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Ablation of VEGFA following a lumbar intervertebral disc injury attenuates intradiscal neurovascular features and prevents chronic low back pain symptoms

Authors: *Ryan S. Potter[1],*Hong J. Moon[3], *Sade W. Clayton[2], Remy E. Walk[2], Addison L. Liefer[2], Liufang Jing[2], Amber N. Stratman[3], Lori A. Setton[1,2,4], Munish C. Gupta[3], Simon Y. Tang[1,2,4]

*These authors led the study equally.

Affiliations:
[1] Department of Mechanical Engineering and Materials Science, Washington University in St. Louis
[2] Department of Biomedical Engineering, Washington University in St. Louis
[3] Department of Cell Biology and Physiology, Washington University School of Medicine
[4] Department of Orthopaedic Surgery, Washington University School of Medicine

Corresponding author: Simon Tang (simon.tang@wustl.edu)

One Sentence Summary:
VEGFA critically mediates the infiltration of neurovascular features into the degenerating intervertebral disc, and the ablation of VEGFA prevents chronic low back pain behavior.

Keywords:
VEGFA, intervertebral disc injury, low back pain, neurovascular features, preclinical model

Abstract

There are currently no therapies for the staggering disability and public health costs of chronic low back pain (LBP). Innervation of the degenerating intervertebral disc (IVD) is suspected to cause discogenic LBP, but the mechanisms that orchestrate the IVD's neo-innervation and subsequent symptoms of LBP remain unknown. We hypothesize that Vascular Growth Endothelial Factor-A (VEGFA) critically mediates the neurite invasion in the IVD and contributes to prolonged LBP. Initiating IVD degeneration through a mechanical injury, we evaluated the progression of neurovascular features into the IVD, as well as resultant LBP symptoms and locomotive performance at acute (3-weeks) and prolonged (12-weeks) time points following IVD injury. To determine the role of VEGFA, we used a mouse model with ubiquitous inducible recombination of the floxed VEGFA allele (UBC-Cre^{ERT2}; VEGFA$^{fl/fl}$). The ablation of VEGFA attenuated the neurite and vessel infiltration into the degenerating IVD, and the VEGFA-null animals exhibited alleviated mechanical allodynia and improved locomotive performance. To determine the effects of IVD-derived VEGFA on endothelial cells and neurons, we cultured HMEC-1 endothelial cells and SH-SY5Y neurons using conditioned media from VEGFA-silenced (siRNA) human primary IVD cells stimulated by IL1β. The endothelial cells and neurons exposed to the secretome of the VEGFA-silenced IVD cells exhibited reduced growth, suggesting that the inhibition of IVD-derived VEGFA may be sufficient to attenuate intradiscal neurovascular features. Together, we show that VEGFA orchestrates the growth of intradiscal vessels and neurites that cause low back pain and impaired function, and the inhibition of VEGFA can prevent prolonged low back pain.

INTRODUCTION

Low back pain (LBP) is a prevalent, debilitating, costly, and multifactorial condition worldwide, imposing substantial global and economic burdens [1-3]. LBP affects 30–80% of the population and increases with age [4]. It is the leading cause of years lived with disability and contributes an estimated $50 to $100 billion USD annually in direct healthcare costs [5,6] Chronic LBP is commonly associated with intervertebral disc (IVD) degeneration, which occurs due to aging, genetic predisposition, and mechanical stress [7-10]. Degeneration leads to structural and biochemical changes—such as loss of hydration, inflammation, and matrix breakdown—that impair disc function [11-14]. However, due to the complexity of these changes, it remains unclear which specific features of degeneration are responsible for pain [15].

Innervation has been identified in painful human intervertebral discs, suggesting a role for aberrant nerve growth in discogenic pain [16]. VEGFA (Vascular Endothelial Growth Factor A) has canonical roles in neurovascular growth and its enhanced expression during injury may thus facilitate this pathological growth [17]. VEGFA is a potent signaling protein known for its role in promoting angiogenesis and neurogenesis. It stimulates endothelial and neural cells, influencing processes such as cell proliferation, migration, and survival [18,19]. While VEGFA has known roles in both vascular and neural support, its function in IVD injury models and its potential to initiate or sustain discogenic pain remain unresolved. We hypothesized that the ablation of VEGFA attenuates neurite growth and vascularization into the injured IVD, mediates structural and functional deficits of the IVD, and alleviates ensuing low back pain symptoms.

To investigate the role of VEGFA in IVD degeneration and pain, we utilized a mouse model with an

inducible ablation of VEGFA and we created an injury in the lumbar IVD to model degeneration. We

subsequently quantified pain-related symptoms such as mechanical sensitivity and locomotor deficits

[OBJ][OBJ][OBJ][20][OBJ]. Because VEGFA is also a critical regulator of vascular growth, we assessed neo-

vascularization into the injured IVD along with neural infiltration to better understand the relationship

between vascular and neural response in disc degeneration. In human cells, we performed co-culture

experiments determine the role of IVD-produced VEGFA on neuron and endothelial cell growth.

Our findings demonstrate that VEGFA plays a central role in promoting neurovascular ingrowth into

the degenerating intervertebral disc, contributing to chronic low back pain. The ablation of VEGFA

significantly reduced neurite and blood vessel infiltration into the injured disc and alleviated mechanical

allodynia and locomotor deficits in mice. In vitro, endothelial cells and neurons exposed to conditioned

media from VEGFA-silenced IVD cells exhibited diminished growth, reinforcing VEGFA's role as a

key mediator of discogenic pain pathways. These results demonstrate that targeted inhibition of VEGFA

can suppress intradiscal vessels and neurites that cause low back pain and impaired function, and this

results in the prevention of chronic low back pain.

RESULTS

Removal of VEGFA Blunts Intradiscal Invasion of Neurovascular Structures

In WT animals, the injured IVD was infiltrated by PGP9.5+ neurons and by endomucin+ vessels VEGFA ablation significantly blunted these neuronal and vascular features. Immunofluorescence images from 50μm thick sections (Figure 1A-B) showed reduced neural and vascular structures in VEGFA-null mice compared to WT particularly in the region around the outer AF. Quantitative measurements of structural length and depth (Figure 1C-F) revealed significant differences in neurite length (p=0.01), vessel length (p=0.004), neurite depth (p=0.002), and for vessel depth (p=0.047) due to VEGFA removal. Post hoc analysis reveals differences between the WT and VEGFA-null IVDs in neurite depth at 3 weeks (p=0.001) and in vessel length at 12 weeks (p=0.01). These findings show that VEGFA is crucial for instigating the neural and vascular structures in IVDs following injury. VEGFA ablation can limit the penetration into- and the propagation within- of these structures of the disc.

Fig. 1: Removal of VEGFA Blunts Intradiscal Invasion of Neurovascular Structures Following IVD Injury. A) IHC of dorsal sides of the injured lumbar IVD with the nuclei (DAPI) in blue, neuronal

features (PGP9.5) in green and vascular features (Endomucin) in red. The ROI is boxed with dotted lines. B) Outer AF ROI with key features notated C-F) VEGFA loss reduces neurite lengths (p=0.01) and vessel lengths (p=0.004) across both post-injury time points. Likewise, the neurite penetration depth (p=0.002) and vessel penetration depth (p=0.047) are blunted by VEGFA removal. Data are presented as mean ± standard deviation.

VEGFA Is Essential for the Spatial Colocalization of Intradiscal Nerves and Vessels

The spatial proximity of intradiscal nerves and vessels are disrupted by VEGFA ablation. Immunofluorescence images reveal the distribution of neural (PGP9.5, green) and vascular (Endomucin, red) structures highlighting the regions of interest around the outer annulus fibrosus (AF) and overlapping lengths relative to neurites, due to their rarer presence in the disc Figure 2A-B. The colocalization analysis show that the WT injured IVDs have neurites with a large amount of vessels in its proximity. The removal of VEGFA in the IVDs 3 weeks following injury prevented neurites from infiltrating the IVD (Figure 2-2C), but this difference between KO and WT animals did not persist at 12 weeks after injury. However, the loss of VEGFA did not deter Endomucin+ vessels from penetrating the injured IVD at both time points (Figure 2D). The spatial overlap analysis between neurites and vessels (Figure 2E) show that VEGFA ablation dramatically reduces the amount of neurites near vessels (p=0.007). After adjusting for the length of the neurites, this observation remains robust independent of the basal level of neurites in each animal (p=0.008; Figure 3F).

The lack of difference between the depth and length ratios between neurites and vessels indicates that once initiated, VEGFA does not appear to affect the propagation of these features (Figure 2G; 2H). These findings highlight the critical role of VEGFA in the initiation of neural and vascular features in the IVD.

