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Corrigendum: The mast cell: A Janus in kidney transplants

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A Corrigendum on

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In the published article, there was an error in the legend for **Figures 1, 2** as published. The legends of **Figures 1, 2** were switched. The corrected legend appears below.

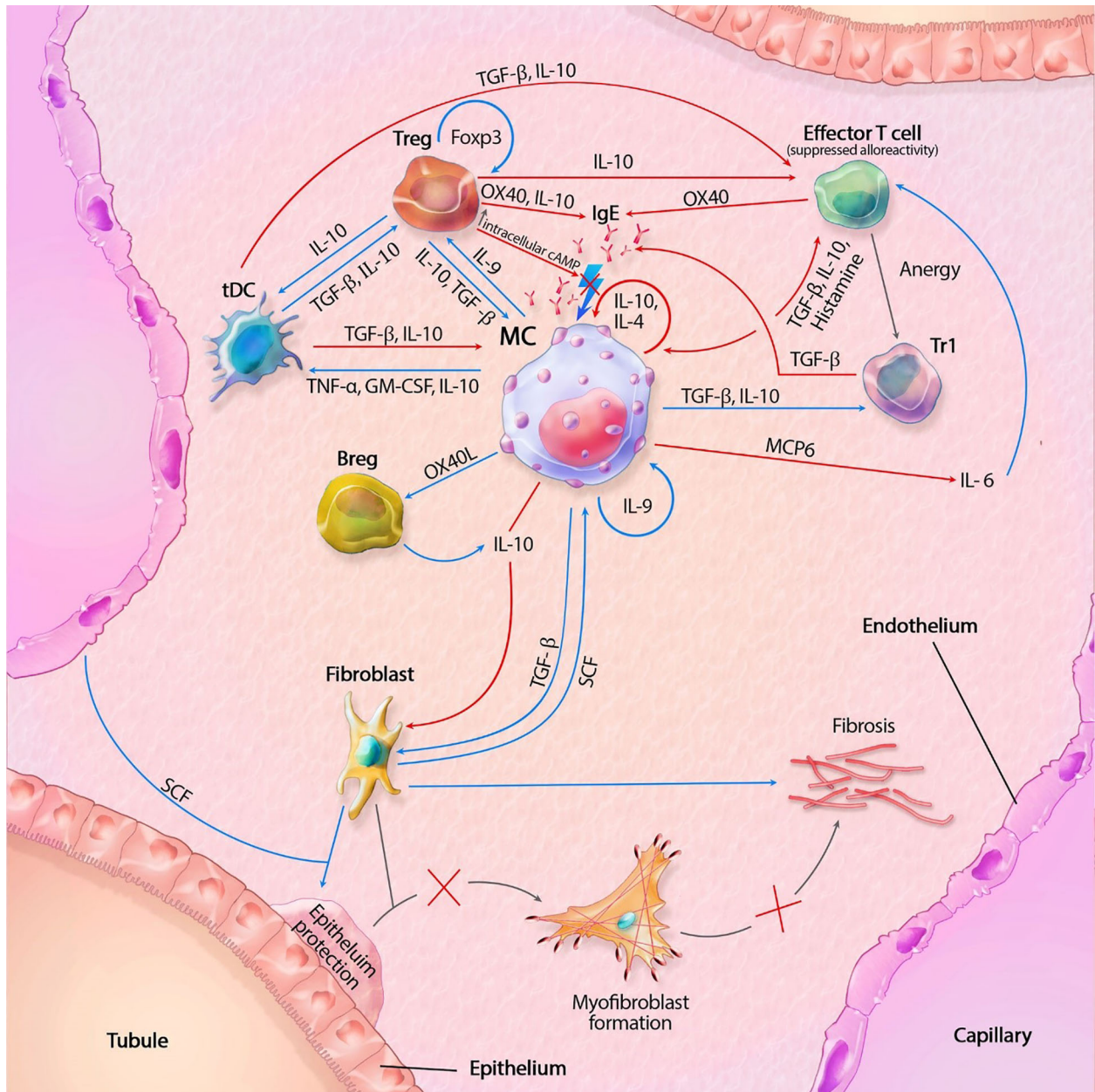


FIGURE 1
 Mast cell (MC) interactions within the transplant during tolerance. FcεRI activity is inhibited by TGF-β, IL-10 and OX40 ligation. Tregs also inhibit degranulation by lowering intracellular Ca²⁺ levels through increased cAMP. IL-10 suppresses alloreactivity within CD4+ and CD8+ T cells and promote anergy and regulatory functions of CD4+ T cells. IL-10 mediated inhibition of fibroblasts also inhibit subsequent formation of myfibroblasts. IL-10 with co-stimulation of IL-4 decrease MC proliferation, while IL-9 increases proliferation. GM-CSF, granulocyte-macrophage colony-stimulating factor; IL, interleukin; MCP6, mat cell protease 6; SCF, stem cell factor; tDC, tolerogenic dendritic cell; TGF-β, tissue growth factor beta; TNF-α, tissue necrotic factor alpha; Tr1, regulatory T cell type 1 (induced); Treg, regulatory T cell (natural); Blue lines symbolize activating pathways, red lines inhibitory pathways, gray lines symbolize subsequent events. Lighting icons are used in the most profound activation patterns, which are inhibited in tolerogenic environments.

