

Supplementary Materials for  
**Nuclear factor  $\kappa$ B overactivation in the intervertebral disc leads to  
macrophage recruitment and severe disc degeneration**

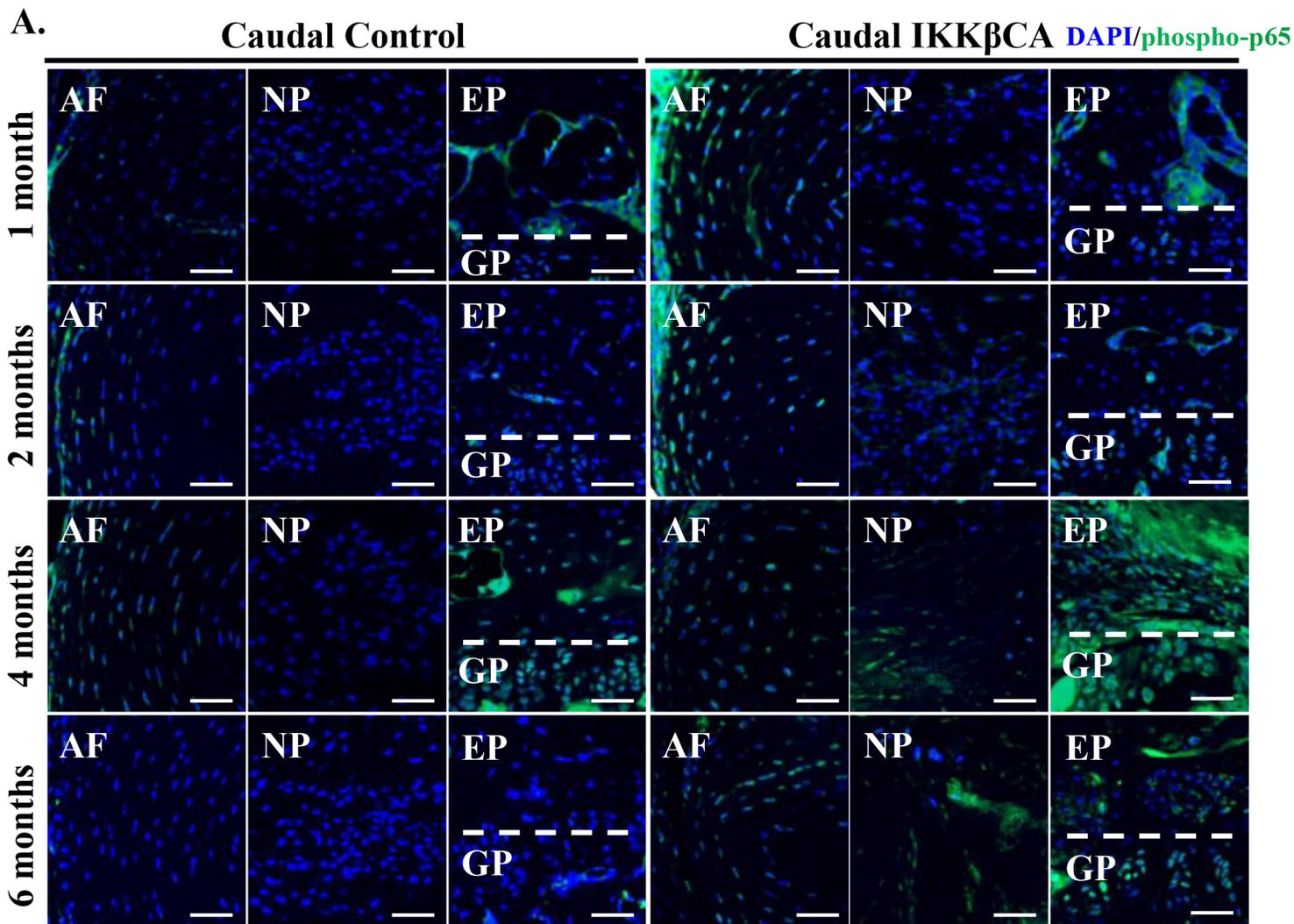
Kevin G. Burt *et al.*

Corresponding author: Nadeen O. Chahine, noc7@columbia.edu

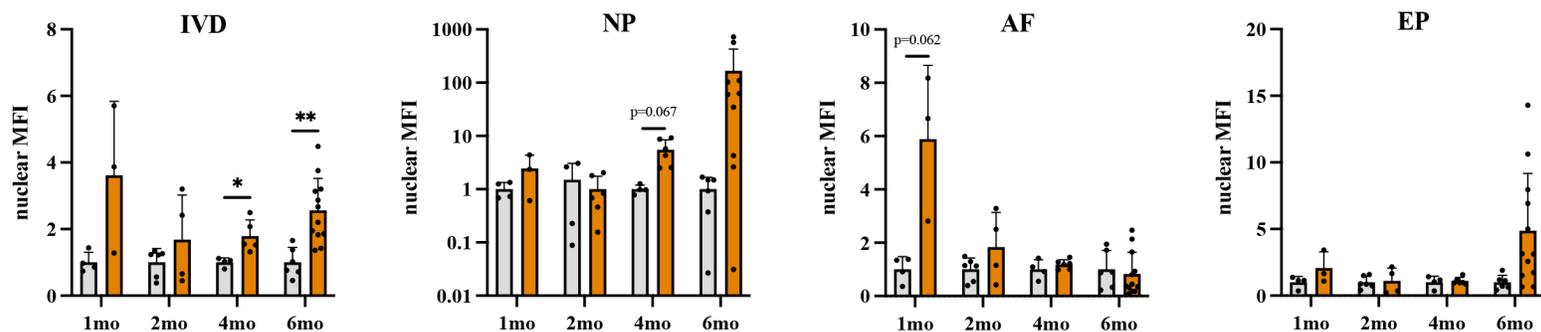
*Sci. Adv.* **10**, eadj3194 (2024)  
DOI: 10.1126/sciadv.adj3194

