Christian Baerlocher and Lynne B. McCusker Laboratory of Crystallography, ETH Zurich

#### Powder Diffraction and Electron Microscopy

a powerful combination



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Introduction

powder diffraction and electron microscopy structure envelope

#### Combinations of XPD and electron microscopy

TNU-9	(FOCUS + HRTEM)
IM-5	(pCF + HRTEM)
SSZ-74	(pCF + HRTEM)

#### Precession electron diffraction

weak reflection elimination phase retrieval

#### FAKED electron diffraction data

SSZ-82

#### Conclusions

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	Powder	Electron
Data collection	easy	difficult
Crystallite size	μm	nm
Dataset	complete	incomplete
Lattice parameters	precise	approximate
Systematic absences	difficult	easy
Intensities	kinematic	dynamical
Multiple diffraction	no	yes
Overlap	yes	no
Phase information	no	yes

PowderElectronData collectioneasydifficult

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1) to help in indexing powder pattern by

- determine lattice parameters from SAED patterns
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2) determine crystal structure usings electron crystallography and then refining the structure with powder data

Solving crystal structures by integrating the two data sets

High resolution transmission electron microscopy

High resolution transmission electron microscopy



High resolution transmission electron microscopy



High resolution transmission electron microscopy



High resolution transmission electron microscopy



CTF corrected symmetry averaged

High resolution transmission electron microscopy



CTF corrected symmetry averaged

High resolution transmission electron microscopy



CTF corrected symmetry averaged

Fourier transform → Structure factor amplitudes and phases

Zeolite Analcime



Zeolite Analcime



A structure envelope is a curved surface that separates regions of high electron density from those of low electron density



A structure envelope is a curved surface that separates regions of high electron density from those of low electron density

 $\rightarrow$  Isosurface of a low resolution electron density map



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Reflection 2 1 1
# Structure Envelope

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Reflection 2 1 1

100, 101, 102, 002

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Reflection 2 1 1

phase can be fixed

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Reflection 2 1 1

phase can be fixed

100, 101, 102, 002

Phases can be obtained from HRTEM images

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#### Zeolites

3-dimensional, 4-connected framework structure



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3-dimensional, 4-connected framework structure

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approximate unit cell contents individual minimum distances selected reflection intensities

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assign random starting phases

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assign random starting phases electron density map

approximate unit cell contents individual minimum distances selected reflection intensities assign random starting phases electron density map

peak search

approximate unit cell contents individual minimum distances selected reflection intensities assign random starting phases electron density map peak search

















Unit Cell	
Space Group	<b>C</b> 2/m
a	28.2219 Å
b	20.0123 Å
С	19.4926 Å
β	92.33°
Reflections powder pattern (d <sub>min</sub> =1.17Å) overlapping (0.3*FWHM)	3705 3154
FOCUS	
reflections used (65% strongest)	1481

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Unit Cell	
Space Group	<b>C2/m</b>
a	28.2219 Å
b	20.0123 Å
С	19.49 <b>2</b> 6 Å
β	92.33°
Reflections powder pattern (d <sub>min</sub> =1.17Å) overlapping (0.3*FWHM)	<ul> <li>3705</li> <li>3154</li> <li>85% overlap</li> </ul>
FOCUS reflections used (65% strongest)	1481

 $\rightarrow$  no solution













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Reflections powder pattern ( $d_{min}$ =1.17Å) overlapping (0.3*EW/HM)	3705 3154
	3134
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 $\rightarrow$  solution after 16 days





24 Si atoms in the asymmetric unit



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Unit Cell	
Space Group	Стст
a	14.2088 Å
b	57.2368 Å
С	19.9940 Å
Reflections	
powder pattern (d <sub>min</sub> =1.05 Å)	4120
overlapping (0.3*FWHM)	3499





4120

3499

#### Reflections

powder pattern (*d<sub>min</sub>* =1.05 Å) overlapping (0.3\*FWHM)





#### Reflections

powder pattern ( $d_{min}$  =1.05 Å) overlapping (0.3\*FWHM) 4120 3499 85% overlap

#### Unit Cell Space Group a b c

*Cmcm* 14.2088 Å 57.2368 Å 19.9940 Å

#### Reflections

powder pattern ( $d_{min}$  =1.05 Å) overlapping (0.3\*FWHM) 4120 3499 85% overlap

Superflip no phases

promising, but...