Fig. 2: VEGFA is Essential for the Colocalization of Intradiscal Nerves and Vessels Initiation. A) WT injured disc showing key features of the IVD and a ROI for analysis B) Graphic representation of how spatial colocalization analyses quantified the amount of neurites that were in proximity of vessels with a 30 μm region. C-D) Loss of VEGFA significantly reduced the number injured IVDs that experienced infiltrating neurites 3 weeks after injury (p=0.01) in VEGFA-null at 3 weeks. Neurite penetration (p=0.02) and vessel penetration (p=0.03) both show a significant decrease in VEGFA-null groups. E) VEGFA ablation reduced the amount of vessel-adjacent neurites (p=0.007), and was dependent on amount of time following injury (p=0.04). F) The effect of VEGFA loss on preventing neurite-vessel spatial coupling persevered even when accounting for varying intradiscal neurite lengths between different animals (p=0.008), confirming that this is a local phenomenon conserved across animals. G-H) The lack of difference in the depth penetration and length elongation ratios suggest that, on average, once the neurites have initiated, the loss of VEGFA did not appear to affect their propagation. Statistical model C-D: Fisher's exact test. Statistical model E-H: 2 factor ANOVA with Fisher LSD post-hoc comparisons. E-H Data are presented as mean ± standard deviation.

Depletion of VEGFA Suppresses the Expression of Pain-related Ion Channels in the DRG

Quantitative measurements of CTCF of mice DRGs (Figure 3B-D) revealed a significant reduction in TRPA1 fluorescence (p=0.04) due to VEGFA removal following injury. No significant changes were measured in TRPV1 CTCF levels from VEGFA ablation indicating that TRPV1 is unaffected by VEGFA levels.

Fig. 3: VEGFA Inhibition Reduces the Expression of Ion Channels Associated with DRG pain. A) TrKA (top), TRPV1 (middle), and TRPA1 (bottom) in green and DAPI in blue were quantified. B-D) Corrected total cell fluorescence (CTCF) for each pain related ion channel shows VEGFA ablation regulated TRPA1 expression (p=0.037).

Ablation of VEGFA Alleviates Mechanical sensitivity and Locomotive Function

Loss of VEGFA Attenuates Mechanical, but Not Hot or Cold, Sensitivity

The WT animals exhibited mechanical sensitivity measured by the Von Frey filament test at both 3 and 12 weeks after intervertebral disc injury. Ablation of VEGFA alleviated this impairment at the 12-week time point (p=0.001, Figure 4A). The electronic Von Frey filament test also revealed that there is a greater mechanical sensitivity on the side of the surgical approach. VEGFA removal specifically alleviated the mechanical sensitivity of the injured side, without modifying the contralateral side. No notable changes observed in heat or cold sensitivity.

VEGFA Ablation Improves Rotarod Performance, but Does Not Affect Passive Function

We also evaluated locomotive performance through an active challenge of locomotive function, measured through the rotarod assay, longitudinally, every three weeks up to 12 weeks after injury. The longitudinal Rotarod test (Figure 4D) reveals that VEGFA-null mice exhibit significantly maintained latency to fall at 9 and 12 weeks compared to impaired WT (p=0.004), highlighting impaired motor coordination over time (p=0.003). In the Open Field test, total distance traveled (Figure 4E) and rearing counts (Figure 4F) are not significantly different between genotypes, suggesting comparable levels of exploratory behavior and general activity.

Fig. 4: **VEGFA Ablation Improves Mechanical Sensitivity and Performance** A) Mechanical

sensitivity shows a statistically significant interaction between genotype and time (p=0.001) with a

worsening effect seen in WT leading to a chronic sensitivity and an improving effect in VEGFA-null

over time that recovers near baseline. B-C) Temperature sensitivity changes were not statistically

significant. D) Longitudinal measures of performance included rotarod which shows genotype (p=0.004) and time (p=0.003) statistical significance, indicative of a VEGFA rescue of a more chronic progression of performance impairment. E-F) Open field measures of total distance and rearing are not statistically significantly different. G-H) Inverted screen endurance at 3 and 12 weeks was not statistically significantly different. I) Hind paw grip strength remained unchanged. Statistical model A-F, I: Linear mixed effects with genotype and time as factors with post-hoc LSM with Sidak correction. G-H Mantel-Cox test. A-F, I Data are presented as mean ± standard deviation.

Assay	Sex M/F	Tamoxifen +/-	Naïve/Sham
Von Frey	0.94	0.63	0.66
Grip Strength	0.09	0.92	0.07
Inverted Screen	0.65	0.33	0.64
Cold Sensitivity	0.52	0.06	0.40
Heat Sensitivity	0.44	0.47	0.57
Rotarod	0.34	0.43	0.25
Distance (Open Field)	0.33	0.27	0.18
Rearing (Open Field)	0.25	0.32	0.59
Neurite Length	0.08	NA	NA
Vessel Length	0.19	NA	NA
Neurite Depth	0.51	NA	NA
Vessel Depth	0.15	NA	NA

Table S4: Behavioral and Neurovascular Significance Values for Sex, Tamoxifen, and Naïve/Sham P-values showing the lack of significant differences (p>0.05) between Male and Female, Tamoxifen +/-, and Naïve and Sham surgery for the behavioral, neuronal, and vascular assays. P-values from the Linear Mixed Effects statistical model. Factors without samples for comparison are shown as NA.

VEGFA Removal Does Not Affect Measures of Strength

Strength assessments include inverted screen endurance and hind paw grip strength. The inverted screen endurance test, indicated by the survival analysis, showed that VEGFA-null mice have similar probabilities of hanging on compared to WT at both 3 weeks (Figure 4G) and 12 weeks (Figure 4H). Hind paw grip strength (Figure 4I) shows no differences, indicative of no compromise in muscle function from VEGFA ablation.

Overall, these findings show that VEGFA-null mice have improved recovery of mechanical sensitivity and avert the transition to chronic sensitivity seen in the WT while also avoiding the performance deficits in WT at chronic time points.

Ablation of VEGFA Does Not Affect structure, function, or degeneration of the IVD

Loss of VEGFA Does Not Affect the Histopathic Degeneration of the IVD

CEμCT analysis of the IVDs 3 weeks after injury reveals significant deterioration of the IVD structure. By using a contrast enhancing agent, Ioversol, the injury can be observed on the ventral side of the outer AF as a progressively higher attenuating regions that mark the fibrous scar tissues (Figure 5A). The injured IVDs also exhibit a loss of NP attenuation, measured by NI/DI, demonstrating the loss of the water retaining ability of the NP (p=0.03, Figure 5B). The ablation of VEGFA does not affect these injury-mediated changes in degeneration. A narrowing of the intradiscal space measured by Disc height index (DHI), a hallmark of IVD degeneration, is apparent in the injured discs (p=0.01, Figure 5C). Surprisingly, there was not a change in NP volume either due to injury or VEGFA ablation. Structurally, injury increases degenerate features including decreased NP hydration and a narrowing DHI while VEGFA ablation shows promising trends to recover these injury induced changes. Noticeably, there was

a spectrum of structural adaptations that may be due to variations in the surgical injury due to needle

depths, approach angles differences, and surgical conditions.

Injury to the IVD Does Not Impair Its Mechanical Function

Dynamic compression of IVDs 3 weeks post injury show no distinct responses to injury or impact from

VEGFA ablation. The stiffness measurements show no significant differences between WT and

VEGFA-null groups, both in sham and injured conditions (Figure 5D). Likewise, the dissipation factor

indicates a similar response with no appreciable changes from injury or VEGFA loss (Figure 5E). These

findings show that the degeneration that occurs at 3 weeks after injury does not yet impair its mechanical

function.

VEGFA Loss Does Not Rescue the Degenerative Changes Caused by Injury

Histopathological scoring (Figure 5H) revealed the destructive effect of a needle stab on the

compartments in the disc, including decreased NP cell density and changes in cellular morphology,

disruption of the AF concentric lamellar structures with increased cellularity and presence of clefts,

along with thinning and disruptions in the endplate cartilage, and structural disorganization in the

boundary regions. Scores for injured groups averaged near 26 out of 36 maximum while sham controls

were near 11 with the greater degeneration in the injured discs (p < 0.0001) compared to the sham

animals with no IVD injury. Statistical differences from VEGFA removal were not present. Mechanical

injury dramatically changed all compartments of the disc with the largest effects on the AF followed by

NP as expected based on the AF stab model while sham scores indicate a moderate degeneration from

the exposure of the disc and likely disruption of surrounding soft tissues.

Fig. 5: Loss of VEGFA Does Not Alter Structural, Mechanical, or Histopathologic Degeneration to the Injured IVD. A) IVD structure and compositional changes via CEμCT show that B) injury decreased NP hydration (p=0.03), indicated by the decreased attenuation of the NP, C) injury decreased disc height index, DHI, (p=0.01) compared to sham, and D) no effect of NP volume was observed. VEGFA removal showed a trend towards Sham levels on the DHI indicating a potential protection against hydration loss. Changes to mechanical behavior were not observed in E) stiffness and F) dissipation factor, indicating that tissue level mechanics were not changed by injury or VEGFA removal at this 3-week post injury time point. G) Histological images stained with Safranin-O and Fast Green show IVD degeneration H) with histopathologic scoring showing a highly statistically significant effect of injury (p<0.0001) with no appreciable differences due to VEGFA loss in nucleus pulposus, annulus fibrosis, end plate, or the interface/boundary demonstrating the pathophysiological response from needle puncture. Data are presented as mean ± standard deviation.