**This PDF file includes:**

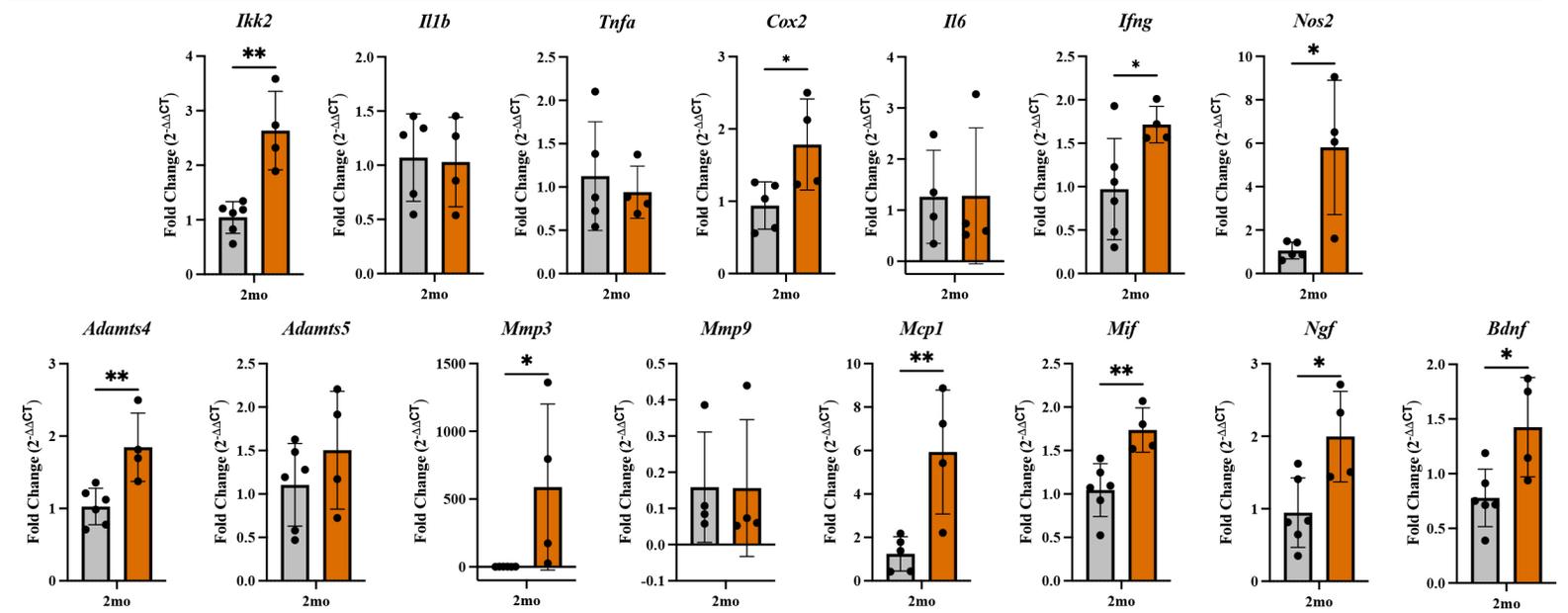
Figs. S1 to S6  
Tables S1 and S2



