

Characterization of the phenotype and function of monocyte-derived dendritic cells in allergic conjunctivae

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| 1 | CHARACTERIZATION OF THE PHENOTYPE AND FUNCTION OF MONOCYTE- |
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| 2 | DERIVED DENDRITIC CELLS IN ALLERGIC CONJUNCTIVA |
| 3 | Bita Manzouri ^{*†} , Masaharu Ohbayashi [*] , Andrea Leonardi [§] , Dilniya Fattah [¶] , D. Frank P. |
| 4 | Larkin ^{*†} & Santa J. Ono [*] |
| 5 | * UCL Institute of Ophthalmology, London, UK. |
| 6 | † Moorfields Eye Hospital, London, UK. |
| 7 | § Ophthalmology Unit, Department of Neuroscience, University of Padua, Italy. |
| 8 | ¶ Medicines Research Centre, GlaxoSmithKline, Stevenage, UK |
| 9 | |
| 10 | Corresponding author: |
| 11 | Bita Manzouri |
| 12 | Department of Cornea & External Disease |
| 13 | Moorfields Eye Hospital |
| 14 | City Road |
| 15 | London EC1V 2PD |
| 16 | bita.manzouri@moorfields.nhs.uk |
| 17 | Tel: 07711 357 211 |
| 18 | Fax: 020 8447 0353 |
| 19 | |
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| 21 | assay. |
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| 23 | |

25 ABSTRACT

| 27 | Background: Dendritic cells (DCs) are the most potent antigen-presenting cells involved in |
|----|---|
| 28 | initiating the immune response, presenting antigens to T cells and leading to T cell proliferation. |
| 29 | In an immature state, DCs lack accessory signals required for T cell stimulation but are highly |
| 30 | specialized to capture antigens. Full DC maturation changes the cell surface phenotype and |
| 31 | facilitates stimulation of T cell proliferative responses. To examine the degree of DC maturity |
| 32 | associated with vernal keratoconjunctivitis (VKC), we examined the phenotype and antigen- |
| 33 | presentation capability of blood derived DCs from VKC patients and from normal controls. |
| 34 | Methods: Flow cytometry was used to identify the cell surface expression of markers of DC |
| 35 | maturity (CD83, CD86, MHC class II) and mixed leukocyte reactions to assess DC induction of |
| 36 | T cell proliferation. |
| 37 | Results: DCs derived from VKC patients were of more mature phenotype than those from |
| 38 | normal controls. However, these VKC DCs had reduced capability for induction of T cell |
| 39 | proliferation compared to DCs from controls. |
| 40 | Conclusion: The increased maturity of DCs in VKC patients correlates with the heightened |
| 41 | immune responsiveness associated with this disorder. A number of mechanisms may underlie |
| 42 | the impaired ability of DCs in atopy to stimulate T cell proliferation. This impairment of DC |
| 43 | induction of T cell activation is likely to be one factor which contributes to the modified |
| 44 | inflammatory response seen in VKC patients and the recognised susceptibility of these patients |
| 45 | to viral infection. |

47 INTRODUCTION

48 Approximately 20% of the Western world suffers from allergic diseases, such as allergic 49 rhinoconjunctivitis, allergic asthma and the atopic asthma / dermatitis syndrome.[1] DCs are the 50 sentinel cells that ensure that an immune response is appropriate in specificity, intensity, and 51 type to the nature of the threat posed by the antigen bearing entity. DCs are divided into 2 major 52 subtypes in human: the myeloid DCs (including Langerhan's cells (LCs)) and plasmacytoid DCs. 53 DCs are located in most tissues where they capture and process antigens, and where they exist 54 in an 'immature' state: they lack the signals required for T cell stimulation but they are highly 55 specialized to capture antigens. Following encounter with and uptake of an antigen, DCs 56 process this antigen into smaller fragments, couple it to the major histocompatibility complex 57 (MHC) class II molecule and present it on their surface to T cells. Additionally, a second co-58 stimulatory signal is also required to initiate T cell proliferation.[2] DCs that display the MHC 59 class II molecule and the co-stimulatory molecules (CD83, CD86) on their surface are 60 considered to be mature.

