SUPPLEMENTAL FIGURE 1

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SUPPLEMENTAL TABLES 1 - 2

Article Title:	Pathways analysis of differential gene expression induced by engrafting doses of total body irradiation for allogeneic bone marrow transplantation in mice
Journal Name:	Immunogenetics
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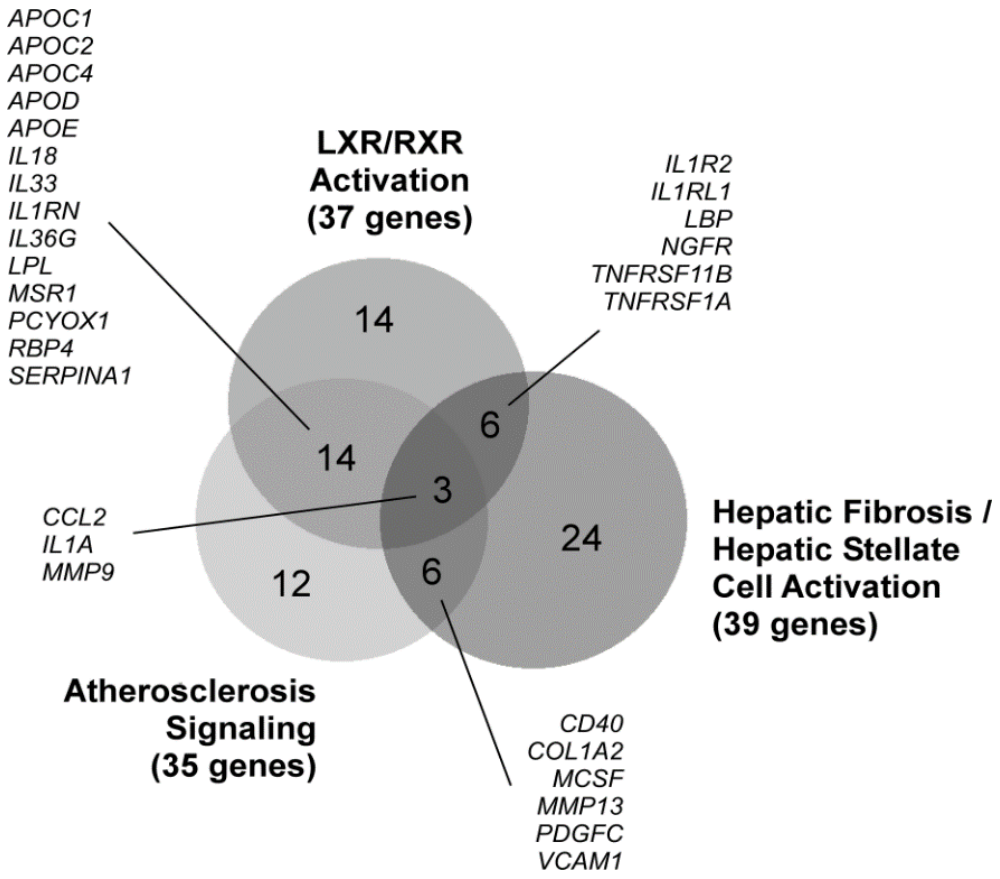
Supplemental Figure 1



Supplemental Fig. 1. Three-way Venn-diagrams of genes enriched in the LXR/RXR activation, atherosclerosis signaling, and hepatic fibrosis / hepatic stellate cell activation canonical pathways.

Supplemental Table 1. Real-time PCR primers.

Primer	Sequence (5' to 3')
H2-Ab1	
Forward	GCGACGTGGGCGAGTACC
Reverse	CATTCCGGAACCAGCGCA
H2-Oa	
Forward	AGACCAGCTTCTACTCTCAGCCTAAC
Reverse	ACAGATCAGGGTCTCTATGGTGTC
H2-Ob	
Forward	AGAGAATTTTGTGATTCAGGC
Reverse	TCTGGATACACTGTCACCTCTG
H2-DMb1	
Forward	TTGGGGTGCTGTATCCATGGGCCGAAAATT
Reverse	GAAGCCCCAGACGTAGCAGGCCAGCATCACC
H2-DMb2	
Forward	TTGGGGTGCTGTCTAGATTGGCTGAAATAA
Reverse	GAAGCCCCAGACGTAGCAGGCCAGCATCACG
Cd80	
Forward	AAGTGGTGCTGTCTGTCATTG
Reverse	CGGAAGCAAAGCAGGTAATC