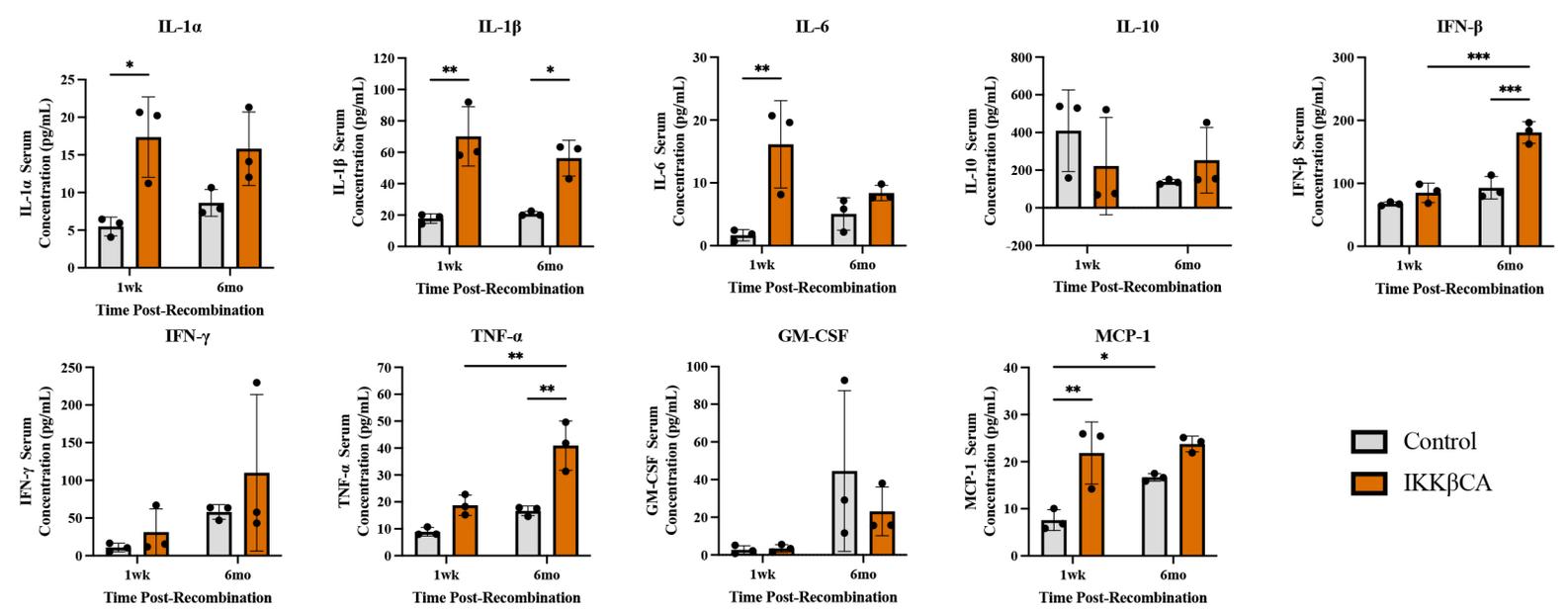
**B.** Caudal phospho-p65 expression  Control  IKK $\beta$ CA