Fig. 6: VEGFA Inhibition Reduces *in vitro* Neurite Activity and Alters Gene Transcription A) Neurite extension assay after treating neuronal cells with media collected from IL-1B stimulated and siRNA transfected human disc cells shows inhibition of neurites from VEGFA siRNA. B-D) Quantification of neurite morphology showing functional suppression with VEGFA siRNA E) RNA-seq showing 146 differentially expressed genes and F) key biological processes regulated by VEGFA siRNA. NCs co-cultured with IL1β- NAIVE DISC CELLS, (IL1β- NAIVE DISC CELLS)NC showed undifferentiated cell possessing few short projections and cluster together (Figure 5A, open arrows), while NCs co-cultured with IL1β+SCRAM siRNA, (IL1β+SCRAM siRNA)NC and NCs co-cultured with IL1β+VEGFA siRNA, (IL1β+VEGFA siRNA)NC showed differentiated cells having many extensive projections (Figure 5A, closed arrows). Total neurite number from 1. (IL1 β-NAIVE DISC

CELLS)NC, 2. (IL1β+SCRAM siRNA)NC and 3. (IL1β+VEGFA siRNA)NC was 194.7 ± 52.6, 325.3

± 76.5 and 281.0 ± 61.8 respectively. 1 vs 2 and 2 vs 3 showed significant difference (p=0.01 and

p=0.04). Total neurite length was 4094.2 ±1185.0, 16283.5 ± 2149.6 and 10376.5 ± 813.4 respectively.

1 vs 2 and 2 vs 3 showed significant difference (p=0.02 and p=0.02).

Silencing VEGFA RNA in Human IVD Cells Attenuates the Growth of Co-Cultured Neuronal

and Endothelial Cells

Transfecting human IVD cells with Vegfa-specific small interfering RNA (siRNA) suppressed their

subsequent production of the VEGFA protein expression by 57.4% when stimulated with IL-1β (Figure

S8). Coculturing these Vefga-siRNA transfected, IL-1β stimulated IVD cells with SH-SY5Y neurons

maintained the neuron morphology but blunted the growth of the axonal extensions (Figure 6A).

Although the number of neurites and their averaged lengths did not change for co-cultured neurons

(Figure 6B), the total neurite length increased in the group cultured with Vegfa-intact IVD cells (Figure

6C, D). There was no change in average neurite length (Figure 6D). RNA sequencing revealed that

neurons cultured with Vegfa-deficient IVD cells expressed 39 downregulated and 107 upregulated

DEGs (Figure 6E). Gene ontology analysis determined the most enriched biological process from the

DEGs involved the regulation of general cell processes, e.g. GCPR signaling pathway involved on hear

process (*Ednra, Chga, Ramp3*), Negative regulation of growth (*DCC, GREM1, STC2, MYOZ1, MT1F,*

BST2, MT1X, BCL2), and the Negative regulation of cell population proliferation (*GREM1, PTPN14,*

DHRS2, SSTR2, SCG2, SERPINF1, IGFBP3, OVOL1, ADM, IGFBP7, BCL2, S100A11). Notably, a

number of DEGs were specific to cell growth inhibition (*SSTR2, S100A11), immune* cells and

inflammation (*CXCR4, FGFB2, IL32)*, and neuronal cell functions (*DCC, SDK2*, *ASTN2*).

We also co-cultured siRNA transfected IVD cells with HMEC-1 cells. Endothelial cells (EC) cocultured

with Vegfa-silenced human IVD cells were less capable of forming closed tubes, a measure angiogenesis

(Figure 7A). ECs cultured with Scram siRNA significantly developed a higher number of closed loops

than ECs cultured with Vegfa siRNA (16.3 ± 4.4 vs 7.3 ± 3.1, p=0.04) after 6 hours culturing. (Figure

7B). RNAseq uncovered 175 downregulated and 0 upregulated DEGs due to Vegfa siRNA transfection

when compared to human IVD cells transfected with Scram siRNA (Figure 7C). Gene ontology analysis

determined the most enriched biological process from the DEGs involved in vessel and vasculature

related process, e.g. Lymph vessel development (*NR2F2, TBX1, SOX18, EFNB2,HEG1*), Blood vessel

morphogenesis (*TNFAIP2, EFNB2, HEG1, PRKACA, NRARP, TBX1, SOX4, NR2F2, AKT1, QKI,*

SMAD7, SOX18, VEGFB), and Regulation of neuron projection development (*EFNB2, CDK5R1, RYK,*

RAPGEF1, CARM1, AKT1, PLZNA1, NCS1, DVL1, SLK, STK24, BRSK2, ADNP). Interestingly,

Canoincal Wnt signaling pathway was also an enriched biologically pathway and Wnt signaling plays

an important role in endothelial cell proliferation, survival, differentiation and response to mechanical

forces due to shear stress from blood flow.[20,21]

Figure S8. Validation of VEGFA-siRNA Transfection in Disc Cells A) Control siRNA (Fluorescein

Conjugate)-A revealed efficient transfection of cationic lipid-based transfection reagents in disc cells.

B) Stimulated disc cells by IL-1β with Scrambled siRNA shows a significant increase in the VEGF

secretion compared to naïve disc cells (158.9 ± 27.6 vs 69.1 ± 27.9, p=0.044). Stimulated disc cells with

IL-1β after being transfected with VEGFA siRNA showed a significant silencing (57.4%) of VEGFA

secretion compared to Scram siRNA (67.6 ± 16.0 vs 158.9 ± 27.6, p=0.034).

Fig. 7: VEGFA Inhibition Reduced *in vitro* Endothelial Cell Function and Regulates Gene Transcription A) Matrigel tube forming assay images of endothelial cells cocultured with human disc cells from degenerate discectomy tissue treated with siRNA showing initiation of vessel formation after 6 hours. B) VEGFA siRNA significantly reduced tube formation compared to Scrambled siRNA C) VEGFA siRNA regulated 176 differentially expressed genes. There is a suppression of key angiogenic genes including *Heg1*, *Narap*, and *Ncs1*. There is also a suppression of endothelial cell-derived neurogenic genes including *Brsk2* and *Crlf1*. D) The RNAseq analyses of the HMEC-1 also show that a ontological shift to lymphangiogenesism, matrix remodeling, and suppression of neurovascular regulating pathways.

Fig. 8: The secreted factors from HMEC-1 (aka ECs) co-cultured with IVD cells were analyzed using the Luminex multiplex ELISA. Despite having a functional VEGFA gene, the ECs produced less VEGFA and CCL3 when co-cocultured VEGFA-silenced IL-1β–stimulated IVD cells. The ECs also produced more VEGFC and PDGF-AA that are both positive regulators of lymphangiogenesis; MMPs (MMP1, MMP3, MMP10, MMP13); chemokines including G-CSF, GM-CSF, CXCL1, and CCL5 (RANTES); and PDGF-AA, a positive relator of was also Units are pg/ml.

Group	Factor	Scrambled siRNA	VEGFA siRNA	p-value (paired)
Angiogenesis	Angiopoietin-2	29.1±2.1 (pg/ml)	28.9±2.4 (pg/ml)	0.53
Angiogenesis	Endoglin	84.8±10	92.9±5.9	0.06
Angiogenesis	Endothelin-1	337.3±42.5	336.7±34.9	0.84
Angiogenesis	Follistatin	436.8±485.1	915.9±1105.5	0.12
Angiogenesis	Leptin	70.4±11	70.4±8.4	0.79
Angiogenesis	PLGF	127.9±4	126.2±7.6	0.88
MMP	MMP2	8775±1081.8	9309.3±1816.7	0.55
MMP	MMP7	613.5±45.3	759.9±120.3	0.24
MMP	MMP8	138.6±116.4	239.4±120	0.30
MMP	MMP9	163±30.5	249.7±44.1	0.05
MMP	MMP12	582.4±65.1	604.9±59.9	0.60
MMP	TIMP1	100638.6±13530.7	89032.7±6718.9	0.13
MMP	TIMP2	30098.3±2192.1	31829.1±3278.2	0.07
MMP	TIMP3	3770.7±1298.2	2823.7±677.1	0.42
MMP	TIMP4	56.8±6.4	54.1±3.1	0.55
Cytokines	EGF	3.2±0.8	3.5±0.7	0.64
Cytokines	Eotaxin	7.7±0.6	7.6±0.7	0.89
Cytokines	FGF-2	187±72.4	200.5±58.4	0.23
Cytokines	FLT-3L	0.6±0.1	0.6±0.1	0.74
Cytokines	M-CSF	246.1±74.1	222.9±61	0.60
Cytokines	IL-6	2481.3±183	2518±201.4	0.94
Cytokines	IL-8	14958±894.4	16325.9±275.8	0.09
Cytokines	CCL2/MCP1	3441.9±1491.7	3898.9±1275.8	0.18
Cytokines	PDGF-BB	32.9±7.5	20.9±5.8	0.45

Supplemental Table 2. Panel of measured cytokines produced from the EC after co-culturing with the VEGFA-silenced IL-1β–stimulated IVD cells.

