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HLA-B27 Alters the Response to TNFα and Promotes Osteoclastogenesis in Bone Marrow Monocytes from HLA-B27 Transgenic Rats

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Abstract

Objective—To determine whether HLA-B27 expression alters the response of bone marrow monocytes (BMMo) from HLA-B27/human β_2 -microglobulin transgenic (B27-Tg) rats to tumor necrosis factor- α (TNF α), and whether this affects cells involved in bone homeostasis.

Methods—BMMo were treated with receptor activator of NF- κ B ligand or TNF α to promote osteoclast formation. Osteoclasts were quantified by counting. Gene expression was measured using quantitative polymerase chain reaction, and protein was detected by enzyme-linked immunosorbent assay, immunoblotting, or immunofluorescence. Effects of endogenously produced cytokines on osteoclast formation were determined with neutralizing antibodies.

Results—TNF α enhanced osteoclast formation 2.5-fold in HLA-B27-expressing cells compared to either wild type or HLA-B7/human β_2 -microglobulin expressing monocytes. TNF α induced approximately 4-fold upregulation of HLA-B27, which was associated with accumulation of misfolded heavy chains, binding of the ER chaperone BiP, and activation of an ER stress response, which was not seen with HLA-B7. No differences were seen with RANKL-induced osteoclastogenesis. Enhanced interleukin-1 α (IL-1 α) production from ER stressed B27-Tg BMMo was found to be necessary and sufficient for enhanced osteoclast formation. However, B27-Tg BMMo also produced more interferon- β (IFN β), which attenuated the effect of IL-1 α on osteoclast formation.

Conclusions—HLA-B27-induced ER stress alters the response of BMMo from B27-Tg rats to TNF α , which is associated with enhanced production of IL-1 α and IFN β , cytokines that exhibit opposing effects on osteoclast formation. The altered response of cells expressing HLA-B27 to pro-inflammatory cytokines suggests that this MHC class I allele may contribute to the pathogenesis of spondyloarthritis and its unique phenotype through downstream effects involving alterations in bone homeostasis.

Keywords

Rodent; MHC; inflammation

INTRODUCTION

The role of HLA-B27 in the pathogenesis of spondyloarthritis remains poorly understood (1). Most hypotheses focus on HLA-B27 as an upstream factor that is envisioned to trigger an aberrant immune response – either adaptive or innate - resulting in chronic inflammation.

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For example, peptide-loaded HLA-B27- β_2 -microglobulin (β_2 m) complexes could be targeted by autoreactive CD8+ T cells (2), or β_2 m-free heavy chain homodimers might trigger the survival and activation of CD4+ Th17 T cells (3). HLA-B27 and human β_2 m expression in transgenic (B27-Tg) rats has been linked to dendritic cell (DC) dysfunction (4) and increased apoptosis (5, 6), that could contribute to enhanced DC-mediated activation of CD4+ Th17 T cells (7) and/or failure to induce tolerance. In addition, the propensity of HLA-B27 heavy chains to misfold and generate endoplasmic reticulum (ER) stress (8, 9), initiates intracellular signals through the unfolded protein response (UPR) that converge with Toll-like receptor (TLR) signaling pathways to promote cytokine production, including interleukin (IL)-23 (IL-23) (10–12). Systemic IL-23 expression has recently been shown to activate an unusual population of enthesis-resident T cells in mice that produce proinflammatory cytokines such as IL-17 and IL-22, providing an explanation for some of the unique anatomical features of the spondyloarthritis phenotype (13). The relative contribution of these mechanisms to human disease remains an area of intense investigation.

Despite evidence implicating HLA-B27 as an upstream factor in spondyloarthritis, its tendency to misfold and generate ER stress, particularly when upregulated (8, 9, 14), raises the possibility that HLA-B27 may have a downstream impact on disease pathogenesis by altering the response of cells to pro-inflammatory cytokines or other factors.