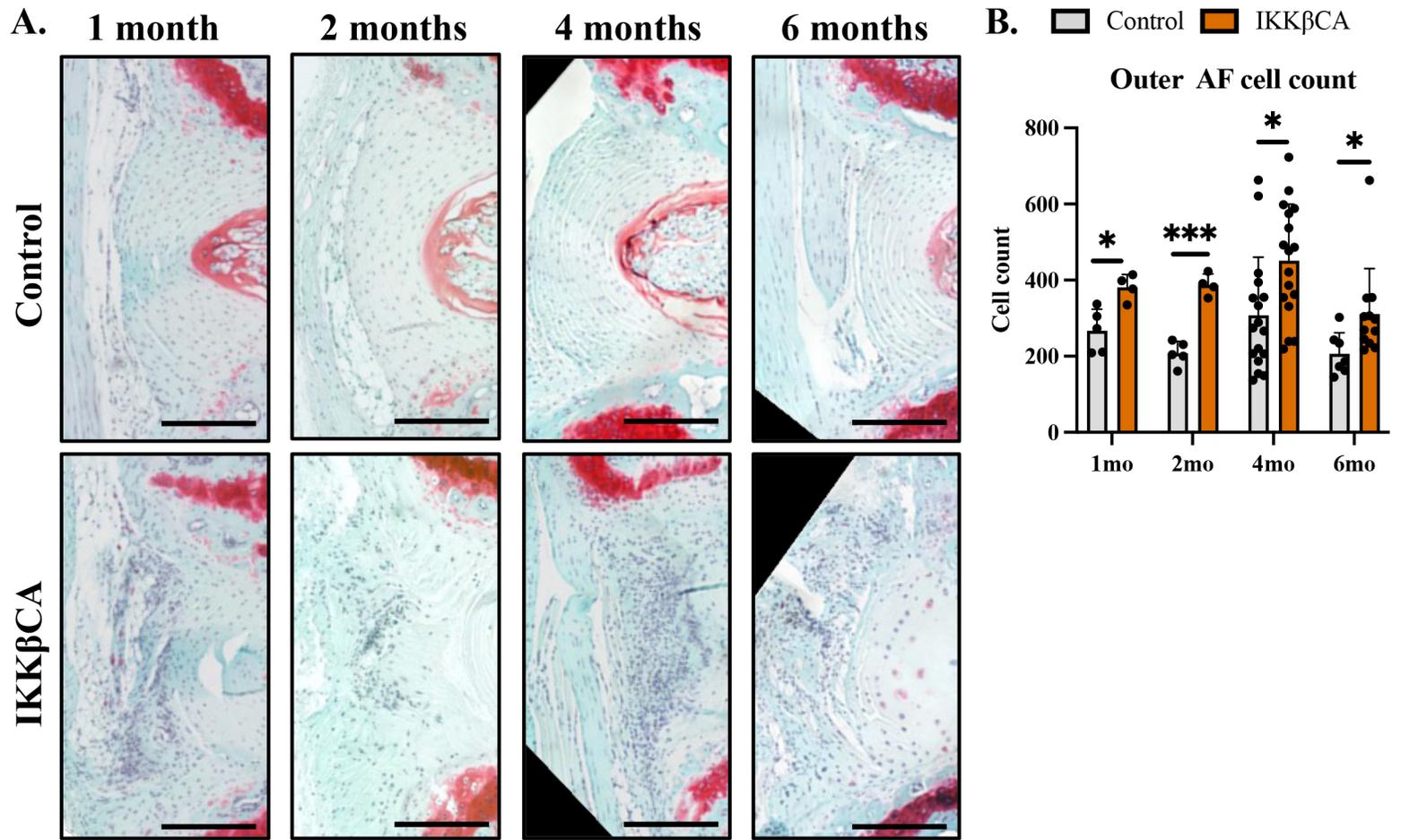
**Fig. S1. NF- $\kappa$ B activation within IKK $\beta$ CA caudal IVDs.** (A) Representative IF staining for phosphorylated p65 (green) within mid sagittal sections of control (*AcanCre<sup>-/-</sup>Ikk2ca<sup>fl/fl</sup>*) and IKK $\beta$ CA caudal IVDs. Scale bar = 50 $\mu$ m. (B) Nuclear MFI quantification (normalized to control within time point) of phosphorylated p65. \*p<0.05, \*\*p<0.01 (n=3-6 mice/group, 1-2 discs/mouse).