61 KleinJan et al. [3] studied nasal mucosal biopsy specimens taken from patients with 62 perennial allergic rhinitis (AR) and found that the number of DCs in the nasal mucosa was 63 higher and displayed a more mature phenotype in symptomatic patients than in healthy controls. 64 This increase in LCs of the nasal mucosa was also observed during the pollen season in work 65 done by Till et al. [4]. Allam et al. [5] performed detailed phenotypical comparison of nasal and 66 oral myeloid DC (mDC) from atopic and non-atopic donors. In atopic donors, these mDCs 67 showed elevated expression of the high affinity receptor for IgE (FccRI) with receptor occupation 68 by IgE but FccRI was present only in low amounts in these cells in non-atopic donors. Semper 69 et al. [6] studied the phenotype of DCs on bronchial biopsy specimens from normal and 70 asthmatic airways and demonstrated the presence of FccRI-bearing CD1a⁺ LCs in both types of 71 specimens. Total expression of FccRI on all cells was slightly higher in the airways of 72 asthmatics compared to normal controls, but this failed to reach statistical significance. They did

find—in keeping with other studies [7, 8]—that the numbers of CD1a⁺ DCs was higher in
asthmatics when compared to normal controls.

75 We have previously demonstrated that the number of DCs expressing $Fc \in RI$ in the 76 substantia propria of the conjunctiva in the atopic disorder vernal keratoconjunctivitis (VKC) 77 patients is increased, indicating an increased ability of DCs to capture and process specific 78 antigens for presentation to CD4⁺ T cells, thereby initiating the immune response.[9] In this 79 study, we sought to further define the phenotypic and functional characteristics of DCs 80 associated with VKC. We examined DCs derived from the peripheral blood of VKC patients and 81 normal patients, testing the hypothesis that DCs in VKC are more mature, and have increased 82 capacity for T cell proliferation.

83

84 METHODS

85 Selection of donors

86 Donors had a clinical history and signs of VKC, and were selected from patients 87 attending clinic follow-up appointments. The control group donors did not have any history of 88 atopy and did not display any signs of ocular allergy (table 1). A difference did exist between 89 the mean age of the donors in the VKC group compared to the control group but this was 90 unavoidable and was consequent to the difficulties in obtaining blood from children for the 91 control group and in adequate volume quantities to perform the experiments. Local ethics 92 committee approval was obtained for the research and each patient gave written informed 93 consent. All research procedures adhered to the guidelines established in the Declaration of 94 Helsinki.

95

96

| | | Pati | ients for flow | cytometry s | tudy | | |
|-------------------|-------------------------------|----------|---------------------|-------------------|-------------|-----|---------------------|
| | Control | subjects | | VKC patients | | | |
| Patient number | Age (yr) | Sex | lgE level (kU/l) | Patient number | Age (yr) | Sex | IgE level (kU/l) |
| 1 | 47 | F | 18 | 1 | 17 | F | 911 |
| 2 | 27 | F | 11 | 2 | 8 | М | 87 |
| 3 | 47 | F | 17 | 3 | 12 | М | 210 |
| 4 | 28 | F | 21 | 4 | 31 | М | 60 |
| 5 | 46 | F | 21 | 5 | 10 | М | 150 |
| 6 | 39 | F | 32 | 6 | 14 | М | 96 |
| 7 | 29 | F | 15 | 7 | 37 | F | 81 |
| 8 | 49 | М | 44 | 8 | 7 | М | 120 |
| 9 | 47 | F | 334 | 9 | 11 | М | 170 |
| 10 | 28 | М | 257 | 10 | 8 | М | 375 |
| 11 | 23 | F | 254 | 11 | 13 | М | 62 |
| 12 | 28 | М | 129 | 12 | 12 | М | 153 |
| 13 | 30 | М | 203 | | | | |
| | | | Patients for | MLR assays | | | |
| | Control subjects VKC patients | | | | | | |
| Patient number | Age (yr) | Sex | lgE level (kU/l) | Patient number | Age (yr) | Sex | lgE level (kU/l) |
| 1 | 41 | М | 11 | 1 | 10 | М | 80 |
| 2 | 21 | F | 31 | 2 | 29 | М | 16 |
| 3 | 50 | М | 13 | 3 | 13 | М | 69 |
| 4 | 42 | F | 18 | 4 | 9 | М | 216 |
| 5 | 49 | F | 37 | 5 | 31 | М | 10 |
| 6 | 38 | F | 9 | 6 | 14 | F | 42 |
| 7 | 21 | F | 212 | 7 | 11 | F | 48 |
| 8 | 25 | F | 55 | | | | |