One aspect of spondyloarthritis that is poorly understood is the excessive bone loss prominent in vertebral bodies, juxtaposed with pathologic bone formation that occurs along the spine of many patients with ankylosing spondylitis (AS) (15–17). One of the several features of spondyloarthritis reproduced in B27-Tg rats is trabecular bone loss in vertebral bodies and long bones (18). Since bone integrity is regulated by osteoclasts and osteoblasts we asked whether HLA-B27 expression could influence osteoclast formation in response to known stimulators of osteoclastogenesis, receptor activator of NF- κ B ligand (RANKL) and tumor necrosis factor-alpha (TNF α).

MATERIALS and METHODS

Animals

Rats were housed in AAALAC-approved facilities on the Bethesda NIH campus, and all animal experiments were pre-approved by the NIAMS Animal Care and Use Committee. Lewis HLA-B*27:05 and human β_2 m transgenic rats (B27-Tg) were generated by backcrossing the 33-3 transgene locus from Dark Agouti (DA) B27-Tg, onto the Lewis background for over 10 generations. The 33-3 transgene locus contains 55 copies of HLA-B27 and 66 copies of human β_2 m (19). Lewis HLA-B*07:02 and human β_2 m transgenic (B7-Tg) rat breeders carrying the 120-4 transgene locus were obtained from Joel Taurog (Dallas, TX). The 120-4 locus contains 26 copies of HLA-B7 and 5 copies of human β_2 m (20). Hemizygous B27-Tg rats were used in this study, while homozygous B7-Tg rats were used (i.e. 52 copies of HLA-B7 and 10 copies of human β_2 m). Transgene negative littermates or Lewis rats purchased from Taconic (Rockville, MD) were used as wild type (WT).

Bone marrow-derived monocytes and osteoclast development

Bone marrow was obtained from tibias and femurs of euthanized 6–9 week old rats. Mononuclear cells were isolated using Lympholyte-Rat (Cedarlane Laboratories Inc., Burlington, NC), washed twice, and re-suspended at 5×10^6 cells/ml in Gibco DMEM high glucose medium (Invitrogen, Grand Island, NY) containing 10% FCS (Invitrogen), 10^{-8} M 1α ,25-dihydroxyvitamin D₃ (Enzo Life Sciences, Inc., Farmingdale, NY), and 20 ng/ml rat macrophage colony stimulating factor (M-CSF) (Peprotech Inc., Rocky Hill, NJ), and

cultured for two days at 37°C in a 5% CO₂ humidified atmosphere. To evaluate effects of cytokines on osteoclast formation, non-adherent bone marrow monocytes (BMMo) were collected, adjusted to 10^6 cells/ml in media, and seeded at 10^5 cells per well in 96 well plates with M-CSF (20 ng/ml final concentration) to examine the effects of RANKL (receptor activator of NF- κ B ligand) and TNF α (tumor necrosis factor-alpha) on osteoclast differentiation. Cytokines were obtained from Peprotech Inc. (Rocky Hill, NY), and used at final concentrations indicated in the figures. BMMo were treated with RANKL for 3 days, or with TNF α for 5 days, which were established in preliminary experiments as optimal conditions for osteoclast formation. Mature osteoclasts were visualized by tartrate resistant acid phosphatase (TRAP) staining (Kamiya Biomedical Company, Seattle, WA), photographed, and quantitated by counting as TRAP-positive cells with 3 nuclei normalized to area.

Reagents

Neutralizing antibodies were goat anti-rat IL-1 α (AF500, R&D Systems, Minneapolis, MN), hamster anti-rat IL-1 β (4-7012, eBioscience, San Diego, CA), and rabbit anti-rat IFN β (CL 9143AP, Cedarlane Laboratories Inc., Burlington, NC). All antibodies were used at a final concentration of 1 µg/ml during the entire 5 days of TNF α induced osteoclast development. Thapsigargin (Sigma-Aldrich, St. Louis, MO) was used at a final concentration of 300 nM and tunicamycin (Sigma-Aldrich) at 10 µg/ml.