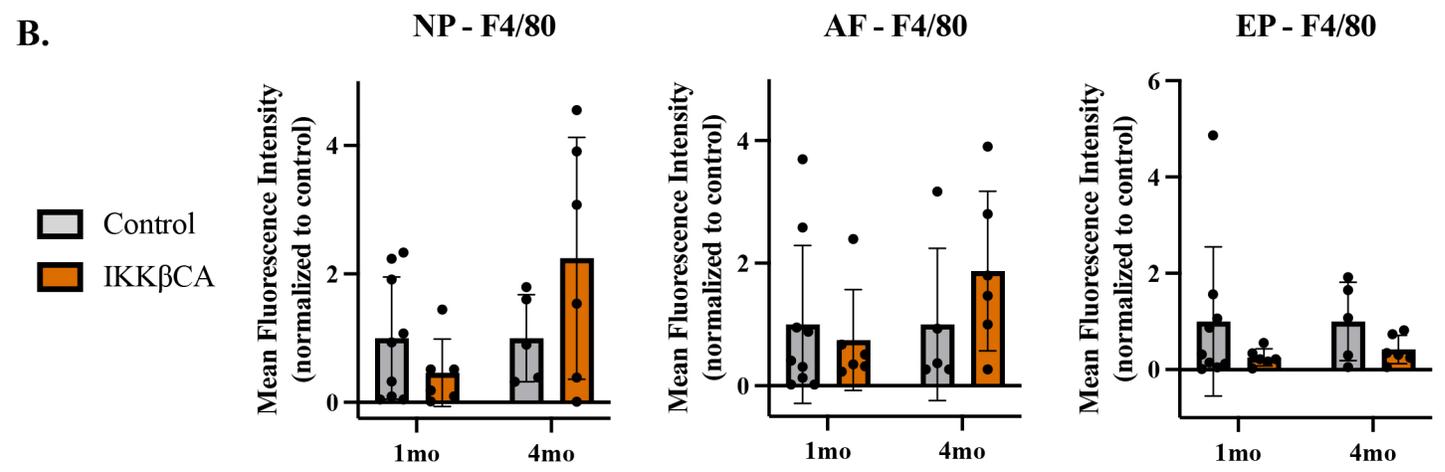
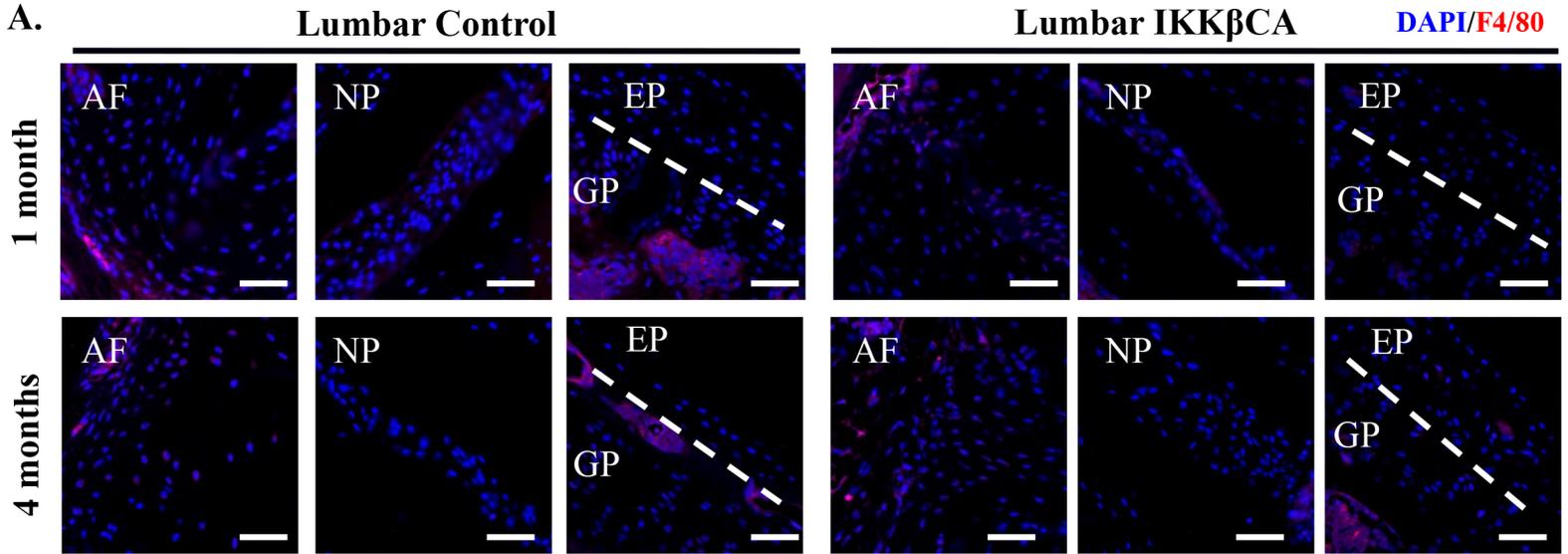
**Fig. S2. IKKβ over-expression upregulates caudal IVD inflammatory cytokine, chemokine, catabolic enzyme, and neurotrophic factor gene expression.** Gene expression changes (relative to control) from total RNA isolated from control (*AcanCre<sup>-/-</sup>;Ikk2ca<sup>fl/fl</sup>*) and IKKβCA whole caudal IVDs containing NP, AF, and EP, 2-months post recombination. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001. (n=3 mice/genotype per timepoint, 1-2 discs/mouse).



