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Cell death mechanism in an isolated wood smoke inhalation induced-ARDS large animal model

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Abstract

Acute respiratory distress syndrome (ARDS) is a lethal disease condition in critically ill patients with a reported mortality rate reaching 45%. The current treatment modalities available for severe ARDS are invasive and carry significant risk for patients. Most published studies involving smoke inhalation utilize another simultaneous injury (such as cutaneous burn) to increase pathology burden of their animal models. This introduces confounding variables to investigations which aim to concentrate on inhalation injury. In this study, we evaluated the potential molecular targets associated with isolated smoke inhalation-induced ARDS. We observed an increase in lung injury score and wet/dry ratio 48h post smoke inhalation together with upregulation of inflammatory markers, ILand IL-6 levels. Furthermore, there was a decrease in phosphorylation of cell survival marker Akt and an increase in proapoptotic protein BAX at 48h post smoke inhalation. These results indicate that smoke inhalation induced inflammatory processes resulting in increased apoptosis and decreased cell survival in lung parenchymal cells. Use of this unique model may be of benefit in studying the pathophysiology of inhalation injury and for the development of novel therapeutic strategies.

Materials and Methods

<u>Smoke inhalation</u>: Duroc pigs $(50\pm 5 \text{ kg})$ were exposed to oak wood smoke for 2 hours while intubated. Continuous monitoring and serial blood collection was performed via catheters placed in the carotid artery, internal jugular vein, pulmonary artery, and femoral artery. Pigs were euthanized at 48h post smoke inhalation after final blood sampling, and lung tissue was collected for analysis.

H&E: Analysis was performed in Tissue Science Facility, UNMC to determine lung injury.

<u>Wet/Dry ratio:</u> Lung tissues were dried in an incubator at 60°C for 5 days and weighed again (dry weight).

Cytokine analysis: ELISA was performed using Quantikine® ELISA kit from R&D systems.

Western Blots: Analysis was performed using actin as loading control.

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