Supplemental Table 2. Differentially expressed genes in canonical pathways shared between BALB.K and B10.BR mice in the spleen after conditioning with engrafting TBI doses. [a]

LXR/RXR Activation (37 genes)		Complement System (16 genes)		Atherosclerosis Signaling (35 genes)		Hepatic Fibrosis / Hepatic Stellate Cell Activation (39 genes)		Heme Biosynthesis II (8 genes)	
Gene Symbol	Fold Change	Gene Symbol	Fold Change	Gene Symbol	Fold Change	Gene Symbol	Fold Change	Gene Symbol	Fold Change
ABCG1	↑ 2.61	C2	↑ 2.80	ALOXE3	↑ 2.02	BAX	↑ 2.06	ALAD	↓ -6.34
ABCG4	↓ -57.13	C3	↑ 2.36	APOC1	↑ 4.65	CCL2	↑ 9.90	ALAS2	↓ -4.12
AHSG	↓ -3.07	C6	↑ 4.27	APOC2	↑ 3.32	CD40	↓ -2.36	CPOX	↓ -7.62
APOC1	↑ 4.65	C1QA	↑ 3.39	APOC4	↑ 2.13	COL1A2	↑ 2.13	FECH	↓ -3.99
APOC2	↑ 3.32	C1QB	↑ 3.67	APOD	↑ 3.95	EDNRA	↑ 2.44	HMBS	↓ -12.52
APOC4	↑ 2.13	C1QC	↑ 2.81	APOE	↑ 2.29	EDNRB	↑ 3.28	PPOX	↓ -3.19
APOD	↑ 3.95	C1R	↑ 3.10	CCL11	↑ 9.90	EGFR	↑ 2.69	UROD	↓ -6.20
APOE	↑ 2.29	C1S	↑ 3.09	CCL2	↑ 2.70	FGFR1	↑ 2.51	UROS	↓ -8.63
APOH	↑ 4.29	C4B	↑ 3.40	CD40	↓ -2.36	FLT1	↑ 2.38		
ARG2	↑ 3.25	C5AR1	↑ 2.26	COL18A1	↑ 2.27	FLT4	↑ 2.10		
C3	↑ 2.36	CD59	↓ -7.45	COL1A2	↑ 2.13	HGF	↑ 2.80		
C4B	↑ 3.40	CFB	↑ 2.38	IL18	↑ 2.12	IGF1	↑ 2.73		
CCL2	↑ 9.90	CFH	↑ 2.53	IL1A	↑ 2.39	IGFBP3	↑ 3.29		
FDFT1	↑ 2.16	CFI	↑ 4.07	IL1RN	↑ 2.22	IGFBP4	↑ 2.57		
IL18	↑ 2.12	CR2	↓ -2.43	IL33	↑ 2.22	IGFBP5	↑ 2.34		
IL33	↑ 2.39	SERPING1	↑ 3.07	IL36G	↑ 2.32	IL1A	↑ 2.32		
IL1A	↑ 2.22			ITGA4	↓ -2.50	IL1R2	↑ 2.22		
IL1R2	↑ 4.80			LPL	↑ 2.72	IL1RL1	↑ 4.80		
IL1RL1	↓ -14.24			MCSF	↑ 2.32	IL4	↓ -14.24		
IL1RN	↑ 2.22			MMP13	↑ 11.54	KDR	↑ 2.02		
IL36G	↑ 2.32			MMP3	↑ 2.07	LBP	↑ 2.61		
ITIH4	↑ 3.94			MMP9	↑ 6.19	LEPR	↑ 3.35		
KNG1	↑ 2.18			MSR1	↑ 4.56	MCSF	↑ 2.32		
LBP	↑ 2.61			PCYOX1	↑ 2.27	MMP13	↑ 7.41		
LPL	↑ 2.72			PDGFC	↑ 4.22	MMP2	↑ 2.07		
MMP9	↑ 2.07			PLA2G1B	↑ 3.53	MMP9	↑ 6.19		
MSR1	↑ 4.56			PLA2G2C	↓ -3.57	MYH11	↑ 6.85		
NGFR	↑ 2.88			PLA2G2D	↑ 3.56	MYH7	↑ 2.47		
NR1H3	↑ 2.29			PLA2G4F	↑ 2.28	NGFR	↑ 2.88		
PCYOX1	↑ 2.27			PLA2G7	↑ 5.12	PDGFC	↑ 4.22		
PON3	↑ 2.10			RBP4	↑ 2.83	PDGFRA	↑ 3.04		
RBP4	↑ 2.83			SELP	↑ 2.20	PDGFRB	↑ 2.26		
SAA2	↑ 26.88			SERPINA1	↑ 2.43	PGF	↑ 2.77		
SERPINA1	↑ 2.43			TNFSF12	↑ 2.53	STAT1	↑ 2.62		
TF	↑ 2.22			VCAM1	↑ 3.93	TIMP1	↑ 2.47		
TNFRSF11B	↑ 2.00					TIMP2	↑ 3.16		
TNFRSF1A	↑ 2.38					TNFRSF11B	↑ 2.00		
						TNFRSF1A	↑ 2.38		
						VCAM1	↑ 3.93		