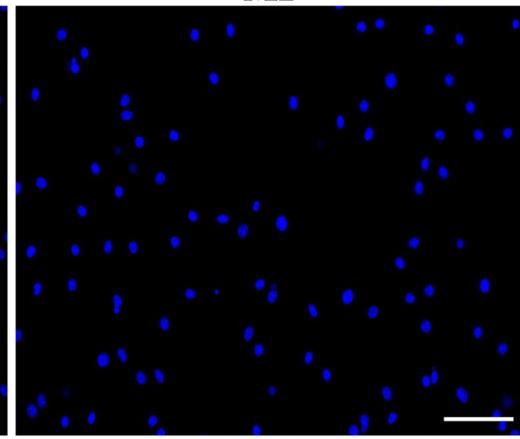
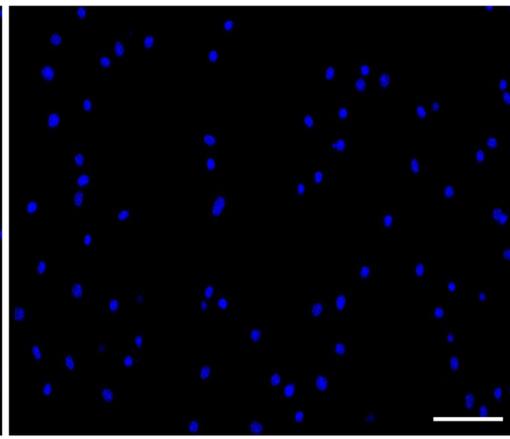
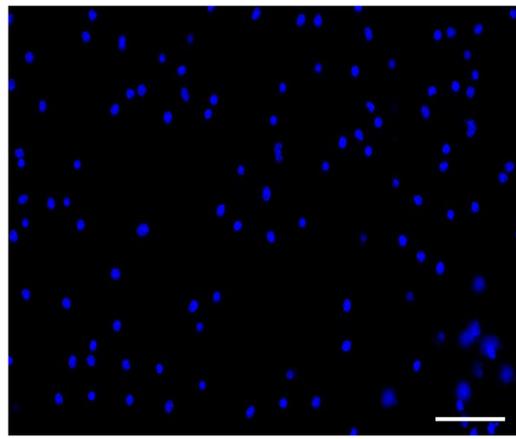
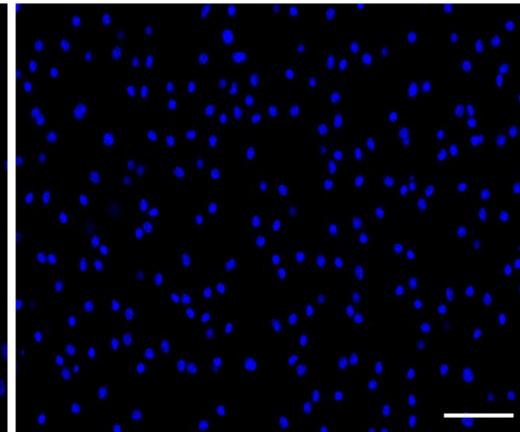
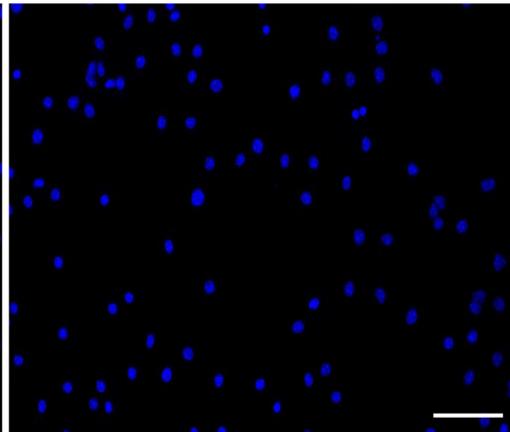
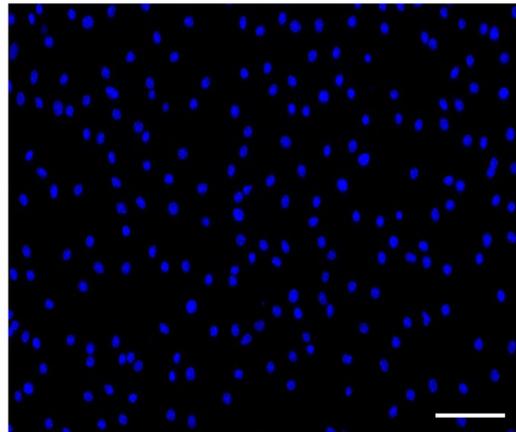
**Fig. S3. IKK $\beta$  over-expression upregulates circulating inflammatory cytokine and chemokines in the serum.** Cytokine and chemokine levels within serum isolated from control (*AcanCre<sup>-/-</sup>Ikk2ca<sup>fl/fl</sup>*) and IKK $\beta$ CA mice 1 week and 6 months post recombination. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001 (n=3/genotype per timepoint).



