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Kidney International (2012) 81, 598-599; doi:10.1038/ki.2011.441

CKD-associated atherosclerosis and monocyte heterogeneity

To the Editor: We noticed with great interest the mini review 'Novel inflammatory mechanisms of accelerated atherosclerosis in kidney disease' by Swaminathan and Shah.¹

Although we commend the authors for their efforts to highlight the substantial impact of monocyte heterogeneity in chronic kidney disease-associated atherosclerosis, we are surprised that the authors exclusively relied on data obtained from murine experiments, ignoring the limits of homology between murine and human monocyte heterogeneity.

Notably, in mice, Gr1⁺/Ly6C^{high} monocytes (the putative homologs of human CD14⁺⁺CD16⁻ monocytes) are proinflammatory, and they are more prone to enter the atherosclerotic plaque than Gr1⁻/Ly6C^{low} monocytes.¹ In contrast, in humans, CD16-positive monocytes (CD14⁺⁺CD16⁺ and CD14⁺CD16⁺⁺ cells, which are counterparts of murine Gr1⁻/Ly6C^{low} monocytes) are considered proinflammatory cells by most groups (as reviewed in Ziegler-Heitbrock²). Moreover, ample circumstantial evidence (as discussed in Rogacev et al.³) and clinical data demonstrate that CD14⁺⁺CD16⁺ monocytes^{3,4}—but not CD14⁺⁺CD16⁻ monocytes-are the relevant monocytes in human atherosclerosis. The importance of the CD14⁺⁺CD16⁺ monocyte subsets in nephrology is underscored by the profoundly elevated counts of these cells in dialysis patients⁴ and their activation during dialysis (referenced in Rogacev et al.³).

Taken together, we agree with the authors that knowledge of monocyte heterogeneity is of major relevance to both clinicians and researchers in nephrology. We kindly disagree with the uncritical transfer of murine data to human (patho)physiology.

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Kidney International (2012) 81, 599; doi:10.1038/ki.2011.433

The Authors Reply: We thank Rogacev *et al.*¹ for their interest in our mini review 'Novel Inflammatory Mechanisms of Accelerated Atherosclerosis in Kidney Disease.'² While we appreciate and acknowledge the important points raised by the authors regarding homology of monocyte heterogeneity between mice and humans, we suggest several additional aspects that are worth considering.

Although we agree that Gr1^{+/}/Ly6C^{hi} are pro-inflammatory monocytes and are putative homologs of human CD14⁺CD16⁻ monocytes, in many settings, depending on local and systemic cues, they can also differentiate into a variety of macrophage and dendritic cell (DC) subtypes that could inhibit immune response.³

Pertaining to atherosclerosis, although Ly6C^{hi} monocytes more efficiently accumulate in atherosclerotic plaques, Ly6C^{lo} anti-inflammatory monocytes were particularly prone to developing into plaque cells that express the DC marker CD11c.⁴ Similarly, in human atheroma, macrophages express c-fms (macrophage colony-stimulating factor receptor) but not inflammatory cytokines such as interleukin-6,⁵and CD14⁺CD16⁻ rather than CD16⁺ monocytes correlate with worse myocardial salvage after myocardial infarction.⁶

More recently, gene expression profiling has identified close similarities between murine and human monocyte subsets in more than 100 genes, suggesting similarity between $Ly6C^{hi}$ monocytes and human $CD14^+CD16^-$ monocytes, and between $Ly6C^{lo}$ anti-inflammatory mouse monocytes and $CD16^+$ human monocytes.⁷ This has led the authors to refer to monocytes as $CD14^+CD16^-$ and $CD16^+$ subsets in both species. However, it is important to point out some notable differences that the authors observed between the homologous human and mouse monocyte subsets, especially in PPAR- γ and phagocytic receptor expression.

Thus, origin, differentiation, and plasticity of different monocyte subsets in mice, and the corresponding cell populations in humans and their independent role in health and disease, remain to be fully elucidated.

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Kidney International (2012) 81, 599-600; doi:10.1038/ki.2011.436