

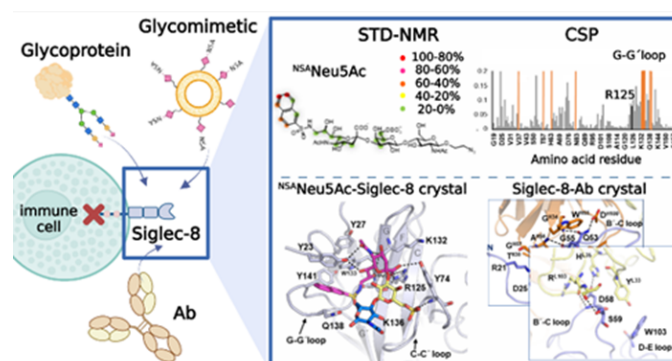
## Siglec-8 complex structures with a therapeutic antibody and a high-affinity sialoside analog

Maria PIA LENZA [1], Unai ATXABAL [2], Corwin NYCHOLAT [3], Iker OYENARTE [2], Antonio FRANCONETTI [2], Jon Imanol QUINTANA [2], Sandra DELGADO [2], Reyes NÚÑEZ-FRANCO [2], Carmen Teresa GARNICA MARROQUÍN [4], Helena COELHO [5], Luca UNIONE [2], Gonzalo JIMÉNEZ-OSÉS [2] [6], Filipa MARCELO [5], Mario SCHUBERT [7], James C. PAULSON [3], Jesús JIMÉNEZ-BARBERO [2,4,6,8], June EREÑO-ORBEA [2] [6]

[1] Department of Chemical Sciences, University of Naples "Federico II", Naples, Italy, [2] CIC bioGUNE, Spain [3] The Scripps Research Institute, La Jolla, United States [4] Department of Inorganic & Organic Chemistry University of the Basque Country, Spain [5] UCIBIO, REQUIMTE, Universidade de Nova de Lisboa, Portugal [6] IKERBASQUE, Basque Foundation for Science and Technology, Bilbao, Spain [7] Department of Biosciences, University of Salzburg, Hellbrunn, Austria [8] Centro de Investigación Biomedica En Red de Enfermedades Respiratorias, Madrid, Spain

mariapia.lenza@unina.it

Siglec-8 is an inhibitory receptor that induces eosinophil apoptosis and inhibits mast cell degranulation when bound by monoclonal antibodies (mAbs) or sialylated ligands. Consequently, Siglec-8 has emerged as a crucial negative regulator of inflammatory responses in various diseases, including allergic airway inflammation [1]. Herein, the molecular recognition features of the interaction of Siglec-8 with the monoclonal antibody (mAb) lirentelimab (2C4) and a sialoside mimic with the potential to reduce mast cell degranulation have been deciphered [2]. The X-ray crystallographic solution of the structure of Siglec-8 and the fragment antigen-binding (Fab) component of 2C4 shows that the mAb binds close to the carbohydrate recognition domain on Siglec-8. Additionally, the STD-NMR experiment demonstrates the inhibition of the binding between Siglec-8 and natural ligand in the presence of 2C4. Moreover, using a combination of NMR spectroscopy and X-ray crystallography, we have also deduced the binding mechanism of a high-affinity analog of its sialic acid ligand (9-N-naphthylsulfonimide-Neu5Ac, <sup>NSA</sup>NeuAc). Our data demonstrate that <sup>NSA</sup>NeuAc sialoside ring binds to the classic sialyl binding pocket of the Siglec receptor family, with the high affinity resulting from the accommodation of the NSA aromatic group in a contiguous hydrophobic patch provided by the N-terminal tail and the unique G-G' loop (Figure 1). These results provide pointers for the rational design of the next generation of Siglec-8 inhibitors and explain the foundation for this ligand's observed high affinity [2].



The binding of a sialic acid mimetic with a 9-N aromatic substituent, and antibody with therapeutic potential to Siglec-8 are revealed by a synergic c

Bibliographic references:

[1] S. Duan, B.M. Arlian, C.M. Nycholat, Y. Wei, . Tateno, S.A. Smith, M.S. Macauley, Z. Zhu, B.S. Bochner, J.C. Paulson (2021), *J Immunol* (206), 2290-2300.

M.P. Lenza, U. Atxabal, C.M. Nycholat, I. Oyenarte, A. Franconetti, J. I. Quintana, S. Delgado, R. Núñez-Franco, C. T. Garnica Marroquín, H. Coelo, L. Unione, G. Gimenez-Oses, F. Marcelo, M. Schubert, J.C. Paulson, J. Jimenez-Barbero, J. Ereno-Orbea (2023), *JACS Au* (3), 204-215.