

Supplementary Informations

Kidney inflammaging is promoted by CCR2+ macrophages and tissue-derived micro-environmental factors

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Supplementary Methods

Supplementary Tables

Table S1

Table S2

Supplementary Methods

Histochemistry

Kidneys were fixed for 6h in 3.4% PFA then in 70% Ethanol and included in paraffin. Tissue sections were stained with sirius red and digitized with a Hamamatsu NanoZoomer. Stained areas and total surface from each section were determined using color-based thresholding and quantified using Fiji software as previously described (1).

Biochemistry

Creatinin and blood urea nitrogen (BUN) were measured in plasma samples with ABX Pentra reagents using the Pentra C400 analyzer (HORIBA Medical).

References

1. Laroumanie F, *et al.* (2014) CD4+ T cells promote the transition from hypertrophy to heart failure during chronic pressure overload. *Circulation* 129(21):2111-2124.

Table S1. List of the antibodies used for DE cyt analysis

name	alternative name	clone	purchased from
Set A			
CD45 APC-Cy7		30-F11	Biologend
Ly6C PB		HK1.4	Biologend
MHCII FITC	I-A/I-E	M5/114.15.2	Biologend
CCR2 PE		FAB5538	R&D Systems
CD14 PercPCy5.5		Sa-2-8	eBioscience
CD64 APC	Fc gamma RI	X54-5/7.1	Biologend
Set B			
Sca-1 PB	Ly6A/E	D7	Biologend
CD31 PercPCy5.5	PECAM1	390	Biologend
CD45 APC-Cy7		30-F11	Biologend
CD326 FITC	EpCAM	G8.8	Biologend
mEFSK4 PE	anti-Feeder Cells	mEF-SK4	Miltenyi Biotec
CD73 PE-Cy7	5'ecto-nucleotidase	TY/11.8	Biologend
CD140b APC	PDGFR β	APB5	Biologend

Table S2. Sequences of the primers used for RT-QPCR

gene	Primer sequences (5' -> 3')	Ref_Seq
<i>Nt5e</i>	F: CTTCATGAACATCCTGGGCT R: AACGTTTCTGAGGAGGGGAT	NM_011851
<i>Cdkn1c</i>	F: ATCACCAATCAGCCAGCAGAA R: CTACGCGCTATCACTGGGAAG	NM_009876.4
<i>Cdkn2c</i>	F: GGGGGACCTAGAGCAACTTAC R: CTCCGGATTTCCAAGTTTCA	NM_007671.2
<i>Cx3cl1</i>	F: TGGCTTTGCTCATCCGCTATCAG R: CGTCTGTGCTGTGTCGTCTCC	NM_009142.3
<i>Cxcl2</i>	F: CGCTGTCAATGCCTGAAG R: GCGGTCACACTCAAGCTCT	NM_009140.2
<i>Ccl2</i>	F: AACCTGGATCGGAACCAAAT R: TACGGGTCAACTTCACATTCAA	NM_011333.3
<i>Il6</i>	F: GAGGATACCACTCCCAACAGACC R: AAGTGCATCATCGTTGTTTCATACA	NM_031168
<i>Il1a</i>	F: TTGGTTAAATGACCTGCAACA R: GAGCGCTCACGAACAGTTG	NM_010554.4
<i>Il10</i>	F: TTCAGCCAGGTGAAGACTTTCT R: GCTTGGCAACCCAAGTAACC	NM_010548.2
<i>Tgfb1</i>	F: AGGGCTACCATGCCAACTTCT R: CCGGGTTGTGTTGGTTGTAGA	NM_011577.2
<i>Vegfa</i>	F: CAGCAGATGTGAATGCAGACCAA R: CTTTCTCCGCTCTGAACAAGGC	NM_001025250.3
<i>Cxcl12</i>	F: CTCAACTCCAACTGTGCC R: TTGGGCTGTTGTGCTTACTTG	NM_021704.3
<i>Cxcr4</i>	F: CCATGGAACCGATCAGTGTGA R: CAGGGTTCCTTGTGGAGTCA	NM_009911.3
<i>Igf1</i>	F: GCTCTTCAGTTCGTGTGTGGAC R: AGCCTGTGGGCTTGTGAAGTA	NM_010512.4
<i>Ccr2</i>	F: CTTGGGAATGAGTAACTGTGTGA R: AATGACAGGATTAATGCAGCAGTGT	NM_009915.2
<i>Tnfa</i>	F: AGCCGATTTGCTATCTCATACCA R: GGGCTCATACCAGGGTTTGA	NM_013693
<i>Il1b</i>	F: CTGCACTACAGGCTCCGAGAT R: TGTTGGTTGATATTCTGTCCATTG A	NM_008361.4
<i>Il12b</i>	F: CGGACGGTTCACGTGCTC R: CACATGTCCTGCCCCGAGAGT	NM_001303244.1

<i>Cdkn2a</i>	F: CCGAACTCTTTCGGTCGTACCC R: CTGCTACGTGAACGTTGCCCA	NM_001040654.1
<i>Rplp0</i>	F: GCTTCATTGTGGGAGCAGAC R: ATGGTGTTCTTGCCCATCAG	NM_007475.5
<i>Trp53</i>	F: TCCTGGCTGTAGGTAGCGACT R: ATCCGACTGTGACTCCTCCAT	NM_001127233.1 NM_011640.3