Abbreviations: IPA, Ingenuity Pathways Analysis; TBI, total body irradiation.

[a] TBI doses were 400 cGy for BALB.K and 500 cGy for B10.BR mice.

Supplemental Table 2 (Continued). Differentially expressed genes in canonical pathways shared between BALB.K and B10.BR mice in the spleen after conditioning with engrafting TBI doses. [a]

B Cell Development (13 genes)		Altered T Cell and B Cell Signaling in Rheumatoid Arthritis (21 genes)		Matrix Metalloproteases (13 genes)		Heme Biosynthesis from Uroporphyrinogen-III I (4 genes)		Communication between Innate and Adaptive Immune Cells (18 genes)	
Gene Symbol	Fold Change	Gene Symbol	Fold Change	Gene Symbol	Fold Change	Gene Symbol	Fold Change	Gene Symbol	Fold Change
CD19	↓ -2.95	CD40	↓ -2.36	HSGP2	↑ 2.11	CPOX	↓ -7.62	CD40	↓ -2.36
CD40	↓ -2.36	CD80	↑ 2.29	LRP1	↑ 3.53	FECH	↓ -3.99	CD80	↑ 2.29
CD43	↓ -2.38	CD79A	↓ -2.88	MMP2	↑ 7.41	PPOX	↓ -3.19	CD8A	↓ -2.78
CD80	↑ 2.29	CD79B	↓ -2.62	MMP3	↑ 11.54	UROD	↓ -6.20	CD8B	↓ -5.17
CD79A	↓ -2.88	FCER1G	↑ 2.44	MMP9	↑ 2.07			CXCL10	↑ 3.39
CD79B	↓ -2.62	H2-DMB2	↓ -2.75	MMP13	↑ 6.19			FCER1G	↑ 2.44
DNTT	↓ -2.77	H2-OA	↓ -2.01	MMP14	↑ 2.40			IL4	↑ 2.32
H2-DMB2	↓ -2.75	H2-OB	↓ -3.92	MMP15	↑ 2.39			IL15	↑ 3.18
H2-OA	↓ -2.01	IL4	↑ 2.32	MMP25	↑ 2.81			IL18	↑ 2.12
H2-OB	↓ -3.92	IL15	↑ 3.18	SDC2	↑ 2.24			IL33	↑ 2.39
IGKC	↓ -2.58	IL18	↑ 2.12	THBS2	↑ 3.09			IL12B	↑ 2.43
IL7	↑ 2.53	IL33	↑ 2.39	TIMP1	↑ 2.47			IL1A	↑ 2.22
RAG1	↓ -3.62	IL12B	↑ 2.43	TIMP2	↑ 3.16			IL1RN	↑ 2.22
		IL1A	↑ 2.22					IIL36G	↑ 2.32
		IL1RN	↑ 2.22					TLR7	↑ 2.64
		IL36G	↑ 2.32					Tlr13	↑ 2.27
		MCSF	↑ 2.32					TNFRSF13C	↓ -3.35
		TLR7	↑ 2.64					TNFSF13B	↑ 3.59
		Tlr13	↑ 2.27						
		RNFRSF13C	↓ -3.35						
		TNFS13B	↑ 3.59						

Abbreviations: IPA, Ingenuity Pathways Analysis; TBI, total body irradiation.

[a] TBI doses were 400 cGy for BALB.K and 500 cGy for B10.BR mice.