

How to curate Morphology Data?

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What morphology data to curate?

- Needle
- Acicular
- Aspect Ratio ~ 5
- Needle oriented along x axis
- Needle bounded by {011} planes





Curate morphology data that are linked to a unit cell

- Needle
- Acicular
- Aspect Ratio ~ 5
- Needle oriented along x axis of BPHEN012
- Needle bounded by {011} planes of BPHENO12





Curating morphology data – benzophenone example





Groth (1919)

Black & Seton (2024)

One of >2000 examples



"Groth"

Volume	Ι	II	III	IV	V
Year of	1906	1908	1910	1917	1919
publication					
Contents	Inorganics		Organics		
	Elements	Salts	Aliphatics	one phenyl	> 1phenyl
	and simple	containing		ring	ring,
	Compounds	oxygen			heterocycles
Pages	914	930	804	801	1063
Figures	390	522	648	828	955
Figure	1-389	390-911	912 - 1559	1560 - 2387	2388-3342
numbers					

Table 1: The five volumes of P. Groth's "Chemische Kristallographie"

Morphologies for > 2000 organic compounds



"Groth" - benzophenone

Summarises 5 references

Benzophenon (Diphenylketon) = $(C_6H_5)_2CO$.

Schmelzpunkt 48°,0-48°,5. Rhombisch bisphenoïdisch.

a:b:c = 0.8511:1:0.6644 Wickel²⁵).

Von Linnemann²⁶) aus wässerigem Alkohol erhaltene Krystalle waren nach Handl lange Prismen $m\{110\}$, an den Enden vorherrschend $d\{101\}$ und $q\{011\}$, ferner $r\{021\}$, $c\{001\}$ und $o\{111\}$ in meist unregelmäßiger Entwicklung; außer den Elementen a:b:c=0.8496:1:0.6535 wird nur ein gemessener Winkel

Fig. 2486.



angegeben: $m: m = (110): (1\bar{1}0) = 80^{\circ} 42'$. Arzruni²⁷) untersuchte ein Präparat von Merz²⁸), aus Ligroïn krystallisiert, wasserhelle Prismen $m\{110\}$, deren Endflächen aus der Combination des Bisphenoïds $o\{111\}$ mit $r\{101\}$ bestanden. Wickel beobachtete an Krystallen, welche Jannasch aus Benzol erhalten hatte, die gleichen Formen, wie Handl, meist ohne $o\{111\}$, welches nur untergeordnet und unregelmäßig auftrat (Wickel übersah offenbar den bisphenoïdischen Charakter der Krystalle, welchen er durch Ätzfiguren leicht hätte feststellen können, und teilte die hier wiedergegebene

idealisierte Fig. 2486 mit). Dagegen wurde die bisphenoïdische Symmetrie bestätigt durch Nacken²⁹), welcher aus dem Schmelzflusse die Combination $m\{110\}, d\{101\}, \omega\{1\overline{1}1\}$ erhielt.



"Groth" – benzophenone symmetry

Benzophenon (Diphenylketon) = $(C_6H_5)_2CO$. Schmelzpunkt 48°,0-48°,5. Rhombisch bisphenoïdisch. a:b:c = 0.8511:1:0.6644 Wickel²⁵).

"rhombisch bisphenoïdisch"

= disphenoidal (International Tables Vol. A, Table 3.1.2.4, p. 728)

= Point Group "222"



"Groth" – benzophenone unit cell

Benzophenon (Diphenylketon) = $(C_6H_5)_2CO$. Schmelzpunkt 48°,0-48°,5. Rhombisch bisphenoïdisch. a:b:c = 0.8511:1:0.6644 Wickel²⁵).

One unit cell

b always = 1

ratios of unit cell lengths determined from angles between faces

b > a > c



"Groth" – benzophenone morphology

• Standard orientation just off x -axis

- Faces {forms} identified by letters
- For example, *m* is always {110}





"Groth" – benzophenone morphology

- Notes in the text
- In this case, o = {111} is different from w = {-111}
- "Verifiable from etch pits"





Benzophenone in CSD - 13 options

- BPHENO02, -03, -11 are the other polymorph
- BPHENO01, -10 have *b* > *a* > *c*, no hydrogens
- BPHENO12, 13-20 have *c* > *b* > *a*
- BPHENO12 as 'best representative'

BPHENO01 P212121 BPHENO02 C2/c BPHENO03 C2/c BPHENO10 P212121 BPHENO11 C2/c BPHENO12 P212121 BPHENO13 P212121 BPHENO14 P212121 BPHENO15 P212121 BPHFNO17 P212121 BPHENO18 P212121 BPHENO19 P212121 BPHENO20 P212121