#### taken by Zhanbing He Stockholm University





# Stockholm University

taken by Zhanbing He







#### introduced in Superflip $\rightarrow$ still no good solutions

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approximated model derived from HRTEM data in C2cm

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approximated model derived from HRTEM data in C2cm



36 Si atoms

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36 Si atoms

#### geometry strained

#### introduced in Superflip $\rightarrow$ still no good solutions

approximated model derived from HRTEM data in C2cm



36 Si atoms

geometry strained calculated powder diffraction pattern does not fit measured one

Superflip run with

*Superflip* run with

1000 starting phase sets generated from HRTEM model (each  $\phi_{hkl}$  allowed to vary by up to 25%)

Superflip run with 1000 starting phase sets generated from HRTEM model (each  $\phi_{hkl}$  allowed to vary by up to 25%) structure envelope used to enforce channel system

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Five best electron density maps averaged C2cm symmetry imposed

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Five best electron density maps averaged <u>C2cm</u> symmetry imposed

Peaks interpreted in C2cm geometry not strained

Superflip run with 1000 starting phase sets generated from HRTEM model (each  $\phi_{hkl}$  allowed to vary by up to 25%) structure envelope used to enforce channel system

Five best electron density maps averaged C2cm symmetry imposed

Peaks interpreted in C2cm geometry not strained main features of powder pattern reproduced

Structure successfully refined in Cmcm





24 Si-atoms in the asymmetric unit



24 Si-atoms in the asymmetric unit unusual 2-dimensional 10-ring channel system



#### 24 Si-atoms in the asymmetric unit unusual 2-dimensional 10-ring channel system



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24 Si-atoms in the asymmetric unit unusual 2-dimensional 10-ring channel system 288 Si and 576 O = 864 atoms in the unit cell
Unit Cell	
Space Group	<b>C2/</b> c or <b>C</b> c
a	20.507 Å
b	13.394Å
С	20.099 Å
β	102.2°
Reflections	
powder pattern (d <sub>min</sub> = 0.95 Å)	3258
overlapping (0.3*FWHM)	2717

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Space Group	C2/c or Cc
a	20.507 Å
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Reflections

powder pattern ( $d_{min}$  = 0.95 Å) overlapping (0.3\*FWHM)

<sup>3258</sup> 83% overlap





HRTEM



# HRTEM

29 reflections





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with structure envelope	partial solution

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Superflip		
with structure envelope above used as seed for 100 phase sets	partial solution	

#### Superflip



Superflip

Superflip symmetry ambiguous, C2/c assumed

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11 Si atom positions taken from electron density map

Superflip

symmetry ambiguous, C2/c assumed 11 Si atom positions taken from electron density map 12th Si atom position added to create a fully 4-connected net

Molecular modelling

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**C2/**c

geometry of 12th Si distorted

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C2/c Cc geometry of 12th Si distorted24 Si atoms in the asymmetric unit

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to estimate the position of the structure directing agent

**Rietveld refinement** 

C2/c Cc

Occupancy refined Final structure geometry of 12th Si distorted 24 Si atoms in the asymmetric unit improvement in profile fit distortion still present (2 Si) one Si disappeared 23 Si + 1 vacancy

Superflip

symmetry ambiguous, C2/c assumed 11 Si atom positions taken from electron density map 12th Si atom position added to create a fully 4-connected net

#### Molecular modelling

to estimate the position of the structure directing agent

#### **Rietveld refinement**

<b>C2/</b> c	geometry of 12th Si distorted
Сс	24 Si atoms in the asymmetric unit
	improvement in profile fit
	distortion still present (2 Si)
Occupancy refined	one Si disappeared
Final structure	23 Si + 1 vacancy

#### Structure solution gave the right answer!

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# **Electron Diffraction**

#### Precession electron diffraction



R. Vincent and P. Midgley,Ultramicroscopy 53,271-281 (1994)

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#### reduced multiple scattering