RNA isolation and quantitation

For evaluation of gene expression, 10^6 cells per sample were collected by centrifugation ($400 \times g$ for 5 min), lysed in Trizol (Life Technologies, Grand Island, NJ) and stored at -80° C. RNA was isolated according to the manufacturers protocol. Complementary DNA (cDNA) was generated using iScript (Biorad, Hercules, CA), and quantitative polymerase chain reaction (qPCR) measurements of specific genes were carried out using a MyiQ cycler with the SsoFast EvaGreen Supermix (Bio-Rad Laboratories, Hercules, CA) with primers (500 nM) and cDNA corresponding to 20–50 ng of starting RNA. Three house keeping genes were used for normalization of target gene expression levels: peptidylprolyl isomerase (*Ppia*), hypoxanthine phosphoribosyl transferase (*Hprt1*), and TATA binding protein (*Tbp*). Fold changes were calculated by comparing normalized CT values of treated to normalized untreated samples using the qBasePlus software (Biogazelle, Ghent, Belgium). For determining *Xbp1* mRNA splicing, reverse transcriptase PCR products were separated on 3% agarose gels and visualized with ethidium bromide (Bio-Rad) under ultraviolet light. Primer sequences are listed below.

Gene	Forward (5'-3')	Reverse (5'-3')
Hprt1	GAC TTT GCT TTC CTT GGT CA	AGT CAA GGG CAT ATC CAA CA
Ppia	CTG GTG GCA AGT CCA TCT AC	CCC GCA AGT CAA AGA AAT TA
Tbp	CGA TAA CCC AGA AAG TCG AA	AGA TGG GAA TTC CAG GAG TC
HLA-B	GTC CAC CGT CCC CAT CG	ACG CAG CCT GAG AGT AGC
Xbp1	ACA CGC TTG GGG ATG AAT GC	CCA TGG GAA GAT GTT CTG GG
Hspa5 (Bip)	TCA GCC CAC CGT AAC AAT CAA G	TGT CTT CCT CAG CAA ACT TCT CG
Il1a	ACA GTT CTG CCA TTG ACC AT	TGT TTG ATG TTG CTG ACA CC
Ifnb1	TGA ATG GAA GGC TCA ACC TCA G	CGT GGA TGT CAC CCA AGT CAA TC

Immunoblotting and immunoprecipitation

BMMo $(2.5 \times 10^7 \text{ cells})$ were harvested, re-suspended in 1 ml ice cold PBS with 10 mM methyl methanethiosulfonate (MMTS) (Thermo Scientific, Rockford, IL) on ice for 20 min to prevent spontaneous sulfhydryl bond formation and reduction. Cells were then spun down and lysed (500 µl per 10⁷ cells) for 30 min on ice in 20 mM Tris, pH 7.8, 100 mM NaCl, 10 mM EDTA, 1% Triton and the Halt Protease Inhibitor Cocktail (Thermo Scientific, Rockford, IL). Nuclei were removed by centrifugation at $13,000 \times g$ for 5 min, and supernatants were collected and used for immunoprecipitation or SDS-polyacrylamide gel electrophoresis (SDS-PAGE) and immunoblotting. For SDS-PAGE, lysates were diluted 1:1 with Tris-Glycine SDS sample buffer (Novex, Invitrogen, Carlsbad, CA) with or without NuPage Sample Reducing Agent (Invitrogen), and subjected to electrophoresis on 4–20% Tris-glycine gels (Novex) followed by immunoblotting. The following primary antibodies were used: HC10 to detect HLA class I heavy chains, rabbit anti-BiP (ab21685, Abcam, Cambridge, MA), and mouse anti-GAPDH (sc-32233, Santa Cruz). Appropriate speciesspecific secondary antibodies were alkaline phosphatase-conjugated (Southern Biotech, Birmingham, AL), and the 5-bromo-4-chloro-3-indolyl phosphate/nitro blue tetrazolium substrate (Thermo Scientific, Rockford, IL) was used to visualize proteins. HLA-B27 immunoprecipitation from 500 µl of cell lysate was performed as described previously (21). Immunoprecipitates were subjected to SDS-PAGE and immunoblotting as described above, except that HLA-class I heavy chains were detected using 3B10.7 (22).