Benzophenone – predicted morphology

- BPHENO12
- "Visual Habit"
- Default view in Mercury





- 1. (011) and (110)
- 2. (011) and (01-1)
- 3. (011) and (0-1-1)
- 4. (111) and (1-1-1)
- 5. (111) and (-1-1-1)







(103)

(002)

(-103)



- 1. (011) and (110)
- 2. (011) and (01-1) {011}
- 3. (011) and (0-1-1) {011}
- 4. (111) and (1-1-1) {111}
- 5. (111) and (-1-1-1)





- 1. (011) and (110)
- 2. (011) and (01-1)
- 3. (011) and (0-1-1)
- 4. (111) and (1-1-1)
- 5. (111) and (-1-1-1)





benzophenone – matching unit cells





benzophenone – matching unit cells





benzophenone – Groth morphology in Mercury

- Approximation to standard orientation
- Forms identified by colours
- Easy to recognise which faces are symmetry related
- {111} different from {-111}





Groth's experimental morphology displayed in Mercury





Groth (1919)

Black & Seton (2024)



Groth's experimental morphology compared with prediction







How to store these morphologies?







	data BPHENO12			
The "morphology cif"	symmetry cell setting	orthorhombic		
	symmetry space group name H-M _ P 21 21 21'			
	symmetry Int Tables number	19		
	space group name Hall	'P 2ac 2ab'		
	loop			
	_symmetry_equiv_pos_site_id			
	_symmetry_equiv_pos_as_xyz			
	1 x,y,z			
CSD Reference	2 1/2-x,-y,1/2+z			
CSD Reference	3 -x,1/2+y,1/2-z			
	4 1/2+x,1/2-y,-z			
	_cell_length_a	7.7378(2)		
 Space Group 	_cell_length_b	10.2421(3)		
	_cell_length_c	12.0395(3)		
	cell_angle_alpha	90		
• Unit Coll	_cell_angle_beta	90		
• Unit Cell	_cell_angle_gamma	90		
	_cell_volume	954.146		
	loop_			
 Centreface distances 	_exptl_crystal_face_index_h			
	_exptl_crystal_face_index_k			
	_exptl_crystal_face_index_1			
	_exptl_crystal_face_perp_dist			
Durham	1 0 2 10.5			
University				

<b

The "morphology .cif" - what's not there

Point group deduce from Space Group

Multiplicity of each form Deduce from point group

Display Changes Orientation Changes to face labels Changes to face colours

data_BPHEN012	
_symmetry_cell_setting	orthorhombic
_symmetry_space_group_name_H-M	'P 21 21 21'
_symmetry_Int_Tables_number	19
_space_group_name_Hall	'P 2ac 2ab'
loop	
symmetry equiv pos site id	
symmetry equiv pos as xyz	
1 x,y,z	
2 1/2-x,-y,1/2+z	
3 -x,1/2+y,1/2-z	
4 1/2+x,1/2-y,-z	
_cell_length_a	7.7378(2)
_cell_length_b	10.2421(3)
_cell_length_c	12.0395(3)
_cell_angle_alpha	90
_cell_angle_beta	90
_cell_angle_gamma	90
_cell_volume	954.146
loop_	
_exptl_crystal_face_index_h	
_exptl_crystal_face_index_k	
_exptl_crystal_face_index_l	
_exptl_crystal_face_perp_dist	
0 1 1 10	
1 1 1 15	
1 1 0 15	
1 1 -1 15	
2 0 0 16	
1 0 1 16	
1 0 2 16.5	



The "morphology .cif"

A standardised way of recording morphology data

<u>The</u> standardised way of recording morphology data?

data BPHENO12 symmetry cell setting orthorhombic symmetry space group name H-M 'P 21 21 21' symmetry Int Tables number 19 space group name Hall 'P 2ac 2ab' loop symmetry equiv pos site id _symmetry_equiv pos as xyz 1 x,y,z 2 1/2-x,-y,1/2+z 3 -x, 1/2+y, 1/2-z 4 1/2+x, 1/2-y, -z cell length a 7.7378(2) cell length b 10.2421(3)cell length c 12.0395(3)cell angle alpha 90 cell angle beta 90 cell angle gamma 90 cell volume 954.146 loop _exptl_crystal face index h exptl crystal face index k exptl crystal face index l exptl crystal face perp dist 0 1 1 10 1 1 1 15 1 1 0 15 $1 \ 1 \ -1 \ 15$ 2 0 0 16 1 0 1 16 1 0 2 16.5



Another example - glycine

Only 143 CSD entries for glycine

GLYCIN02 matched the unit cell in Groth

>2000 other examples
of organic crystal
morphologies in Groth





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Link the other ~2000 morphologies in "Groth" to the CSD?

A standardised way of recording Morphology Data?





Thank you

Comments?