99 Table 1: Background patient information.

100

101 Measurement of IgE concentration

102 Serum (6 ml) was collected for measurement of IgE levels by a normal sandwich

103 enzyme-linked immunosorbent assay (ELISA; ImmunoCAP 100, Phadia, Uppsala, Sweden).

104 **MoDC** isolation and culture

105 Peripheral blood mononuclear cells were isolated from EDTA-treated blood using 106 Lymphoprep (Sigma-Aldrich, Poole, United Kingdom) centrifugation. Isolation of CD14⁺ 107 monocytes from the resultant buffy coat layer was performed using anti-CD14 mAb conjugated 108 to magnetic MicroBeads and the autoMACS system as described in the manufacturer's 109 instructions (Miltenyi Biotech, Bergisch Gladbach, Germany). The derived pellet of purified 110 monocytes was cultured into dendritic cells as per the method described by Sallusto and 111 Lanzavecchia.[10] To generate monocyte-derived DCs (MoDCs), IL-4 (4 µg/ml, R & D systems, 112 Abingdon, United Kingdom) and GM-CSF (6 µg/ml, GlaxoSmithKline Research Institute, 113 Stevenage, United Kingdom) were added to each well of purified monocytes.

114 Flow cytometry

On day 6 or 7 the MoDC cultures from each well were combined for fluorescent staining.
 The cells were blocked for 10 minutes with human Fcγ fragments (Miltenyi Biotech) and then
 incubated on ice with the appropriate fluoroscein isothiocyanate (FITC)- or phycoerythrin (PE) conjugated antibody or with the respective isotype-matched control immunoglobulins.

119 Monoclonal antibodies directed against CD83 (FITC, clone HB15e), CD11c (PE, clone 120 B-ly6), CD86 (FITC, clone 2331), HLA-DR (FITC, clone Tü39), CD23 (PE, clone M-L233), CD58 121 (PE, clone 1C3) and CD11c (PE, clone D12) were purchased from Becton Dickinson (Cowley, 122 United Kingdom). The antibody against $Fc \in RI-\alpha$ (PE, clone AER-37) was purchased from 123 eBioscience (Insight Biotechnology Ltd, Wembley, United Kingdom). FITC- and PE-conjugated 124 isotype control mouse immunoglobulins (Igs; IgG_1 , IgG_{2a} , IgG_{2b}) were obtained from Becton 125 Dickinson and eBioscience. Additional FACS staining of T cells was performed using CD3 126 (FITC, clone HIT3a) (BD Pharmingen) and CD209 (DC-SIGN, PE, clone eB-h209, eBioscience). 127 The α CD28/ α CD3 (GlaxoSmithKline, Stevenage, United Kingdom) was used as a positive 128 control.

Events were acquired using the Beckman Coulter flow cytometry machine and analyzed with Expo32 (Beckman Coulter Ltd, High Wycombe, United Kingdom). Cells were acquired and manually gated on the physical properties of the cell by FSC versus SSC. The measurements taken were the mean fluorescence intensity (MFI) and the percentage of positive cells as defined by the proportion expressing higher levels of fluorescence than those of the isotype controls.