Immunofluorescence

Immunofluorescence staining was used to identify BiP and HLA-B27-expressing cells during osteoclast development. Four days after the addition of RANKL or TNF α to BMMo, cells were fixed with 4% formaldehyde, washed twice with BD Perm/Wash buffer (BD Biosciences, San Jose, CA) and then incubated in the same buffer for 15 min at 4°C to permeabilize the cells. Primary antibodies (anti-BiP, ab21685 [Abcam, Cambridge, MA] or HC10 (23)) were added at 1 µg/ml and cells were incubated for an additional hour at 4°C. Cells were then washed 3 times with Perm/Wash buffer and incubated with secondary antibodies (Alexa Fluor 594-labeled chicken anti-rabbit; Invitrogen, Grand Island, NY, or FITC-labeled chicken anti-mouse IgG; LifeSpan Biosciences Inc., Seattle, WA) diluted 1:100 in Perm/Wash buffer for 30 min at 4°C in the dark. Cells were washed again and visualized with an Axiovert Zeiss 40 CFL fluorescence microscope at 20× magnification.

Statistical analysis

Values are expressed as means with standard deviation (SD) depicted by error bars, unless otherwise noted. Statistical analysis was performed using Student's *t* test. *P* values less than 0.05 were considered significant. For ratios, the mean and 95% confidence intervals are shown.

RESULTS

HLA-B27 expression promotes osteoclast formation

To determine whether HLA-B27 has an influence on osteoclast development, BMMo from WT and B27-Tg rats were differentiated with RANKL or TNFα. There was no difference in the number of osteoclasts formed in the presence of RANKL (Figure 1A, B), although we observed a trend toward larger osteoclasts with more nuclei in the presence of HLA-B27 and HLA-B7 (Figure 1A and unpublished observations). In contrast, TNFα treatment consistently resulted in greater osteoclast formation in BMMo derived from B27-Tg animals (Figure 1C–F). Representative experiments are shown in Figure 1C–E, with the average fold change from several experiments shown in Figure 1F. The average increase in

osteoclastogenesis in HLA-B27 expressing cells treated with TNF α was 2.5 fold (Figure 1F). Experiments with BMMo derived from B7-Tg rats used as a control for HLA class I overexpression revealed no difference in osteoclast formation compared with WT rats (Figure 1E, F), indicating the effect is specific for HLA-B27. Osteoclasts formed from WT and B27-Tg BMMo were active and exhibited similar resorption patterns on calcium phosphate-coated slides, indicating they were functional (unpublished observations).

The roles of IL-1 α and IFN β in HLA-B27-induced osteoclast formation

IL-1 α and IFN β have been shown to regulate osteoclast formation. IFN β is a potent inhibitor (24), while IL-1 α promotes TNF α induced osteoclastogenesis (25). To determine whether these cytokines were involved in mediating the effect of HLA-B27 expression on TNF α -induced osteoclastogenesis, each cytokine was blocked with a neutralizing antibody. Blocking IL-1 α completely inhibited the effect of HLA-B27 on TNF α -induced osteoclastogenesis, whereas an IL-1 β neutralizing antibody had no effect (Figure 2A). The addition of HC10, which recognizes HLA-B27 homodimers as well as free heavy chains, also had no effect on osteoclastogenesis, IL-1 α was added to TNF α -treated WT BMMo, where it enhanced osteoclast formation approximately 2-fold (Figure 2B). HLA-B27-expressing cells produced more immunoreactive IL-1 α protein (Figure 2C) and exhibited greater induction of *Il1a* mRNA in response to TNF α (Figure 2D), consistent with this cytokine being responsible for the effect of HLA-B27 on osteoclastogenesis.

In contrast to the results with IL-1 α , neutralizing IFN β further enhanced osteoclastogenesis in cultures of HLA-B27 expressing cells (Figure 2A). This effect was associated with a several-fold induction of *Ifnb1* mRNA in HLA-B27-expressing cells following exposure to TNF α , whereas WT and HLA-B7-expressing BMMo exhibited only a small increase (Figure 2E). We have not found a reliable ELISA for rat IFN β , and thus are unable to quantitate protein production in BMMo cultures. Taken together with the results for IL-1 α , these data suggest that HLA-B27-expressing cells produce more IL-1 α and IFN β with opposing effects on osteoclastogenesis, although under these experimental conditions the net result is greater osteoclast formation.

