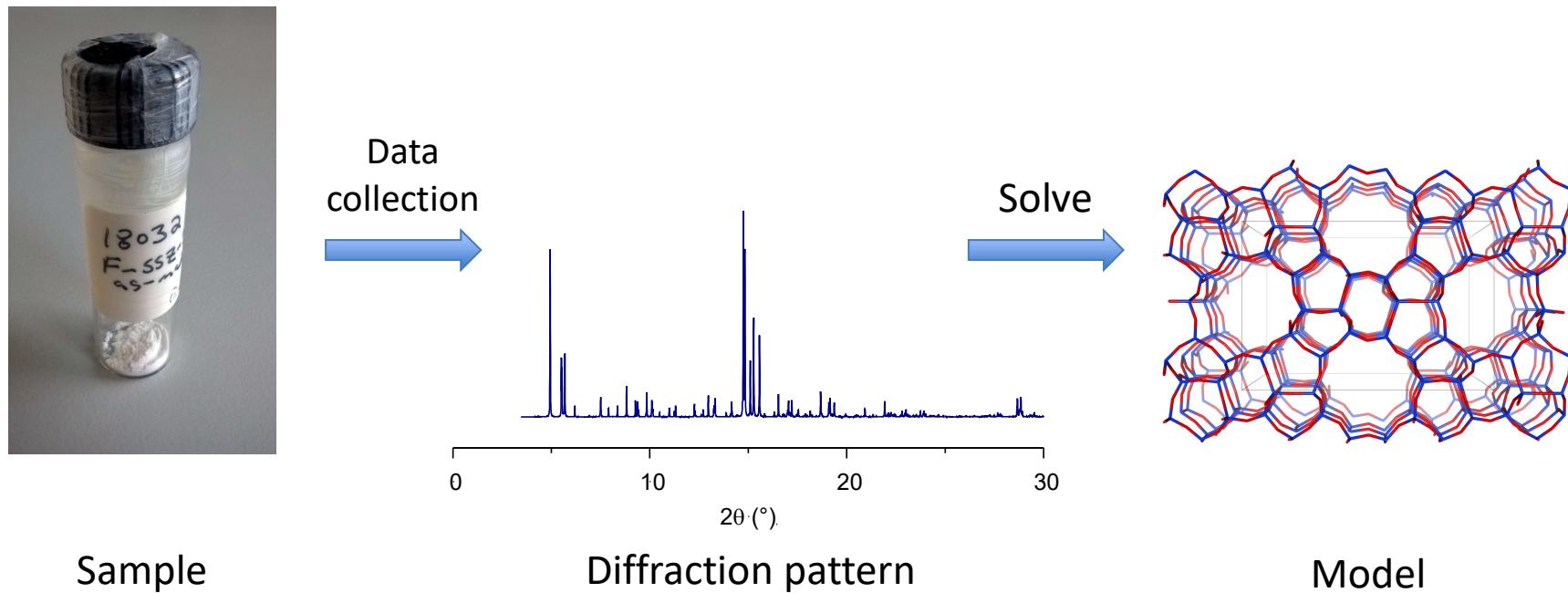


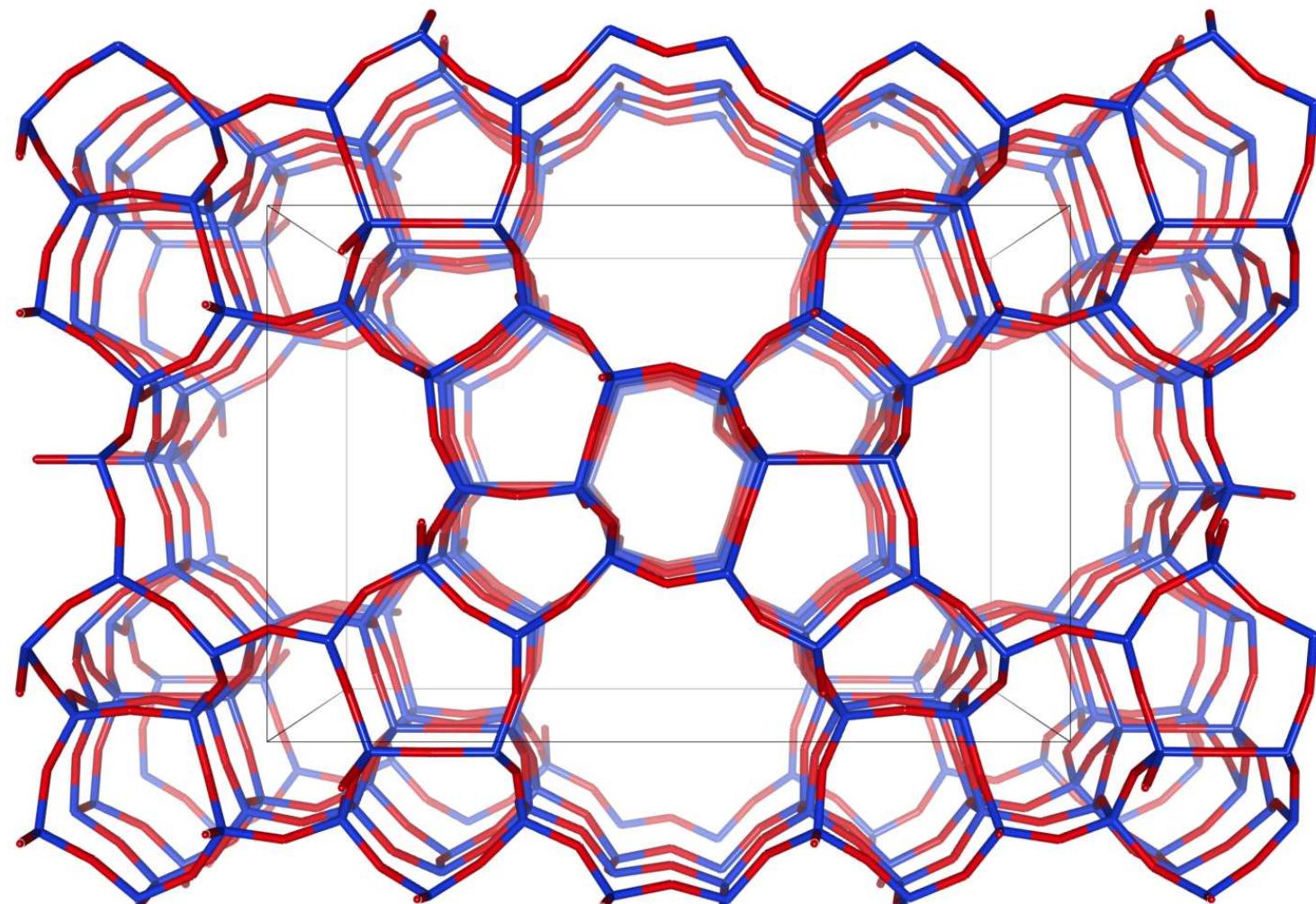
HOW DO THE POSITIONS OF ORGANIC SDAs DETERMINED FROM POWDER DIFFRACTION DATA COMPARE WITH THOSE OBTAINED FROM MOLECULAR MODELING?