The factors secreted by endothelial cells co-cultured co-culturing with the VEGFA-silenced IL-1β- stimulated IVD cells reveals compensatory activity in the absence of VEGFA

After removal from the co-cultured conditions with the VEGFA-silenced, IL-1β-stimulated IVD cells, the HMEC-1 also secreted less VEGFA and CCL3, suggesting that the factors produced by the VEFA-silenced IVD cells contained less stimuli. Yet these HMEC-1s also produced more VEGFC, which is important for lymphangiogenesis; active matrix metalloproteinases (MMPs -1, -3, -10, and -13); G-CSF, GM-CSF, CXCL1, and CCL5 (RANTES) (Figure 8). The production of PDGF-AA suggests that there are compensation mechanisms to stimulate VEGFA. In all, IL-1β -stimulated IVD cells caused the HMEC-1s to form more mature tubular structures with more robust production of VEGFA. Silencing the VEGFA from these IVD cells prior to IL-1β -stimulation attenuated the tube formation

331   and resulted in reduced inflammatory factors but more MMPs, immune recruiting chemokines and

332   lymphoangiogenic factors, that indicates greater cellular emphasis on matrix remodeling and

333   lymphoangiogensis rather than angiogenesis. There are also a number of chemokines that are not

334   changed due to VEGFA-silencing of the IVD cells (Table S2).

DISCUSSION

This is the first study to demonstrate that intradiscal innervation contributes to locomotive performance and pain sensitivity in a model of intervertebral disc degeneration. While many studies in the past have shown progressive associations where neurovascular structures increase with progressive degeneration and pain,[22-26] none have been able to block or prevent these structures to prevent chronic pain and maintain locomotive performance. From this work, we also demonstrate that VEGFA may be a therapeutic target for a disease modifying therapy of low back pain. While VEGFA ablation did not significantly impact the structural and mechanical properties of the IVDs, it showed a pronounced effect on neural and vascular infiltration and the subsequent mechanical sensitivity and locomotive performance following degeneration.

VEGFA plays a crucial role in limiting nerve and vessel infiltration post-injury. IVD degeneration in VEGFA-null mice exhibited significantly reduced neurite and vessel lengths and depths, particularly in the outer annulus fibrosus (AF). VEGFA ablation inhibits the elongation and penetration of neural and vascular structures in the normally avascular IVD, potentially preventing painful sensory neurons from entering injured areas. The synergistic relationship between nerve and vessel infiltration highlights VEGFA's importance in neurovascular coupling within the IVD, with its absence leading to diminished neurite and vessel growth, which may limit painful neurological signals and affect overall disc health. These findings emphasize VEGFA's critical role in supporting neural and vascular initiation post-injury. Yet, the loss of VEGFA does not significantly influence IVD tissue degeneration, 3D structure, or mechanical properties, and it is not a major contributor to structural or functional integrity at the mechanical level during IVD degeneration.

The loss of VEGFA also appeared to reduce the neuropeptide-related signaling in TrKA, TRPV1, and TRPA1 in the Dorsal Root Ganglion (DRG). The increase expression of these factors in

the DRG have been observed in chronic low back pain, as well as in DRG that are sensitive to capsaisin stimulation. The decrease in TrKA activation could be responsible for the reduction in pain sensitivity demonstrated by the Von Frey mechanical test. The results of the Von Frey mechanical test revealed that VEGFA-null mice were less sensitive to pain than the wild-type mice. VEGFA removal also showed a significant decrease in TRPA1 fluorescence, however, no cold sensitivity was observed. TRPV1 levels did not have significant changes between VEGFA-null mice and wild-type mice. This data suggests that there would be no observable difference in heat sensitivity between VEGFA-null mice and wild-type mice, which is consistent with the hot plate experiment.

The behavioral assessments revealed that despite exhibiting acute pain post-injury, VEGFA-null mice had improved recovery from mechanical sensitivity and avoided the chronic sensitivity observed in WT mice. While VEGFA-null mice did not develop the performance impairments seen in WT mice at later points (9-12 weeks), indicating a protective effect of VEGFA ablation against chronic pain and impaired performance. Mechanical sensitivity showed imbalanced injury induced sensitivity while VEGFA inhibition balanced this asymmetry demonstrating there is no global effect of pain alleviation, and that VEGFA loss rescued the IVD injury effects but did not change systemic pain sensing.

Elongation vs Sprouting

Injuries to the intervertebral disc primarily trigger nerve and blood vessel sprouting as an adaptive response to inflammation and hypoxia [27,28]. This sprouting—stimulated by factors like NGF and inflammatory cytokines—leads to increased neurovascular infiltration into normally aneural and avascular disc regions, contributing to chronic pain and ongoing inflammation.[29-31] Consistent with these, we observed neurites and vessels infiltrating into the degenerating IVD at 3-weeks following the injury. However, once sprouted, the neurites did not appear to elongate further, as evidenced by the relative similar between the neurites between 3- and 12- weeks. In this animal model, VEGFA is

ubiquitously ablated from the whole animal, and it is not possible to determine the source of the VEGFA.

Our in vitro experiments however, support that blocking IVD-produced VEGFA was sufficient to

prevent growth of the neurites and the maturation of the blood vessels. RNAseq analysis suggest that

the reduced growth of neural outgrowths is driven suppression of cell proliferation. We observed similar

behavior in the endothelial cells, where the HMEC-1s co-cultured VEGFA-silenced IL-1β–stimulated

secreted VEGFC and PDGF-AA, both positive regulators of lymphangiogenesis, more MMPs, and more

immune cell recruiting chemokines. These immune cell recruiting chemokines are likely to be important

for the intrinsic healing of the IVD. [32] The RNAseq analyses of the HMEC-1 also show that a ontological

shift to lymphangiogenesism and matrix remodeling with a suppression of key angiogenic genes

including Heg1 Narap, and Ncs1. There is also a suppression of endothelial cell-derived neurogenic

genes including Brsk2 and Crlf1.

Needle stab injury models in animals are frequently utilized to study the pathophysiology of

intervertebral disc (IVD) degeneration and to develop treatments for chronic lower back pain (CLBP).

These models effectively replicate critical aspects of human disc degeneration, such as mechanical

damage, inflammatory responses, and biochemical changes. For example, they simulate the mechanical

damage caused by traumatic events and the inflammatory cytokine upregulation observed in human

discs, which are crucial for understanding the progression of disc degeneration.[33] However, these models

have limitations, especially in replicating the chronic, multifactorial nature of human CLBP and the

complexity of human pain perception.[34,35] Human IVD degeneration occurs over years and decades,

with the herniation of the IVD being one of the most apparent instigating factors. Despite the varied

etiologies of discogenic pain, the chronic symptoms and eventual degenerative presentation, such as the

deterioration of IVD structure, are quite similar. Therefore, the needle injury model in murine IVD is

employed to understand the mechanisms following IVD injury that eventually lead to low back pain

symptoms.

While these models are beneficial for basic research and initial therapeutic testing, their findings

must be interpreted with caution and validated in more complex models and ultimately in human clinical

studies. Differences in disc anatomy and biomechanics between humans and animals can affect the

translational relevance of these models, underscoring the need for complementary approaches to fully

capture the complexity of human IVD degeneration.[35] The needle injury in a mouse induces progressive,

but relatively rapid degeneration, occurring over weeks and months, compared to human degeneration,

which spans years and decades. Despite being traumatic relative to the human scale, this model is a

necessary caveat for developing feasible and mechanistic experiments. A single level lumbar injury

utilized here may also have underestimated the magnitude of our findings, since multiple groups have

shown a more pronounced pain behavior with 3 levels of injury in rodents. [36]

The multifactorial nature of the injury and VEGFA ablation at 3 weeks complicated the detection

of significant changes acutely, but by chronic timepoints, painful alterations were apparent.

Additionally, the WT sham group displayed elevated disc degeneration scores, suggesting that sham

procedures may induce a mild phenotype. Notably, neurovascular ingrowth was more pronounced on

the dorsal side opposite the ventral injury location, as seen by others,[22] likely due to its proximity to the

spinal cord and increased availability of neural and vascular sources. These findings are likely clinically

translatable as the treatment is compatible with the sequalae chronic low back pain development. Since

patients typically seek treatment only after an injury or an acute episode of pain, the timely post-injury

inhibition of VEGFA may be able to prevent the transition to chronic low back pain.

Overall, the loss of VEGFA appears to be protective against mechanical sensitivity, with

decreased innervation and no major impact on IVD degeneration. These findings, for the first time to

date, show a causative relationship between neurovascular pathoanatomy and pain while implicating

VEGFA as a critical mediator these processes following lumbar injury and highlight its potential as a

therapeutic target for preventing lower back pain through neurovascular inhibition. These findings are

likely clinically translatable as the treatment is compatible with the sequalae of chronic low back pain

development. Since patients typically seek treatment only after an injury or an acute episode of pain, the

timely post-injury inhibition of VEGFA may be able to prevent the transition to chronic low back pain.

MATERIALS AND METHODS

Experimental Design

The Postnatal Inducible Ablation of VEGFA in Adult Mice

Young adult, 4–6-month-old, ubiquitin (UBC)-Cre[ERT2] ; VEGFA^{fl/fl}; Ai9 tomato reporter mice which were previously described [37-39] were used on a C57BL/6J background and allow for a temporally controllable ablation of VEGFA upon tamoxifen administration and report recombination via an Ai9 fluorescence reporter. This ubiquitous model systemically ablates VEGFA in all tissue. An estrogen responsive Cre element is activated when a mutant estrogen receptor (ERT2) renders Cre active upon tamoxifen binding and initiates a conformational change that allows Cre to enter the nucleus and catalyze recombination at LoxP sites. These Cre+ mice allow for precise temporal control over gene modification. Moreover, this approach can avoid developmental defects and embryonic lethality, especially in pathways that are essential to organogenesis. All procedures (breeding, drug administration, procedures, and tissue extraction) were performed with WUSM IACUC (Institutional Animal Care and Use Committee) approval.

