Bright expression of CD91 identifies highly activated human dendritic cells that can be expanded by defensins

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Introduction

CD91 is a member of the low-density lipoprotein receptor family that recognizes more than 40 structurally and functionally distinct ligands.¹ Particularly, CD91 has been

Summary

CD91 is a scavenger receptor expressed by different immune cells and its ligands defensins have been demonstrated to contribute to immune responses against infections and tumours. We previously demonstrated that CD91 is expressed on human monocyte-derived dendritic cells (moDCs) and that human defensins stimulate in vitro the activation of these cells. In this study, we observed that CD91 is expressed at different levels on two distinct moDC subsets: CD91^{dim} and CD91^{bright} moDCs. Although CD91^{bright} moDCs represented a small proportion of total moDCs, this subset showed higher levels of activation and maturation markers compared with $CD91^{\text{dim}}$ moDCs. The frequency of $CD91^{\text{bright}}$ moDCs increased by \sim 50% after *in vitro* stimulation with recombinant human neutrophil peptide-1 (rHNP-1) and recombinant human β defensin-1 (rHBD-1), while lipopolysaccharide (LPS) stimulation decreased it by \sim 35%. Both defensins up-regulated moDC expression of CD80, CD40, CD83 and HLA-DR, although to a lower extent compared with LPS. Notably, upon culture with rHNP-1 and rHBD-1, CD91bright moDCs maintained their higher activation/maturation status, whereas this was lost upon culture with LPS. Our findings suggest that defensins promote the differentiation into activated $CD91^{bright} DCs$ and may encourage the exploitation of the CD91/defensins axis as a novel therapeutic strategy to potentiate antimicrobial and anti-tumour immune response.

Keywords: CD91; defensins; dendritic cells; human β defensin-1; human neutrophil peptide-1.

> demonstrated to bind and internalize α_2 -macroglobulin,² $C1q³$ and defensins.⁴ Reflecting this diversity of ligands, CD91 displays a wide tissue distribution and participates in a variety of physiological responses, including lipoprotein metabolism, proteinase homeostasis and cell migration.

Abbreviations: hsp 90, heat-shock protein 90; LPS, lipopolysaccharide; MFI, mean fluorescence intensity; moDCs, monocytederived dendritic cells; rHBD-1, recombinant human β defensin-1; rHNP-1, recombinant human neutrophil peptide-1

Subsequent studies defined CD91 as an immunologically relevant receptor for heat-shock proteins and molecular chaperones.^{5,6} It is expressed on different cell types, including dendritic cells, monocytes, macrophages, B and T cells as well as splenocytes and thymocytes.⁷ In addition, CD91 is expressed on monocyte-derived dendritic cells (moDCs) and it plays a role in the internalization of CD91-targeted antigens and their presentation to T cells.⁸ Moreover, the importance of this molecule in the immune response has been strengthened by the observation that CD91 represents an important route for stimulating $CDS⁺ T$ cell responses by MHC class I-restricted presentation.⁹

The scavenger receptor CD91 is involved in a wide variety of diseases and infections. Kebba et al.,¹⁰ showed that a significantly higher surface expression of CD91 on monocytes of HIV-1-infected long-term non-progressors may contribute to host anti-HIV-1 defence and play a role in protection against HIV-1 infection. Further, the up-regulation of CD91 in patients affected by cancer may represent an additional strategy to improve DC activation and subsequent stimulation of a specific CD8⁺ T-cell response towards tumour cells dying an immunogenic cell death.¹¹ Of note, CD91 ligands heat-shock proteins, α_2 -macroglobulin and defensins are currently under investigation in pre-clinical and clinical trials exploring new therapeutic strategies for the treatment of infectious diseases and cancers.^{12–14}

In a previous study, we demonstrated that moDCs express CD91 and that moDC treatment with human defensins up-regulates the surface expression of CD91 as well as the activation/maturation markers on these cells.¹⁵ In this study, we describe two subsets of moDCs, characterized by variable levels of CD91 expression and different states of activation/maturation. We further investigate the effects of recombinant human α and β defensins, human neutrophil peptide-1 (HNP-1) and human β defensin-1 (rHBD-1), on these two subsets of moDCs.