Stef Smeets
Laboratory for Crystallography
ETH Zurich, Switzerland

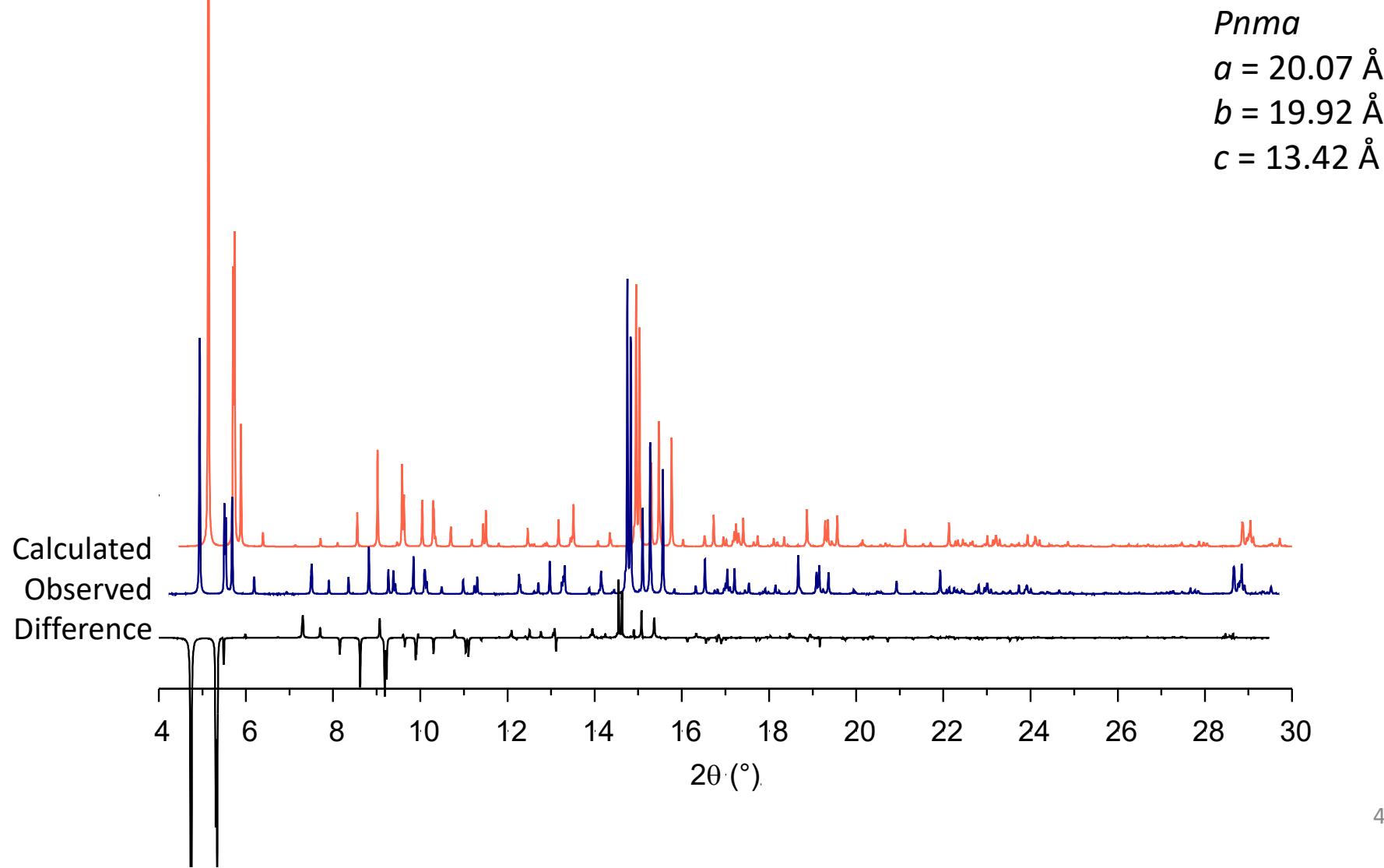
X-ray Powder diffraction



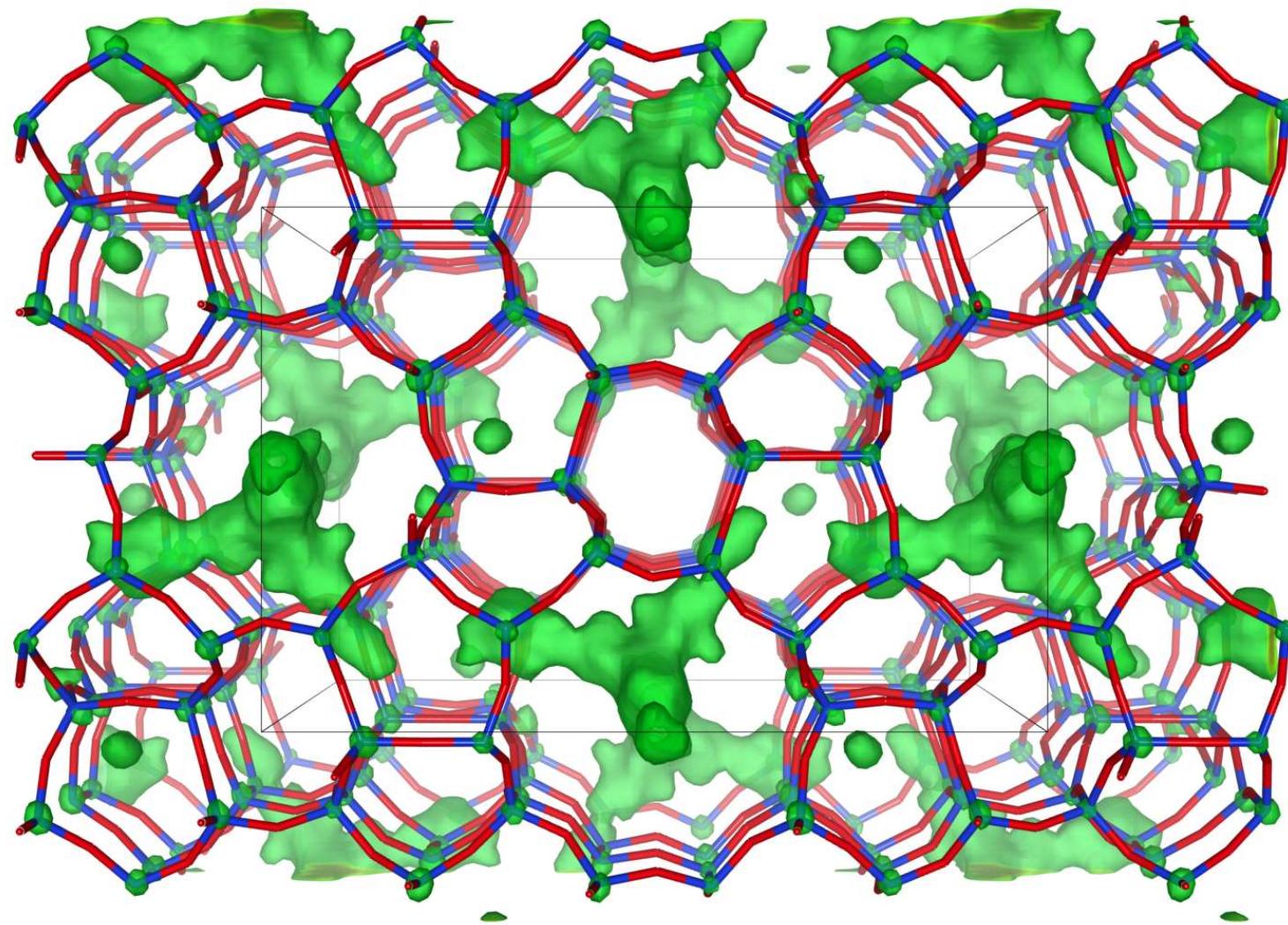
ZSM-5



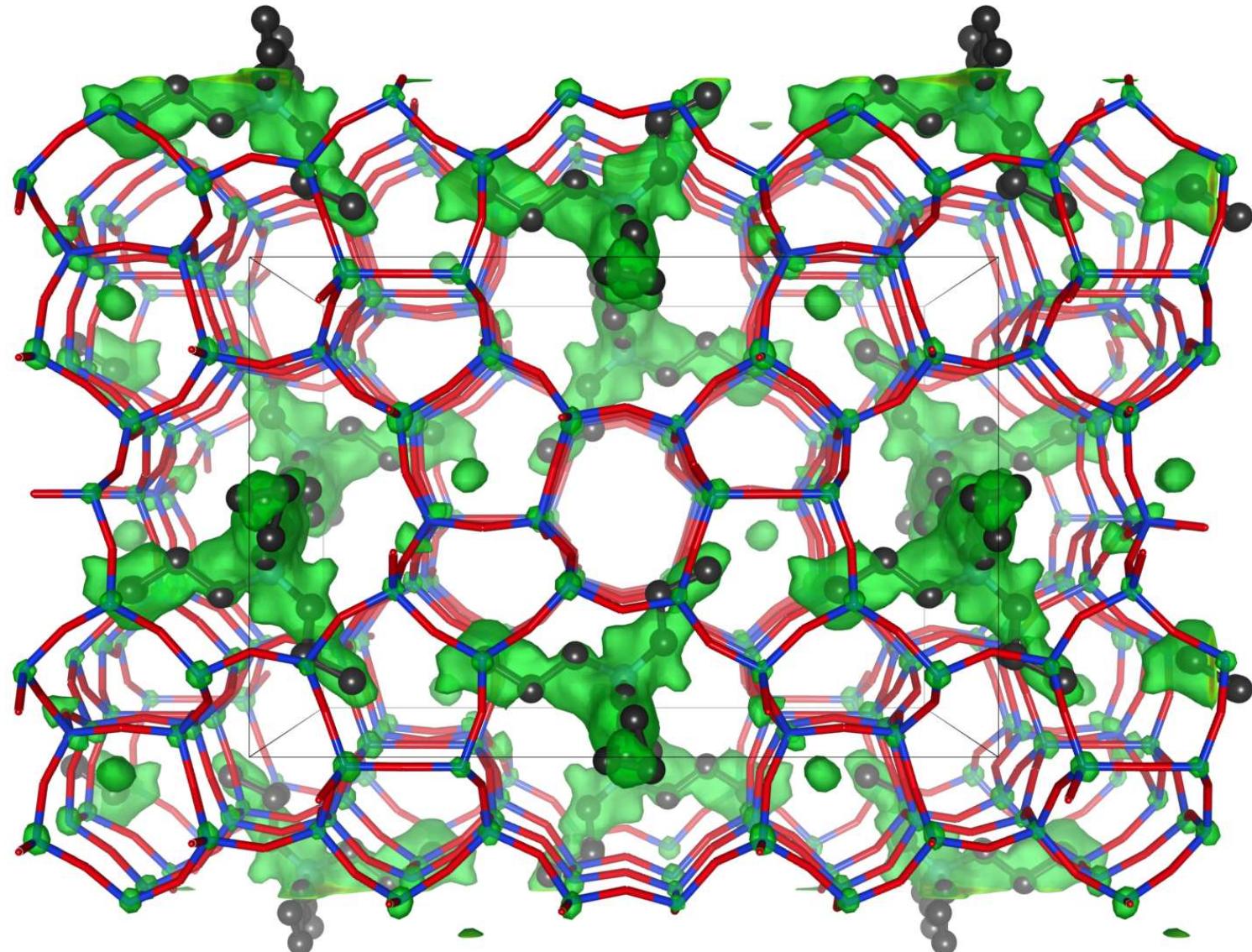
Difference plot ZSM-5



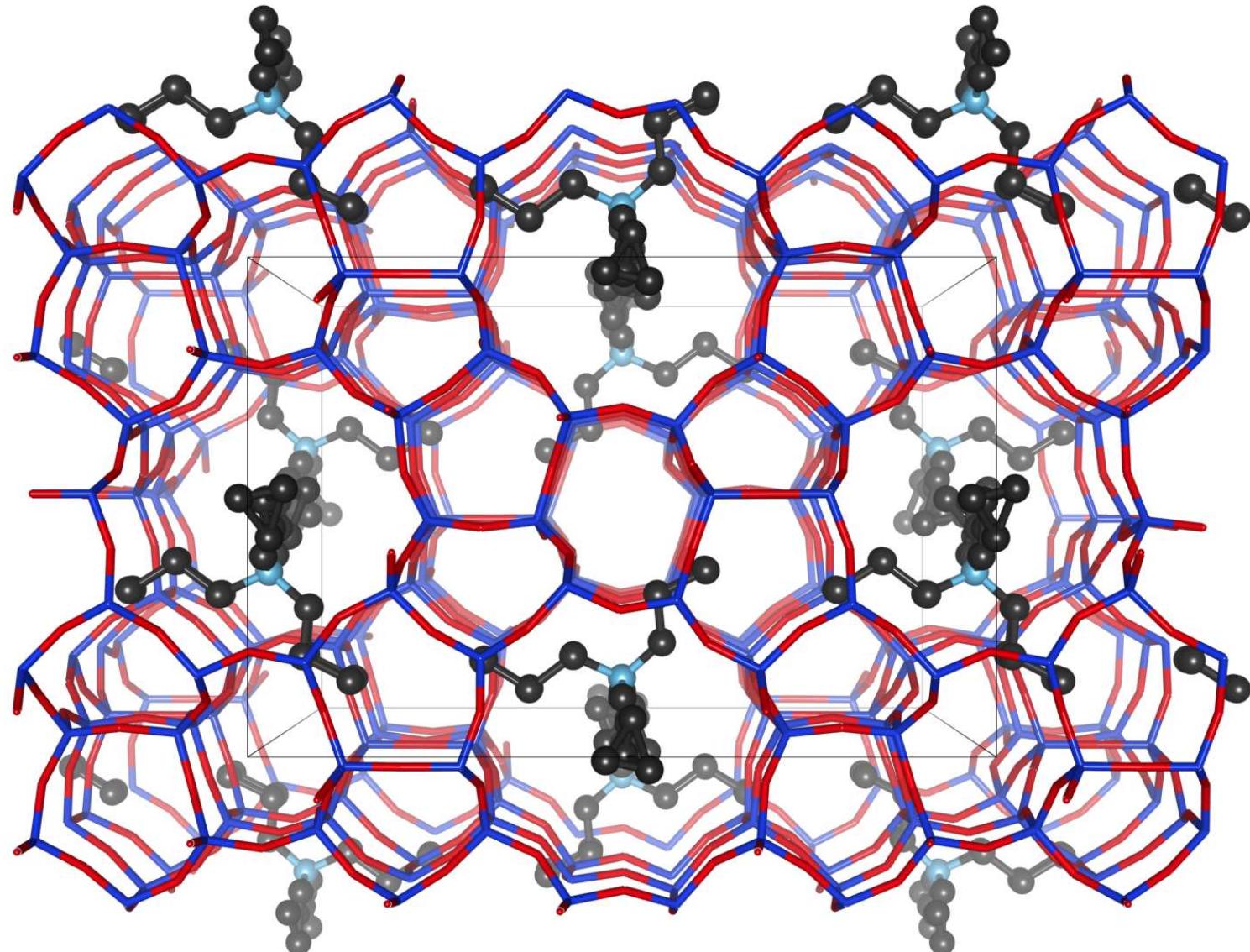
Difference map



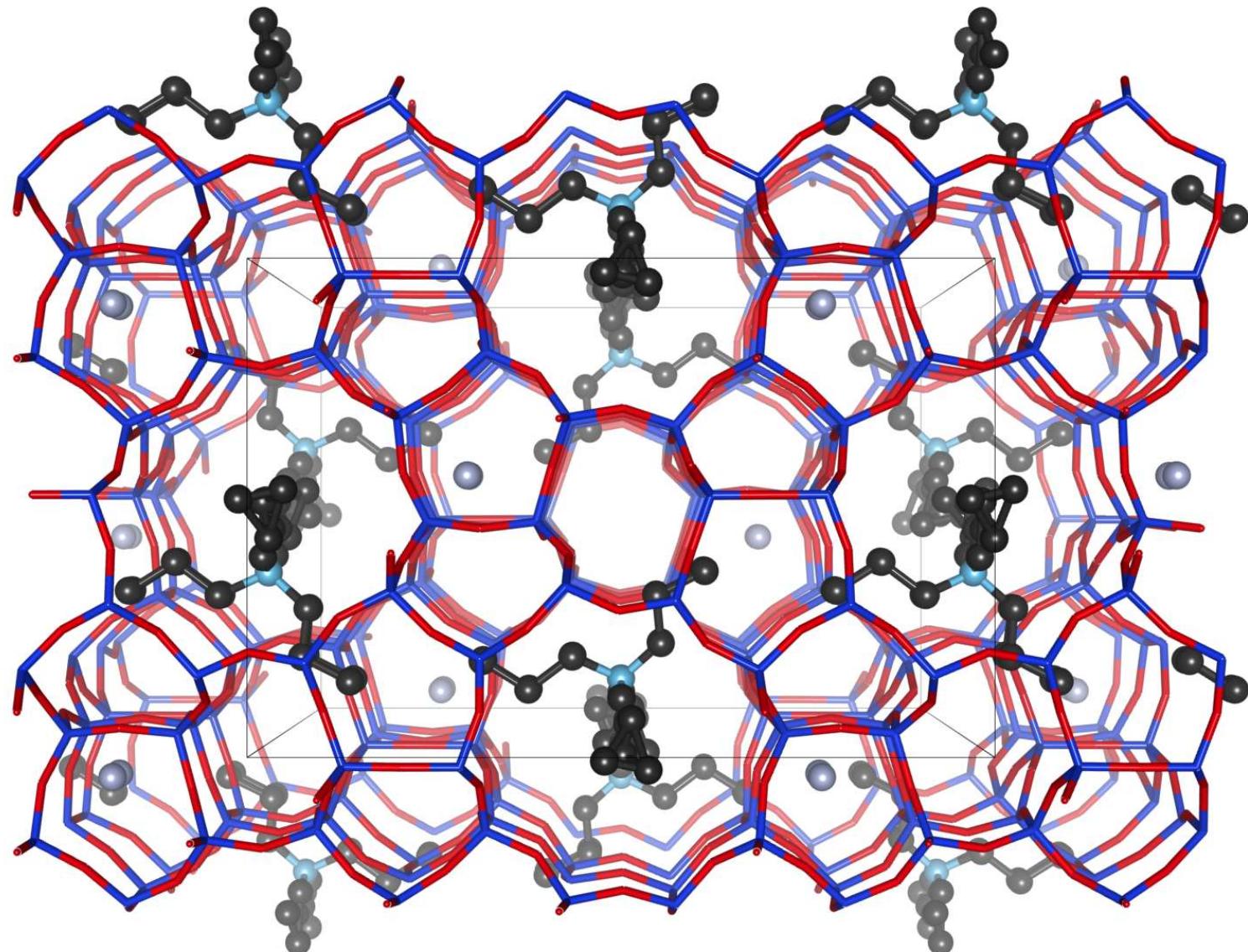
Difference map



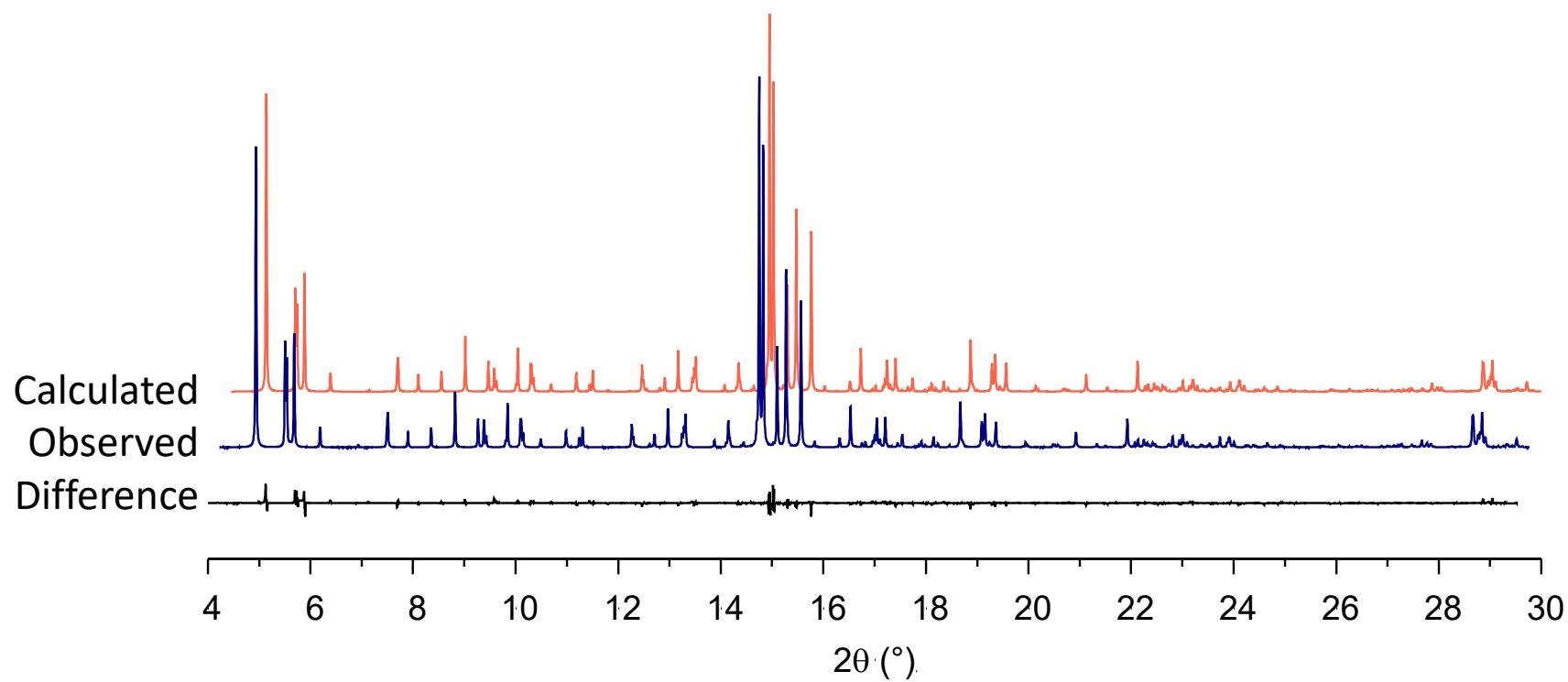
Refinement



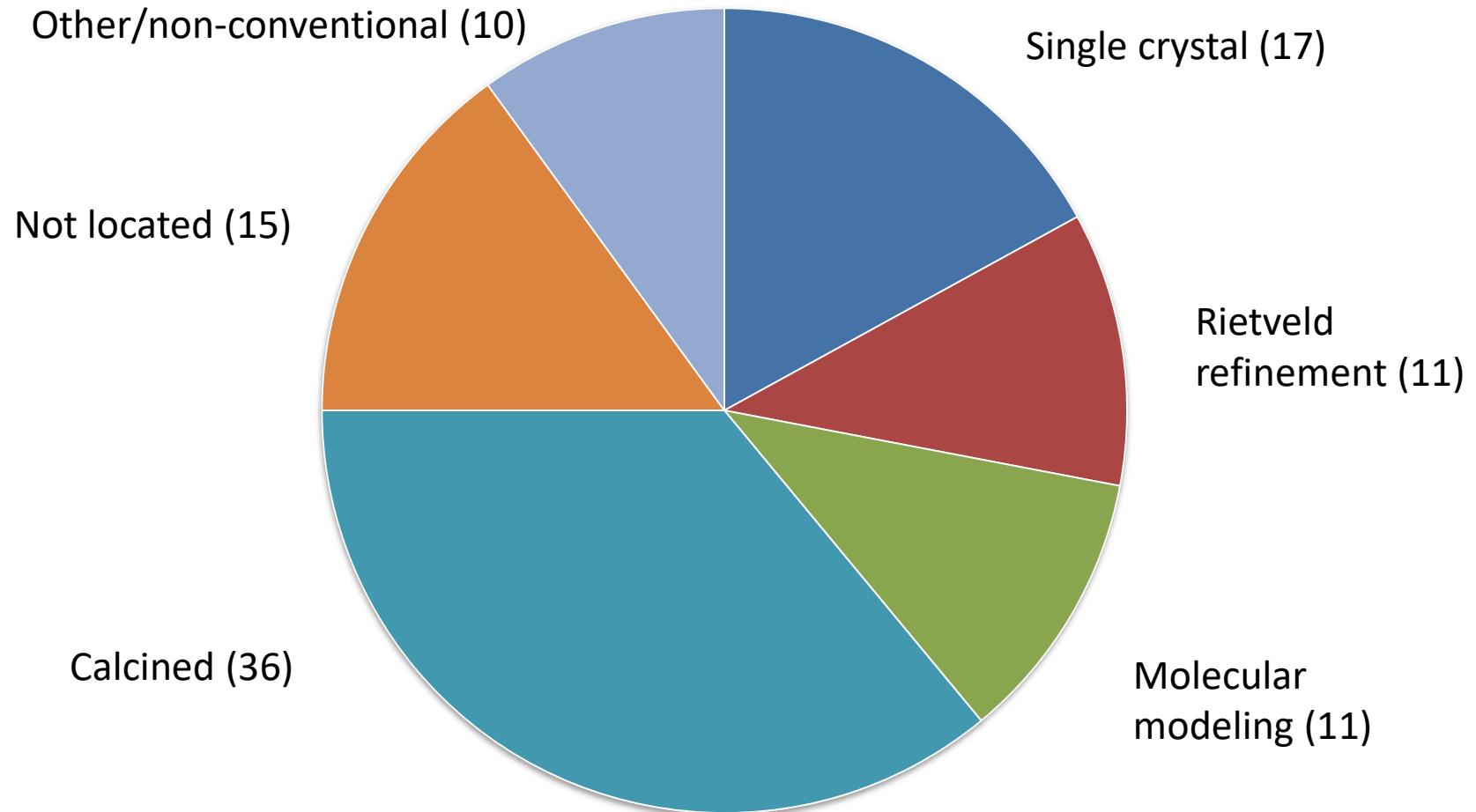
Refinement



Refinement of ZSM-5

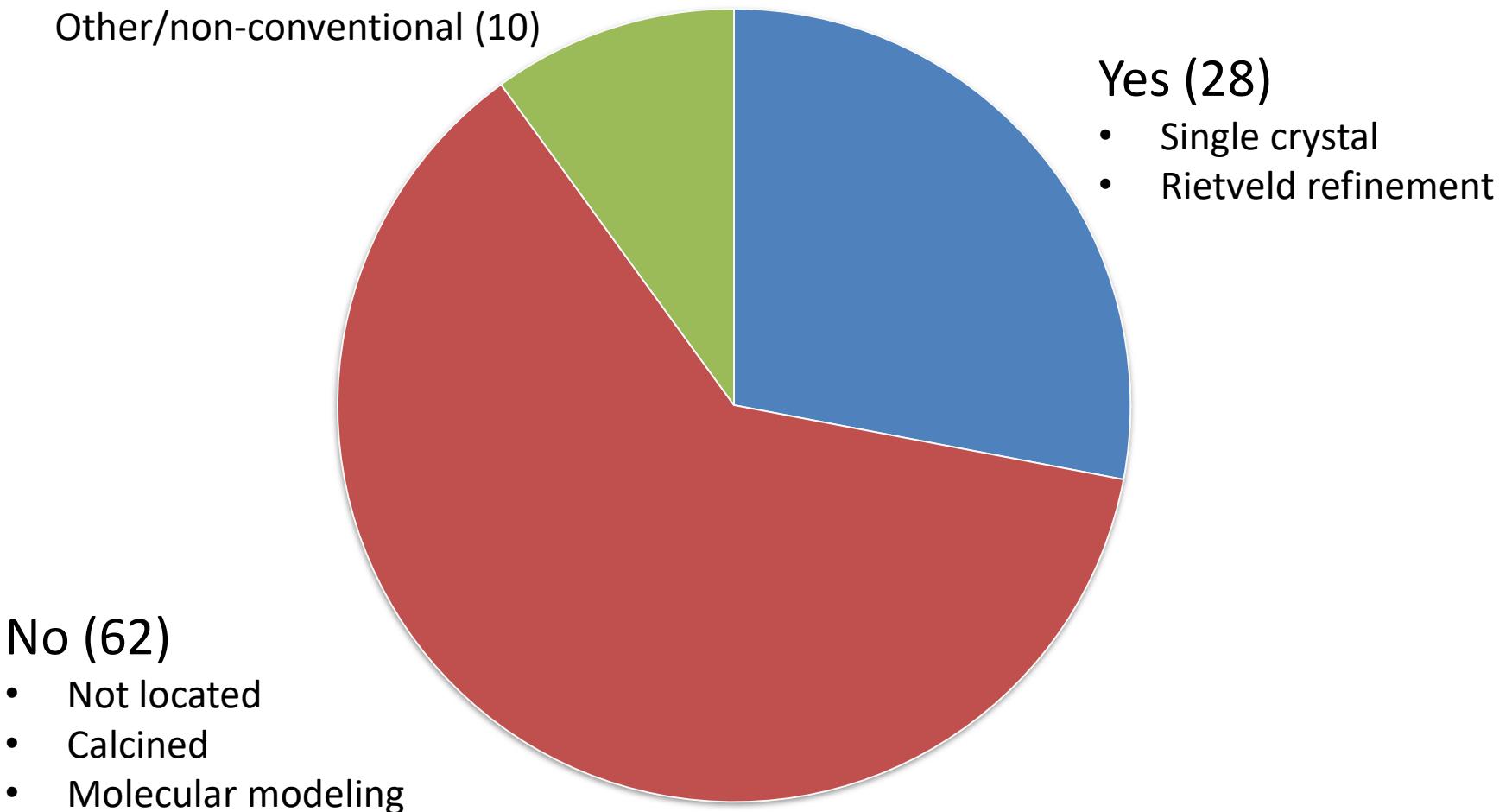


Organic SDA in zeolites 1998-2015



Number of structures: 100
(Database of zeolite structures)

