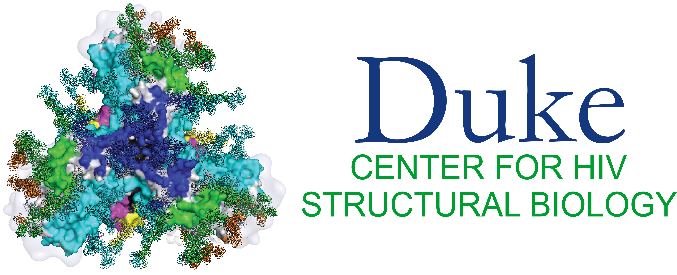
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4. Liu HF, Zhou Y, Huang Q, et al. nextPYP: a comprehensive and scalable platform for characterizing protein variability in situ using single-particle cryo-electron tomography. Nat Methods 2023. DOI: 10.1038/s41592-023-02045-0.

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11. Do HN, Zhao M, Alam SM, Gnanakaran S. Dynamics and Activation of Membrane-Bound B Cell Receptor Assembly. bioRxiv 2024:2024.07.10.602784. DOI: 10.1101/2024.07.10.602784.

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14. Watson AJI, Bartesaghi A. Advances in cryo-ET data processing: meeting the demands of visual proteomics. Curr Opin Struct Biol 2024;87:102861. DOI: 10.1016/j.sbi.2024.102861.