

Sex-dependent spinal expression of *SERPINA1*/Alpha-1 antitrypsin may predispose women to inflammation

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SIGNIFICANCE/CLINICAL RELEVANCE: Low expression of spinal alpha-1 antitrypsin (AAT-1) may partially account for the heightened frequency of inflammation and back pain in women.

INTRODUCTION: Women exhibit stronger inflammation-driven pain responses compared to men, suggesting a distinct interplay between sex and inflammation; potentially indicating a need for unique treatment options for women⁽¹⁾. Several bulk- and single-cell bioinformatics approaches^(2,3) have identified *SERPINA1* (human) / *serpinA1a-e* (mouse) as a prominent gene in intervertebral disc (IVD) cells during IVD degeneration, but the role of *SERPINA1* in the IVD and adjacent bone are unclear. *SERPINA1* (SERine Proteinase INhibitor A1) is a gene that encodes AAT-1 protein, which suppresses recruitment of myeloid-derived cells (macrophages, neutrophils) to the site adjacent of injury⁽⁴⁾ and is a protease inhibitor of elastase. AAT-1 deficiency exhibits a greater occurrence and prevalence in males, yet it is associated with a lower mortality rate compared to females⁽⁵⁾. We find by RNA-sequencing that IVD injury activates more inflammatory pathways in female mice than male mice (unpublished). Therefore, we hypothesized that *SERPINA1*/AAT-1 is differentially expressed in spinal tissues/cells by sex of mice and humans.

METHODS: This study is approved by IACUC. Tissues (liver, lungs, IVD, bone) were harvested from 4-month-old male and female mice (n=5/sex). The tissues were subsequently pulverized, and processed into mRNA, which was subsequently converted into cDNA to determine *serpinA1a* gene expression by qPCR. Lumbar L1-L3 was used for histological staining of SafraninO (IVD degeneration score) and AAT-1 (immunofluorescent), L3-L5 IVDs for gene expression, L6-S1 for IVD mechanics and L6 vertebra for μ CT. Bone marrow was extracted from tibia, femur, and ilium of C57BL6 mice and used as a cell source for osteoclast induction. While M-CSF facilitated the differentiation of marrow cells into macrophages, the combination of M-CSF+RANKL promoted osteoclast formation, and M-CSF+RANKL+IL-1 β supported both osteoclast formation and differentiation⁽⁶⁾. Ten human IVDs from male (n=5) and female (n=5) individuals within the age range of 28-85 years, were collected. The IVDs were categorized based on their degeneration level using the Thompson grading system, ranging from 1 (healthy) to 5 (most degenerated). Subsequently, these IVDs were processed and stained for AAT-1. A 2-way ANOVA was used to compare *serpinA1a* expression by tissue and sex or osteoclast expression of *serpinA1a* expression by osteoclast-differentiation by treatment (RANKL, RANKL+IL-1 β) and sex. A Student's t-test compared sex for all other comparisons, with statistical significance defined at a p<0.05.

RESULTS: *SerpinA1a* expression is differentially expressed across mouse tissues, where the order from most to least expression of the tissues harvested is liver, lung, IVD and bone (Fig. 1A). Further, female tissues expressed less *serpinA1a* than males in lung, IVD and bone cells (mostly osteocytes) by 6-fold, 4-fold and 2-fold, respectively. Compared to male murine IVDs, females expressed 97% less AAT-1 protein (green color) in the IVD, particularly annulus fibrosus (AF) cells (Fig. 1B, B'). Next, we focused on bone marrow-derived osteoclast differentiation because the histological expression of AAT1 was intense in the bone marrow of vertebrae (data not shown). In males, treatment to induce osteoclast differentiation (RANKL or RANKL+IL-1 β) reduced *serpinA1a* gene expression by ~0.4 fold but, in females, the treatment-induced osteoclast differentiation inconsistently reduced *serpinA1a* expression (interaction: p<0.05) (Fig. 1C). In human IVD, the AAT-1 protein expression in male IVD is less than that of women by 76% (Fig. 1D, D'). Investigation of the remaining outcomes is currently ongoing.

DISCUSSION: Males consistently exhibited higher *serpinA1a*/AAT-1 expression levels than females in various tissues. Notably, the annulus fibrosus of the IVD and bones expressed less *serpinA1a*/AAT among all tissues tested, which could promote greater recruitment of neutrophils and macrophages as occurs in the lungs of people with a genetic variant in *SERPINA1*⁽⁷⁾. In human IVDs, AAT1 protein expression was greater in male IVDs than female IVDs, highlighting a conserved sexual dimorphism across species. In bone, we harvested a predominately osteocyte-rich tissue for qPCR and the sexual disparity in the expression of *serpinA1a* was mild. However, the expression of *serpinA1a* with osteoclast differentiation was strikingly different between biological sex. In myeloid-derived osteoclasts, RANKL promotes osteoclast differentiation and the presence of RANKL downregulated *serpinA1a* in males and the inclusion of IL-1 β further promoted differentiation and suppression of *serpinA1a*. By contrast in bone marrow-derived female macrophages, RANKL appeared to upregulate *serpinA1a* during osteoclastic differentiation and IL-1 β downregulated *serpinA1a* with further differentiation. Overall, the sexual dimorphic expression of *serpinA1a* in the spine and regulation of *serpinA1a* during osteoclast differentiation suggest that inflammatory signaling may yield greater inflammatory-related consequences to the musculoskeletal system of women than men.

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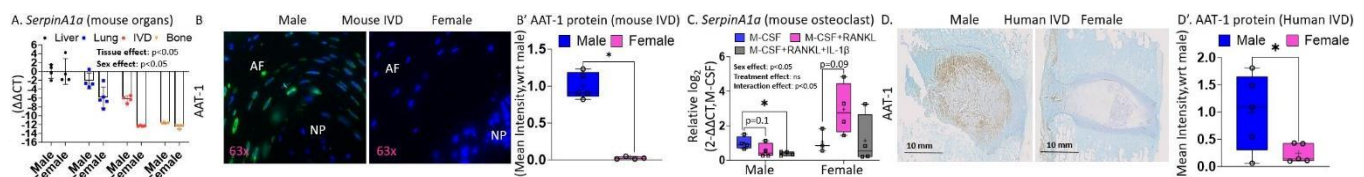


Fig. 1. (A) Gene expression of *serpinA1a* in male (n=5) and female (n=5) mouse tissues (Liver, Lung, IVD, bone), (B) Representative immunofluorescent images of AAT-1 (magnification 63x) in CON male (n=5) and female (n=5) IVDs. Blue color (DAPI) represents cell nuclei while green is the AAT-1 (B') Quantification of AAT-1 protein in the mouse male (n=5) and female (n=5) IVD, (C) Gene expression of *serpinA1a* in the macrophages (M-CSF), pre-osteoclast (M-CSF+RANKL) and osteoclast (M-CSF+RANKL+IL-1 β) (n=4/group), (D) Representative immunohistochemistry images of AAT-1 in the human male and female IVD, (D') AAT-1 protein expression in the human IVDs of male and female (N=5/sex), *: p<0.05, ns-not significant