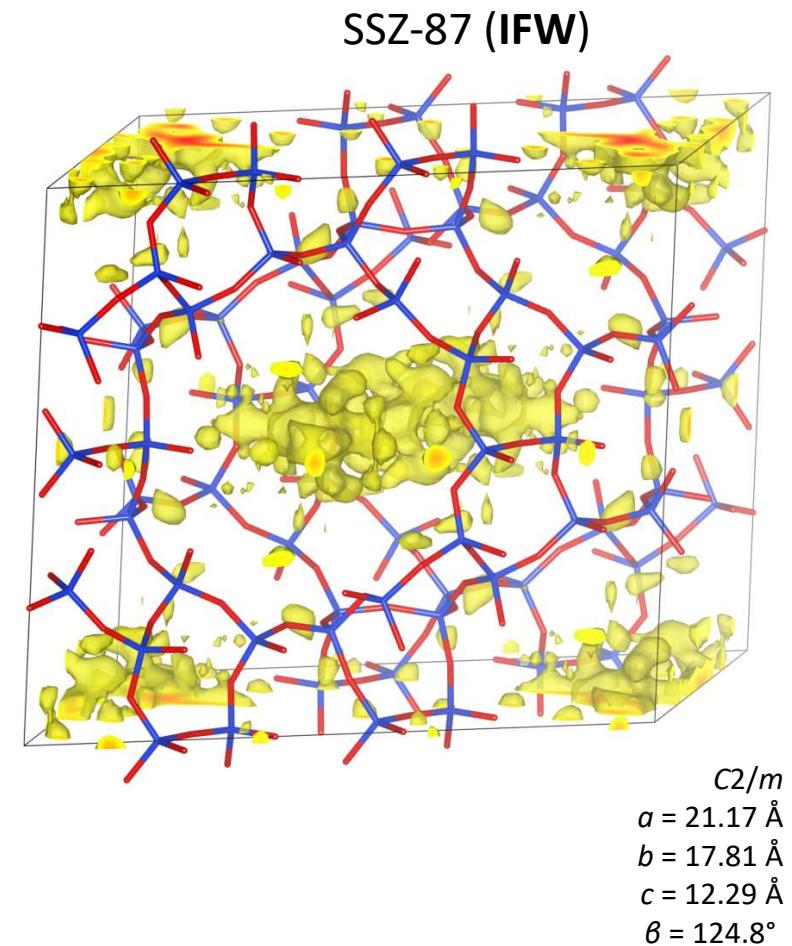
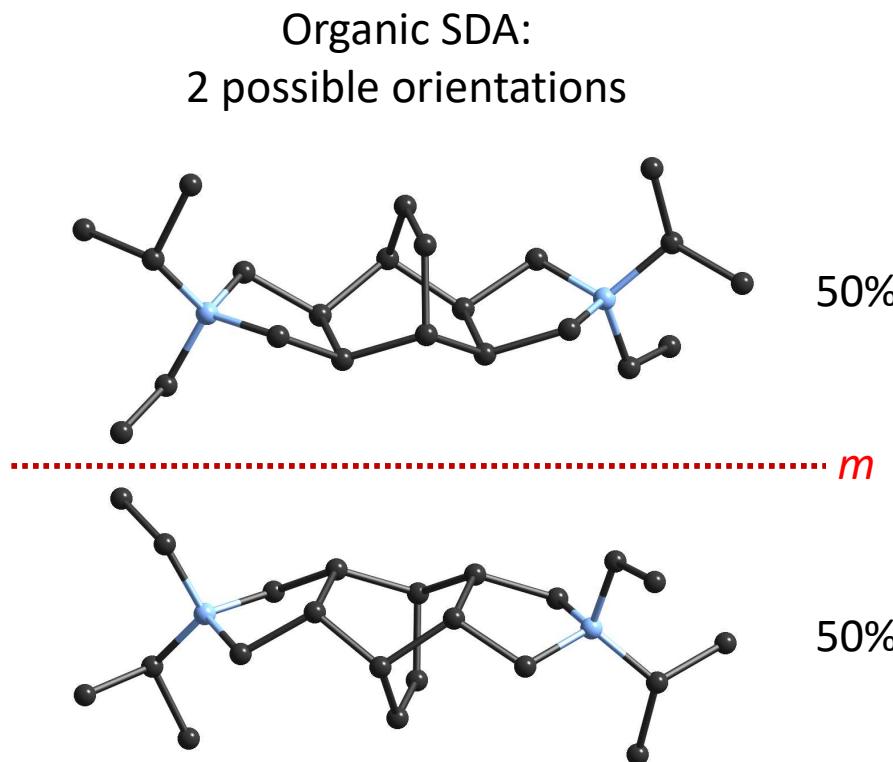
Organic SDA located from data



Reasons for locating the SDA

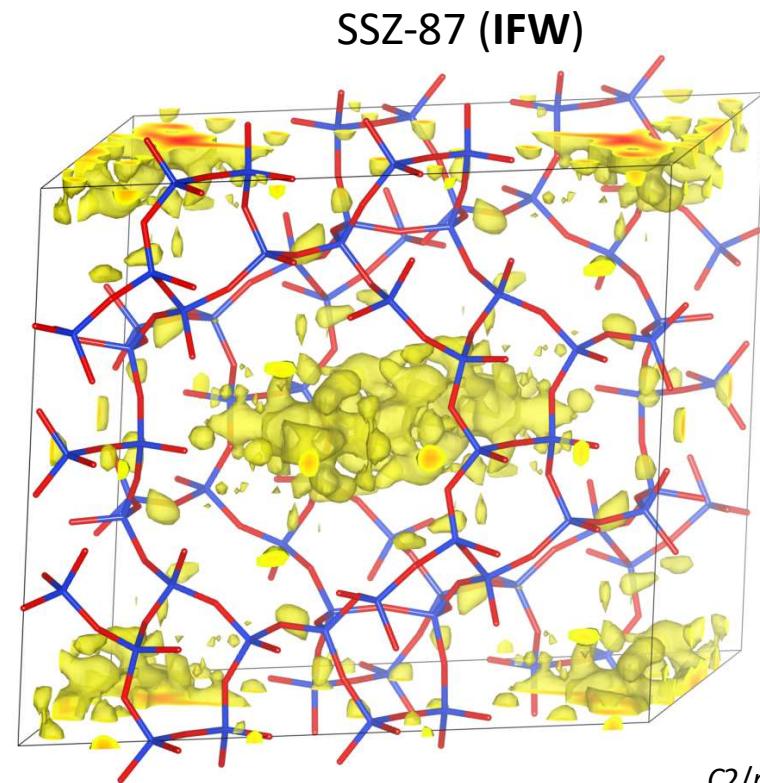
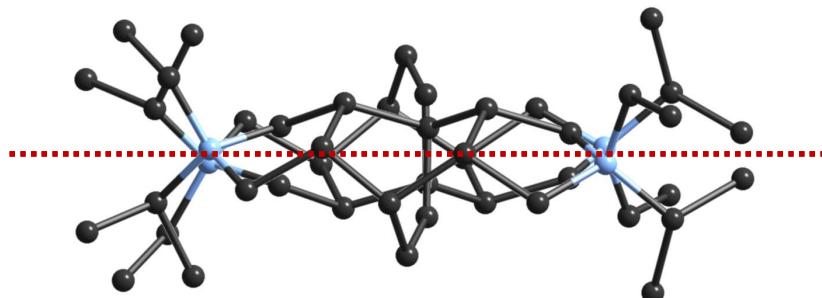
- Understanding zeolite formation
- Data quality affected by calcination
 - Better data => more accurate model
- Structure validation (SDA is part of synthesis)
- Obstacles
 - Disorder
 - Data limitations (overlap, resolution)

Obstacle 1: Disorder



Obstacle 1: Disorder

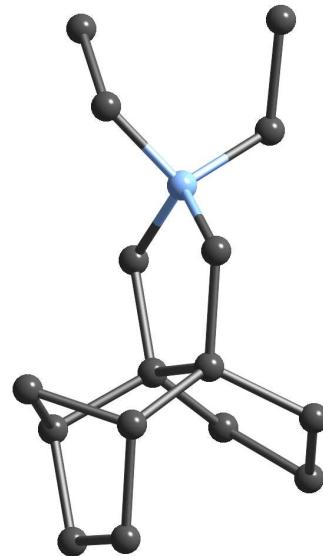
Organic SDA:
2 possible orientations



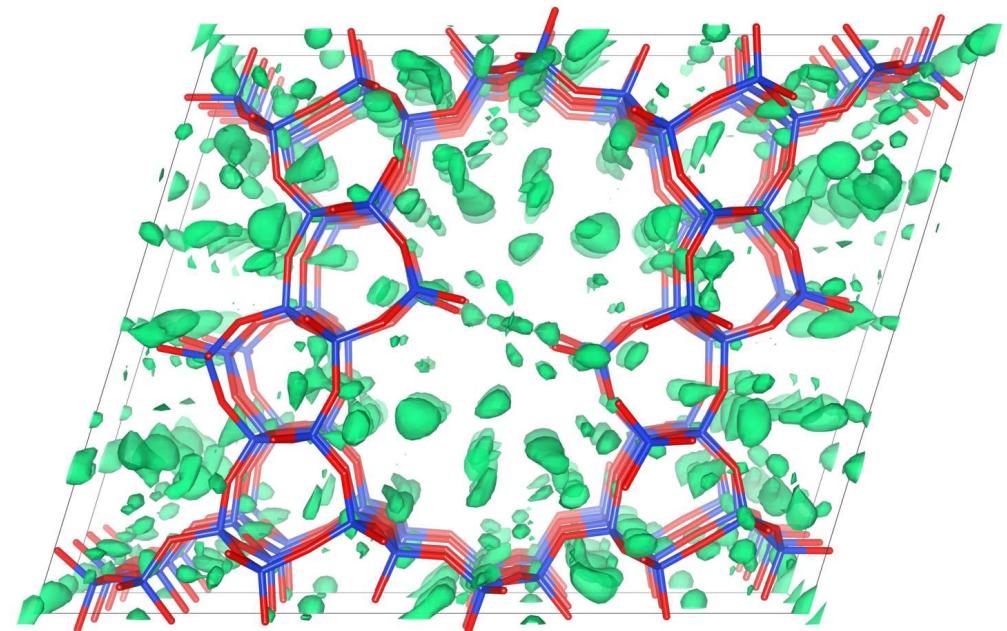
$C2/m$
 $a = 21.17 \text{ \AA}$
 $b = 17.81 \text{ \AA}$
 $c = 12.29 \text{ \AA}$
 $\beta = 124.8^\circ$

Obstacle 2: Data quality

Organic SDA:



SSZ-61 (**SFO**)



$P2_1/c$

$a = 19.76 \text{ \AA}$

$b = 10.07 \text{ \AA}$

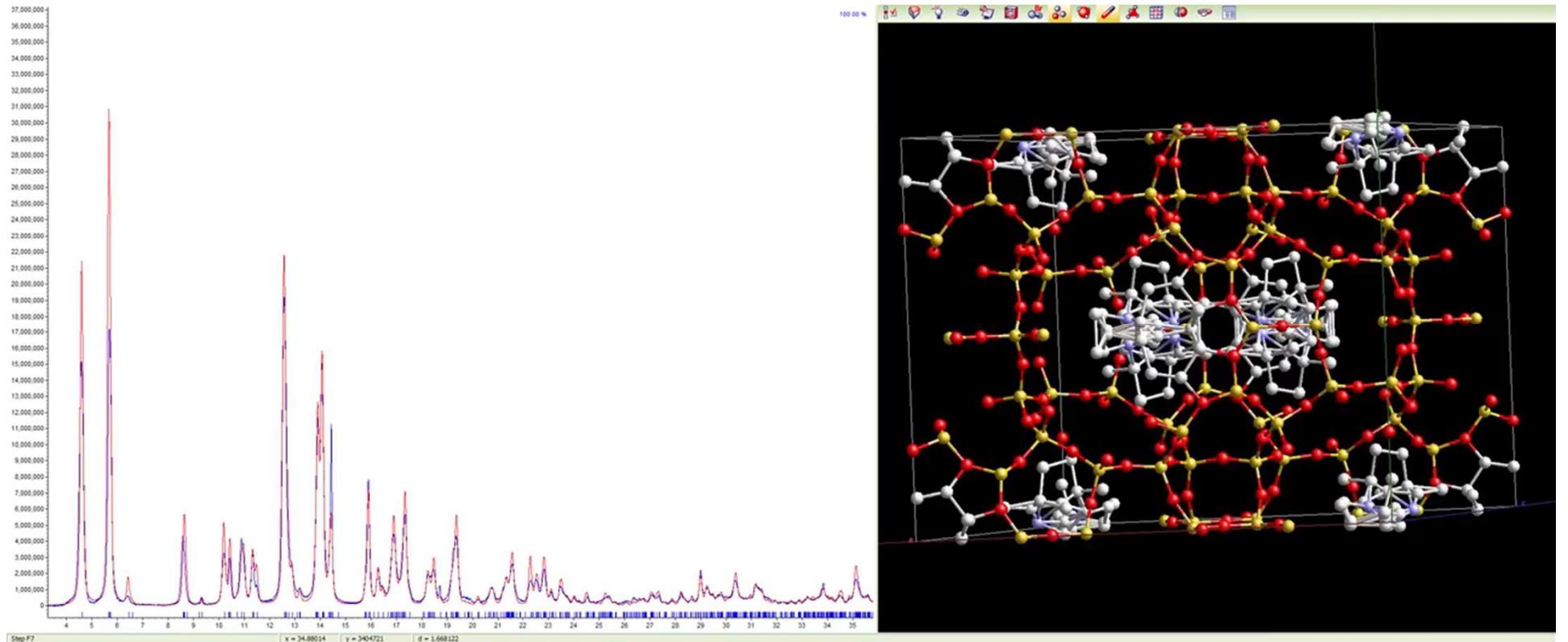
$c = 25.21 \text{ \AA}$

$\beta = 106.9^\circ$

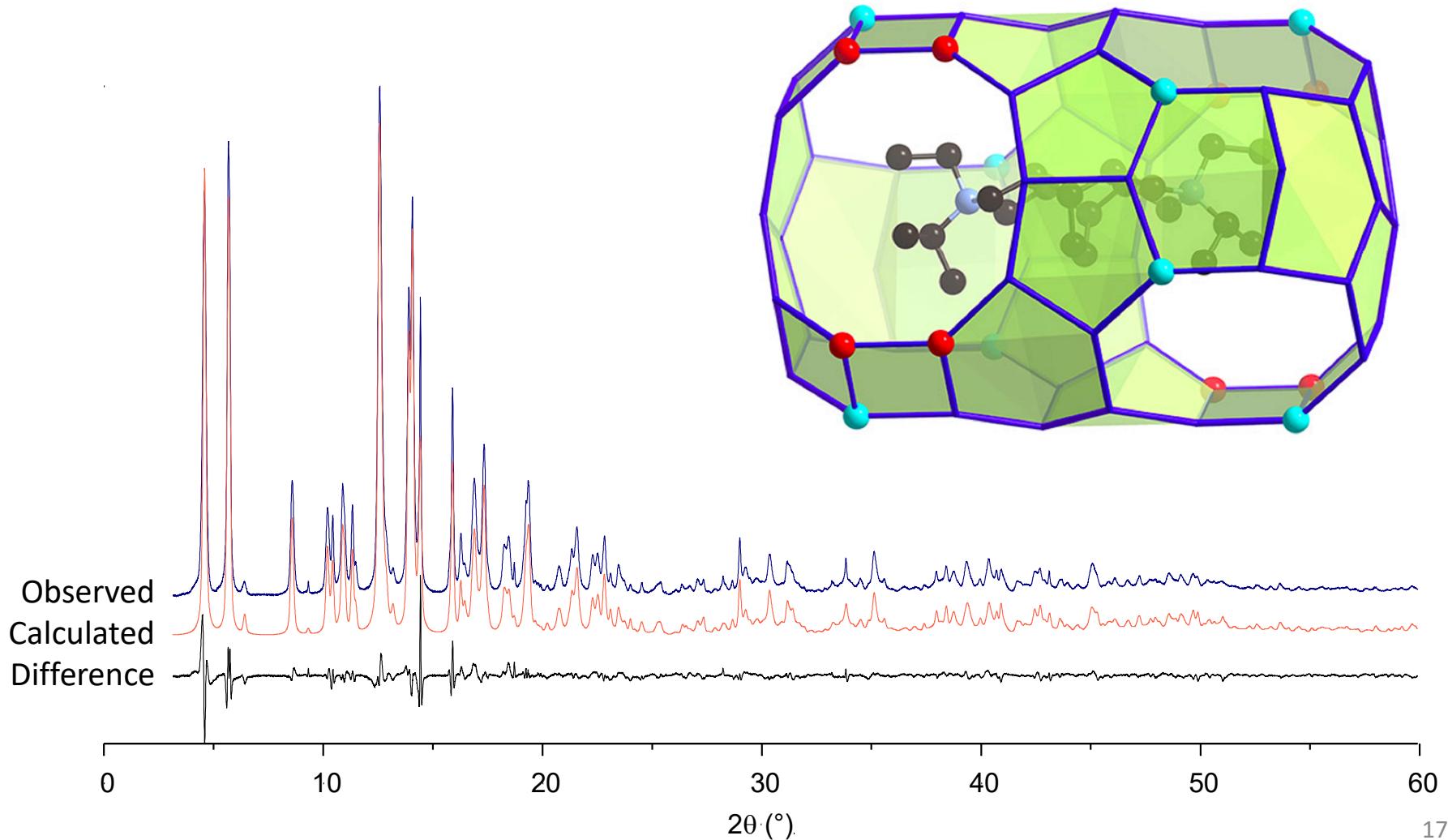
How to overcome these obstacles?