135 **T cell isolation and culture**

T cells were isolated and purified from the buffy coat following layering of blood over
Lymphoprep (Sigma-Aldrich) and centrifugation. The T cells were isolated from the resultant
PBMC cell pellet using the pan T cell isolation kit II from Miltenyi Biotech (negative selection)
according to the manufacturer's instructions. T cell purity was always greater than 99%.

140 Mixed leukocyte reaction (MLR) assays

141 MoDCs were harvested on day 6 and purity of population assessed with flow cytometry 142 for CD11c +/- CD209. The MoDCs were washed and resuspended in T cell medium at a 143 concentration of 1 x 10⁴ DCs/100 µl T cell medium. In all allogenic MLR assays, T cells from the 144 same donor were used, thereby maintaining consistency between the MLR results obtained 145 from the normal control MoDCs and the VKC-derived MoDCs. Each of the MLRs for each 146 patient was performed in triplicate at five different concentrations of MoDC to T cells (1:10, 1:20, 147 1:40, 1:80 and 1:160). Graded doses of MoDC were added to T cells in 96-well U-bottom tissue 148 culture plates (200 µl/well)-the T cell number remained constant whilst serial dilution of MoDCs 149 was performed to obtain the appropriate DC:T cell ratios for the MLR assays. The control wells, 150 consisted of two sets of negative controls (1 x 10⁴ MoDC only in T cell medium and 1 x 10⁵ T cells in T cell medium) and one positive control $(1 \times 10^5 \text{ T cells only in T cell medium in wells})$ 151 152 lined with α CD28/ α CD3). The plates were incubated for 5 days at 37°C in 5% CO₂. T cell

| 155 | last 16 hours of the 5-day culture. Cells were harvested on a filter and radioactivity measured in |
|-----|---|
| 156 | a beta-plate liquid scintillation counter. |
| 157 | Statistical analysis |
| 157 | |
| 158 | The Mann Whitney U test was used to determine if there was significant up/down |
| 159 | regulation of the measured cell surface markers when comparing VKC MoDC to those derived |
| 160 | from control subjects and to determine if the proliferation of T cells observed between VKC and |
| 161 | control subjects at the different ratios tested was significant. A value of $p \le 0.05$ was considered |
| 162 | to be significant. |
| 163 | |
| 164 | RESULTS |
| 165 | Flow cytometric analysis of circulating DC phenotype |
| 166 | MoDCs were isolated from the peripheral blood of 12 VKC patients and 13 normal |
| 167 | controls. For each of the control group donors, a yield of $3 - 16.2 \times 10^{6}$ MoDCs was obtained |
| 168 | after culture as compared with the VKC donors where a yield of $0.5 - 13.2 \times 10^6$ MoDCs was |
| 169 | obtained. No morphological differences were readily identifiable between the two groups. DCs |
| 170 | were identified as staining brightly for CD11c, the widely-used marker for human myeloid DCs. |
| 171 | The cells were gated on their FSC and SSC properties with the MoDCs within the gate being $>$ |
| 172 | 95% CD11c ⁺ (figure 1, (a)). |
| 173 | MoDCs of the control group individuals displayed constitutive cell surface expression of |
| 17. | MHC class II, low expression of CD86, and no expression of CD83, a typical profile for DCs in |
| 17: | the immature state. In MoDCs from VKC patients in comparison, there was a statistically |
| 176 | significant upregulation of MHC class II, CD83 and CD86 (figure 1 (b) and table 2), an |
| 177 | expression profile consistent with a more-mature state of development. |
| | |

proliferation was quantified by incubating the cells with 1 μ Ci of methyl-titrated thymidine

deoxyriboside (10 µl; Amersham Life Sciences, Buckinghamshire, United Kingdom) during the

153

154

The cell surface adhesion proteins CD58, CD11b, and CD11c were constitutively
expressed in the control MoDCs (figure 1 (c) and table 2). Of the two IgE receptors, only CD23
was constitutively expressed on control MoDCs, and little or no expression of FccRI was
detected on these cells. There was statistically insignificant upregulation of surface FccRI on
VKC-derived MoDCs, but significant upregulation of CD11b, CD11c and CD58. CD23 was
down-regulated in VKC-derived MoDCs (p = 0.02).