Materials and methods

In vitro culture of moDCs

Human peripheral blood mononuclear cells were isolated by Ficoll density gradient centrifugation from standard buffy coat preparations of healthy donors. Monocytes were obtained from adherent peripheral blood mononuclear cells ($> 90\%$ pure CD14⁺, as assessed by flow cytometry) and cultured in RPMI-1640 (Euroclone, Wetherby, UK) supplemented with 10% heat-inactivated fetal calf serum (Gibco, Invitrogen, Carlsbad, CA), in the presence of 800 U/ml recombinant human granulocyte–macrophage colony-stimulating factor (Strathmann Biotech, Hannover, Germany) and 10 ng/ml recombinant human interleukin-4 (R&D Systems, Minneapolis, MN) at 37° in 5% $CO₂$ for 5 days. Immature moDCs were characterized by long dendrites and expression of high levels of CD1a (> 90%) and low levels of CD14 $(< 5\%)$. Cell viability was $> 90\%$ in all experiments, as assessed by trypan blue exclusion. These immature moDCs were stimulated for 18 hr by adding full-length 100 ng/ml rHNP-1 (30 amino acids), 500 ng/ ml rHBD-1 (36 amino acids ; both from AlphaDiagnostic, San Antonio, TX), 20 μ g/ml heat-shock protein 90 (hsp 90; Enzo Life Sciences, Lausen, Switzerland), or 100 ng/ml lipolysaccharide (LPS; serotype 055:B5; Sigma Chemicals Co., St Louis, MO). The concentration and duration of culture stimulation were established based on preliminary experiments.¹⁵ Endotoxin contamination in culture media and defensins was excluded by Limulus amoebocyte lysate assay (BioWhittaker, Walkersville, MD) following the manufacturer's protocol.

Flow cytometric analysis

Phenotypic analysis was performed by five-colour flow cytometry. Briefly, 1×10^5 cells were incubated for 20 min at 4° with different monoclonal antibodies: FITC-conjugated anti-CD91 (Biomac, Leipzig, Germany), phycoerythrin (PE) -conjugated anti-CD80, allophycocyanin (APC) -conjugated anti-CD86, PE-conjugated anti-CD1a, APC-conjugated anti-CD83, APC-conjugated anti-CD14, PE-conjugated anti-CD40, PE-conjugated anti-CCR7, APC-Cy7-conjugated anti-HLA-DR (all from BD Biosciences, San Diego, CA). To optimize the staining, monoclonal antibodies were titrated in preliminary experiments. 7-Amino-actinomycin D (Sigma Aldrich, St Louis, MO) was used to exclude non-viable cells from the analysis. Fluorescence minus one samples were used as negative controls.

Cells were collected and analysed using a FACSCanto II (Becton Dickinson, San Jose, CA) flow cytometer, and data analysis was performed using FACSDIVA (Becton Dickinson) and FLOWJO software (TreeStar Inc., Ashland, OR).

Statistical analysis

Statistical analyses were performed using GRAPHPAD PRISM 5.0a (GraphPad Software, San Diego, CA). All statistical analyses assumed a two-sided significance level of $P < 0.05$. The Student's paired t-test was used for comparisons between groups.

Results

Different levels of CD91 identify two moDC subpopulations

It has been shown that CD91 is expressed on moDCs. $9,15-17$ However, whether CD91 is expressed differently in moDC subpopulations is still unknown. To investigate the expression of CD91 receptor, we cultured human monocytes with granulocyte–macrophage colony-stimulating factor and interleukin-4 for 5 days to generate immature moDCs that were subsequently characterized by flow cytometry. As expected, 15 we observed that during the differentiation of monocytes into moDCs characterized by down-regulation of CD14 and up-regulation of CD1a (Fig. 1a), a partial down-regulation of CD91

occurred in parallel (Fig. 1b). Moreover, our results clearly showed that $CD91⁺$ moDCs were segregated into two distinct populations: CD91 was highly expressed in a small proportion of total moDCs (CD91bright moDCs) (mean \pm SE, 1.19 \pm 0.28%), whereas the majority of

Figure 1. The expression of CD91 is partially down-regulated during differentiation of monocytes into monocyte-derived dendritic cells (moDCs). Monocytes obtained by plastic adhesion from freshly isolated peripheral blood mononuclear cells were stained for surface expression of CD14, CD1a and CD91 before and after their differentiation into moDCs obtained by 5 day-culture with recombinant human granulocyte– macrophage colony-stimulating factor and recombinant human interleukin-4. (a) Dot plots showing the gating strategy and the expression of CD14 and CD1a on monocytes (upper row) and moDCs (lower row). (b) Histogram showing the mean fluorescence intensity (MFI) of CD91 expression on monocytes (grey line), moDCs (black line), and fluorescence minus one (FMO) negative control (dotted line). One representative experiment is shown.