**Fig. S4. IKK $\beta$  over-expression increases cellularity presence within the caudal AF.** (A) Representative images of safranin-O stained mid sagittal sections of control (*AcanCre<sup>-/-</sup>;Ikk2ca<sup>fl/fl</sup>*) and IKK $\beta$ CA caudal IVDs 1-, 2-, 4-, and 6-months post recombination. Scale bar = 250 $\mu$ m. (B) Quantification of outer AF cellularity using haematoxylin nuclear stain (n=4-6 mice/genotype and timepoint, 1-4 IVDs/mouse).



**Fig. S5. Macrophage presence within the lumbar IVDs.** (A) Representative images of IF staining for F4/80, in mid sagittal sections of control (*AcanCre<sup>-/-</sup>Ikk2ca<sup>fl/fl</sup>*) and IKK $\beta$ CA lumbar IVDs at 1- and 4-months post recombination. Scale bar = 100 $\mu$ m. (B) MFI quantification of F4/80 expression within individual NP, AF, and EP compartments. Letters (a,b,c) indicating statistically significant ( $p < 0.05$ ) different groupings. (n=3-6 mice/genotype, 1-2 discs/mouse).

**M0****M1****M2****Control-CM****IKKβCA-CM**

**Fig. S6. Caudal IKK $\beta$ CA-CM increases macrophage migration *in vitro*.** Representative images of DAPI (nuclear) stained M0, M1, and M2 macrophages on transwell membranes following caudal IKK $\beta$ CA-CM stimulation. Cell counting used for quantification of migration through transwell membranes. Scale bar = 100 $\mu$ m (n=3).