184

185 Table 2: The mean fluorescent intensity (MFI) recorded for each marker measured during flow

| 186 | cytometric analysis. * | Denotes a statistically significant result. |
|-----|------------------------|---|
|-----|------------------------|---|

| | Control subject MFI | | | VKC patient MFI | | | Fold | р |
|-----------------|---------------------|-------|-----------|-----------------|-------|-----------|--------|----------|
| | Mean | SD | Range | Mean | SD | Range | change | value |
| MHC class II | 40.97 | 14.01 | 20.8-66.9 | 71.41 | 29.70 | 14.4-122 | 1.74 | <0.01 * |
| CD86 | 2.42 | 1.85 | 0.45-5.68 | 5.98 | 3.40 | 1.71-11.7 | 2.47 | <0.01 * |
| CD83 | 0.34 | 0.09 | 0.2-0.48 | 0.68 | 0.15 | 0.46-0.98 | 2.00 | <0.001 * |
| CD58 | 4.18 | 2.59 | 1.97-12 | 9.28 | 3.17 | 4.77-15.6 | 2.22 | <0.001 * |
| CD11c | 7.86 | 4.29 | 3.63-18 | 12.09 | 6.77 | 5.07-30.7 | 1.54 | 0.05 * |
| CD11b | 76.92 | 33.56 | 38.7-152 | 111.05 | 48.28 | 40.4-192 | 1.44 | 0.05 * |
| CD23 | 6.26 | 3.78 | 3.03-15.4 | 5.55 | 11.37 | 0.94-41.5 | -1.13 | 0.02 * |
| FcεRI | 0.52 | 0.24 | 0.27-0.92 | 0.54 | 0.18 | 0.23-0.91 | 1.04 | 0.77 |

188 Functional analysis of circulating DCs in VKC patients

Having determined that VKC patient-derived MoDCs had altered cell surface marker profiles compared to the controls, we then examined these cells for altered function by testing T cell stimulatory capacity. The responder T cells used were from a subject without any history of allergy or medication, and an IgE level of 61 kU/l (table 1).

For each of the control group donors, a yield of $2.4 - 7.5 \times 10^6$ MoDCs was obtained after culture as compared with a yield of $8 - 11 \times 10^5$ MoDCs for the VKC donors. No morphological differences were observed between the MoDCs from the VKC and the control donors. The MoDCs were labelled with anti-CD11c-PE and the T cells with anti-CD3-FITC. The yield of CD11c⁺ cells from both normal and VKC cultured MoDC populations was >90% (figure 2 (a)), whereas the yield of T cells from the T cell donor was consistently >99% pure (figure 2 (b)).

199 As expected, MoDCs from both the control group and VKC patients stimulated T cell 200 proliferation. The mitogenic effect was statistically significantly greater with MoDCs from control 201 group individuals than with MoDCs from VKC patients (figure 2 (c)) and was observed for all 202 ratios of MoDC to T cells that were tested. Since all T cells used were from the same donor, the 203 observed differences are likely to be explained by differing stimulatory capacity of MoDCs in the 204 two groups. In a trial MLR using MoDCs derived from a non-atopic donor outside of our group of 205 experimental subjects and T cells from the same donor as for the experiments described in this 206 study, adequate proliferative responses were observed (data not shown). This effect on 207 stimulation is therefore not limited to the patients in our control group.

