



# Nucleotide-Binding Domain and Leucine-Rich Repeat Protein Inflammasome may Modulate Sleep Induced by Both Increased Wakefulness and Bacterial Components in the Brain

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## INTRODUCTION

Molecules involved in innate immunity influence sleep and circadian oscillators and vice versa. Sleep-inducing inflammatory molecules are activated by wakefulness and increased pathogens. Pathologies that alter inflammatory molecules. Traumatic brain injury, cancer, cardiovascular disease, and stroke are often associated with sleep disturbances and the EEG power spectrum. Furthermore, sleep disorders such as insomnia and sleep disturbances are associated with increased dysregulation of inflammatory processes. Inflammatory molecules both in the central nervous system and in the periphery can alter sleep. Inflammation may also modulate cerebrovascular hemodynamics associated with changes in the EEG power spectrum. However, further studies are needed to determine the interplay between sleep-regulating inflammatory molecules and the circadian clock.

## DESCRIPTION

Both sleep deprivation and pathogens can increase brain inflammation, sleep and sleep intensity, as shown by EEG delta power. The proinflammatory cytokine interleukin-one beta, which is increased in the cortex after sleep deprivation and in response to lipopolysaccharide, a cell wall component of Gram-negative bacteria, does not suppress these effects. The nucleotide-binding domain and leucine rich repeat protein three inflammasome protein complexes is formed in response to changes in the local environment and activates caspase one to convert interleukin-one beta to its active form. Sleep deprivation enhances the cortical expression of the somnogenic cytokine interleukin-one beta, although the underlying mechanism is, as yet, unidentified. Using nucleotide-binding domain and leucine rich repeat protein gene knockout mice, we provide evidence that inflammasome activation is a crucial mechanism for the downstream pathway leading to increased inter-

leukin enhanced sleep. Nucleotide binding domain and leucine rich repeat protein knockout mice exhibited reduced non-rapid eye movement sleep during the light period. We also found that sleep amount and intensity were drastically attenuated in knockout mice following sleep deprivation (homeostatic sleep response), as well as after administration, although they were enhanced by central administration of interleukin. Nucleotide binding domain and leucine rich repeat protein, and caspase one activity were greater in the somatosensory cortex at the end of the wake-active period when sleep propensity was high and after sleep deprivation in wild type but not nucleotide binding domain and leucine rich repeat protein mice. Our new convergent results therefore suggest that activation of the nucleotide-binding domain and the leucine rich repeat protein inflammasome may modulate sleep induced by both increased wakefulness and bacterial components in the brain.

## CONCLUSION

Minocycline has beneficial effects in early brain injury after subarachnoid hemorrhage. This study was conducted to determine the effects of minocycline on inflammation and neuronal apoptosis and possible mechanisms for these effects in early brain injury after subarachnoid hemorrhage. Subarachnoid hemorrhage was induced by his filament perforation model of subarachnoid hemorrhage in male sprague dawley rats. One hour after subarachnoid hemorrhage induction, minocycline or vehicle was administered by intraperitoneal injection. Minocycline treatment significantly reduced cerebral edema secondary to blood-brain barrier dysfunction by inhibiting activation of the inflammasome, which regulates the maturation and release of inflammatory cytokines, particularly interleukin. Proteins containing nucleotide-binding oligomerization domains are known to play important roles in innate immunity, but their relationship to the inflammatory response in endometriosis remains unclear.

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