- Simulated annealing (global optimization)
 - Search model that best matches XPD data
 - Get starting location SDA => Rietveld refinement

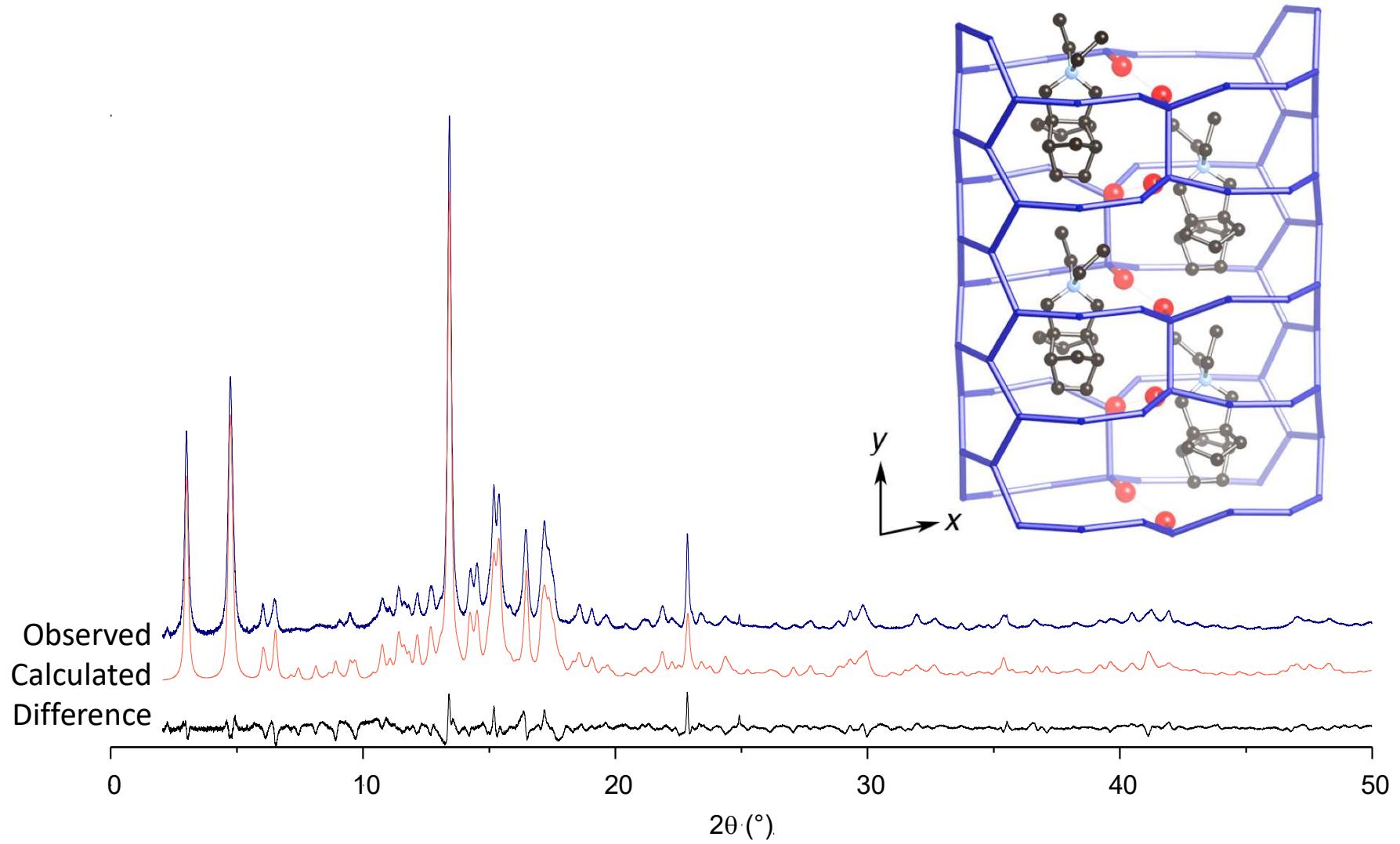
SSZ-87



Rietveld refinement of SSZ-87

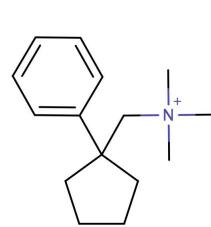


Rietveld refinement of SSZ-61

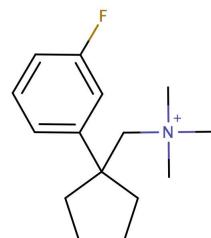


Can simulated annealing be applied in a routine manner to locate the SDA?

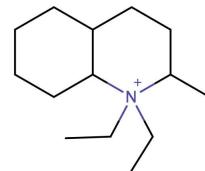
	ftc	a	b	c	α	β	γ	Spgr.
SSZ-53	SFH	5.02	33.74	21.17		90.5		<i>C2/c</i>
SSZ-55	ATS	12.95	21.85	5.08				<i>Cmc2</i> ₁
SSZ-56	SFS	13.95	19.90	12.36		106.7		<i>P2</i> ₁ / <i>m</i>
SSZ-58	SFG	25.11	12.50	12.86				<i>Pmma</i>
SSZ-59	SFN	5.02	12.74	14.72	103.4	90.5	100.9	<i>P-1</i>
SSZ-60	SSY	21.95	13.70	5.01				<i>P2</i> ₁



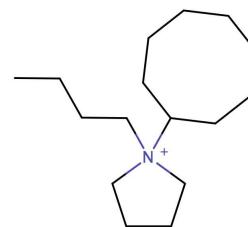
SSZ-53



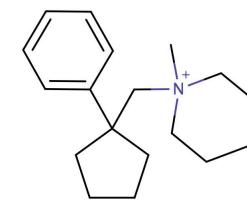
SSZ-55



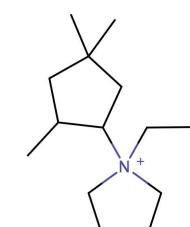
SSZ-56



SSZ-58



SSZ-59

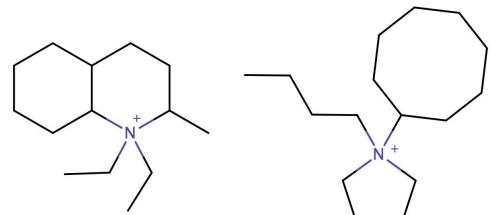


SSZ-60

Can simulated annealing be applied in a routine manner to locate the SDA?

- Six borosilicate zeolites (Saleh Elomari, Chevron)
- Flexible SDAs
- Location SDA originally simulated with molecular modeling

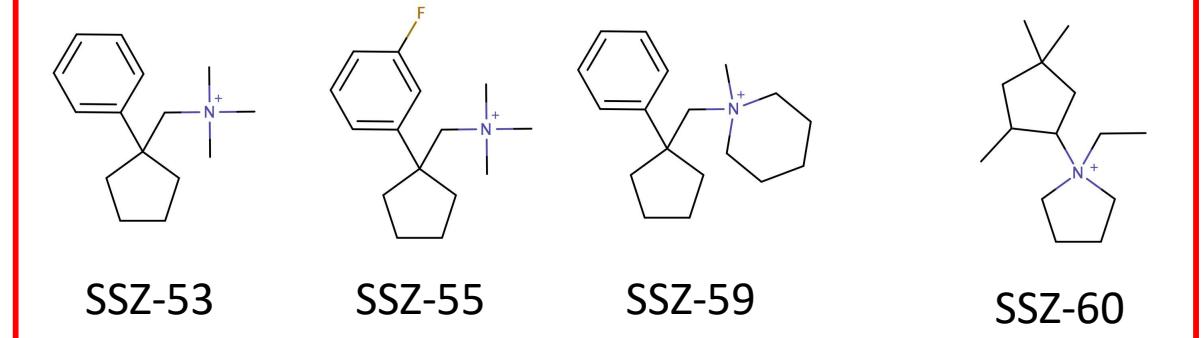
Two-dimensional
channel system



SSZ-56

SSZ-58

One-dimensional
channel system



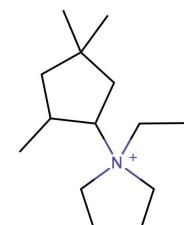
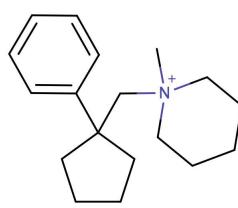
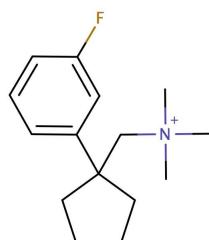
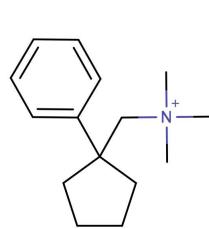
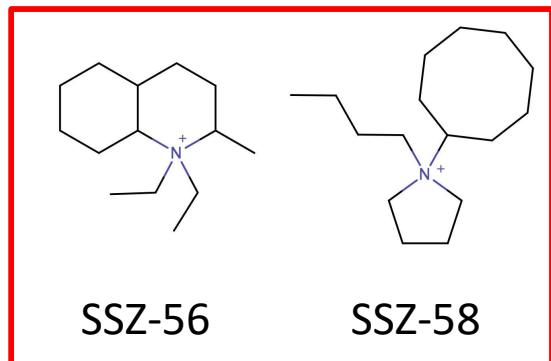
SSZ-60

SSZ-56 & SSZ-58

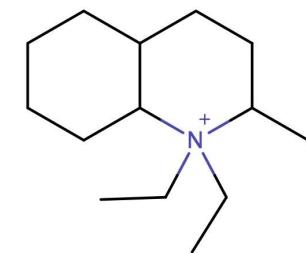
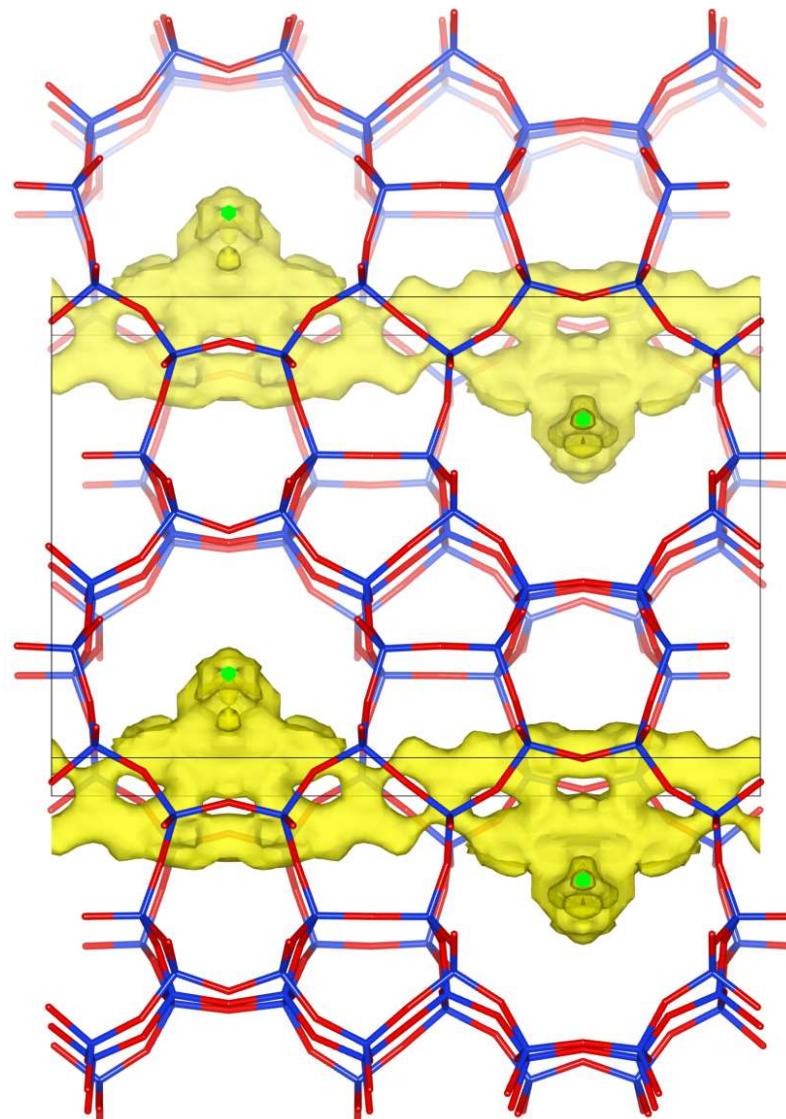
	ftc	a	b	c	α	β	γ	Spgr.
SSZ-56	SFS	13.95	19.90	12.36		106.7		<i>P</i> 2 ₁ /m
SSZ-58	SFG	25.11	12.50	12.86				<i>Pmma</i>

12-ring,
10-ring

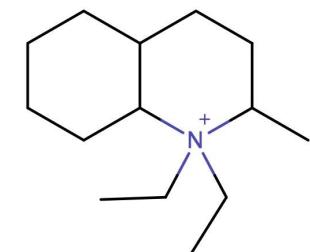
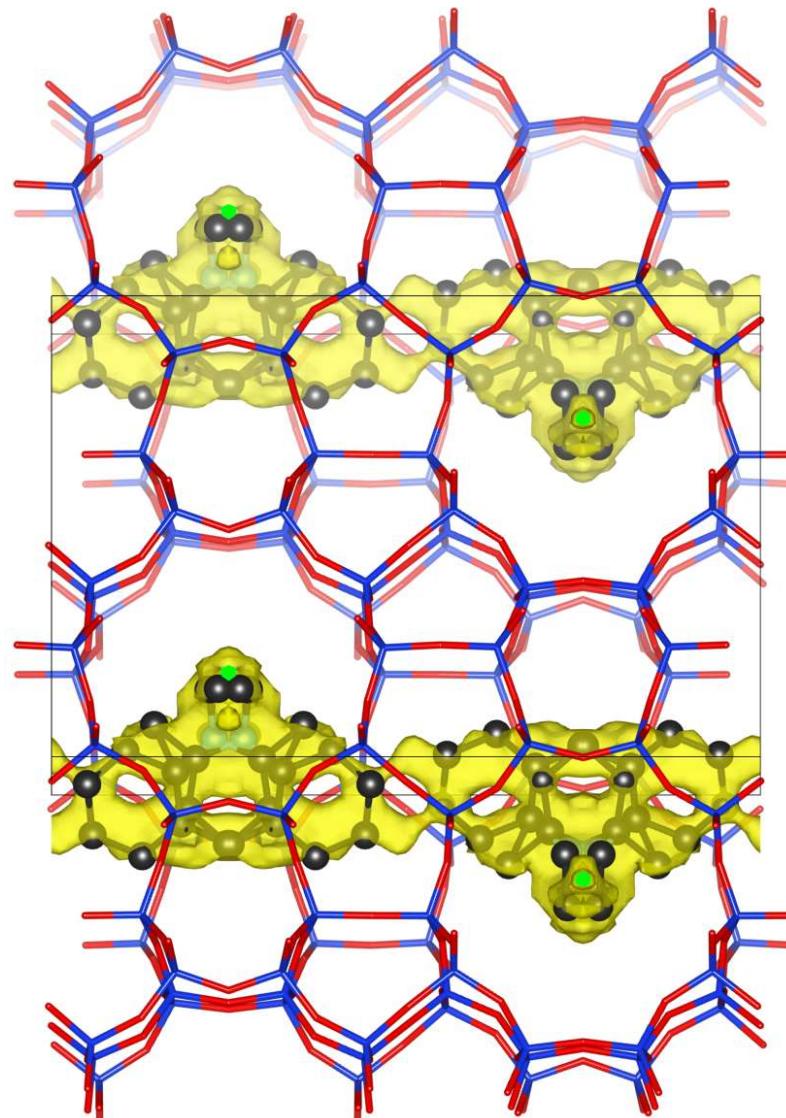
10-ring,
10-ring