208

209 **DISCUSSION**

210 Phenotypic alterations in MoDCs derived from VKC patients

The MoDC derived from VKC patients are a more mature phenotype than those from
 control subjects as indicated in the statistically significant upregulation of MHC class II and the

213 co-stimulatory molecules CD86 and CD83. This finding is comparable to those of McCarthy et 214 al. [11] who found that DCs in atopic asthma patients expressed higher levels of CD80 and 215 CD86. The upregulation of MHC class II and co-stimulatory molecules is likely to increase the 216 antigen presenting capacity of DCs in VKC patients. CD83 expression on the cell surface is a 217 marker of DC maturation.[12] Zhou and Tedder have demonstrated that most CD83⁺ cells 218 express high levels of CD86:[13] the VKC MoDCs in this study displayed upregulation of both 219 molecules. As CD86 is upregulated in respiratory tract DCs and in circulating blood B cells in 220 allergic asthma [11, 14] and on LC's in the conjunctiva of patients with VKC, [15] it is clear that 221 this co-stimulatory molecule may contribute to the T-cell driven pathology of VKC.

Adhesion proteins upregulated on VKC MoDCs included CD58, CD11c and CD11b. These molecules facilitate the interaction of DCs with T cells and influence the ability of DCs to migrate to regional lymph nodes upon maturation.

225 Expression of FcεRI on MoDCs

226 DCs can express the high-affinity IgE receptor FccRI, which, in the presence of specific 227 IgE, facilitates the uptake of allergen, leading to increased activation of allergen-specific T cells. 228 We found that MoDCs from VKC patients and normal subjects displayed similar levels of FccRI 229 expression. These results are consistent with those of Holloway et al., [16] who found that the 230 total expression of FccRI on the surface of DCs isolated from the peripheral blood of healthy 231 and asthmatic subjects were not significantly different. Other studies have associated elevated 232 levels of DC FccRI with atopic conditions.[17, 18] There are at least three possible explanations 233 for the results in our study. First, the mature state of the VKC MoDCs may have led to 234 downregulation of the FccRI receptor: upon *in vitro* maturation, LCs completely and irreversibly 235 downregulate their transcripts for FccRI subunits, leading to a rapid loss of intracellular and 236 surface receptor expression.[19] Second, although a direct correlation between serum IgE 237 levels and expression of $Fc \in \mathbb{R}$ on human basophils from allergic subjects has been shown, [20]

this control of FccRI expression may not apply to DCs, since FccRI on these cells exists as a trimeric complex lacking the β subunit, in contrast to the tetrameric form expressed by mast cells and basophils . Finally, DCs in different tissues may have different phenotypes which may be influenced by the microenvironment. Indeed we have found increased FccRI expression in conjunctiva in allergic conjunctivitis.[9]

243 Mechanisms underlying reduced T cell proliferation in response to VKC MoDCs

244 Mature DCs are better able to prime T cells than immature DCs, an important early step 245 in cell-mediated immunopathology. We found the expected potent T cell proliferation in 246 response to MoDCs derived from control subjects, but it was surprising to find that our mature 247 VKC MoDCs were less effective at stimulating T cell proliferation. This, however, is not the first 248 report of a reduction in T cell function in atopic disease—the literature in support of this finding 249 is well summarized in the review article by Strannegård and Strannegård.[21] In the setting of 250 allergic disorders, this reduced capacity to cause T cell proliferation might be one reason why 251 those with allergic disease are so prone to viral and fungal infections, in particular herpes 252 simplex virus infection. Upon presentation of viral antigen from virus-infected cells it may be that 253 the ineffective ability of DCs to stimulate T cell proliferation leads to impaired clearance of viral 254 infection in atopic patients.