TNFα induces ER stress in HLA-B27-expressing BMMo

To explore differences between B27-Tg and WT BMMo that might contribute to greater TNF α induced osteoclastogenesis, we compared effects of TNF α and RANKL on HLA-B27 expression and misfolding. TNF α treatment resulted in 4-fold upregulation of HLA-B27 mRNA, with a comparable increase in heavy chain expression, while RANKL had no effect (Figure 3A, B, E and data not shown). TNF α -induced HLA-B27 heavy chains accumulated as disulfide-linked dimers and multimers, as well as conventional monomers (Figure 3A). There was also an increase in co-immunoprecipitation of BiP with HLA-B27 heavy chains (Figure 3B).

The accumulation of disulfide-linked HC10-reactive forms of HLA-B27 with enhanced BiP binding in response to TNF α is indicative of heavy chain misfolding, and reminiscent of previous results demonstrating that IFN γ upregulation of HLA-B27 exacerbates misfolding in macrophages (9). This prompted us to look for evidence of ER stress. HLA-B27-expressing BMMo exhibited increased *Bip* mRNA and protein, and *Xbp1* mRNA splicing (Figure 3C–E) when treated with TNF α , but not RANKL. These data demonstrate that HLA-B27-expressing BMMo exhibit ER stress and activate the UPR when treated with TNF α , but not RANKL.

ER stress upregulates IL-1 α and IFN β

To determine whether ER stress alone is sufficient to induce IL-1 α and IFN β expression, BMMo from WT rats were treated with thapsigargin, an inhibitor of the sarco/endoplasmic reticulum Ca²⁺ adenosine triphosphatase (SERCA) that prevents autophagosome-lysosome fusion and induces ER stress (26). UPR activation resulting in *Bip* mRNA upregulation and *Xbp1* splicing was readily apparent within 3 hours of treatment of cells (Figure 4 A, B), and was accompanied by strong upregulation of *Il1a* and *Ifnb1* mRNAs (Figure 4 C, D).

Cellular localization of TNFα-induced HLA-B27 and BiP

To determine whether BMMo (osteoclast precursors) or mature osteoclasts were exhibiting effects of TNF α on HLA-B27 expression and UPR activation, BMMo treated with RANKL or TNF α were assessed by immunofluorescence microscopy for BiP and HLA class I heavy chain expression. Increased BiP and HLA class I heavy chain staining was noted only with TNF α treatment and not RANKL, and was most prominent in small round/oval cells with single nuclei (BMMo; white arrows), rather than large osteoclasts with multiple nuclei (blue arrows) (Figure 5). In contrast, RANKL had no effect on HLA-B27 or BiP staining in B27-Tg cells, and neither TNF α nor RANKL had any effect on BiP staining in WT BMMo (Figure 5). These data are consistent with the immunoblot analyses shown previously (Figure 3), and further indicate the predominant effect of TNF α on HLA-B27-induced ER stress is in osteoclast precursors.

DISCUSSION

The propensity of HLA-B27 to misfold, generate ER stress, and activate the UPR has the potential to exert biological effects in many different cell types expressing MHC class I molecules. We previously documented effects of HLA-B27 misfolding in rat macrophages where UPR activation after upregulation of HLA-B27 with IFN γ promotes increased expression of IL-23 and IFN β in response to TLR agonists (1, 8, 10). In earlier work we found no biochemical evidence, such as heavy chain oligomerization or prolonged BiP binding, to suggest that HLA-B7 misfolds (1). Furthermore, in these studies upregulation of HLA-B27 (8, 10). Since these earlier comparisons were made with IFN γ -treated macrophages, we also examined HLA-B7-expressing monocytes treated for 20 hours with TNF α (e.g. Fig. 2D, E), and found no UPR activation (unpublished observations). Thus, comparable overexpression in rat cells of another HLA-B allele that is closely related to HLA-B27 but does not misfold, does not appear to cause ER stress, at least under the conditions examined to date.