Difference map SSZ-56

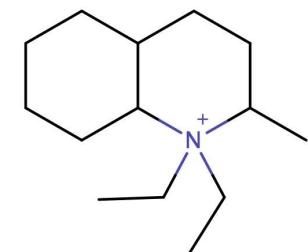
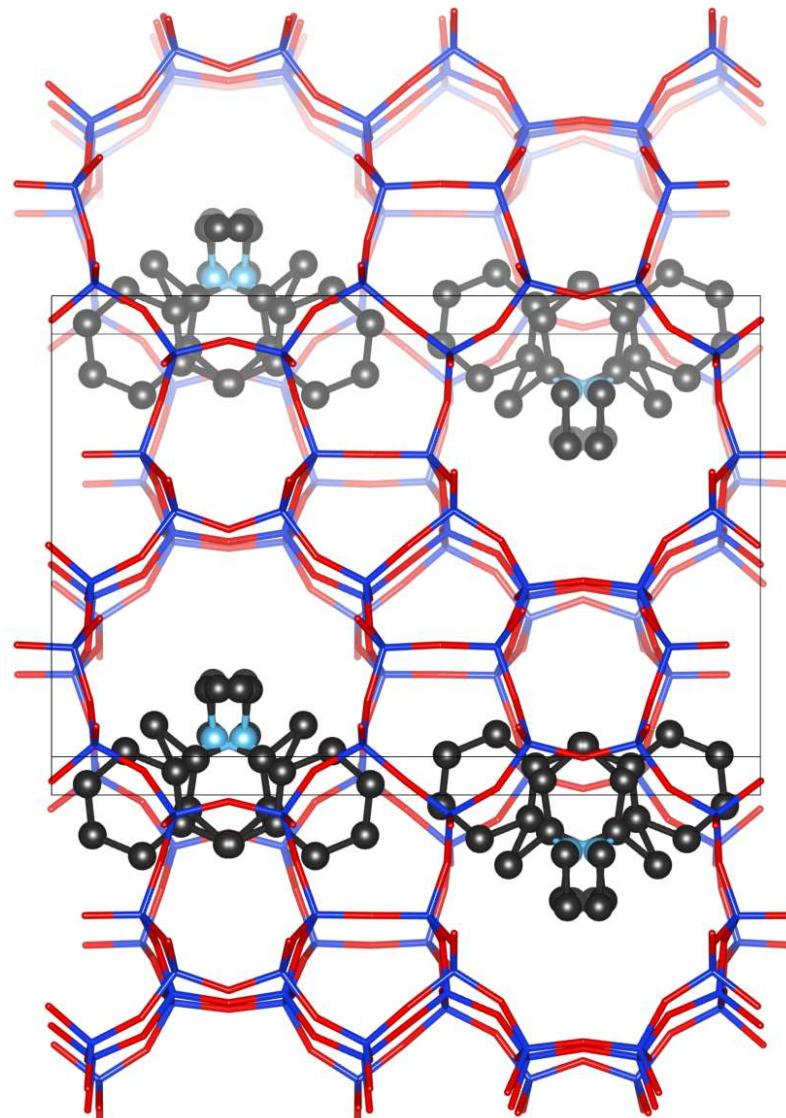


Difference map SSZ-56



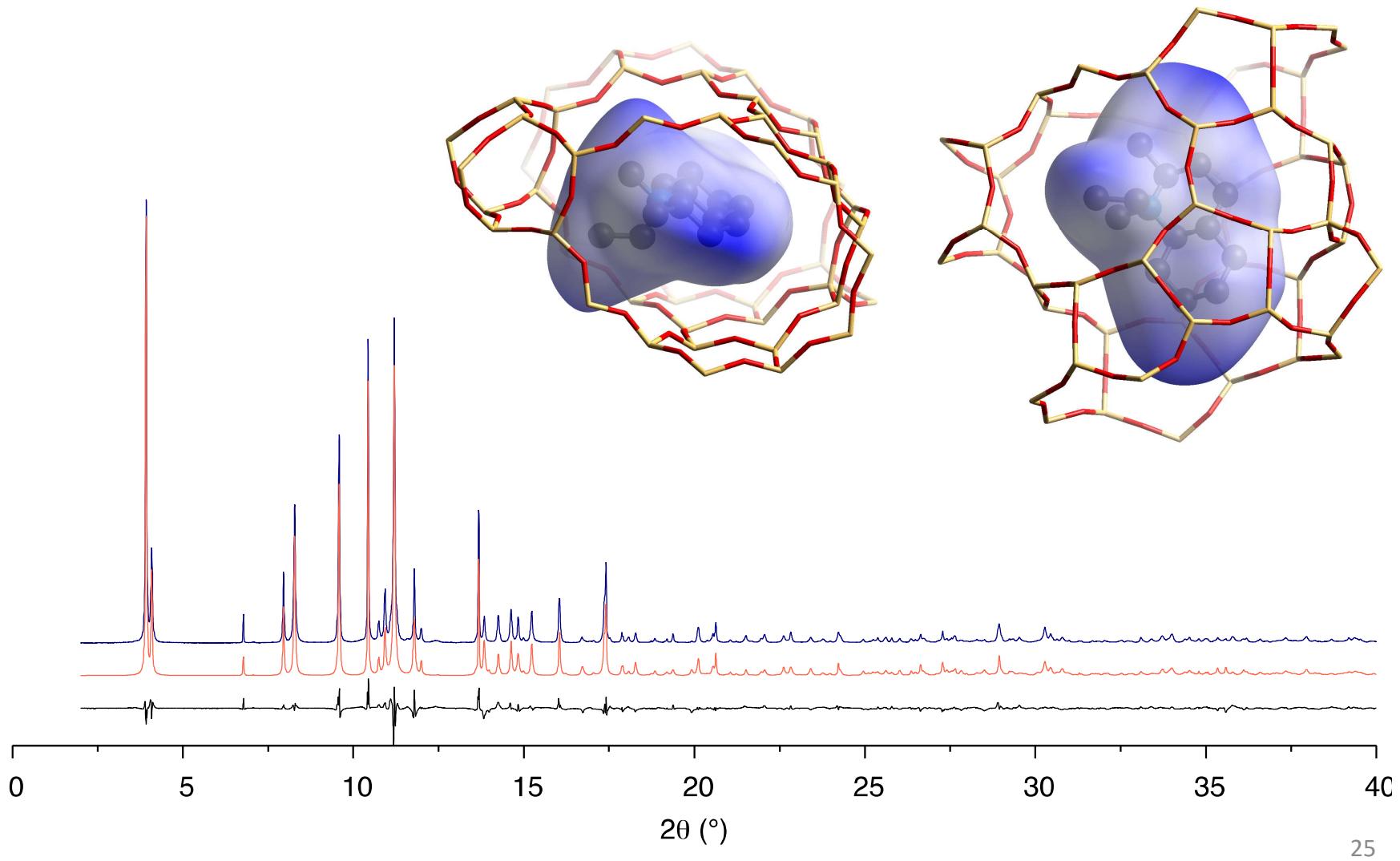
Two-fold disorder

Difference map SSZ-56

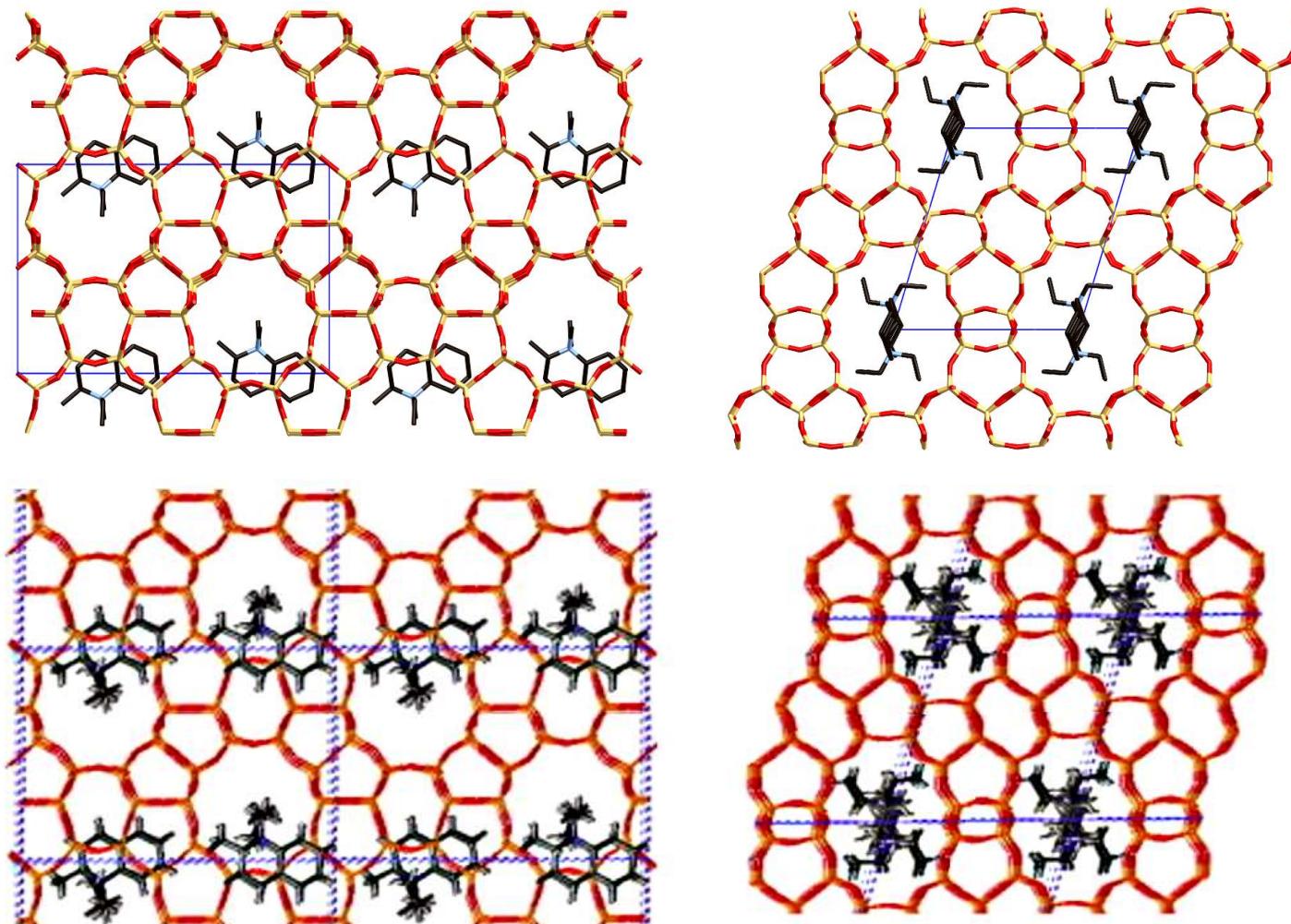


Two-fold disorder

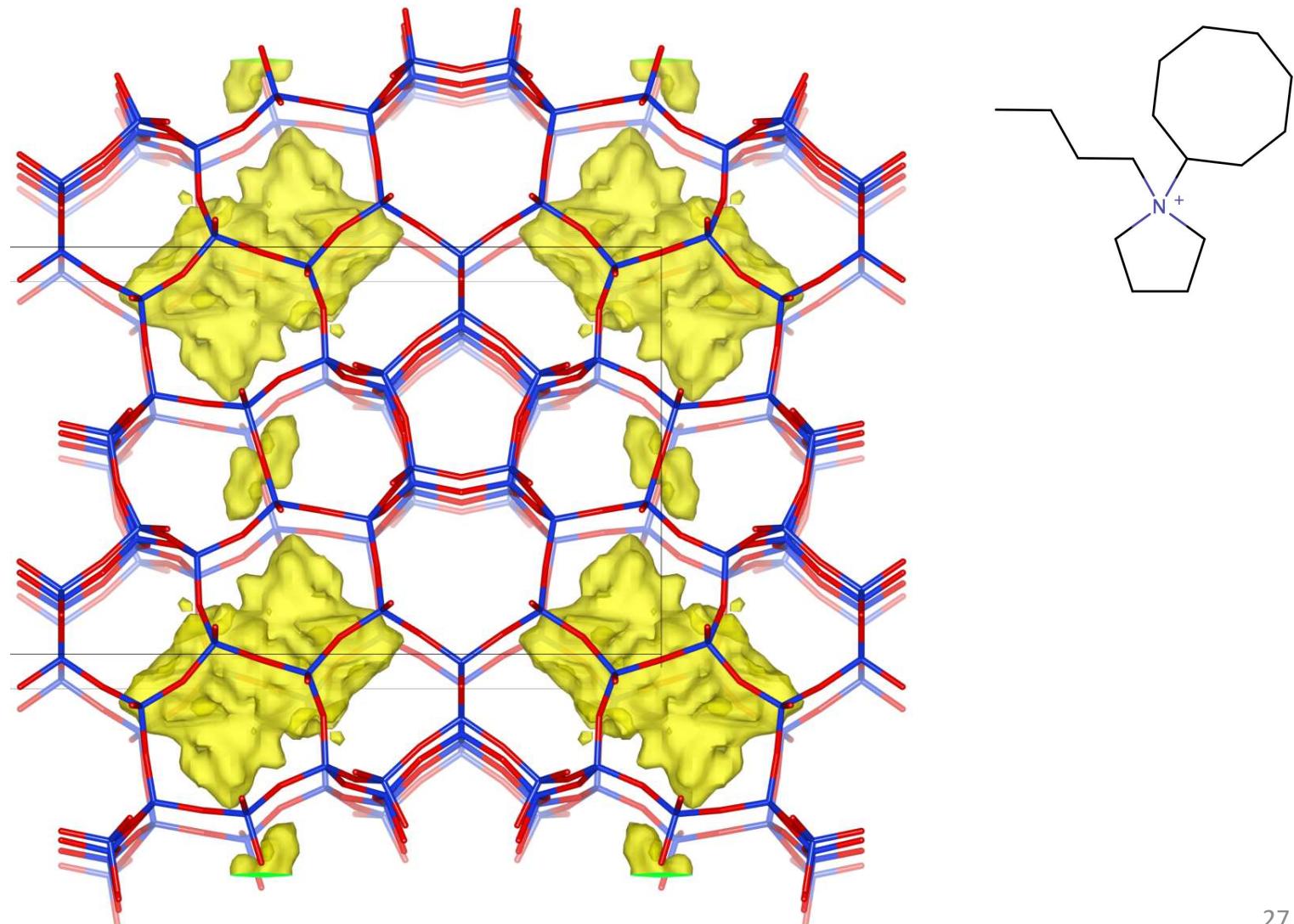
Rietveld refinement of SSZ-56



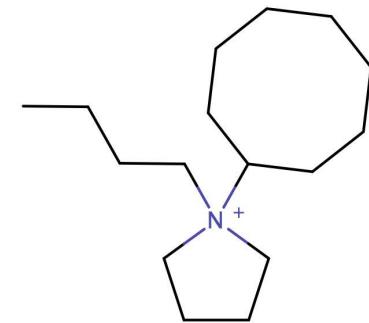
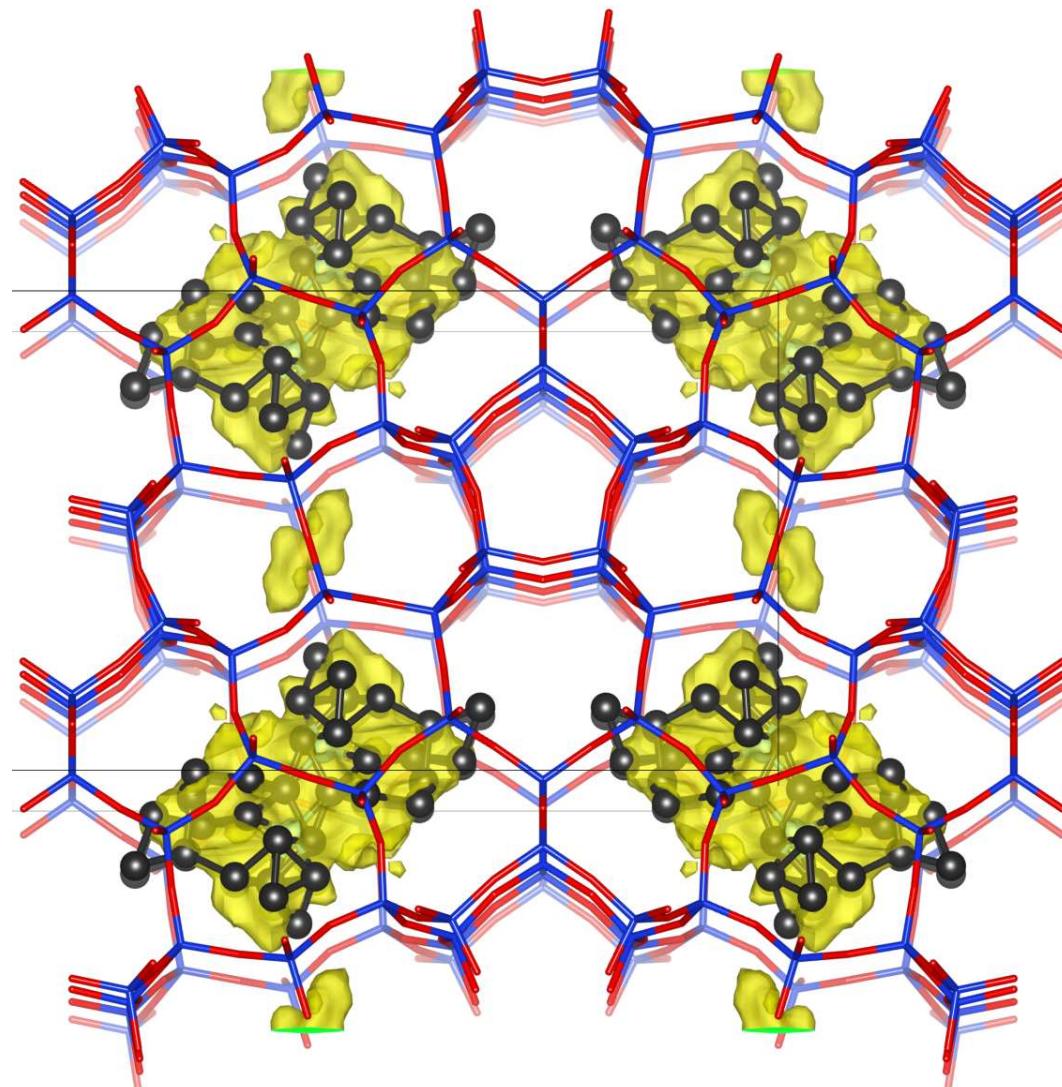
Comparison with molecular modeling



