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www.chem.uzh.ch/en/research/services/xray.html

Department of Chemistry X-ray Crystallography Facility

Fill in ALL sections. Label your sample clearly with your name and sample code.

Name:	Date:
Group:	Laboratory:
E-mail: a UZH e-mail address is preferred	
Sample code or name:	
Molecular formula:	Melting point:°C
Solvent(s):	Colour:
Safety precautions & hazards:	
Are you confident of the structure of your molecule?	
	_ 1 ,
Chirality: none racemic If chiral: all sites known all sites unknown Clearly indicate the chirality of known sites on you	single enantiomer own some sites unknown ur diagram opposite.
Crystal stability:	
stable moisture sen keep cold hygroscopic loses solvent thermally un Speed of decomposition: instant	_ °
Special requirements:	
temperature of study: K (160 K) investigation of classic hydrogen bonding interactorsion angles involving H atoms determine absolute configuration (possible only if a other:	an element heavier than N is present)

Diagrams: (these will be e-mailed	I to you)		
You will normally receive the follow			
One displacement ellipsoid plot o A packing diagram if intermolecu	of the molecule ready for publication. ular hydrogen bonds are present.		
If you are interested in the packing, it is often best to visualise this yourself using Mercury and the provided CIF			
Indicate additional requirements below (e.g. stereoview, view directions, style, etc.):			
Format of report & data files:	(these will be e-mailed to you)		
Report & tables document:			
File of atomic coordinates:	CIF (for structure visualisation with Mercury, Diamond, Suitable for deposition in the CSD or with a publication)		
other file formats, tables, o	calculations, special needs:	,	
Structural formula:			
If the composition is uncertain, give	the reaction starting materials and all likely products.		
Please indicate the desired atom nu Please indicate what you hope to le	umbering if it is important to you. Clearly show chirality, if known earn from this analysis.	•	
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