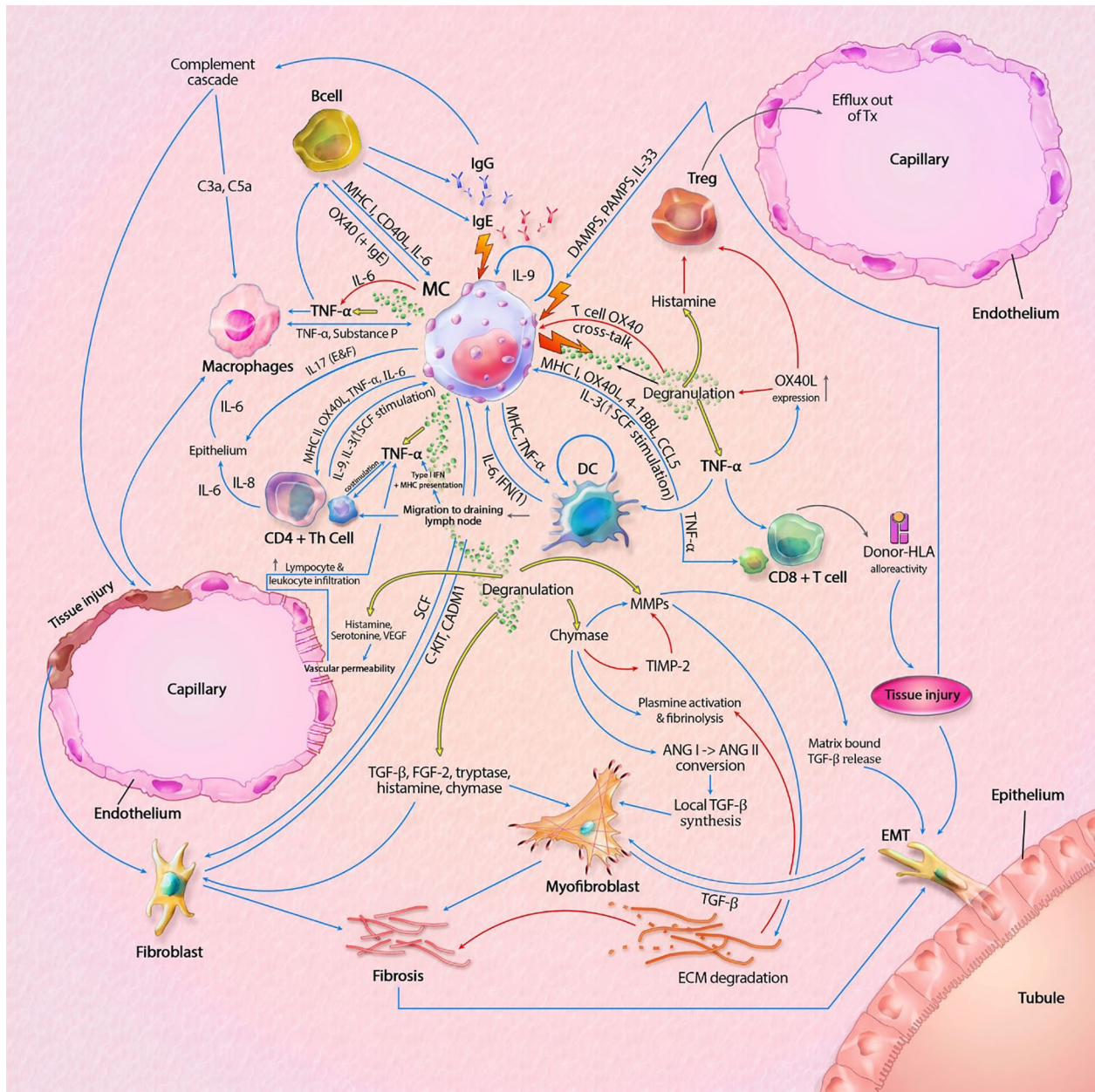


FIGURE 2

Mast cell (MC) interactions within the graft during rejection. Pathways can include both cytokines (like TNF- α) and membrane bound interaction (like MHC I-TLR interaction). MC-T cell interaction through OX40L-OX40 cross-linking inhibits MC degranulation, represented by the inhibitory pathway towards degranulation. Innate immune cells can also result in tissue injury, which is not shown in this model. Interaction between APCs, T cells and B cells, resulting in antigen production is also not shown in this model. The model shows almost no inhibitory pathways, explaining the progressive state of fibrosis within KTx even when immunosuppressive drugs are taken. Detailed description of the model can be found within the text. ANG, angiotensin; C3a/C5a, complement component; ECM, extracellular matrix; EMT, epithelial-mesenchymal transition; FGF-2; fibroblast growth factor-2; Ig, immunoglobulin; IL, interleukin; MHC, major histocompatibility complex; MMPs, matrix metalloproteinase; SCF, stem cell factor; tDC, tolerogenic dendritic cell; TGF- β , tissue growth factor beta; Th cell, T helper cell; TIMP-2, tissue inhibitor of metalloproteinase-2; TNF- α , tissue necrotic factor alpha; Treg, regulatory T cell (natural); VEGF, Vascular Endothelial Growth Factor. Blue lines symbolize activating pathways, red lines inhibitory pathways, yellow lines represent pre-formed mediators within MCs. Grey lines represent subsequent events. Lighting icons are used in the most profound activation patterns.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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