Figure 2. Different levels of CD91 identify two monocyte-derived dendritic cell (moDC) subpopulations. Immature moDCs were stained for surface expression of CD91 and activation/maturation markers. (a) Representative flow cytometric analysis showing the gating strategy used to define CD91^{dim} and CD91^{bright} moDCs. (b) Graphs show the mean fluorescence intensity (MFI) of CD80, CD86, CD40, CD83, CCR7 and HLA-DR molecules in CD91^{dim} (white bars) and CD91^{bright} (black bars) moDCs from 11 independent experiments. Data shown as mean \pm SE. $*P < 0.05, **P < 0.01.$

cells expressed low levels of CD91 receptor (CD91^{dim} moDCs) (Fig. 2a).

$CD91^{bright}$ moDCs show an activated phenotype

The ability of DCs to activate immune responses depends on their maturation status and the expression of co-stimulatory molecules. To investigate whether CD91^{bright} and CD91^{dim} subsets differ in term of activation/maturation, we evaluated their surface expression of CD80, CD86, CD40, CD83, HLA-DR molecules. As shown in Fig. 2(b), $CD91^{bright}$ moDCs expressed significantly higher levels of all the activation and maturation markers compared with CD91^{dim} moDCs: CD80 [mean fluorescence intensity (MFI), mean \pm SE, 905 \pm 116 versus 647 \pm 95, P < 0.05), CD86 (MFI, 4864 \pm 861 versus 3093 \pm 473, P = 0.05), CD40 (MFI 1287 \pm 157 versus 813 \pm 125, $P < 0.01$), CD83 (MFI, 1235 \pm 99 versus 865 \pm 120, $P < 0.01$) and HLA-DR (MFI, 32 273 \pm 6815 versus 20 578 \pm 6862, $P < 0.05$). Also shown in Fig. 2(b), CD91^{bright} moDCs expressed significantly higher levels of CCR7 than CD91^{dim} moDCs (MFI 3814 \pm 649 versus 1520 ± 185 , $P = 0.01$).

rHNP-1 and rHBD-1 increase the proportion of the CD91^{bright} moDC subset

Based on our previous findings indicating that moDC treatment with human defensins up-regulates the surface

expression of CD91, we next evaluated whether in vitro stimulation with rHNP-1 and rHBD-1 may indeed positively affect the frequency of CD91bright moDCs. Interestingly, we observed that the proportion of CD91^{bright} moDCs increased after rHNP-1 stimulation by $45.4 \pm 15.4\%$ (mean \pm SE, $P < 0.01$) and after rHBD-1 stimulation by $47.0 \pm 17.4\%$ ($P < 0.05$). On the contrary, the proportion of CD91^{bright} moDCs decreased after treatment with the CD91 ligand hsp 90 by $-32.7 \pm 6.2\%$ ($P < 0.05$) and after LPS stimulation by $-34.0 \pm 8.9\%$ ($P < 0.05$) (Fig. 3a,b).

rHNP-1 and rHBD-1 increase the activation of CD91^{bright} moDC subset

Next, we performed experiments to test the ability of CD91^{bright} moDCs to respond to defensins. First, we confirmed previously published data that both defensins promoted the activation and maturation of total moDCs: rHNP-1 versus unstimulated CD80 (MFI, mean \pm SE, 739 \pm 37 versus 611 \pm 39, P < 0.05), CD86 (3482 \pm 1162 versus 2984 \pm 557, P = ns), CD40 (1432 \pm 101 versus 1018 ± 174 , $P < 0.05$), CD83 (1288 ± 252 versus 738 \pm 165, P < 0.05) and HLA-DR (24 026 \pm 3209 versus 14 490 \pm 3905, $P < 0.05$; rHBD-1 versus unstimulated CD80 $(830 \pm 64$ versus 611 ± 39 , $P < 0.05$), CD86 $(5327 \pm 1589 \text{ versus } 2984 \pm 557, P = \text{ns})$, CD40 (1549 ± 145 versus 1018 \pm 174, $P < 0.05$), CD83 (1229 \pm 253 versus 738 ± 165 , $P = \text{ns}$ and HLA-DR (26 580 \pm 4690