Confirmation and Quantification of VEGFA Ablation

The Ai9-tdTomato reporter protein is co-expressed with Cre recombinase presence, and it is visually evident by inspection at the macroscopic scale after two doses of tamoxifen (100 mg/ kg) administered over two consecutive days.

We also evaluated the efficiency of VEGFA ablation in the intervertebral disc by measuring the production of the VEGFA protein secreted by the intervertebral disc (IVD). Mice with Cre- and Cre+ alleles driven by the ubiquitin promotor in animals with the homozygous VEGFA floxed alleles (UBC-

Cre[ERT2]; VEGFA^{fl/fl} Ai9) received 2 consecutive days of tamoxifen administration via oral gavage

(100 mg/kg in corn oil). The animals were sacrificed on the third day, following the two days of

induction. The functional spine units (FSUs), containing the vertebrae-intervertebral disc-vertebrae

structure were cultured as previously described and stimulated with TNFα (5ng/ml) to provoke an

inflammatory response.[40] Media was collected every other day for 7 days and analyzed using a VEGFA

ELISA (R&D Systems Cat: DY493).

Figure S2: Validation of VEGFA Ablation and Mouse Health A) Ubiquitous recombination in a WT

and VEGFA-null showing the pink WT and red VEGFA-null Ai9 tdTomato reporter B) Mouse body

weights at 0 (Baseline), 3 and 12 weeks remain similar between groups irrespective of genotype or

surgery type C) Disc height ratio of injured/adjacent control disc height showing a similar decrease in

heights following injury D) Surgical survival curves show no significant differences between genotype

or surgery E) VEGFA ablation confirmation after initial ablation and 12 weeks from FSU media after

TNFα stimulation ELISA showing a sustained ablation over time F) E-Von Frey shows that WT injured

sensitivity increased over time (decreased withdrawal threshold) while VEGFA-null decreased over

time (increased withdrawal threshold) G) E-Von Frey asymmetry signifies an imbalance in WT mice

that fluctuates over time while VEGFA-null mice remained nearer to symmetry (dotted line). Data are

presented as mean ± standard deviation.

The Lumbar Puncture Intervertebral Disc Injury Model

Male and female mice of 4-6 months of age were randomly allocated into respective experimental

groups. The animals undergoing surgery were first anesthetized using a subcutaneous lidocaine injection

(0.5% at 7mg/kg) followed by isoflurane inhalation (3–4% v/v induction and 2–2.5% v/v maintenance

at 1 L/min flow rate). The left flank was shaved, and the skin sterilized with 70% ethanol and povidone

iodine. Under microscopic guidance, a retroperitoneal dissection was performed to expose the lateral

aspect of the spine using a Penfield dissector. The pelvis and hip were rotated posteriorly to enhance the

working space. The peritoneal wall was gently tugged to expose the psoas muscle, which was retracted

posteriorly to anteriorly using a cotton swab and then held in place with a metal spatula. The spinal

column and IVDs at the L5/6 and L6/S1 levels were exposed by blunt dissection. The superior margin

of the pelvic bone indicated the L6 vertebral body, confirming the locations of L5/6 and L6/S1 IVDs.

The L5/L6 IVD was targeted and either exposed in shams or injured with a 30G needle 3 times through

the annulus fibrosis and partially into the nucleus pulposus.[22,41]

Perioperatively, mice were monitored every 8–12 hours for 4 days post-surgery. Pain

management included intraperitoneal injections of carprofen (5 mg/kg) every 8–12 hours, supplemented

with carprofen tablets (2 mg/tablet) for 48-72 hours. On days 3 and 4 post-operatively, tamoxifen (TAM,

Cat#: T5648, Millipore Sigma) was given via oral gavage (100 mg/kg in corn oil, Cat#: C8267, Millipore

Sigma) to Cre+;VEGFA$^{fl/fl}$ animals (aka 'VEGFA-null') and Cre-;VEGFA$^{fl/fl}$ littermates ('WT')

inducing VEGFA ablation. Data collection occurred at 3 and 12 weeks with behavioral assays including

additional timepoints at baseline, within 7 days of injury, and longitudinal cohorts at 6 and 9 weeks.

Mouse weights were collected at baseline, 3, and 12 weeks to track overall health and survival was

tracked throughout the experiment to create Kaplan-Meier survival curves.

Acute 3-week post-op and chronic 12-week post-op time points were chosen with the acute

injury response allowing for the initial surgical wound healing recovery, while chronic LBP is defined

as pain lasting 12 weeks or longer.[42,43]

Figure S1: Experimental Design A) Lower lumbar disc puncture model showing the relative scale of

a 30G needle and the depth of injury B) Experimental design for various outcomes including injury at

time=0 with VEGFA ablation at 3-4 days Post-op, and acute and chronic terminal time points at 3 and

12 weeks. Baseline behavior within one week or surgery normalized behavioral outcomes.

Histological analyses

Histopathological evaluation of degeneration

To quantify acute IVD degeneration at 3 weeks, frozen 10 μm thick midsagittal sections were stained

with Safranin O / Fast Green and imaged at 10x on a Nanozoomer (Hamamatsu, Japan).

Histopathological scoring followed an established protocol for degeneration in four compartments of

the IVD including the NP, AF, end plate, and interface/boundary regions.[44] Mean degeneration scores

from three independent investigators (n=3/sex/genotype/surgery, N=24 total) were used for analysis.

Structural quantification of Nerve and Vascular features by Immunohistochemistry

Spines from the animals were fixed in 4% paraformaldehyde for approximately 48 hours, washed 6

times for 15 minutes each in 1X PBS, and subsequently demineralized in 14% EDTA for 10-14 days.

After which, samples were washed 8-10 times for 15 minutes in 1X PBS and serially infiltrated with

10%, 20%, and 30% sucrose solutions for 1 hour each prior to optimal cutting temperature (OCT)

infiltration and freezing for cryosectioning. Section thickness was 50 μm to provide a semi-3-

dimensional visualization of the neuronal and vascular morphology. Serial sections were collected with

CryoJane slides to adhere tissue to the slide during IHC staining with tissues being stored at -80°C prior

to antibody staining.

To stain for Protein Gene Product 9.5 (PGP9.5), Endomucin, and DAPI counter stain, a 4-day protocol[45]

was followed whereby slides were dried at room temperature for 30 minutes and washed in PBS 2 times

for 5 minutes. A PAP pen was used to create a hydrophobic perimeter around the tissue and allow for

direct pipetting of reagents onto the slides to avoid tissue loss from vertical slide immersion. Slides were

blocked with 10% donkey serum and 0.3% Triton X-100 in PBS for 1 hour at room temperature. The

primary PGP9.5 antibody (Millipore Cat # Ab1761) at a 1:1000 dilution and Endomucin antibody (Cat

#14-5851-85) at a 1:500 dilution was added to 1% normal donkey serum in a TNT (Tris-NaCl-Tween)

buffer and incubated 48 hours at 4°C in a humidified chamber. Samples were then washed 3 times for 5

minutes in TNT buffer prior to incubation of a PGP9.5 secondary antibody (Jackson Cat #711-545-152)

at 1:1000 concentration and 488nm fluorophore and Endomucin secondary antibody (Jackson Cat #712-

605-153) at 1:1000 concentration and 657nm fluorophore in 1% normal donkey serum in TNT buffer

for 24 hours at 4°C in a humidified chamber. Samples were washed 3 times with TNT for 5 minutes

prior to DAPI (Sigma-Aldrich D9542) at 1:1000 for 5 minutes at room temperature. A final wash with

TNT 3 times for 5 minutes was done prior to cover slipping and subsequent sealing of the slides with

Fluoromount-G a day later (Thermo Cat:00-4958-02).

To confirm that the PGP9.5 positive structures within the IVD are sensory neurons, we conducted a

direct comparison of the PGP9.5 straining of an injured intervertebral disc in a $Na_v1.8$ td tomato reporter

mouse and found the $Na_v1.8+$ features accounted for 88% of the PGP9.5+ features with a spatial

colocalization correlation of R=0.86.   The sodium-sensitive ion channel $Na_v1.8$ is highly specific to

sensory neurons.[46-48] A serial section as described above was obtained and then stained for PGP9.5. We

then examined the localization and specificity of PGP9.5 and $Na_v1.8$ within the same section.

DAPI, PGP9.5, and Endomucin helped visualize, respectively, cell nuclei, broad neural populations, and

endothelial cells of vasculature via a confocal microscope (Leica, USA) at 2.5 μm focal depth in

increments with 10x magnification and 2048x2048 resolution and 600 speed with 2 frame averaging.

Excitation and emission wavelengths were set as follows: DAPI excitation 405nm and emission 414-

450 nm, PGP9.5 excitation 488nm and emission 498-530 nm, and Endomucin excitation 635nm and

emission 650-725 nm. Note that there was no overlap between the Ai9 tdTomato reported with excitation

of 554 and emission of 581nm. Neurite and vessel quantification of the outer AF were performed on

max projections of a Z-stack with Fiji (ImageJ), with lengths on or within the AF and the surrounding

fibrous tissue calculated from the Neuroanatomy SNT plugin and with depths calculated normal to the

peripheral edge of the AF. PGP9.5 staining showed some non-specific regions so was only not included

if neuronal morphology was apparent.