255 This is the first study to show a functional difference between the MoDCs derived from 256 VKC patients and normal control subjects who had no known history of allergy. There are 257 several possible explanations for reduced T cell-stimulating activity. The proliferative signal 258 presented by the MoDCs to the T cells might be defective or insufficient. Indeed, the 'strength of 259 stimulation' model of T cell activation and differentiation is based on the fact that naïve T cells 260 interacting with DCs receive stimulatory signals of differing strengths, depending on the degree 261 of DC maturation, the level of peptide-MHC and co-stimulatory molecule interaction, and the 262 duration of T cell-DC interaction.[22] According to this model, T cells receiving a signal of

263 excessive strength undergo activation-induced cell death. The MoDCs from VKC patients, being 264 in a more mature state of differentiation, may deliver a signal of excessive strength to the T 265 cells, thereby causing cell death and a reduced proliferative respons. Alternatively, inhibitory 266 signals (DC-derived immunoglobulin receptor 2 (DigR2), cytotoxic T lymphocyte-associated 267 receptor 4 (CTLA-4), programmed death-1 (PD-1), and DC-associated heparan sulfate 268 proteoglycan-dependent integrin (DC-HIL)) presented or secreted by the VKC MoDCs hinder an 269 adequate proliferative response from T cells. Finally, the difference in MoDC stimulatory 270 capacity might be due to the differences in age and blood volumes between the donor groups, 271 unavoidable limitations of this study. The normal control patients were older, on average, than 272 the VKC patients, and the blood volumes withdrawn from the normal patients were higher than 273 for the VKC patients.

In conclusion, we have shown that MoDCs derived from circulating monocytes in VKC individuals display a more mature phenotype than their normal counterparts, as evidenced by enhanced expression of MHC II, CD86, CD83, CD58, CD11c and CD11b. Despite this phenotypic maturity, the VKC MoDCs are less-potent T cell stimulators - and potentially lesseffective APCs - than their normal counterparts. The impaired T cell proliferation may contribute to the abnormal inflammatory response, not least to viral or fungal infection, seen in patients with VKC and other allergic disorders.

281

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- 292
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- 294

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358 Figure & Table Legends

359 Figure 1: Flow cytometric analysis of (a) MoDCs cultured for analysis of CD11c surface marker. 360 The FACS pattern was generated based on cells that were gated ('A'), and confirms their 361 phenotype as CD11c+. The same methodology was used for characterization of normal MoDCs and VKC MoDCs. (b) and (c) Staining for surface markers labelled with FITC and PE 362 363 respectively. Expression of each of these markers was determined by flow cytometry and is 364 represented in the form of histograms. A shift of the curve to the right in comparison with the 365 isotype control indicates surface expression of this marker. In (b) MoDCs derived from normal 366 control and VKC patients were labelled with FITC-labelled antibodies against MHC class II. 367 CD83 and CD86. The histograms show significant upregulation of the DC surface markers MHC 368 class II, CD86 and CD83 in VKC subjects compared to normal control subjects; in (c) MoDCs 369 derived from normal control and VKC patients were labelled with PE-labelled antibodies against 370 CD11b, CD11c, CD23, CD58 and FccRI. Significantly higher expression of CD11c, CD11b and 371 CD58 was found on VKC-derived cells with significantly lower expression of the CD23 receptor 372 but not of FceRI. (* denotes p < 0.05.)

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374 Figure 2: Flow cytometric analysis of MoDCs and T cells for MLRs. CD11c labelling (a) of cells 375 derived from normal control and VKC patients that were gated ('A') indicated DC phenotype. 376 CD3 labelling (b) of isolated lymphocytes that were gated ('A') confirmed T cell phenotype. (c) 377 Allogeneic T cell proliferation stimulated by MoDCs derived from normal control and VKC 378 patients. Donor T cells tested were identical for the MLRs involving control MoDCs and VKC 379 MoDCs. T cells were co-cultured with MoDC for 5 days. Triplicate assays were performed at 380 each MoDC:T cell ratio. Results shown indicate mean of triplicate assays for each patient and 381 then mean of all patients at that ratio. Error bars indicate standard deviation. At all cell ratios

| 382 | tested, higher T cell proliferation, as determined by thymidine incorporation, was found in |
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| 383 | response to incubation with MoDCs from normal control subjects than from VKC patients. |
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- **Table 1:** Background patient information.
- **Table 2:** The mean fluorescent intensity (MFI) recorded for each marker measured during flow
- 388 cytometric analysis. * Denotes a statistically significant result.







(c)