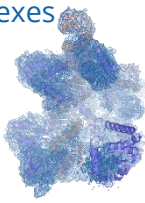
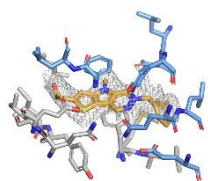
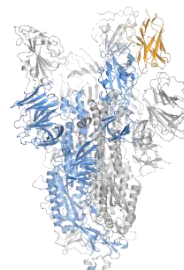
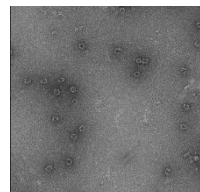
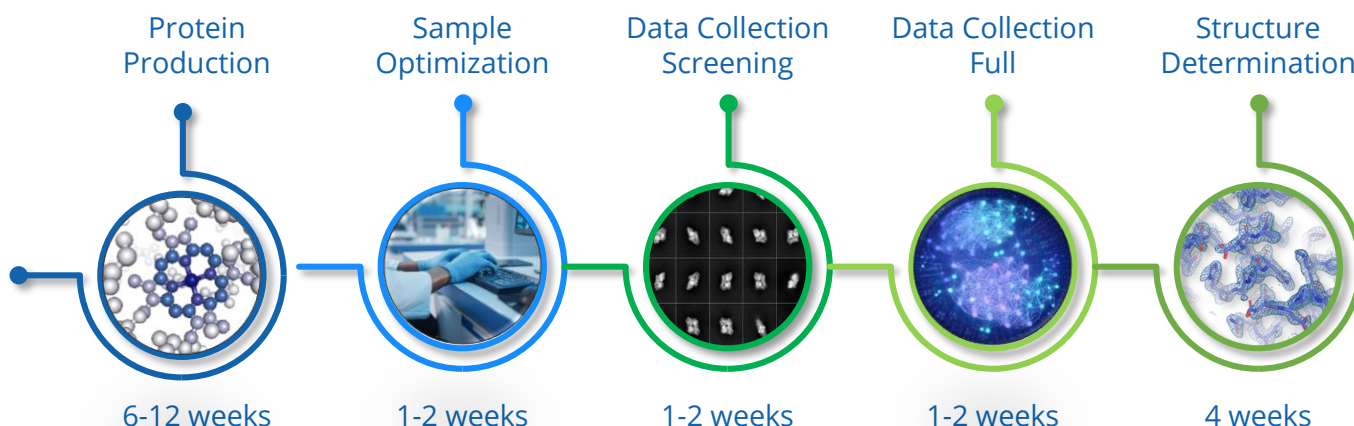


Cryo-EM

<p>De novo Structure Determination</p> <ul style="list-style-type: none"> Membrane proteins Multi-domain proteins Multi-subunit complexes 	<p>Structure-Based Drug Discovery</p> <ul style="list-style-type: none"> Determination of small molecule binding mode Support of hit-to-lead optimization projects 	<p>Epitope Mapping</p> <ul style="list-style-type: none"> Characterization of epitope-paratope interactions 	<p>Formulation Characterization</p> <ul style="list-style-type: none"> LLP and VLP Characterization Formulation screening and optimization 
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Gene-to-Structure Workflow



Experience and Cutting-Edge Facilities

- >20 cryo-EM structures determined in 2023
- 15 years' experience structural biology sample preparation
- 7 PhD-level scientists with proven EM track record
- In-house sample preparation laboratory (Vitrobot IV)
- Rapid access to screening (Glacios) and data collection (Krios) microscopes
- Close integration with protein production, biochemistry, biophysics and screening groups

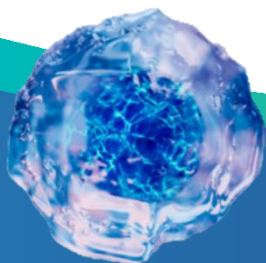
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