Here, we show that HLA-B27-expressing rat monocytes cultured with M-CSF and stimulated with TNF α alone, also exhibit ER stress and UPR activation linked to HLA-B27 misfolding, with enhanced expression of IL-1 α and IFN β , while these effects are not seen with HLA-B7. More importantly, the autocrine/paracrine effects of IL-1 α result in enhanced differentiation of monocytes into osteoclasts. Interestingly, the effect of TNF α is almost doubled when IFN β is neutralized (Figure 2A). Taken together, these results indicate that HLA-B27-expressing rat monocytes produce more pro- and anti-osteoclastogenic cytokines when stimulated with TNF α , with IL-1 α promoting the effects of TNF α , with IFN β attenuating the stimulatory effect of IL-1 α . The counter-regulatory effect of IFN β is insufficient to prevent the pro-osteoclastogenic effect of IL-1 α , at least under current experimental conditions.

ER stress-induction of IFN β has been shown to occur via the UPR transcription factor XBP1 (10). The active transcription factor produced from the spliced *Xbp1* mRNA (XBP1s) binds

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to an enhancer to promote transcription of the *Ifnb* gene (27). The greatest effect of ER stress on *Ifnb* mRNA induction is seen when macrophages are exposed to TLR4/3 agonists that already induce IFN β , such as lipopolysaccharide (LPS) and double stranded RNA (10, 27). However, even in the absence of TLR agonists, HLA-B27-expressing rat macrophages exhibit low-level (~3-fold) *Ifnb* mRNA upregulation when stimulated with IFN γ (1). Our current results extend these previous findings in an important way by showing TNF α can act as an inducer of low-level IFN β expression in monocytes experiencing ER stress due to HLA-B27 expressing cells, where *Ifnb* mRNA is increased only 2–4-fold) in response to TNF α (Figure 2E). Moreover, our results establish that even a low-level of IFN β produced from HLA-B27-expressing cells can be biologically significant as an inhibitor of osteoclastogenesis.

Effects of ER stress on IL-1 α production have not, to our knowledge, been reported previously. We had observed *Il1a* mRNA induction in a pilot experiment using tunicamycin (unpublished observations), which activates the UPR by inhibiting protein glycosylation. We show here that thapsigargin has a similar effect on *Il1a* mRNA, indicating that *Il1a* induction is a consequence of ER stress, and unlikely due to some other effect of HLA-B27 expression. We have not further investigated the mechanism of *Il1a* upregulation, although enhanced ER stress-induced NF- κ B and/or AP-1 activation could contribute. Interestingly, ER stress has been shown to activate the NLRP3 inflammasome and increase IL-1 β production by a mechanism independent of classical UPR activation (28). However, in contrast to IL-1 α , we do not see increased IL-1 β production by HLA-B27-expressing rat monocytes (unpublished observations), so inflammasome activation seems unlikely to be playing a role.

Opposite effects of IL-1 α and IFN β on osteoclast development have been documented previously (24, 25). Kobayashi *et al.* demonstrated IL-1 α was required for the formation of fully functional osteoclasts induced by TNF α in a RANKL/RANK-independent mechanism (25). The inhibitory effect of IFN β on osteoclast formation was demonstrated in the context of RANKL-induced osteoclast formation, where small amounts of IFN β induced by RANKL and signaling through an autocrine/paracrine loop, inhibit c-Fos expression which is essential for osteoclast formation (24). We have not further investigated the mechanism by which IFN β inhibits TNF α -induced osteoclastogenesis in HLA-B27-expressing rat BMMo, but since TNF α also induces c-Fos, it is plausible to expect that the mechanism would be similar. It is important to note that in our system, TNF α -induced IFN β does not appear to be limiting osteoclast formation in WT BMMo, since there is little difference with and without α -IFN β (Figure 2A) and little IFN β mRNA induction in WT or HLA-B7-expressing cells (Figure 2E).