# **Electron Diffraction**

#### Precession electron diffraction



R. Vincent and P. Midgley, Ultramicroscopy 53, 271-281 (1994)

# reduced multiple scattering→ intensities more reliable

Combining PED data with X-ray powder diffraction data

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weak reflection elimination

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identify weak reflection in PED patterns

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True

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Combining PED data with X-ray powder diffraction data

phase retrieval
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phase retrieval

run charge flipping using just the reflections in the plane

Combining PED data with X-ray powder diffraction data

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→ phases for these reflections

Combining PED data with X-ray powder diffraction data

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ca 70% of total  $|F_{hkl}|$  amplitudes are correctly phased

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PED amplitudes + CF phases

Tuesday, June 7, 2011

Combining PED data with X-ray powder diffraction data

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Combining PED data with X-ray powder diffraction data

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→ phases for these reflections

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PED amplitudes + CF phases

Phases are almost as good as those from HRTEM, but easier to obtain

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It is possible to get phases from 2D electron diffraction data



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Extracted intensities from powder data are also not very accurate



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Question:



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### faked electron diffraction



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Yes, we can

### 2D-XPD



[010] Projection with "empty" regions



• Normal LeBail extraction from powder pattern



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- Select reflections of zone [010] (d<sub>min</sub>=3Å)



- Normal LeBail extraction from powder pattern
- Select reflections of zone [010] (d<sub>min</sub>=3Å) (47 reflections, 30% overlapped)



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- Select reflections of zone [010] (d<sub>min</sub>=3Å) (47 reflections, 30% overlapped)
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- Use some more tricks



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3.0Å

1.5Å

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• Select reflections of zone [010] with dmin=1.5Å

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- Run superflip with delta = 0.0
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### TNU-9







[010]



[001]



### TNU-9













### TNU-9



22 reflections, 83% correct

[010]



47 reflections, 72% correct





28 reflections, 85% correct
# TNU-9





22 reflections, 83% correct



47 reflections, 72% correct





28 reflections, 85% correct





# TNU-9





22 reflections, 83% correct

[010]



47 reflections, 72% correct





28 reflections, 85% correct





76 reflections, 79% correct

192reflections, 68% correct

106 reflections, 85% correct

Unit Cell	
Space Group	Pmmn
a	24.278 Å
b	11.466 Å
С	14.113 Å
Reflections	
powder pattern ( <i>d<sub>min</sub></i> = 0.95 Å)	3116
overlapping (0.25*FWHM)	2783
d <sub>min</sub>	0.9 Å

Unit Cell	
Space Group	Pmmn
a	24.278 Å
b	11.466 Å
С	14.113 Å

Reflections powder pattern ( $d_{min}$  = 0.95 Å) overlapping (0.25\*FWHM)  $d_{min}$ 

<sup>3116</sup> 2783 0.9 Å

### 2D-XPD fourier maps



### 2D-XPD fourier maps



[100]



61 Reflections



[010]

124 Reflections

[001]



53 Reflections

best "superflip" solutions



### normal run

best "superflip" solutions





normal run

using phases from 2D-XPD

Electron microscopy is a valuable complement to powder diffraction

Electron microscopy is a valuable complement to powder diffraction SAED for indexing and space group determination

Electron microscopy is a valuable complement to powder diffraction

SAED for indexing and space group determination

HRTEM for phase and symmetry information

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The framework structures of three complex polycrystalline zeolites

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The framework structures of three complex polycrystalline zeolites have been solved by combining powder diffraction with HRTEM data

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The powder charge-flipping (*pCF*) approach works in both real and reciprocal space, and therfore is well suited to combine information from different sources can deal with more than 800 atoms in the unit cell

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2D-X-ray powder diffraction data

can be phased with charge flipping

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#### 2D-X-ray powder diffraction data

can be phased with charge flipping used successfully in the structure determination of SSZ-82

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#### IM-5

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Stacey Zones Fabian Gramm Lars Massüger

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#### IM-5

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# Acknowledgments

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Dan Xie

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