

Activation of the NLRP3 inflammasome by SARS-CoV-2 via mitochondrial dysfunction.

The three SARS-CoV proteins (E, ORF3a and ORF8b) induce the activation of inflammasome.

E protein (**orange**) induces Ca^{2+} efflux from Golgi apparatus, where it stores, to the cytosol. This induces its entry into the mitochondria via VDAC (**2**) to generate mtROS (**3**). ORF3a (**blue**) induces K^+ efflux to the extracellular space and promotes inflammasome assembly through TRAF3-mediated ubiquitination of ASC. On the other hand, TRAF3-ORF3a interaction is required for NF- κ B activation, resulting in transcription of the pro-IL-1 β /IL-18 and NLRP3 genes. ORF8b (**violet**) is able to interact directly with NLRP3 stimulating its activation. Consequently to inflammasome activation, the formation of GSDMD pores on the plasma and mitochondrial (**A**) membranes occurs causing IL-1 β /IL-18 secretion, the cellular swelling associated to pyroptosis and the induction of mitochondrial apoptotic pathway via Bax-dependent release of cytochrome c in the cytosol (**B**). In addition, activation of BAX can trigger NLRP3 activation via apoptotic caspases (**dotted line**) in a K^+ efflux-dependent manner (**C**).

Thus, SARS-CoV-2 triggers NLRP3 inflammasome assembly and activation by damaging the mitochondria and inducing the production of mtROS (**3**) and the loss of mitochondrial membrane potential ($\Delta\Psi_m$) (**4**) to release damaged mitochondrial DNA (mtDNA) in the cytosol through the mitochondrial pore transition (mPT) (**5**). Therefore, mitophagy (**1**) plays an important role as a regulator of NLRP3.

Cardiolipin, MAVS and MFNs can tether NLRP3 to the mitochondria for its activation in a mtROS-dependent manner (**black box**).