Difference map SSZ-58



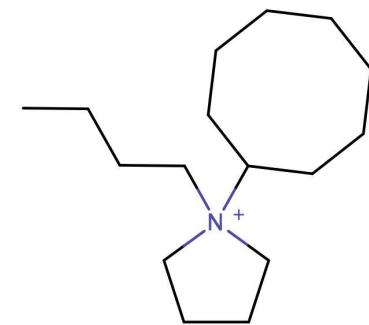
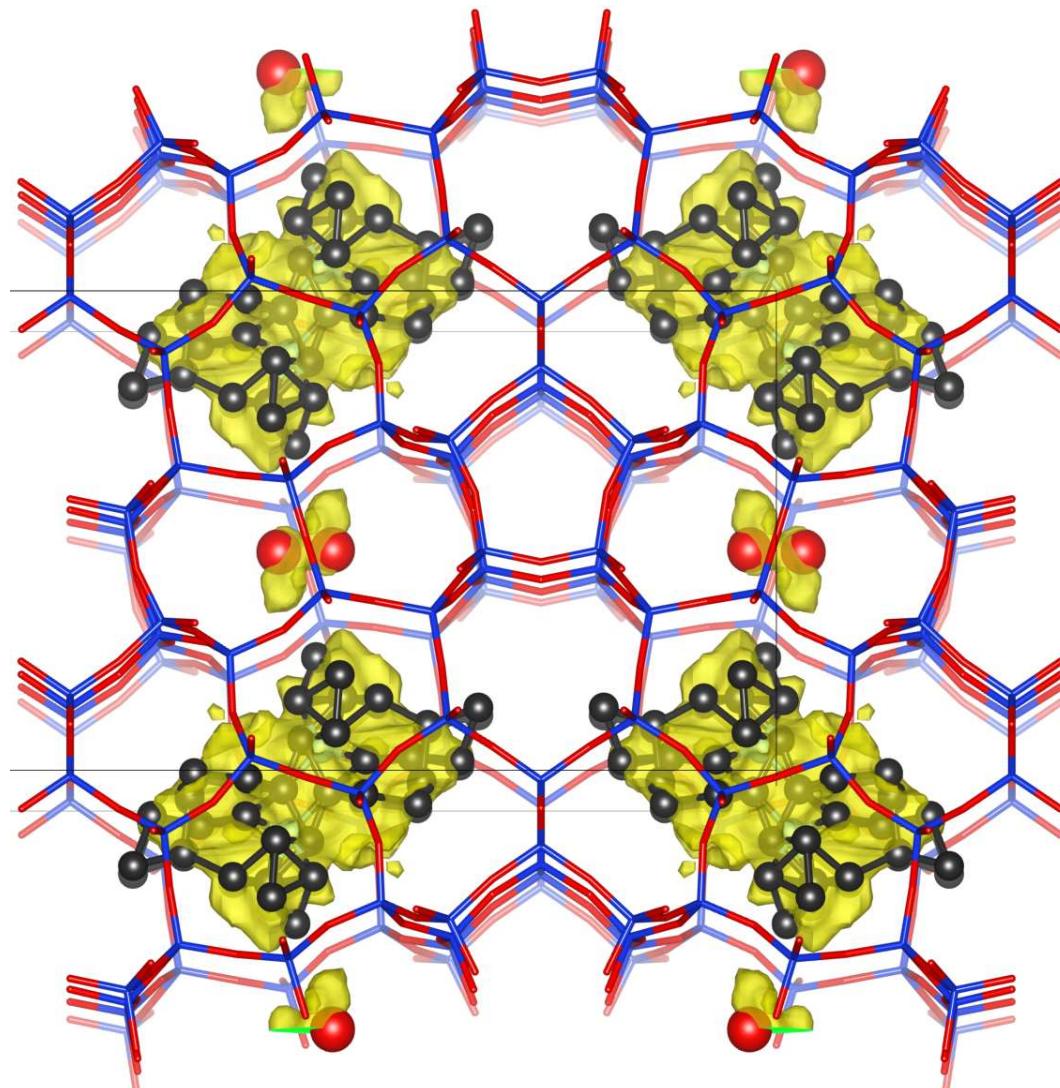
Difference map SSZ-58



Four-fold disorder

Residual electron
density

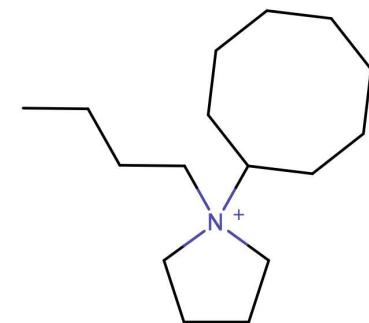
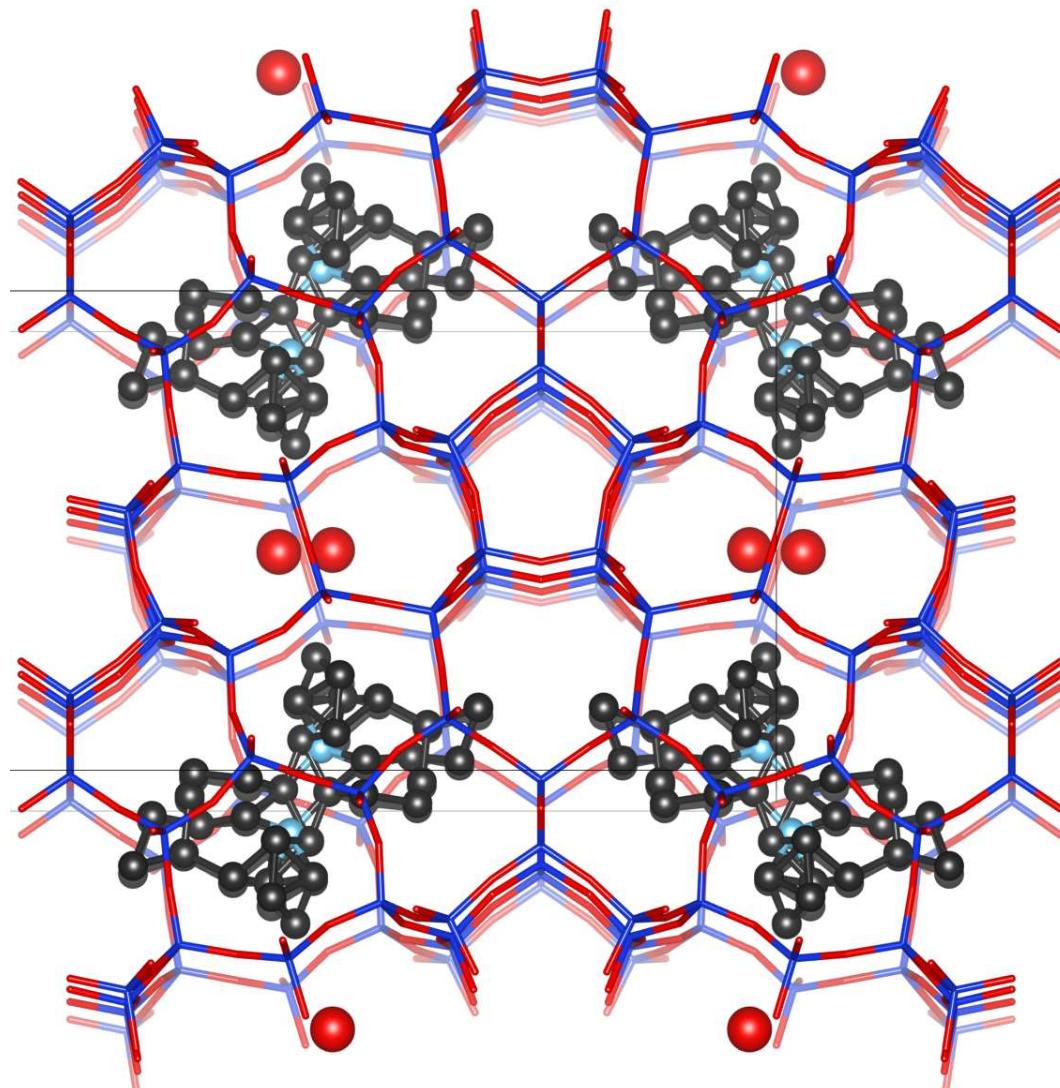
Difference map SSZ-58



Four-fold disorder

Add some water!

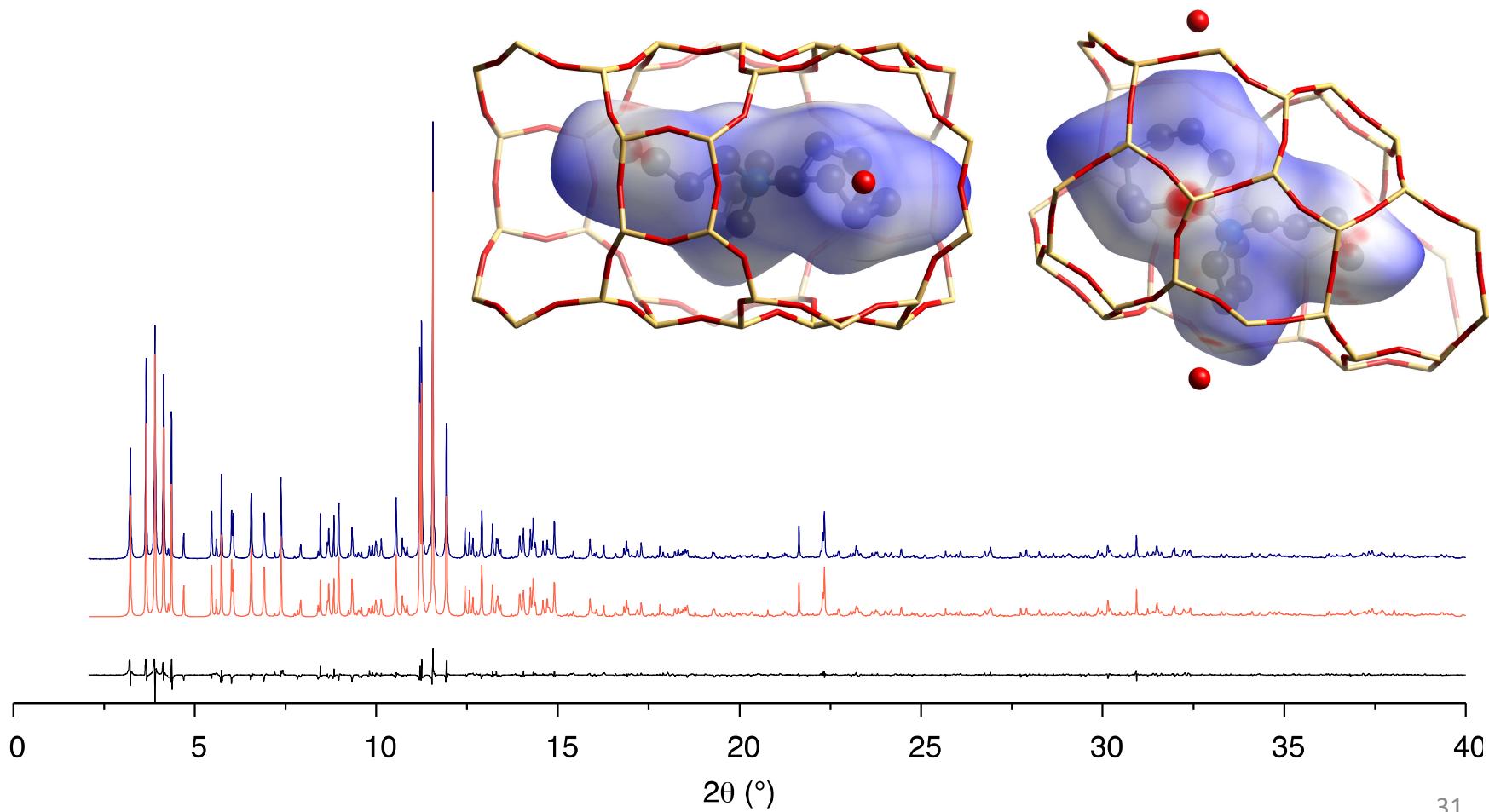
Difference map SSZ-58



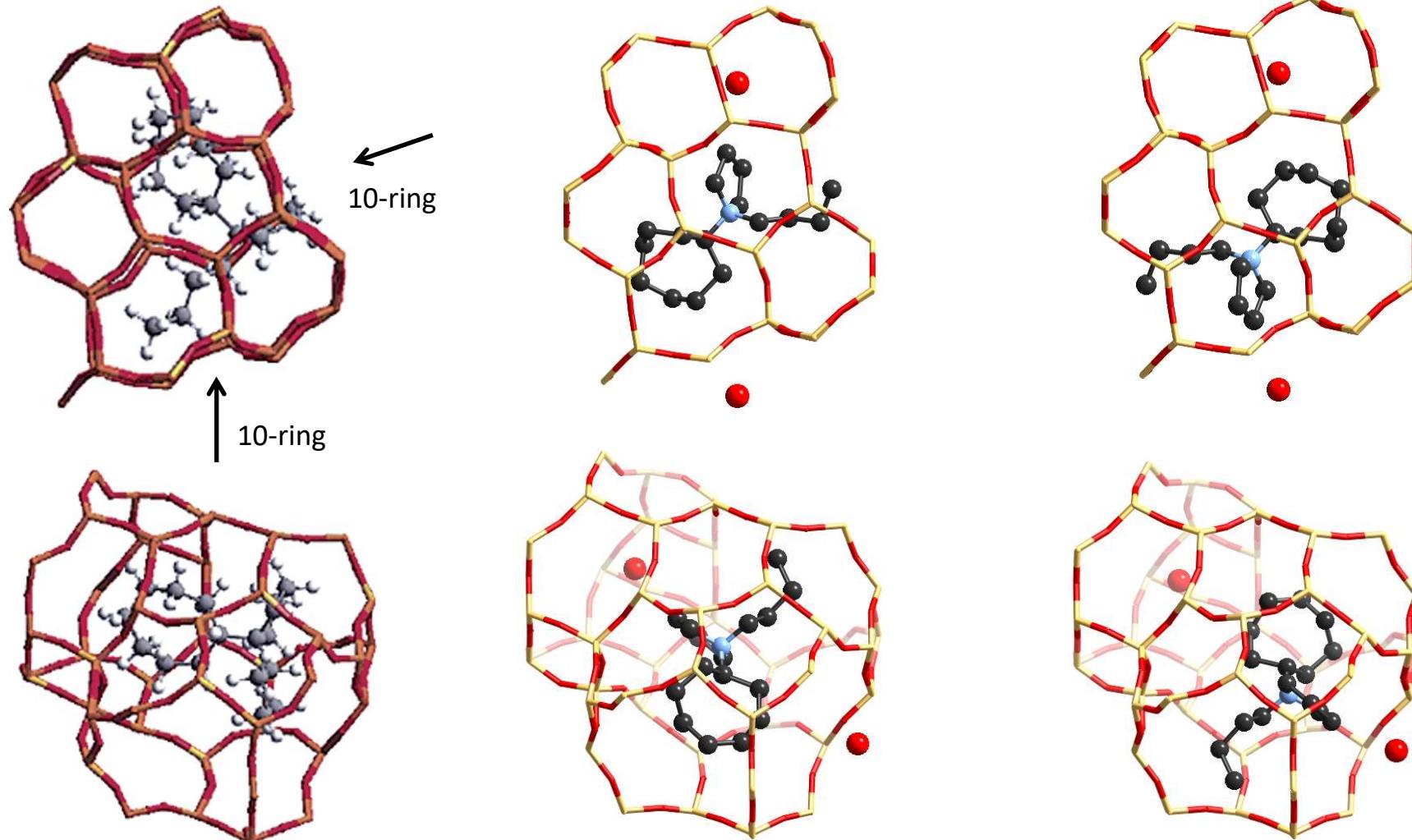
Four-fold disorder

Add some water!