Figure 3. Recombinant human neutrophil peptide-1 (rHNP-1) and recombinant human β defensin-1 (rHBD-1) increase the proportion of the CD91bright monocyte-derived dendritic cell (moDC) subset. Immature moDCs were incubated with 100 ng/ml rHNP-1, 500 ng/ml rHBD-1, 20 lg/ml heat-shock protein 90 (hsp 90) or 100 ng/ml lipopolysaccharide (LPS). After 18 hr, the cells were analysed by flow cytometry to measure cell-surface CD91 expression. (a) Representative flow cytometric analysis showing the increased proportion of CD91^{bright} moDC subset after rHNP-1 and rHBD-1 stimulation compared with the decreased proportion after hsp 90 and LPS challenge. (b) Change of CD91^{bright} moDCs frequency after rHNP-1, rHBD-1, hsp 90 and LPS compared with unstimulated conditions. Data shown as mean \pm SE from six independent experiments. $*P < 0.05$, $*P < 0.01$.

versus $14\,490 \pm 3905$, $P < 0.05$) (Fig. 4). We further analysed the in vitro effects of rHNP-1, rHBD-1 and LPS stimulation on CD91^{bright} compared with CD91^{dim} moDCs. As also shown in Fig. 4, rHNP-1 induced significantly higher levels of all the activation markers on CD91^{bright} moDCs compared with $CD91^{\text{dim}}$ moDCs (MFI on $CD91^{\text{bright}}$ versus CD91^{dim} cells, CD80: 864 \pm 51 versus 721 \pm 45, P < 0.05; CD86: 4373 ± 1260 versus 3463 ± 1154 , $P < 0.05$; CD40: 2164 \pm 289 versus 1367 \pm 132, $P < 0.05$; HLA-DR: 40 483 \pm 5467 versus 23 769 \pm 7002, P < 0.05). Similar results were observed when moDCs were stimulated with rHBD-1 (CD80: 1216 \pm 264 versus 1012 \pm 190, P < 0.05; CD40: 1956 \pm 138 versus 1323 \pm 196, P < 0.05; CD83: 1775 ± 279 versus 1224 ± 253 , $P < 0.05$; HLA-DR:
41 818 \pm 5536 versus 26 457 \pm 3637, $P < 0.05$). $26\;457 \pm 3637$, Conversely, when moDCs were stimulated with LPS, no significant differences in the expression of any activation marker were observed between CD91bright and CD91dim cells. Overall, these results indicate that CD91^{bright} moDCs that had undergone stimulation with rHNP-1 and rHBD-1 maintained their higher state of activation/maturation

compared with CD91^{dim} moDCs while this property was lost upon LPS stimulation.

Discussion

Dendritic cells are potent antigen-presenting cells that sense environmental stimuli through a wide repertoire of receptors expressed on their surface.¹⁸ We and others previously described that moDCs express CD91, a polyfunctional receptor that binds several molecules including defensins. $4,9,15-17$ In this study we confirmed our previous observations and, by using multi-parametric flow cytometry analysis, we showed for the first time to our knowledge the presence of two distinct subsets of moDCs based on their CD91 expression: CD91^{bright} and CD91^{dim} moDCs. These subsets were characterized by a different phenotype. In fact, although $CD91^{bright}$ cells represented a small proportion of moDCs, this subset showed greater expression of activation/maturation markers compared with CD91^{dim} moDCs, suggesting that these two subsets could have different roles in the