**Table S1: FWD and REV murine primers (IDT) for rt-qPCR.**

Gene Target, murine	Forward Primer	Reverse Primer
Glyceraldehyde-3-phosphate dehydrogenase ( <i>Gapdh</i> )	AAC AGC AAC TCC CAC TCT TC	CCT GTT GCT GTA GCC GTA TT
inhibitor of nuclear factor kappa B kinase subunit beta ( <i>Ikk2</i> )	CTG AAG ATC GCC TGT AGA AA	TCC ATC TGT AAC CAG CTC CAG
interleukin-6 ( <i>Il6</i> )	CTT CCA TCC AGT TGC CTT CT	CTC CGA CTT GTG AAG TGG TAT AG
tumor necrosis factor alpha ( <i>Tnfa</i> )	TTG CTC TGT GAA GGG AAT GG	GGC TCT GAG GAG TAG ACA ATA AAG
interleukin-1 beta ( <i>Il1b</i> )	ATG GGC AAC CAC TTA CCT ATT T	GTT CTA GAG AGT GCT GCC TAA TG
nitric oxide synthase 2 ( <i>Nos2</i> )	TCT CCC TTT CCT CCC TTC TT	CTT CAG TCA GGA GGT TGA GTT T
interferon gamma ( <i>Ifng</i> )	GGC CAT CAG CAA CAA CAT AAG	GTT GAC CTC AAA CTT GGC AAT AC
Prostaglandin-Endoperoxide Synthase 2 ( <i>Ptgs2/Cox2</i> )	GAA GAT TCC CTC CGG TGT TT	CCC TTC TCA CTG GCT TAT GTA G
Cluster of differentiation 38 ( <i>CD38</i> )	TCT CTC TCT CTC TCT CTC TCT CT	TCA GCT GTG CTG AGG ATT TAG
Cluster of differentiation 206 ( <i>CD206</i> )	GGA ATC AAG GGC ACA GAG TTA	TTC CAT CTG CTC CAC AAT CC
monocyte chemoattractant protein-1 ( <i>Mcp1</i> )	CTC GGA CTG TGA TGC CTT AAT	TGG ATC CAC ACC TTG CAT TTA
macrophage inhibitory factor ( <i>Mif</i> )	GTT CCA CCT TCG CTT GAG T	CAT CGC TAC CGG TGG ATA AA
Matrix Metalloproteinase 3 ( <i>Mmp3</i> )	GGA CCA GGG ATT AAT GGA GAT G	TGA GCA GCA ACC AGG AAT AG
Matrix Metalloproteinase 9 ( <i>Mmp9</i> )	CTG GAA CTC ACA CGA CAT CTT	TCC ACC TTG TTC ACC TCA TTT
A Disintegrin-Like And Metalloprotease With Thrombospondin Type 1, Motif 4 ( <i>Adamts4</i> )	GGC AGA GAA GGG ATG ATG TAA TAG	CCC AAC ATC ACC CAG GTA ATA A
A Disintegrin-Like And Metalloprotease With Thrombospondin Type 1, Motif 5 ( <i>Adamts5</i> )	GTG CTG TGT TTG CCA TCT TC	GCA CTG CCT TGT TCT GTT TC
nerve growth factor ( <i>Ngf</i> )	CAG TGA GGT GCA TAG CGT AAT	CTC CTT CTG GGA CAT TGC TAT C
brain derived neurotrophic factor ( <i>Bdnf</i> )	CAA GAG TCC CGT CTG TAC TTT AC	GAC TAG GGA AAT GGG CTT AAC A

**Table S2: Primary and secondary fluorescent antibodies used for immunohistochemistry.**

Antigen	Company	Cat #	Dilution
[1°] mCherry	Sicgen	AB0040-200	1:250
[1°] Phosphorylated p-65	Abcam	AB86299	1:500
[1°] F4/80	Bio-rad	MCA497GA	1:100
[1°] IKK $\beta$	Sigma	07-1479	1:500
[1°] CD38	Fisher	PIMA516871	1:100
[1°] CD206	Fisher	PIPA595840	1:100
[2°] Donkey anti-goat (AF 594)	Thermo Fisher	A-11058	1:250
[2°] Donkey anti-rabbit (AF 594)	Abcam	AB150064	1:200
[2°] Donkey anti-rat (AF 488)	Abcam	Ab150153	1:200