

immune response against SARS-CoV-2. These results can contribute to the development of therapeutics, preventive medication against hyperinflammatory syndromes as well as alternative vaccine constructs and adjuvants.

Theobald SJ, Simonis A, [...], Rybniker J. (2021) **Long-lived macrophage reprogramming drives spike protein-mediated inflammasome activation in COVID-19.** *EMBO Molecular Medicine* 13(8): e14150

P033

### **MicroRNA-223 dampens neutrophil-mediated lung inflammation during pneumococcal pneumonia**

C. Gökerj<sup>1</sup>, P. Pennitz<sup>1, 2</sup>, W. Groenewald<sup>1</sup>, U. Behrendt<sup>1</sup>, H. Kirsten<sup>3</sup>, C. Zobel<sup>4</sup>, S. Berger<sup>1, 2</sup>, G. Heinz<sup>5</sup>, M.-F. Mashreghi<sup>5, 6</sup>, S.-M. Wienhold<sup>1</sup>, K. Dieter<sup>7, 8</sup>, A. Dorhoi<sup>9, 10</sup>, A. Gruber<sup>7</sup>, M. Scholz<sup>11</sup>, G. Rohde<sup>12, 13, 14</sup>, N. Suttorp<sup>2, 13, 15</sup>, CAPNETZ Study Group, M. Witzernath<sup>1, 2, 13, 15</sup>, G. Nouailles<sup>1, 2</sup>

<sup>1</sup>Charité-Universitätsmedizin Berlin, Division of Pulmonary Inflammation, Berlin, Germany,

<sup>2</sup>Charité-Universitätsmedizin Berlin, Department of Infectious Diseases and Respiratory Medicine, Berlin, Germany, <sup>3</sup>Universität Leipzig, Institute for Medical Informatics, Statistics and Epidemiology, Leipzig, Germany, <sup>4</sup>Bundeswehrkrankenhaus Berlin, Department of Internal Medicine, Berlin, Germany, <sup>5</sup>Deutsches Rheuma-Forschungszentrum Berlin, Therapeutic Gene Regulation, Berlin, Germany, <sup>6</sup>Berlin Institute of Health at Charité - Universitätsmedizin Berlin, BIH Center for Regenerative Therapies, Berlin, Germany, <sup>7</sup>Freie Universität Berlin, Institute of Veterinary Pathology, Berlin, Germany, <sup>8</sup>Freie Universität Berlin, Veterinary Centre for Resistance Research, Berlin, Germany, <sup>9</sup>Friedrich-Loeffler-Institut, Institute of Immunology, Greifswald, Germany, <sup>10</sup>University of Greifswald, Faculty of Mathematics and Natural Sciences, Greifswald, Germany, <sup>11</sup>Universität Leipzig, Institute for Medical Informatics, Statistics and Epidemiology, Berlin, Germany, <sup>12</sup>Goethe-Universität Frankfurt am Main, Department of Respiratory Medicine, Medical Clinic i, Frankfurt/Main, Germany, <sup>13</sup>CAPNETZ STIFTUNG, Hannover, Germany, <sup>14</sup>Biomedical Research in Endstage and Obstructive Lung Disease Hannover, Hannover, Germany, <sup>15</sup>German Center for Lung Research, Berlin, Germany

Community acquired pneumonia remains a leading cause of communicable disease-related mortality globally, despite the widespread use of protective vaccines and effective antibiotics. Neutrophils play a major role in containing bacterial growth, while also orchestrating detrimental pulmonary inflammation when dysregulated. Here we aimed to elucidate the role of myeloid cell-derived microRNA-223 in regulating pulmonary inflammation during pneumococcal pneumonia.

Serum microRNA-223 was quantified in pneumococcal pneumonia patients and healthy subjects. Wild-type and microRNA-223 knockout mice were intranasally infected with *Streptococcus pneumoniae*, followed by evaluation of pulmonary inflammation in terms of clinical disease course, histopathology, immune cell chemotaxis and inflammatory protein and gene signatures. Single-cell RNA sequencing was utilized to evaluate the transcriptomic repercussions of the absence of microRNA-223 during murine pneumococcal pneumonia.

Cell-free serum microRNA-223 was reduced in pneumococcal pneumonia patients relative to healthy subjects, which was correlated with increased disease severity. MicroRNA-223 knockout mice exhibited enhanced neutrophilic influx into the lungs and bronchoalveolar lavage, leading to histopathological aggravation and increased production of pro-inflammatory mediators following *Streptococcus pneumoniae* infection. MicroRNA-223 was induced in the lungs and sorted lung neutrophils of wild-type mice by *Streptococcus pneumoniae* in a time-sensitive manner, while its absence resulted in a dysregulated transcriptome of pulmonary neutrophils involving anti-microbial and cellular maturation genes.

Altogether, our findings indicate that in the absence of microRNA-223, alterations in the neutrophil transcriptome provoked exacerbated acute lung injury in mice, while reduced levels in the serum correlated to increased disease severity in pneumococcal pneumonia patients.