To quantify spatial coupling of neurites and vessels, an automated MATLAB was developed to measure

lengths and depths and calculate colocalization lengths and a colocalization ratio based on distance from

the rarer neurite features. Spatial coupling was defined as vessels within 30 μm from neurites and based

on the distance between two adjacent lamellae as lamellar structure and proximity likely impact the

transmission of neurovascular coupling signaling.

Figure S3: Validation of Neuronal Stain with Injury A) Validation of PGP9.5 IHC with a Na$_v$ 1.8

showing the coincidence of the more specific sensory Na$_v$ 1.8 with pan neuronal PGP9.5. B-E) Acute (3

week) neurite length and depth and vessel length and depth data showing the increased lengths

associated with injured discs F) IHC of a herniated disc at 3 weeks showing diffuse non-specific green

PGP9.5 staining and red Endomucin vessels. Data are presented as mean ± standard deviation.

Neuropeptide immunostaining of the Dorsal Root Ganglia

Frozen DRG sections were processed for immunohistochemistry using a standardized protocol across

all targets. Briefly, slides were fixed in 4% paraformaldehyde for 10 minutes at room temperature. After

washing in 0.1 M Tris buffer (pH 7.6), non-specific binding was minimized through sequential blocking

steps with 0.005% BSA in Tris and 10% normal goat serum. Sections were incubated overnight at 4°C

with primary antibodies—anti-TrkA (1:100, Alomone Labs #ANT-018), anti-TRPA1 (1:200, Alomone

Labs #ACC-037), or anti-TRPV1 (1:500, Abcam #ab6166)—diluted in Tris-BSA. After rinsing, a goat

anti-rabbit Alexa Fluor 488 secondary antibody (1:300, Invitrogen #A11006) was applied for 4 hours at

room temperature or overnight at 4°C. Nuclear counterstaining was performed using DAPI (1:5 dilution

in 0.1 M Tris), and slides were mounted prior to imaging. DRG immunohistochemical images were

quantified using the custom Python software, DRGVisual. Contrast Limited Adaptive Histogram

Equalization (CLAHE) was applied to enhance the contrast of the images. The processing began with

edge detection using Canny edge detection to refine the segmentation and ensure accurate depiction of

cell boundaries. Following edge detection, seed points within the cells were identified to indicate the

center of the cells, and diameters were drawn to estimate each cell's maximum size. Cell regions were

grown using a projection-based region-growing algorithm that expands the region from each seed point

by examining neighboring pixels and including them in the region if they meet criteria based on intensity

and connectivity. For each cell, the corrected total cell fluorescence (CTCF) was calculated by

subtracting the background intensity from each pixel's intensity within the cell region and summing the

total fluorescence across the cells for each ion channel.

3D Structural Evaluation of Intervertebral Disc Structure Using μCT

At 3 weeks post lumbar intervertebral disc injury, in vivo μCT was used to evaluate structural changes

of the intradiscal space (VivaCT, 45 kVp, 177 uA, 300 ms integration time, 10 μm voxel) with isoflurane

inhalation (2-3% v/v). Disc height was calculated in ImageJ as the average of 5 measures across the

midsagittal view.[41] To confirm that the loss of VEGFA did not affect the structure of the IVD, we

examined the uninjured L4-L5 and L6-S1 IVDs between the VEGFA KO and the WT animals.

Additionally, we examined the injured L5-L6 IVD level to ensure that the VEGFA KO animals were

susceptible to injury-mediated degeneration. An independent 2-tailed t-test was used to determine differences between groups.

The structural changes at this 3-week post injury time point were further evaluated using the e*x vivo* contrast-enhanced micro-computed tomography (CEµCT). Ioversol as the contrast agent was used to highlight the water-rich nucleus pulposus.[49] The fresh IVDs were immersed in 50% ioversol (OptiRay 350, Guerbet Pharma) at 37°C for 24 hours before imaging. The Scanco40 µCT system (Zurich, Switzerland) scanned samples at 45 keVp, 177 µA, 10.5 µm voxel size, and 300 ms integration after the incubation[50]. Groups were n=3 mice/sex/genotype/injury. Total disc ROI and nucleus pulposus ROI were contoured using the MATLAB graphical user interface from Washington University Musculoskeletal Image Analyses (github.com). NP hydration, a ratio of nucleus attenuation to total disc attenuation (NI/DI), was calculated along with volumes and disc height index, DHI, as an average of 5 height measures in a midsagittal slice over the disc width.

Biomechanical Assessment of the Intervertebral Disc

Mechanical behavior of the IVDs using dynamic compression testing under displacement control was conducted using a BioDent reference point indenter (Active Life Scientific).[51] After CEuCT, these samples (n=3 mice/sex/genotype/injury) were mechanically tested with the strain rate calculated from the 5-point average disc height. FSUs were affixed to aluminum platens and immersed in a phosphate-buffered saline bath at room temperature and preloaded to 0.25N. A sinusoidal compressive waveform at 10% strain and 1 Hz frequency was applied for 20 cycles. The average stiffness was calculated from the second through the final loading cycles and the loss tangent, or dissipation factor, was calculated from the phase angle between load and displacement data.

Behavioral Assessments

Habituation measures were taken to reduce animal stress. Acclimation to room environment prior to behavioral assessment was set for 1 hour at the beginning of each behavioral testing week. To avoid additional animal stress, assessments were typically performed twice a day with 1 hour rest between tests between 9am-5pm to minimize the effect of activity changes due to diurnal variations.[52] All tests were performed by the same laboratory personnel. Baseline data was collected one week prior to surgery and post-operative data collection included a cross-sectional group at baseline, 3, and 12 weeks and a longitudinal group at 3, 6, 9, and 12 weeks. Care was taken to ensure a clean and odor-free environment. The procedures were conducted in a quiet room with controlled temperature (20-23°C). Testing was initiated up to one week prior to the 3- and 12-week time points as the full set of assays took 4-5 days to complete with efforts made to reproduce the order and timing of each test to remove stress on the animals and replicate methods.

Mechanical Sensitivity measured using the Von Frey and E-Von Frey

Mechanical sensitivity was measured using the up-down method of the Von Frey test using Semmes Weinstein filaments (Bioseb, France) and an electronic Von Frey (Figure 2-6C, EVF, BIOSEB, France).[53,54] Prior to baseline behavioral testing, mice were habituated to the elevated mesh grid chamber for 1 hr. Calibrated filaments were applied to the plantar surface (Figure 2-5D) of the hind paw to elicit paw withdrawal or filament buckling.[55] Filament Von Frey was done on the left hind paw to focus on the injury response. EVF consisted of maximum force values at withdrawal from the median of the result of 3 trials which were conducted bilaterally, with at least 5 min between testing opposite paws, and at least 10 min between testing the same paw. Post-op measures were normalized to baseline.

Hot and Cold Nociceptive Sensitivity

Mice were subjected to thermal stimulus to measure the temperature sensitivity response.[56,57] A hot/cold plate (Bioseb, France) was set to 55±0.5°C or 5±0.5°C and mice were placed individually onto the plate with a transparent cylinder to prevent escape and allow for visibility. The latency to the first sign of nociceptive behavior, either lick of the hand paws or jumping for hot plate and lifting, licking, or shaking of the hind paws for cold plate was recorded with a cutoff time of 20 seconds for hot or 30 seconds for cold to prevent tissue damage. Three repeated measures were collected with >15 minutes between tests to allow for home cage recovery. Values were averaged and normalized to baseline.

Functional Assessments of Performance

Locomotor Function – Rotarod

Assessing motor coordination and balance, the Rotarod performance test challenged mice to a rotating rod apparatus (Bioseb, France).[54,58,59] Prior to testing, mice were trained on the Rotarod at a slow 4 rpm for 2 minutes to ensure baseline capabilities. Mice unable to achieve this baseline capacity were excluded from assessment. On a subsequent day, mice were tested on an accelerating bar from 4 to 40 rpm over a 2-minute period. Latency to fall was recorded for each mouse. Each mouse underwent 5 trials with a >5-minute rest period between trials to reduce fatigue. The best performance of the 5 trials was used for data analysis and normalized to baseline when possible.

Open Field Testing

In the open field test designed to evaluate exploratory passive behavior, mice were individually introduced into the center of an open field arena (40 cm x 40 cm, with 30 cm high walls) equipped with an automated tracking system (Omnitech Electronics, Columbus, OH).[54,58,59] The apparatus featured an array of infrared sensors for precise movement tracking over a 60-minute test period. Behavioral parameters recorded included total distance traveled and instances of vertical rearing, among others.

Between tests, the arena was cleaned with Cavicide and water to disinfect and eliminate residual odors

and waste. The total distance and rearing events were reported over 60 minutes.

