ESRF	Experiment title: Bacterial and viral neuraminidases/sialidases	Experiment number: LS-2087						
Beamline:	Date of experiment:	Date of report:						
ID14-2	from: 03/02/02 to: 04/02/02							
Shifts:	Local contact(s):	Received at ESRF:						
3	Antoine Royant							
Names and affiliations of applicants (* indicates experimentalists):								
Professor Garry Taylor								
*Dr Viateslav Zaitsev								
*Mr Ibrahim Moustafa								
*Mr Simon Newstead								
Centre for Biomolecular Sciences								
University of St Andrews								
St Andrews. Fife KY16 9ST. Scotland								

Report:

1. Hemagglutinin neuraminidase

Several datasets of substrate analogues. HN_FLUORO is a sialic acid with substitution of F at the 3-position protons. HN_SULFUR is a non-hydrolyesable disaccharide with a sulfur in place of the glycosidic oxygen.

Ident	Nframes	osc	reso	exp	dist (mm)	conditions	compl	Rm
$\overline{1^{st} xtal, or}$	thorhomb	ic 72.8	89 x 77.7	2 x 200.1	27 Å			
hn_fluoro	200	0.5	2.2 A	10sec	205	100 mM, pH 6.57, 3.5h	90.3%	6.03%
2^{nd} xtal, or	thorhom	bic 73.	15 x 72.9	97 x 199.4	42 Å			
hn_fluoro2	145	0.5	2.2 A	10 sec	205	100 mM, 5.5hours		
hn_fluoro3	100	0.5	2.2 A	10 sec	205	100 mM, 5.5hours		
hn_sulfur *	·) 1-72	0.7	2.2 A	5sec	205	25 mM, pH 6.57, 6.5ho	urs 94.89	% 5.5%
hn_sulfur	86-120	0.7	2.2 A	5sec	205			
*) orthor	hombic	73.35	x 78.67 x	200.38 A	Å			
**) gap of	14 frames	s - due	to refil at	: 6-30 am	l			

Following the success of the last visit in November, co-crystals were prepared with sialyl-lactose:

Ident	Nframes	osc	reso	exp	dist (mm)	conditions	compl	Rm	
vc_sialyl	120	1.0	1.8	15sec	162	co-crystals with sialyl-lact	tose	94%	10%

3. Neuraminidase from *Clostridium perfringens*

Recently obtained crystals of a 55kDa fragment of the 77kDa sialidase from C. perfringens were tested for their high resolution limit. Data in-house extended to 1.6Å, and on ID14-2 data were observed to beyond 1.0Å! Unfortunately on the low resolution pass, the crystals were found to be split – but this trial gives us hope to return to obtain very high resolution data from this large protein.

Ident	Nframes	osc	reso	exp	dist (mm)	conditions	compl	Rm
nanH_G12	2 120	1.0	2.0	10 sec	184	native		
nanH_18	130	1.0	2.2	10sec	205	native		