Figure 4. Human neutrophil peptide-1 (HNP-1) and human β defensin-1 (HBD-1) increase the activation of CD91^{bright} monocyte-derived dendritic cell (moDC) subset. Immature moDCs were stimulated with 100 ng/ml recombinant HNP-1, 500 ng/ml recombinant HBD-1, or 100 ng/ ml lipopolysaccharide (LPS). After 18 hr, the cells were analysed by flow cytometry to measure cell-surface activation and maturation markers expression on total moDCs (shaded bars), CD91^{dim} moDCs (white bars) and CD91^{bright} moDCs (black bars). Graphs show the mean fluorescence intensity (MFI) of CD80, CD86, CD40, CD83 and HLA-DR molecules. Data shown as mean \pm SE from six independent experiments and analysed by the paired t-test. $^{*}P$ < 0.05 stimulated total moDCs compared to unstimulated (NS) cells; $^{*}P$ < 0.05: CD91^{dim} moDCs compared with CD91bright moDCs.

immune response. This hypothesis may be further supported by a previous study that correlated the full maturation status of plasmacytoid DCs with a greater increase of CD91 expression and increased CD91 ligandbinding capacity.¹⁹ Interestingly, in our study in vitro stimulation with defensins increased the proportion of CD91^{bright} moDCs and preserved their higher activation/ maturation status compared with $CD91^{dim}$ moDCs, suggesting that CD91^{bright} moDCs may be more sensitive to CD91 stimulation. Notably, defensins appeared unique in their ability to up-regulate their own receptor, as CD91 up-regulation was not shared by hsp 90, another CD91 ligand highly expressed in eukaryotic cells. Indeed, in accordance with one of two previous studies addressing the effects of heat-shock proteins on the expression of CD91 on different antigen-presenting cells, $19,20$ we observed that hsp 90 decreased the frequency of CD91^{bright} moDCs. Not even the ability of defensins to up-regulate CD91 was shared by LPS, a common pro-inflammatory molecule that signals through toll-like receptor 4. Indeed, LPS decreased the frequency of CD91^{bright} moDCs. This finding is in accordance with our previous study, 15 reporting LPS-induced down-regulation of CD91 expression on moDCs, possibly supported by shedding of CD91 from cell surface.²¹

Although there is evidence that CD91-mediated immune activation can represent a powerful mechanism of immune regulation, the pathway(s) regulating expression of CD91 have not been fully defined so far. CD91 is up-regulated on monocytes of people who are HIVpositive long-term non-progressors, it may enhance cross-presentation of HIV antigens and may in turn enhance the stimulation of activated anti-HIV cytotoxic T lymphocytes.²² Moreover, the pivotal role of CD91 regulation on DCs has been demonstrated in the context of primary effusion lymphoma: DC activation was completely inhibited when these cells were pre-treated with a neutralizing antibody directed against $CD91₁¹¹$ suggesting that the up-regulation of CD91 in cancer patients may represent an additional strategy to improve DC activation and subsequent stimulation of a specific CD8⁺ T-cell response. Notably, we observed that CD91^{bright} cells expressed significantly higher levels of CCR7 than $CD91^{\text{dim}}$ cells, suggesting that the CD91^{bright} moDC subset might preferentially migrate to secondary lymph nodes, where they may elicit adaptive immune responses.

Our results are intriguing because of the plasticity of dendritic cells in the setting of different immune responses. Hence, we could hypothesize a differential role of CD91^{bright} and CD91^{dim} moDC subsets in activating the immune response, possibly involving a differential production of pro-inflammatory and regulatory cytokines. These two cell subsets may also differentially privilege MHC-I rather than MHC-II presentation. Because of the low frequency of CD91bright moDCs in our cultures, we could not separate these populations to compare their antigen-presenting ability. Experiments of cell depletion were not undertaken, as they may provide only approximate and indirect functional data. Additional studies will be required in the future to further characterize the role of CD91^{bright} and CD91^{dim} DCs in the immune response.

Taken together, our results suggest that moDCs preferentially respond to defensins by increasing the frequency of CD91^{bright} moDCs and promoting a higher state of activation/maturation of these cells compared with CD91dim moDCs. The identification of a subset of moDCs expressing high levels of CD91 and molecules involved in DC–T-cell interaction suggests that these CD91^{bright} cells may be particularly devoted to the activation of adaptive immune responses against antigens that may be delivered to DCs through CD91 ligands, in particular defensins. These results may provide new insights into the immune potentiating effects of defensins that are currently under intense investigation as possible new therapeutic agents aimed to potentiate antimicrobial and anti-tumour immunity.

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Disclosure

The authors declare no conflicting financial interests.

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