<u>Strength</u>

Inverted wire hang endurance

In the wire hang behavioral test, mice were evaluated for neuromuscular strength and endurance.[60] The

apparatus consisted of a wire mesh (2 mm diameter) suspended above a soft bedding layer to ensure a

safe landing for the mice. Each mouse was gently held by the base of the tail and allowed to grasp the

wire at a 60° incline before the mesh was inverted to 180°. The duration the mouse remained suspended

inverted was recorded up to a maximum of 2 minutes. A minimum 5-minute rest was given between

trails. The maximum time of three trials was recorded.

Grip Strength

Grip strength, a measure of neuromuscular strength, was measured using a uniaxial grip force tester

(Bioseb, France).[61] Hind paw grip strength assessed mechanical strength as the mouse was gently

restrained and allowed to freely grab and hold a bar with both hind paws. Maximum force was recorded

as the tail was pulled in a consistent manner to avoid rate-dependent effects. A brief rest period, >15

sec, was given between each of the 5 repeated trials. Average strength was reported normalized to body

weight at the time of testing and normalized to baseline.

Cell Culture

Human primary IVD Cell Culture

Human intervertebral disc cells were isolated from the disc tissue of eight patients (mean age ± SD =

68.5 ± 8.1; female:male = 5:3; mean Pfirrmann grade ± SD = 3.5 ± 0.7) removed during elective

discectomy surgeries (Supplemental Table 5). Human biospecimens were obtained under IRB

Exemption Category 4, in accordance with institutional and federal guidelines for research using de-identified, previously collected samples, as exemplified in prior studies (PMC7608136) and (PMC5536110). Tissue specimens were placed in sterile Ham's F-12 medium (F-12; Gibco-BRL, Grand Island, NY) containing 1% penicillin/streptomycin (P/S; Gibco-BRL) and 5% fetal bovine serum (FBS; Gibco-BRL). Disc tissue was washed three times with Hank's balance salt solution (HBSS; Gibco-BRL) containing 1% P/S to remove blood and other bodily contaminants prior to isolation. Specimens were minced and digested for 60 min at 37°C under gentle agitation in F-12 medium containing 1% P/S, 5% FBS, and 0.2% pronase (Calbiochem, La Jolla, CA). The specimens were rinsed three times with HBSS and 1% P/S and incubated overnight in a solution of F-12 medium containing 1% P/S, 5% FBS, and 0.025% collagenase I (Sigma Chemical Co, St. Louis, MO). A sterile nylon mesh filter (70 μm pore size) was used to separate suspended cells from the remaining tissue debris. Isolated cells were centrifuged (2000 rpm, 5 min) and resuspended in F-12 medium with 10% FBS and 1% P/S. Cells were plated in a 75-cm^2 culture flask (VWR Scientific Products, Bridgeport, NJ) and incubated at 37°C, 5% CO_2, 20% O_2 in a humidified incubator. Culture medium was changed three times weekly, and cultures grew to 90% confluence. Cells were then trypsinized (0.05% trypsin/ethylene diaminetetraacetic acid; Gibco-BRL) and re-plated (6 x 10^5 cells/mL) in 75-cm^2 culture flasks.

Age	Sex	T12/L1	L1/2	L2/3	L3/4	L4/5	L5/S1	Mean	STD	Endothelial Cell (RNA/ multiplex/ ELISA)	Matrigel Outgrowth Assays	with Neural Cell (RNA/neurite)
70	F		3	3	4	4	4	3.6	0.5		X	
73	F			3	3	4	5	3.8	1.0	X		
73	M			3	3	3	4	3.3	0.5			X
66	F	3	3	3	5	5		3.8	1.1	X	X	
50	M					3	4	3.5	0.7	X		
76	F		2	3	3	4		3.0	0.8			X
68	F					3	3	3.0	0.0			X
72	M	3	3	4	4	4	5	3.8	0.8		X	

Table S5: Patients' Demographics for Disc Cells (DCs): DCs were isolated from the disc tissue of eight patients removed during elective surgical procedures for degenerative spinal disease. Phirman Grades for T12/S1 are shown along with assay distributions for each patient sample.

Immortalized human microvascular endothelial cell (HMEC-1) culture

HMEC-1(CRL-3243; ATCC, Masassas, VA) were cultured in MCDB 131 medium (10372019; Gibco-BRL) containing 10 ng/mL epidermal growth factor (PHG0311; Thermo Fisher Scientific, Waltham, MA), 500 mg/mL hydrocortisone (H0396; Sigma), 1% P/S, 2 mM L-Glutamine and supplemented with 10% FBS. Cells at passage number 5-10 were plated at 75 cm^2 culture flask and grown to 90% confluence.

Immortalized human neuroblastoma cell (SH-SY5Y) culture

The SH-SY5Y human neuroblastoma cell line (CRL-2266; ATCC) was cultured in a 1:1 mixture of Eagle's Minimum Essential Medium (EMEM; ATCC, Manassas, VA) and Ham's F12 medium, supplemented with 10% FBS and 1% P/S in a 5% carbon dioxide humidified incubator at 37°C. Cells at passage number 5-10 were plated at 75 cm^2 culture flask and grown to 90% confluence.

Vascular Endothelial Growth Factor (VEGF)-siRNA Transfection

Isolated human IVD cells were seeded in 6-well plates at a density of 3×10^5 cells per well 24 hours prior

to transfection. VEGF-specific siRNA (sc-29620; Santa Cruz Biotechnology, Dallas, Tx) and scrambled

control siRNA- FITC (sc-36869; Santa Cruz Biotechnology) were transfected into DCs using

Lipofectamine™ RNAiMAX Transfection Reagent (Catalog #: 13778100, Thermo Fisher) following

the manufacturer's protocol. The final concentration of siRNA in the culture medium was 50 pM. The

effectiveness of transfection was evaluated by checking the amount of the secreted VEGF in media using

ELISA (DY293; R&D system, Minneapolis, MN) and confirming transfection of control siRNA by

FITC fluorescence with a Nikon Eclipse Ti2 inverted scope at 10x.

Co-culture between IVD cells and HMEC-1 (ECs) Cells

IVD cells were plated in a 6 well plate and transfected with either control, scrambled siRNA (Scram

siRNA) or with VEGF specific siRNA (VEGF siRNA). Medium with 1% FBS and recombinant human

interleukin-1β (IL-1β; R&D Systems) at 1 ng/mL was added to stimulate Scram siRNA (IL-1β+Scram

siRNA) and VEGF siRNA (IL-1β+VEGF siRNA) transfected IVD cells for 24 hours. After stimulation,

HMEC-1 cells "ECs" that were plated for 24 hours on 1 µm pore co-culture inserts (Thincert cell culture

insert; Greiner, Monroe, NC) at a density of 3×10^5 cells were inserted to the wells containing Scram

siRNA or VEGF siRNA transfected IVD cells with MCDB media plus 1% FBS. ECs co-cultured with

stimulated Scram siRNA treated IVD cells are denoted as $EC_{IL-1β+Scram\ siRNA}$ and ECs co-cultured

stimulated with VEGF siRNA treated IVD cells are denoted as $EC_{IL-1β+VEGF\ siRNA}$. Naïve ECs that did

not undergo co-culturing were used as a control group. ECs were analyzed via a Matrigel tube formation

assay, multiplex, and RNA sequencing. (Supplement Figure 6)

Figure S6. Schematic drawing of Co-culture between Disc Cells (DCs) and HMEC-1(ECs). DCs were plated on a 6 well plate left as naïve or transfected with Scrambled control siRNA (Scram siRNA) or VEGFA specific siRNA (VEGFA siRNA) for 24 hours. Cells were then treated with 1 ng/mL of IL-1β to stimulate the Scram siRNA and VEGF siRNA transfected DCs (IL-1β+Scram siRNA or siRNA IL-1β+VEGF siRNA, respectively) for 24 hours but the naïve DCs were left unstimulated. After stimulation, 1 μm pore co-culture inserts plated with ECs were inserted into each well for direct coculturing for 24 hours.

Co-culture between IVD cells and SH-SY5Y Cells (NCs)

SH-SY5Y neuronal cells "NCs" were plated at a density of 1.5×10^5 cells per well in a 6 well plate and cultured for 48 hours prior to co-culture experiment. For co-culturing, non-stimulated IVD cells "naive", IL-1β+Scram siRNA, and IL-1β+VEGF siRNA transfected IVD cells were plated on the co-culture insert at a density of 3.0×10^5 cells and co-cultured for 48 hours with NCs. NCs were analyzed via a neurite outgrowth assay and RNA sequencing. (Supplement Figure S7)

Figure S7. Schematic drawing of Co-culture between Disc Cells (DCs) and SH SY5Y cells (NCs). NCs were plated on a 6 well plate and cocultured with 1 μm pore co-culture inserts plated with DCs 48 hours. DCs were either left as naïve or transfected with Scrambled control siRNA (Scram siRNA) or VEGFA specific siRNA (VEGFA siRNA) for 24 hours. The naïve DCs were left unstimulated and the Scram siRNA and VEGF siRNA transfected DCs were stimulated with 1 ng/mL of IL-1β (IL-1β+Scram siRNA or siRNA IL-1β+VEGF siRNA, respectively) for 24 hours before coculturing.

