SARS-CoV-2 Spike 1 protein controls Natural killer cells activation via HLA-E/NKG2A pathway.

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Backgroud

Natural killer cells are important in the control of viral infections. NK cells showed Interestingly a functional exhaustion with an increased NKG2A expression during SARS-CoV-2 infection [1]. The CD94/NK group 2 member A (NKG2A) is a heterodimeric inhibitory receptor expressed by NK cells [2]. It binds to the nonclassical HLA class I molecule (HLA-E), which presents peptides derived from leader peptide sequences of other HLA class I molecules, including HLA-G [3]. The ligation of the peptide-loaded HLA-E with NKG2A transduces inhibitory signaling through 2 inhibitory immune-receptor tyrosine-based inhibition motifs, that suppress NK cytokine cytotoxicity and secretion [3]. By now, no data are available on how SARS-CoV-2 might control NK cells. We evaluated the possible role of SARS-CoV-2 spike proteins in modifying NK cell functions [4].



Aim

Peripheral blood NK cells from SARS-CoV and SARS-CoV-2 naïve subjects were evaluated for their activation, degranulation, interferon-gamma expression in the presence of SARS-CoV and SARS-CoV-2 spike proteins.

Results



SP1 induces NK anergy via HLA-E/NKG2A pathway



Effect of SP1 and SP2 on NKs



SP1 and SP2 induced NKs migration and IFN-gamma secretion

SP1 modifies NKs cytotoxicity



SP1 induces NKs anergy increasing HLA-E and NKG2A expression via GATA3

Conclusions

We show for the first time that NK cells are affected by SP1 expression in lung epithelial cells via HLA-E/NKG2A interaction. The resulting NK cells exhaustion might contribute to immunopathogenesis in Sars-Cov-2 infection.

References

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