Trabecular bone loss resulting in osteopenia and osteoporosis is a significant problem in SpA and particularly in AS (17, 29–31). Bone loss often begins early in disease, prior to physical immobilization (15). Osteoclastic bone resorption occurs in the sacroiliac and zygapophyseal joints of patients with AS (32, 33), and is even present in long-standing disease where osteoclasts can be found adjacent to fibrous tissue (34). It is most severe in the spine where it increases the risk of vertebral fractures, and in many individuals is anatomically juxtaposed with areas of abnormal bone formation such as syndesmophytes. TNFa inhibition significantly increases bone mineral density in SpA (35) and AS (36), implicating this cytokine in bone loss, while effects of TNF inhibitors on syndesmophyte growth remain unclear. Indeed, whether or not trabecular bone loss is coupled with bone formation is the subject of much debate (37).

B27-Tg rats carrying the 33-3 transgene locus serving as a model for SpA-like disease also exhibit substantial loss of trabecular bone in vertebral bodies, large bones of the leg, and reduced bone strength (18). Bone loss in this model occurs primarily as a consequence of increased bone resorption rather than reduced formation (38), and is associated with increased RANKL expression relative to OPG (39). TNFa also contributes to the inflammatory disease phenotype in B27-Tg rats, although a role in bone loss has not been established. Our results raise the possibility that HLA-B27 expression contributes to TNFainduced bone loss in this model and in human SpA including AS, by promoting local production of IL-1a. Since these transgenic rats have multiple copies of the HLA-B27 transgene, they overexpress HLA-B27 heavy chain relative to human cells. Although results with HLA-B7 indicate that heavy chain misfolding is required to generate ER stress in this model, it may still occur more readily than in humans due to greater expression. It is noteworthy that Feng et al. have recently shown that peripheral blood mononuclear cells (PBMC) from HLA-B27 positive AS patients, but not healthy controls, generate sufficient ER stress to activate the UPR when treated with IFN_Y (14). In previous studies, HLA-B27expressing PBMC-derived macrophages from AS patients did not exhibit UPR activation when treated with IFN γ (40, 41). This may have been due to a lack of (40) or insufficient (41) upregulation of HLA-B in macrophages derived ex vivo. It is also possible that monocytes and macrophages respond differently to cytokine treatments. Indeed, monocyte to macrophage differentiation is itself associated with UPR activation, which in turn protects cells against further ER stress (42). It will be important to better characterize the response of PBMCs and isolated monocytes from HLA-B27 positive individuals to cytokines that upregulate MHC class I, including TNFa.

In conclusion, we demonstrate that HLA-B27 alters the response of rat bone marrow-derived monocytes to TNF α , and promotes osteoclastogenesis by causing ER stress and enhancing IL-1 α production. This also increases production of the anti-osteoclastogenic cytokine, IFN β . While the net effect under the conditions studied here is to promote osteoclastogenesis, this balance could be altered *in vivo* by other mediators of inflammation. These findings suggest that this unusual MHC class I molecule may exert downstream effects in the pathogenesis of spondyloarthritis with implications for bone remodeling and the unique spondyloarthritis phenotype.

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Non-standard abbreviations used in this paper

$\beta_2 m$	β_2 -microglobulin
BMMo	bone marrow monocyte
ER	endoplasmic reticulum
$h\beta_2m$	human β_2 -microglobulin
UPR	unfolded protein response
TG	thapsigargin
XBP1	X box binding protein-1

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Figure 1.

Augmentation of TNF α -induced osteoclast formation by HLA-B27. BMMo from WT and B27-Tg (B27) rats were treated with M-CSF and (**A**, **B**) increasing concentrations of RANKL for three days, or (**C**, **D**) TNF α for five days. **A**, **C**, TRAP staining from a representative experiment (original magnification 5×). **B**, **D**, Number of osteoclasts, defined as TRAP positive cells with 3 nuclei per unit area. **E**, Representative results from an experiment comparing TNF α -induced osteoclast formation with BMMo derived from WT, B7-Tg (B7) and B27-Tg (B27) animals. *P* < 0.05 for B27 vs. B7 or WT. **F**, Fold increase (compared to WT) in number of osteoclasts induced by TNF α (30 ng/ml) using BMMo derived from B7-Tg (B7) and B27-Tg (B27) animals. Values in **B** and **D** represent means with SD indicated by error bars of triplicate wells from a representative experiment. Results in **F** represent averages, with 95% confidence intervals indicated by error bars, from 3 (B7) and 6 (B27) independent experiments. * = *P* < 0.05.