RNA Sequencing and Gene Ontology

Post co-culture experiments, HMEC-1 and SH-SY5Y cells were lysed (TRIzol reagent, Fisher Scientific), and total RNA was extracted (Direct-zol RNA MicroPrep Kit, Zymo Research). Samples were prepared according to library kit manufacturer's protocol, indexed, pooled, and sequenced on an Illumina NovaSeq X Plus. Basecalls and demultiplexing were performed with Illumina's DRAGEN and BCLconvert version 4.2.4 software. RNA-seq reads were then aligned to the Ensembl release 101 primary assembly with STAR version 2.7.9a1. Gene counts were derived from the number of uniquely aligned unambiguous reads by Subread:featureCount version 2.0.32. Isoform expression of known Ensembl transcripts were quantified with Salmon version 1.5.23. Sequencing performance was assessed for the total number of aligned reads, total number of uniquely aligned reads, and features detected. The

ribosomal fraction, known junction saturation, and read distribution over known gene models were quantified with RSeQC version 4.04. All gene counts were then imported into the R/Bioconductor package EdgeR5 and TMM normalization size factors were calculated to adjust for samples for differences in library size. Ribosomal genes and genes not expressed in the smallest group size minus one sample greater than one count-per-million were excluded from further analysis. The TMM size factors and the matrix of counts were then imported into the R/Bioconductor package Limma6. Weighted likelihoods based on the observed mean-variance relationship of every gene and sample were then calculated for all samples and the count matrix was transformed to moderated log 2 counts-per-million with Limma's voomWithQualityWeights7. The performance of all genes was assessed with plots of the residual standard deviation of every gene to their average log-count with a robustly fitted trend line of the residuals. Differential expression analysis was then performed to analyze differences between conditions and the results were filtered for only those genes with Benjamini-Hochberg false-discovery rate adjusted p-values less than or equal to 0.05. The raw data files are available on the Gene Expression Omnibus database: GSE294577

For gene ontology analyses, biological pathways enriched by the upregulated and downregulated DEGs were identified by using the Panther Classification System with version Panther 19.0 by using the statistical overrepresentation test. For endothelial cells, an adjusted p value of $p < 0.05$ and logFC of 0.41 was used to identify statistically significant DEGs with at least a 50% increase in fold change of the treatment group when compared to the control. For neuronal cells, an adjusted p value of $p < 0.10$ and logFC of 0.41 was used to identify statistically significant DEGs with at least a 50% increase in fold change of the treatment group when compared to the controls. The full list of all GO terms for the endothelial cells comparisons are listed in Supplemental Tables X' and X' and for the neuronal cell comparisons are listed in Supplemental Tables X and X.

Multiplex Protein Assay

The supernatant from naive ECs and those co-cultured with IL-1β+Scram siRNA and IL-1β+VEGF siRNA transfected IVD cells was collected and analyzed by multiplex protein assay (Human Angiogenesis & Growth Factor 17-Plex, Human Cytokine Panel A 48-Plex, Human MMPTIMP 13-Plex, Eve Technologies, Calgary, Alberta, Canada). Out of range values or results < 2pg/ml were not included.

Matrigel Tube Formation Assay

To determine the vessel formation capacity between ECs cultured with conditioned media (CM) collected from IL-1β+Scram siRNA versus IL-1β+VEGF siRNA transfected IVD cells, a Matrigel tube formation assay was performed. Matrigel matrix (356231; Corning Inc., Corning, NY), a gelatinous protein mixture that mimics the extracellular matrix, was used to coat a 24-well plate following the manufacturer's protocol. HMEC-1 cells at the density of 3×10^4 cells per well were suspended in the CM collected from IL-1β+Scram siRNA or IL-1β+VEGF siRNA transfected IVD cells for 48 hours and plated onto each Matrigel coated well. Images were acquired at 6 hours and 12 hours post plating via a Nikon Eclipse Ti2 inverted scope at 10x. To assess the ability of tube formation by each conditioned media and time point, the numbers of closed loops were counted.[62]

Neurite Outgrowth Assay

The number and length of neurites from NCs co-cultured with non-stimulated IVD cells "naive", IL-1β+Scram siRNA, or IL-1β+VEGF siRNA transfected IVD cells were compared. Neurite outgrowth was quantified using ImageJ software. For each neuron, neurite number and length was counted manually. Neurite length was measured from the edge of the cell body to the tip of the neurite. The

average neurite length per neurite was calculated for each treatment group, with all measurements

expressed in microns.

Statistical Analysis

Statistical comparisons on repeated measures, IE longitudinal behavioral assays, were done using a

linear mixed effect model to assess surgery (Sham/Injured), genotype (WT/VEGFA-null), weeks after

surgery (3/12), sex (M/F), and surgery (Naïve/Sham). Linear mixed effect models were initiated with

all factors and reduced in complexity as factors were found to be not significant. Additionally, focused

questions were modeled to assess injured groups with genotype and week. A 2 or 3 factor ANOVA was

used on terminal measures such as Ex Vivo CEµCT, biomechanics, disc degeneration scoring, or

immunohistochemistry to assess effects of injury, genotype, and sex. ANOVA pairwise comparisons

were done with the Fisher's Least Significant Difference method. Survival curves implemented Mantel-

Cox test while binary penetration results used a Fisher's exact test. Simple pairwise comparisons were

done using a Welch's t-test when appropriate. Appropriate adjustments were made in the respective

statistical models for unequal variances and non-normal distributions. The ROUT outlier test with

Q=0.1% removed definitive outliers within groups.[63] R and GraphPad Prism were used to model results.

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List of Abbreviations

AF – Annulus Fibrosus

Ai9 – Cre-reporter expressing tdTomato fluorescent protein (Rosa-CAG-LSL-tdTomato)

ANOVA – Analysis of Variance

CEμCT – Contrast-Enhanced Micro-Computed Tomography

CLBP – Chronic Low Back Pain

CM – Conditioned Media

CTCF – Corrected Total Cell Fluorescence

C57BL/6J – Common inbred mouse strain

DAPI – 4′,6-Diamidino-2-Phenylindole (DNA stain)

DEG – Differentially Expressed Gene

DHI – Disc Height Index

DRG – Dorsal Root Ganglion

EC – Endothelial Cell

EDTA – Ethylenediaminetetraacetic Acid (chelating agent)

ELISA – Enzyme-Linked Immunosorbent Assay

EMCN – Endomucin (endothelial cell marker)

EMEM, ATCC – Eagle's Minimum Essential Medium, from American Type Culture Collection

ERT2 – Modified Estrogen Receptor Ligand Binding Domain (used in tamoxifen-inducible Cre systems)

EVF – Electronic Von Frey

FBS – Fetal Bovine Serum

FITC – Fluorescein Isothiocyanate dye

FSU – Functional Spinal Unit

HBSS – Hanks' Balanced Salt Solution

HMEC-1 – Human Microvascular Endothelial Cell Line

IL1β – Interleukin 1 Beta

IVD – Intervertebral Disc

LBP – Low Back Pain

LoxP – Locus of X-over P1 (recombination sites for Cre-Lox systems)

LSD – Least Significant Difference (statistical post hoc test)

MCDB – Molecular, Cell, and Developmental Biology

Nav1.8 – Voltage-Gated Sodium Channel (gene SCN10A) expressed in nociceptive neurons

NFkB – Nuclear Factor Kappa B

NGF – Nerve Growth Factor

NI/DI – Nucleus Attenuation Intensity/Disc Attenuation Intensity

NP – Nucleus Pulposus

OCT – Optimal Cutting Temperature Compound

PAP – Hydrophobic barrier pen

PBS – Phosphate-Buffered Saline

PGP9.5 – Protein Gene Product 9.5 (neuronal marker)

P/S – Penicillin/Streptomycin

ROI – Region of Interest

SCRAM – Scrambled siRNA (negative control)

SH-SY5Y – Human Neuroblastoma Cell Line

siRNA – Small Interfering RNA

SNT – Simple Neurite Tracer (ImageJ plugin)

TNFα – Tumor Necrosis Factor Alpha

TNT – Tris-NaCl-Tween buffer

TrKA – Tropomyosin Receptor Kinase A (NGF receptor)

TRPA1 – Transient Receptor Potential Ankyrin 1

TRPV1 – Transient Receptor Potential Vanilloid 1

UBC – Ubiquitin

VEGF – Vascular Endothelial Growth Factor

WT – Wild Type

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Author contributions:

Conceptualization: RSP, SYT

Methodology: RSP, REW, HJM, ALL, ANS, SWC, LJ, LAS, MCG, SYT

Data Interpretation: Methodology: RSP, REW, HJM, ALL, ANS, SWC, LJ, LAS, MCG, SYT

Investigation: RSP, HJM, ALL, LJ, LAS, MCG, SYT

Visualization: RSP, HJM, ALL, SWC

Funding acquisition: LAS, MCG, SYT

Project administration: RSP

Supervision: SYT

Writing – original draft: RSP, HJM, ALL

Writing – review & editing: RSP, SWC, SYT

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Co-inventors: RSP, HJM, MCG, SYT

Data and materials availability: All data, code, and materials used in the analysis must be available in

some form to any researcher for purposes of reproducing or extending the analysis. Include a note

explaining any restrictions on materials, such as materials transfer agreements (MTAs). Note accession

numbers to any data relating to the paper and deposited in a public database; include a brief description

of the data set or model with the number. If all data are in the paper and supplementary materials, include

the sentence