Rietveld refinement of SSZ-58

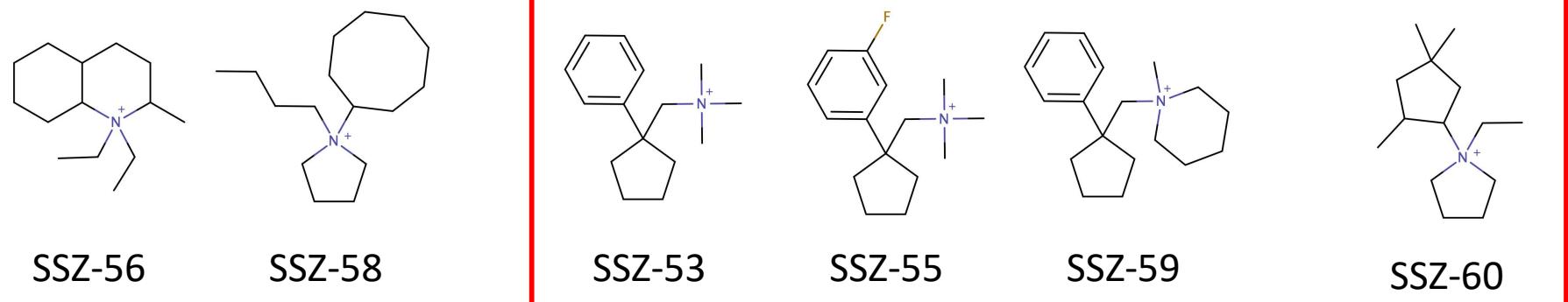


Comparison with molecular modeling

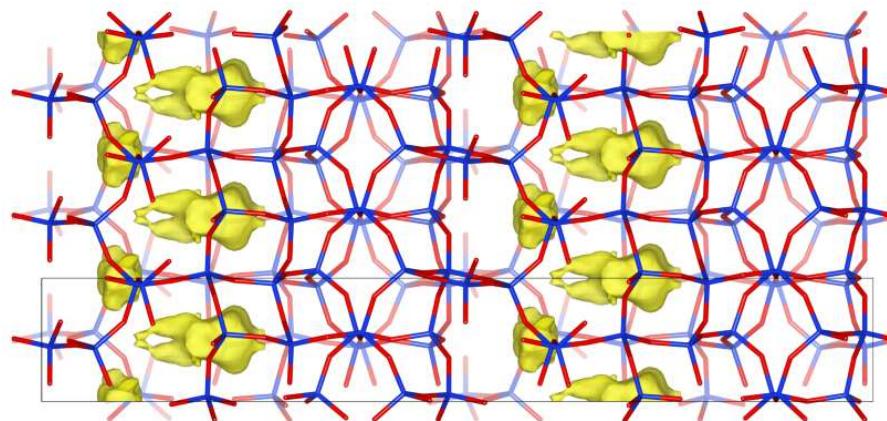
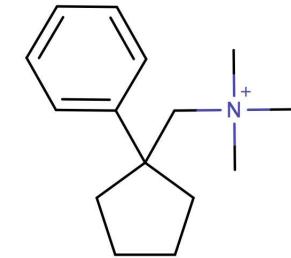
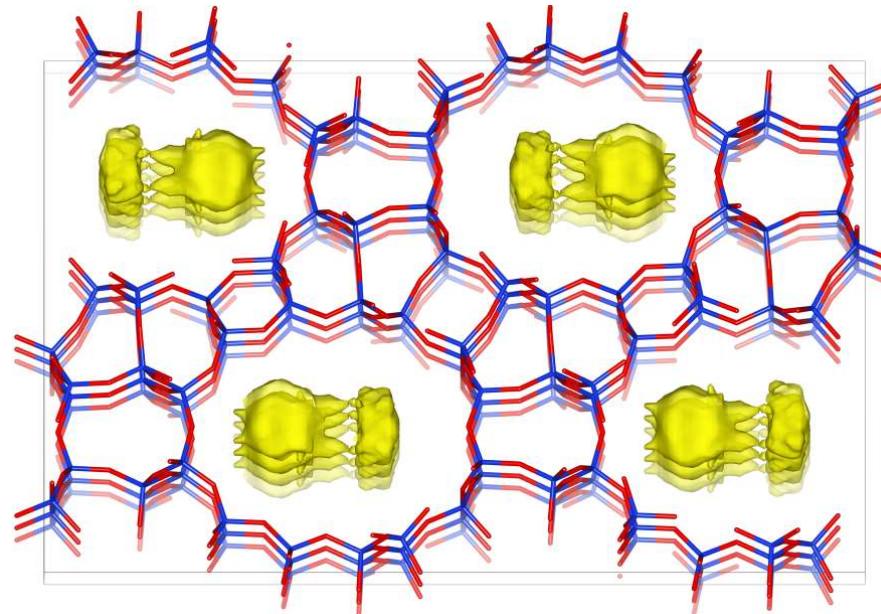


SSZ-53, SSZ-55, SSZ-59, SSZ-60

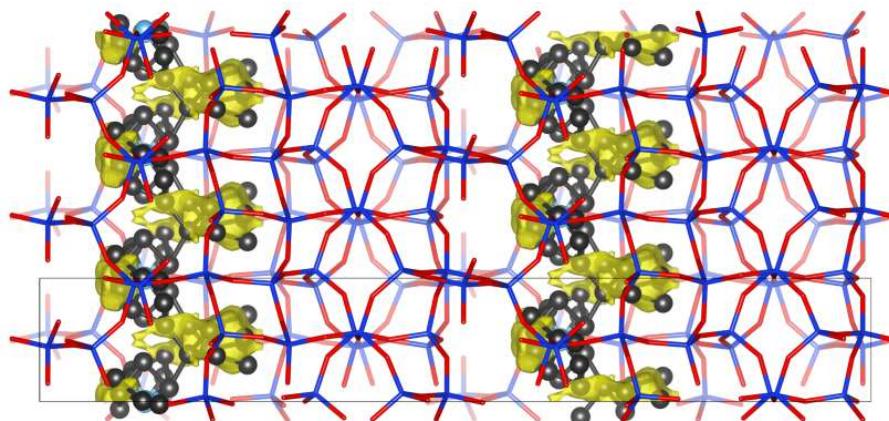
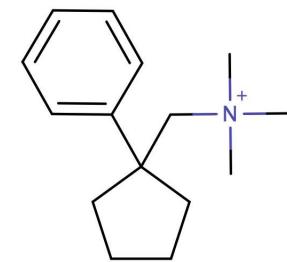
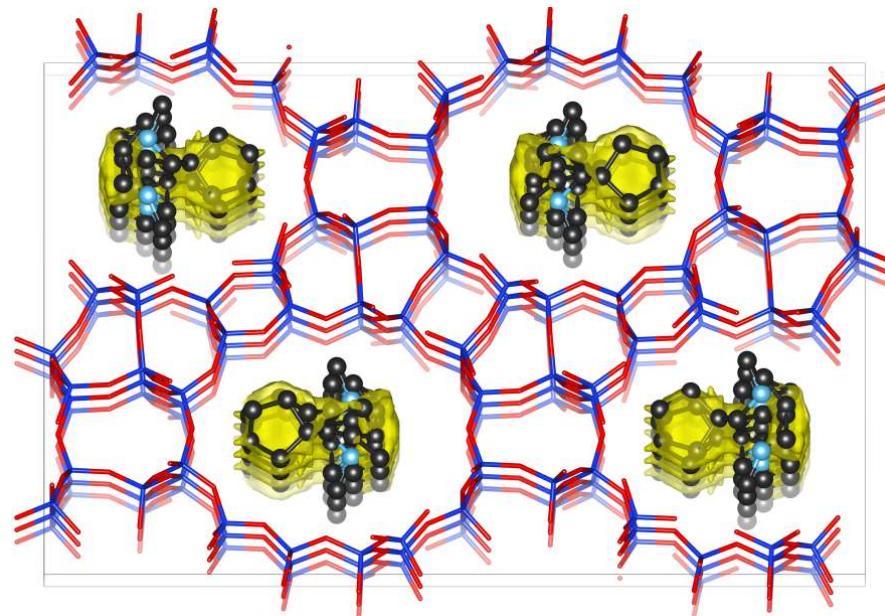
	ftc	<i>a</i>	<i>b</i>	<i>c</i>	α	β	γ	Spgr.
SSZ-53	SFH	5.02	33.74	21.17		90.5		<i>C</i> 2/c
SSZ-55	ATS	12.95	21.85	5.08				<i>Cmc</i> 2 ₁
SSZ-59	SFN	5.02	12.74	14.72	103.4	90.5	100.9	<i>P</i> -1
SSZ-60	SSY	21.95	13.70	5.01				<i>P</i> 2 ₁



Difference map SSZ-53



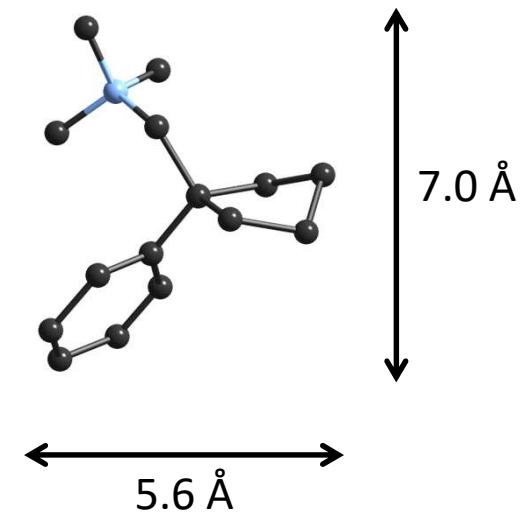
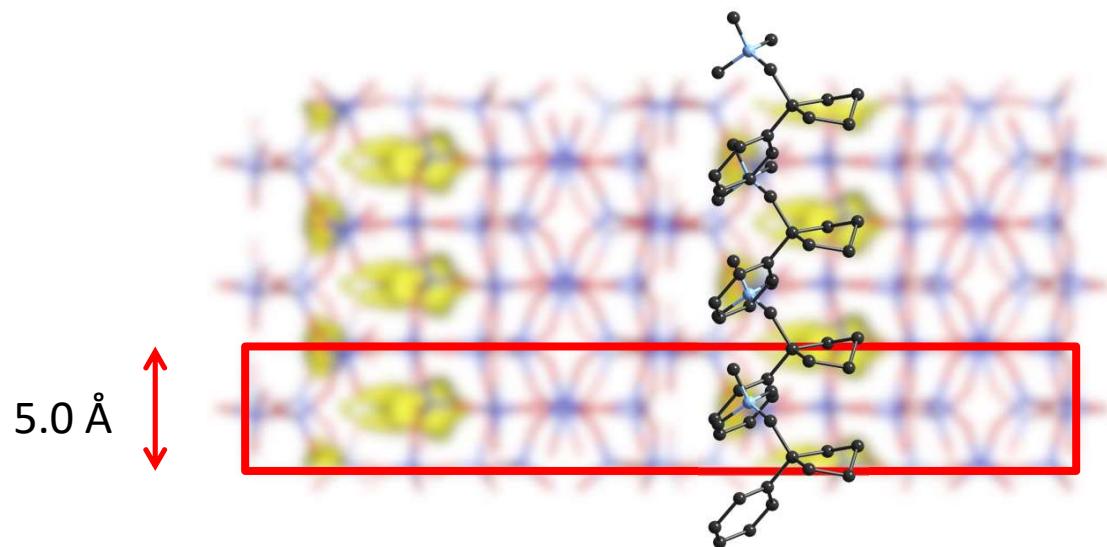
Difference map SSZ-53



SSZ-53

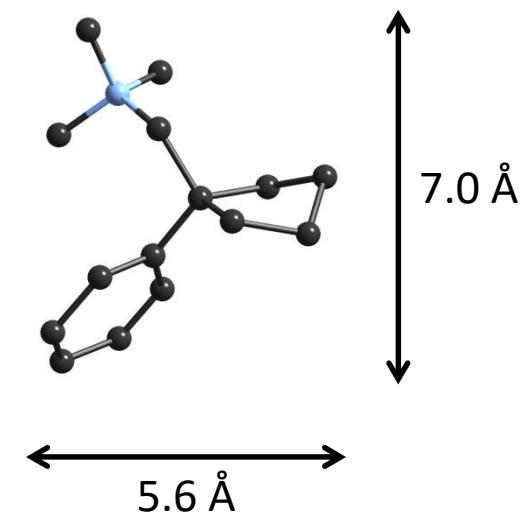
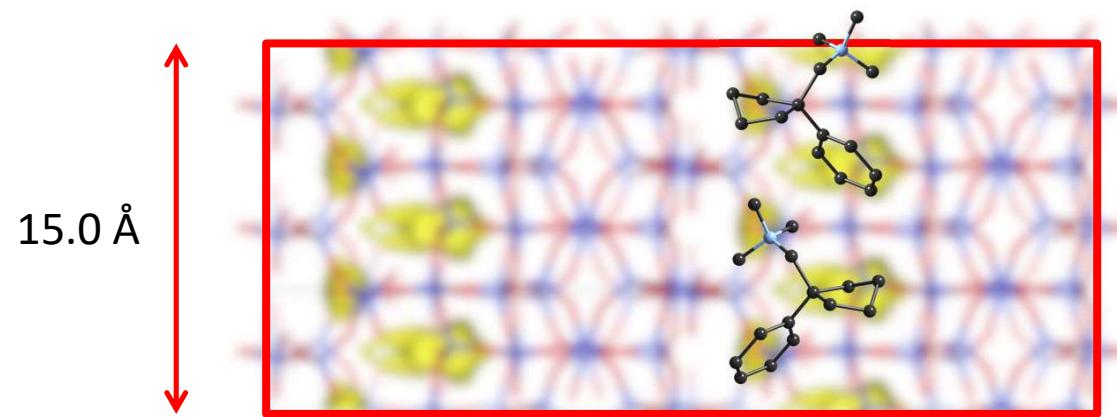
Electron density:

