylated p70-S6K levels were  
 689 normalized relative to  $\beta$ -actin and compared to stimulated control cells. Levels of total p70-  
 690 S6K levels were not significantly different between the different experimental groups. Bars  
 691 represent the mean  $\pm$  SD from five individual experiments with cells from different donors  
 692 (paired t test (two-tailed), \* $p$  < 0.05, \*\* $p$  < 0.01 compared with the respective control).

693 **Footnotes**

694

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