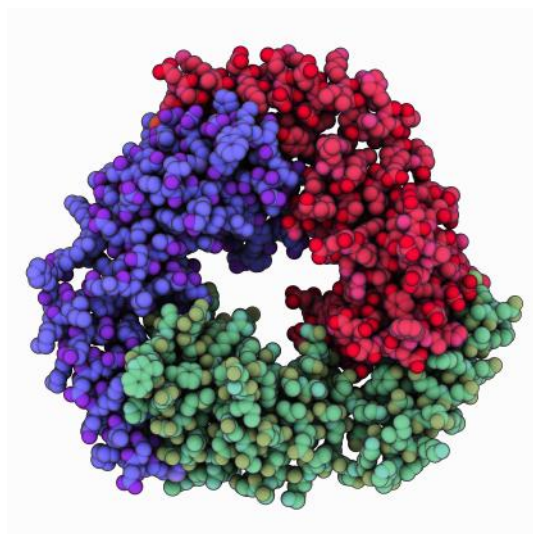


Improved cross-linking models between amino acids with DynamXL software

DynamXL software models cross-link between amino acids by accounting for linker and protein dynamics for improved resolution of protein structure.



To find out more, contact enquiries@innovation.ox.ac.uk and quote the project number.

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