




## CCP-SAS: New developments in computational modelling of X-ray and neutron scattering curves

### Virtual Satellite Meeting at the 60<sup>th</sup> British Biophysical Society Meeting, UK

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A Joint EPSRC-NSF Software Project



Usual Zoom rules apply:


Please keep mikes off.  
Please post questions in the chat throughout the talks.  
The chair will read them to the speaker at the end.  
Auto-captions are available to those wishing to have them.

## (1) What is CCP-SAS?

## (2) Applications?

## (3) Achievements?

## (4) Developments/Future

 **Scattering problem**

Scattering is directly related to the organization of atoms within a material and thus a powerful tool for the elucidation of the detailed microstructure of materials.


BUT


Maximizing the information extracted from the analysis requires:

- 1) Using all a priori information available to constrain the solution.
- 2) Careful posing of the question and design of experiment
- 3) Can become computationally challenging

Is used by a wide variety of scientists from geology to biology, from physics to chemistry etc. – Not all experimentalists in these fields have the skills to extract the most information from the data

**MEANING: The modelling dilemma – to model - or not to model.**




 **CCP-SAS in a nutshell**


Create new and enable existing computational tools to model scattering data in real space using modern MD and MC methods as well as applying constraints from other experimental techniques

AND

dramatically improve their accessibility by non-experts.

**MEANING: Interpret SAS curves at a molecular or atomistic level, not as solid objects or envelopes**



 **The CCP-SAS Solution**

UI – GUI – Web GUI

GenApp

Other?




QuaFit


WillItFit


SASSIE

Build
Simulate
Calculate
Analyze
SCT

HPC – Gateway



 **Good vs. successful software**


The most brilliant software in the world .....

If nobody ever uses it

Or it fails to be maintained long term (after the honeymoon is over)

IS A COMPLETE AND UTTER FAILURE

**MEANING: The need to build a community around CCP-SAS and promote this.**

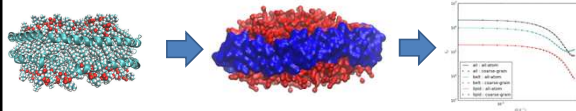


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Session 1 this afternoon will look at applications

## Applications are broad ranging

SOFT MATTER – eg: Polymer + Nanodisc



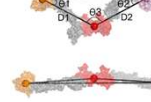
BIOLOGY AND BIOTECHNOLOGY

eg: Antibody + Glycans



G K Hui et al., Biochem. J. (2015), DOI: 10.1042/BJ201506

eg: Complement



R. Nan et al., Structure. (2017), DOI: 10.1016/j.str.2016.12.014

eg: Hfq + RNA



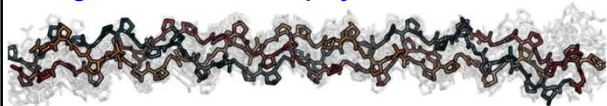
Y Peng et al., PNAS (2014), DOI: 10.1073/pnas.1410114111

Outcomes give molecular structures that clarify biological function

IgG1 antibodies – widespread utilities



Collagen – the classic biopolymer



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## Usage of SASSIE-web on the HPC at Tennessee, USA

May 2016 – 273 registered users  
Sep 2020 – 952 registered users

Period	Hours CPU per year	Active users	Power users >100 jobs
Beta period Jun15-May16	8500		
Jul17-Aug18	24900	134	18
Jul18-Aug19	35000	158	24
Jul19-Aug20	44900	104	12

## CCP-SAS publications in good journals

Year	All	Biology	Techniques	Soft Matter	Biopolymer
2020	4				
2019	12				
2018	5				
2017	8				
2016	14				
2015	9				
2014	10				
2013	7				
Totals	69	39	5	18	7

Biology – includes PNAS, Structure, JBC – a good spread of the stronger journals



**UCL**

- (1) What is CCP-SAS?
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Session 2 this afternoon will look at the future – where are we on the computing side?

**Conclusion – CCP-SAS is on the way to becoming established for the atomistic modelling of macromolecules by scattering**