

Neutrophil extracellular traps (NETs) are part of the innate immune system and are pathogenic in SLE. We therefore investigated 56 lupus patients who met at least 4 SLICC criteria. Results were compared to 20 age, sex, and race matched healthy controls. We found that curli/eDNA induced more NETs in SLE PMNs compared to healthy controls. In SLE, patients who were high inducers of NETs triggered by curli/eDNA complexes were also a high inducer of NETs triggered by LPS and PMA. Interestingly, patients who were anti-dsDNA positive made more NETs in response to curli/eDNA complexes. Moreover, we found patients who are anti-dsDNA positive responded highly to curli/eDNA complexes and LPS. We did not observe this in patients who were anti-dsDNA negative. Mechanistically, we found that curli/eDNA induce NETs via NADPH oxidase. Finally, we found patients who had bacteriuria had a higher amount of NET production in response to curli/eDNA complexes and PMA compared to patients with no bacteriuria. We conclude

1) that lupus PMNs are in a chronic inflammatory state. And 2) that curli/eDNA complexes can activate neutrophils and exposure to UPECs could be a mechanism to sustain autoantigens in the form of neutrophil extracellular traps.

1003

IMPAIRED INTRACELLULAR PROTEIN TRANSPORT AND AN ENDOTHELIAL STRESS RESPONSE REMINISCENT OF COPA SYNDROME IN SLE-ASSOCIATED DIFFUSE ALVEOLAR HEMORRHAGE

Haoyang Zhuang, Erin Hudson, Shuhong Han, Rawad Daniel Arja, Li Lu, Westley Reeves*. *Division of Rheumatology, Allergy, and Clinical Immunology, University of Florida, Gainesville, Florida, USA*

10.1136/lupus-2022-lupus21century.63

Background Human mutations of the coatmer coat protein alpha subunit (*COPA*) affect retrograde Golgi-to-endoplasmic reticulum (ER) protein transport, resulting in endoplasmic reticulum (ER) stress and a clinical syndrome consisting of polyarthritis and diffuse alveolar hemorrhage (DAH) with autoimmune features. Some SLE patients also develop DAH and C57BL/6 (B6) mice with pristane-induced lupus develop monocyte-dependent DAH closely resembling the human disorder. In contrast, BALB/c mice are resistant to DAH. We examined the role of *COPA* and ER stress in lupus mice with DAH.

Methods B6 and BALB/c mice were treated with pristane. Expression of *Copa* and markers of ER stress and vascular injury were assessed by quantitative PCR and immunohistochemistry.

Lung tissue was disrupted by Gentle MACS and ER stress was assessed in CD45+CD146- bone marrow-derived cells and CD45-CD146+ lung microvascular endothelial cells by flow cytometry. *COPA* transcripts were quantified in peripheral blood from 54 SLE patients and 22 controls.

Results DAH in B6 mice was associated with impaired *Copa* mRNA and protein expression and evidence of ER stress (increased *Ddit3* and CHOP protein, *Hspa5* and BiP protein, and other markers). Although DAH did not develop in BALB/c mice treated with either pristane or the ER stress inducer thapsigargin, DAH with impaired *Copa* expression and evidence of ER stress was induced when BALB/c mice were treated with pristane plus low dose thapsigargin. (BALB/c X B6)F1 mice did not develop DAH or ER stress, suggesting that susceptibility was recessively inherited. Flow cytometry of

single cell suspensions of lung tissue revealed increased expression of the ER stress protein BiP in CD45-CD146+ pulmonary endothelial cells and CD45+CD146- myeloid cells from pristane-treated B6 mice and also in CD45- cells from BALB/c mice treated with pristane + thapsigargin. von Willebrand factor (*Vwf*), a marker of endothelial injury, and the monocyte-attractive chemokine *Ccl2* increased in lung from B6 mice with DAH, but not in lung from BALB/c mice treated with pristane (without thapsigargin). These data suggest that pristane-induced ER stress and impaired *Copa* expression promote lung microvascular injury (indicated by increased *Vwf* expression) and the production of monocyte-attractive chemokines, such as *Ccl2*. *COPA* expression also was low in SLE patients with interstitial lung disease or nephritis, suggesting that increased susceptibility to ER stress and impaired retrograde Golgi-to-ER transport also may be associated with a subset of human lupus.

Conclusion DAH in a mouse lupus model appears to be initiated by genetically-regulated susceptibility of the lung microvasculature endothelium to pristane-induced injury, resulting in an ER stress response and culminating in a monocyte-dependent inflammatory response. Lupus-associated DAH in mice and possibly humans may represent an acquired form of COPA syndrome.

Acknowledgments This work was supported by the National Institutes of Health (NIAMS) grant number R01-AR44731.

Lay summary Lung hemorrhage is a severe complication of SLE. Patients with mutations affecting intracellular protein transport ("COPA syndrome") also develop lung hemorrhage along with other clinical features of reminiscent of lupus (arthritis, autoantibodies, and abnormal regulation of interferon production). Using a mouse model, we found that lung hemorrhage in lupus replicates some of the clinical features of COPA syndrome, including low expression levels of *COPA* mRNA and protein and evidence of a stress response in lung microvascular endothelial cells. Additionally, we found that *COPA* expression was low in a subset of SLE patients, raising the possibility that these individuals might be at increased risk to develop lung hemorrhage.

Innate Immunity

1004

ALTERED ER α LOCALIZATION DIFFERENTIALLY MODULATES IMMUNE CELL SUBSETS

¹Cameron Leyers, ²Jena Wirth, ²Melissa Cunningham. ¹*Department of Microbiology and Immunology, Graduate Studies, Medical University of South Carolina, Charleston SC 29407*; ²*Department of Medicine, Division of Rheumatology and Immunology, Medical University of South Carolina, Charleston SC 29407*

10.1136/lupus-2022-lupus21century.64

Background Estrogen is anti- or pro-inflammatory depending on milieu and plays a role in the increased incidence of lupus in reproductive age women. Estrogen's pleiotropic effects are in part due to estrogen receptors (ER) and their variants that localize to different regions in the cell. To understand the role of ER α localization in immune responses, we investigated the effects of altered ER α localization on Toll Like Receptor (TLR7)-stimulated endpoints, often dysregulated in lupus.

Methods Membrane-only ER α (MOER) or Nuclear-only ER α (NOER) mice were used to isolate and culture spleen cells, *ex vivo* bone marrow (BM), and BM-DCs. Cells were phenotyped