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Figure 2.

Enhanced TNF α -induced osteoclast formation in HLA-B27-expressing cells is IL-1 α dependent and inhibited by IFN β . **A**, Neutralizing antibodies (1 µg/ml) against IL-1 α (α -IL-1 α), IFN β (α -IFN β), IL-1 β (α -IL-1 β), or HLA-B class I heavy chains (HC10) were added to WT and B27-Tg BMMo during TNF α (30 ng/ml) induced osteoclastogenesis. Osteoclast formation was quantitated as described for Figure 1. **B**, The effect of IL-1 α (0.1 ng/ml) on TNF α (30 ng/ml) induced osteoclast formation in WT BMMo. **C**, IL-1 α production in culture supernatants from TNF α treated (30 ng/ml for 3 days) WT and B27 BMMo as measured by ELISA. **D**, **E**, WT and B27-Tg BMMo were treated with TNF α (30 ng/ml) for varying lengths of time and (**D**) *Il1a* and (**E**) *Ifnb1* mRNA expression was measured. Data are representative of a minimum of three independent experiments with each performed in triplicate. Data points show the mean ±SD. * = *P* < 0.05.

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Figure 3.

HLA-B27 expression, misfolding, and activation of the unfolded protein response in BMMo cultures. WT and B27-expressing BMMo were treated with M-CSF (20 ng/ml), and then without or with RANKL (100 ng/ml) or TNFa (30 ng/ml) for 3 days (A, B, E) or 20 hours (C, D), followed by harvest. A, Cell lysates were subjected to non-reducing SDS-PAGE and HLA-B27 heavy chains visualized by immunoblotting with the antibody HC10. The lowest band represents monomers, while the higher molecular weight material constitutes dimers and multimers. B, HLA-B27 heavy chains were immunoprecipitated using HC10 and subjected to SDS-PAGE under reducing conditions, then visualized by immunoblotting with 3B10.7 (a-HLA). Co-preciptating BiP was visualized by immunoblotting (a-BiP). C, BiP mRNA expression is shown relative to untreated cells. Bars show the mean ±SD of three experiments. **D**, *Xbp1* mRNA splicing in untreated and RANKL or TNFa treated cells. Tunicamycin (TM) treated cells serve as a positive control. The spliced form of Xbp1 mRNA (XBP1s) runs slightly faster than the unspliced form (XBP1u) due to the removal of 26 nucleotides by activated inositol-requiring 1 alpha (IRE1a). E, Immunoblots of cell lysates subjected to SDS-PAGE under reducing conditions, visualizing BiP (a-BiP), HLA-B27 (α-HLA; HC10), and GAPDH (α-GAPDH) as a loading control.

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Figure 4.

ER stress induces *ll1a* and *lfnb1* mRNA in BMMo. BMMo derived from WT rats were treated with thapsigargin (TG) (300 ng/ml) for 3 and 6 hours. **A**, Expression of BiP mRNA relative to untreated cells. **B**, *Xbp1* mRNA splicing. **C**, **D**, Expression of *ll1a* (IL-1 α) (**C**) and *lfnb1* (IFN β) (**D**) mRNA relative to untreated cells.

Figure 5.

Immunofluorescence visualization of HLA-B27 and BiP-expressing cells. BMMo were treated with M-CSF (20 ng/ml) alone (NT), or M-CSF plus RANKL (100 ng/ml) or TNF α (30 ng/ml) for 3 days. Cells were fixed with formaldehyde and permeabilized, then stained with antibodies against BiP (α -BiP) or HC10 (α -HLA). Texas Red or FITC-labeled secondary antibodies were used to detect α -BiP and HC10, respectively. White arrows identify monocytes showing increased HC10 and BiP staining with TNF α treatment, and blue arrows identify an osteoclast. Original magnification 20×.