- 0.67 SDAs per channel per unit cell
- or
- 2 SDAs per channel per 3 unit cells

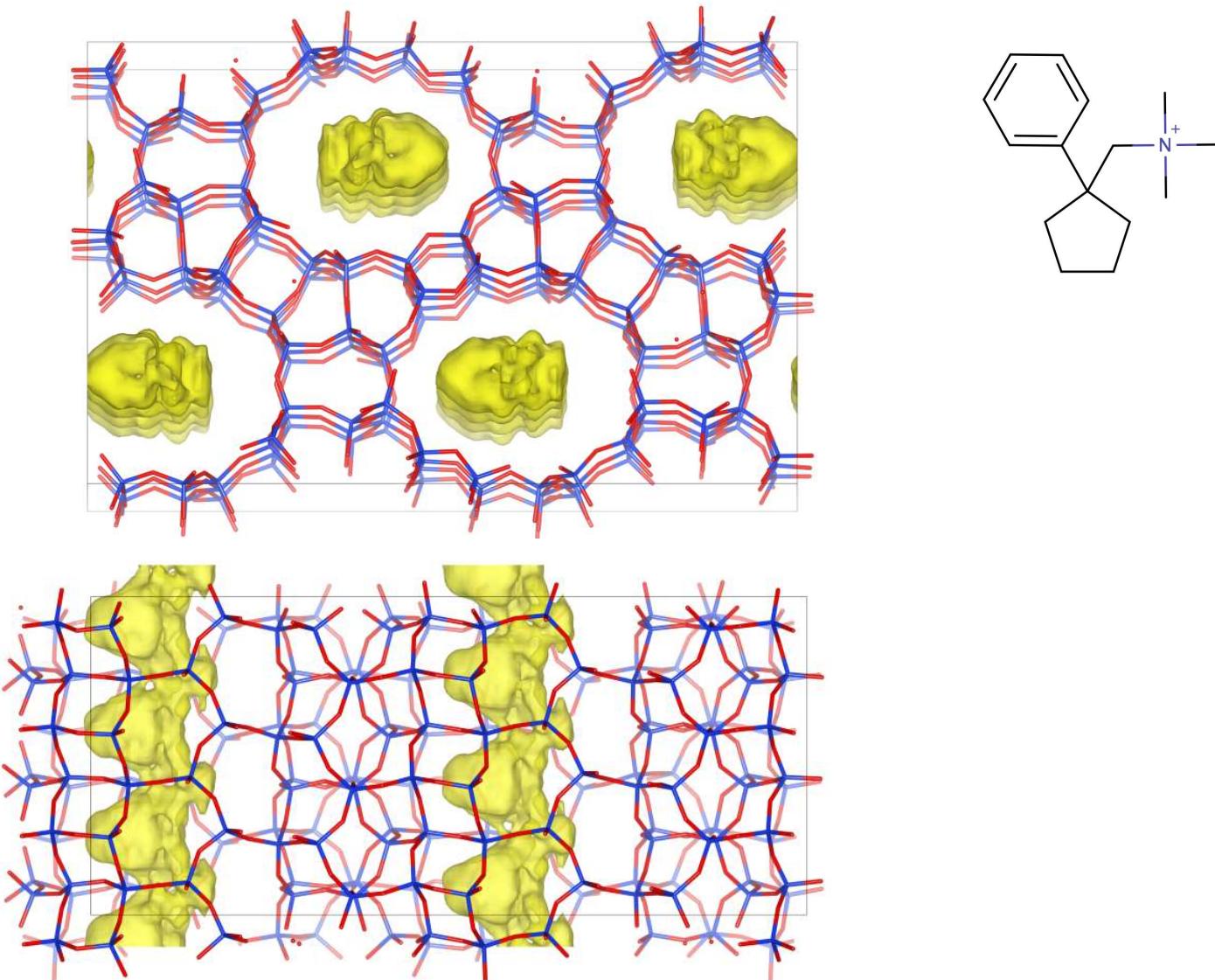


SSZ-53

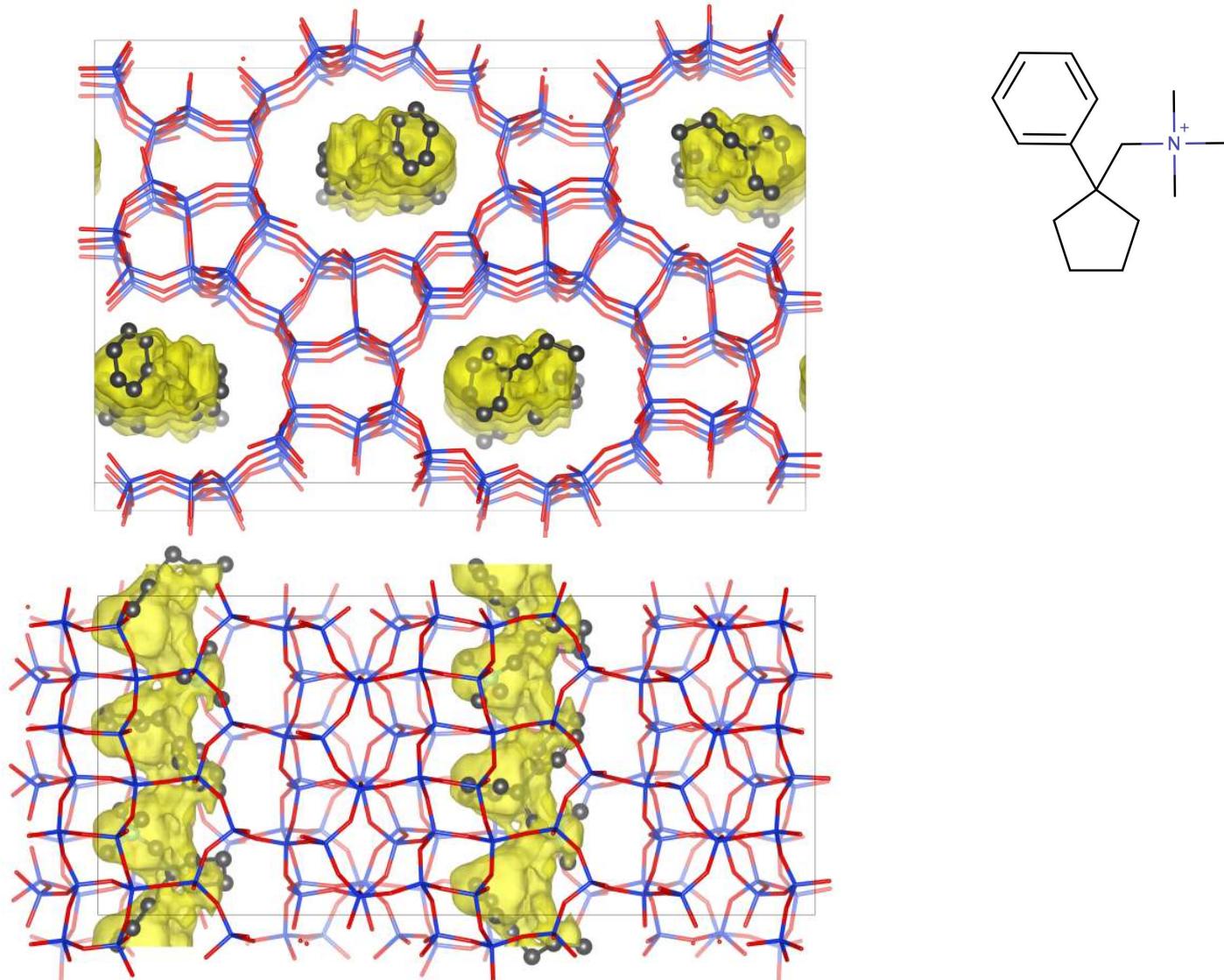
Expand unit cell x3 along channel direction



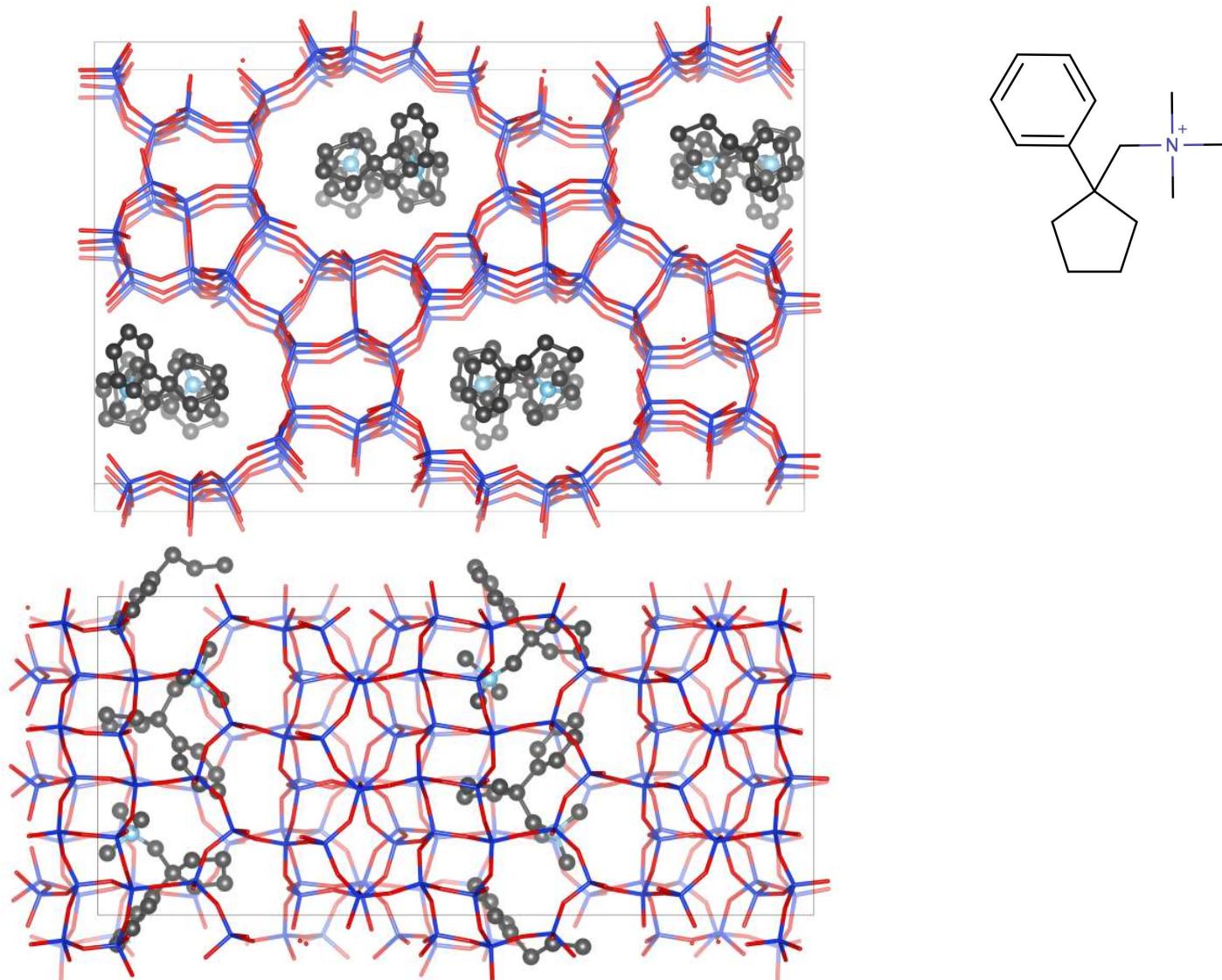
Difference map SSZ-53



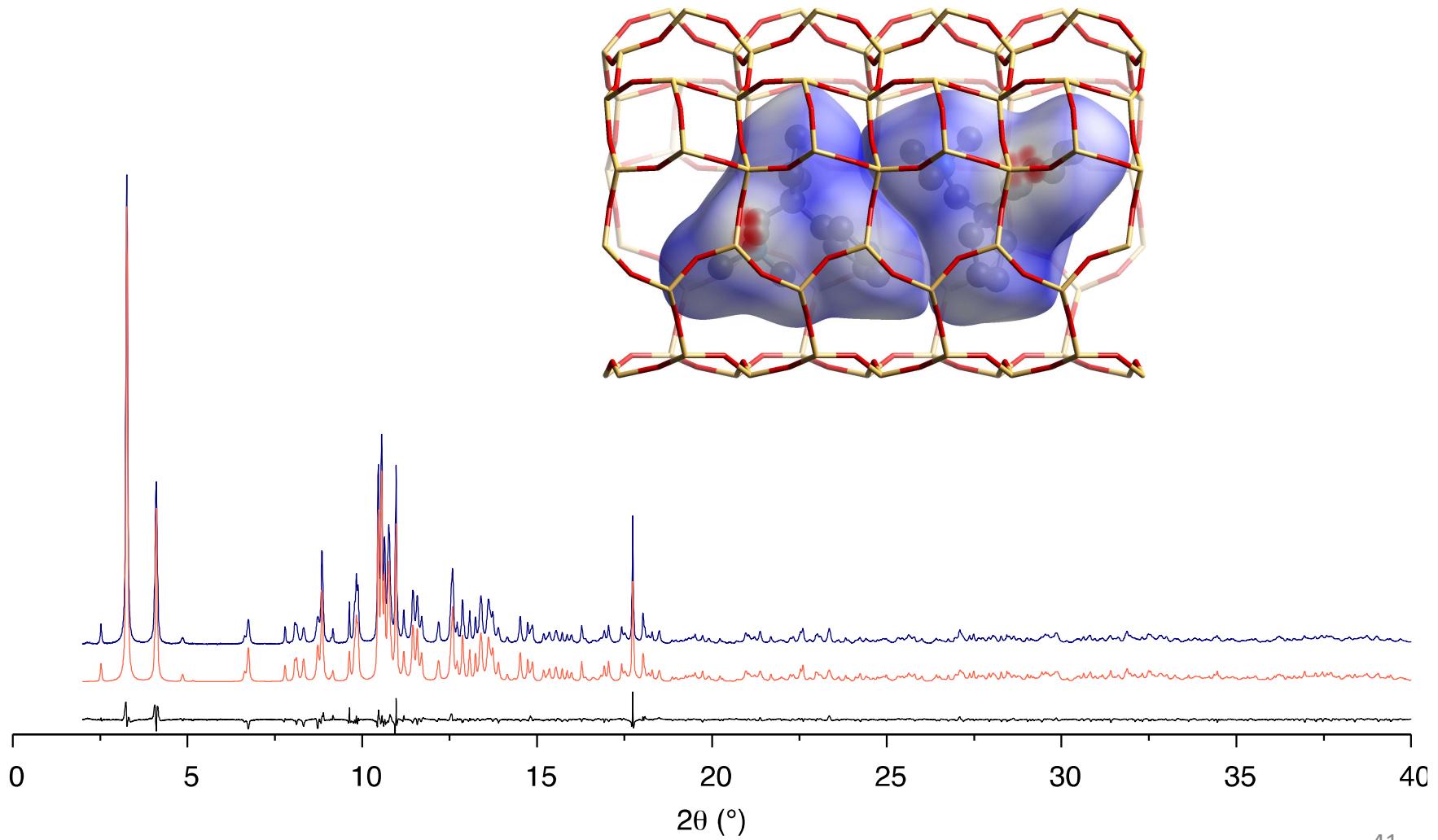
Difference map SSZ-53



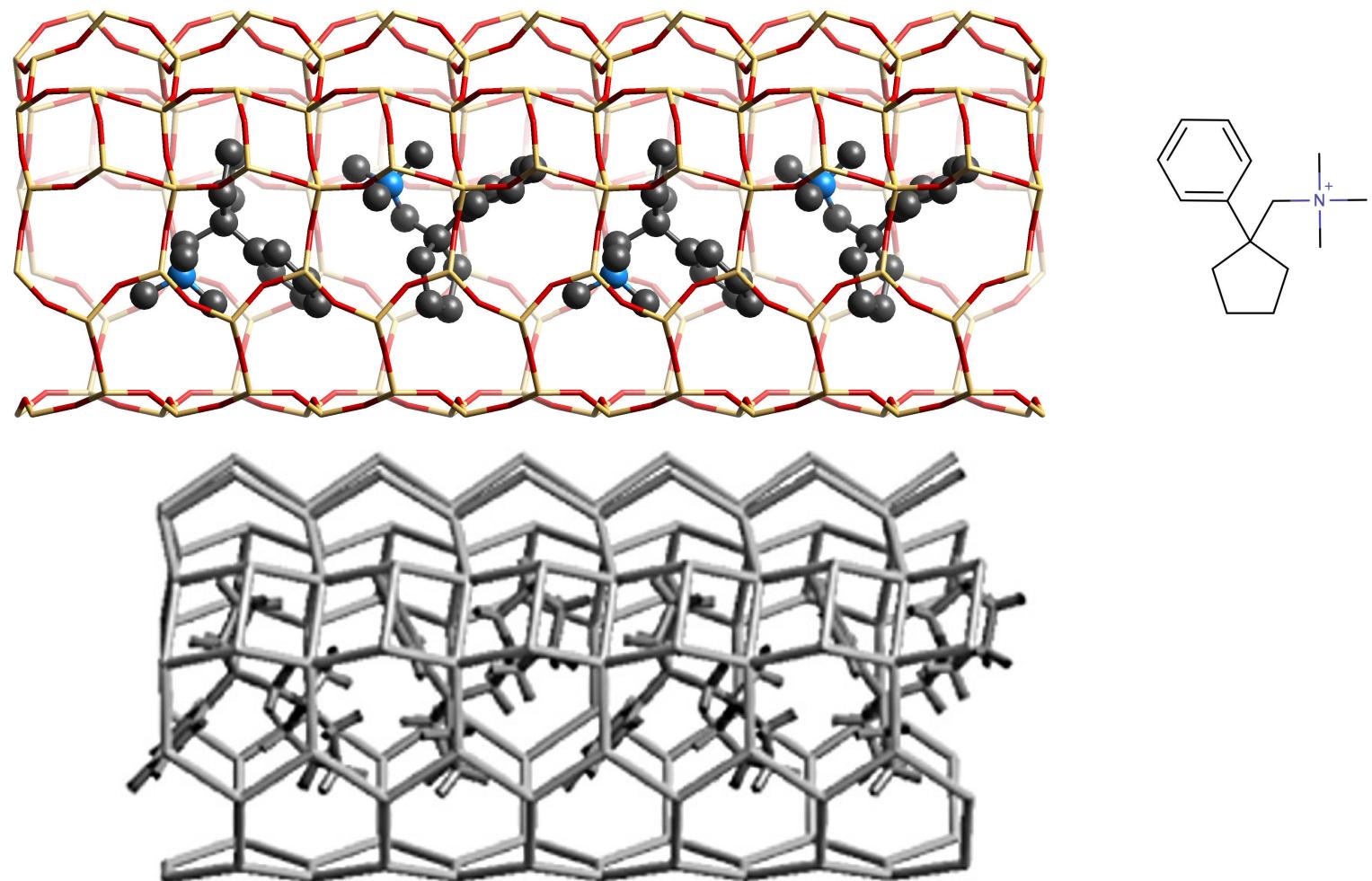
Difference map SSZ-53



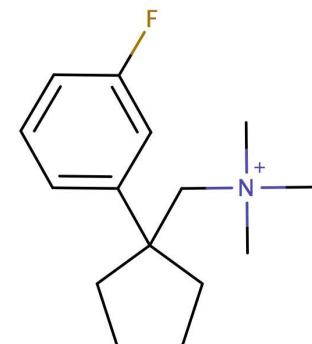
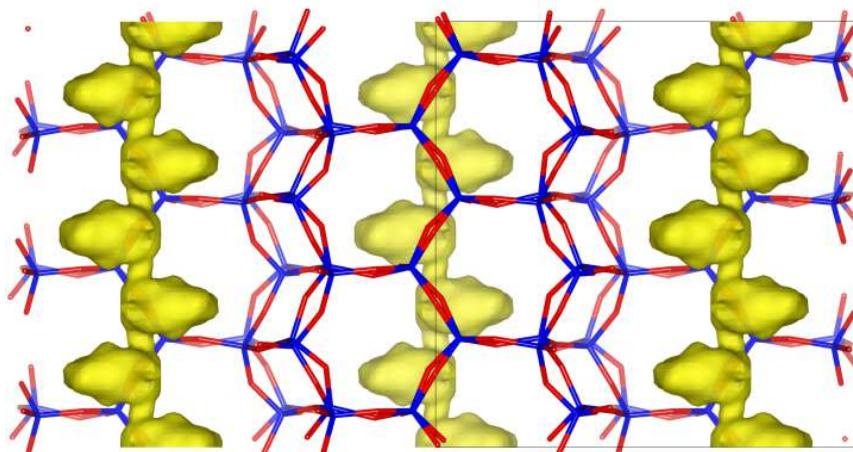
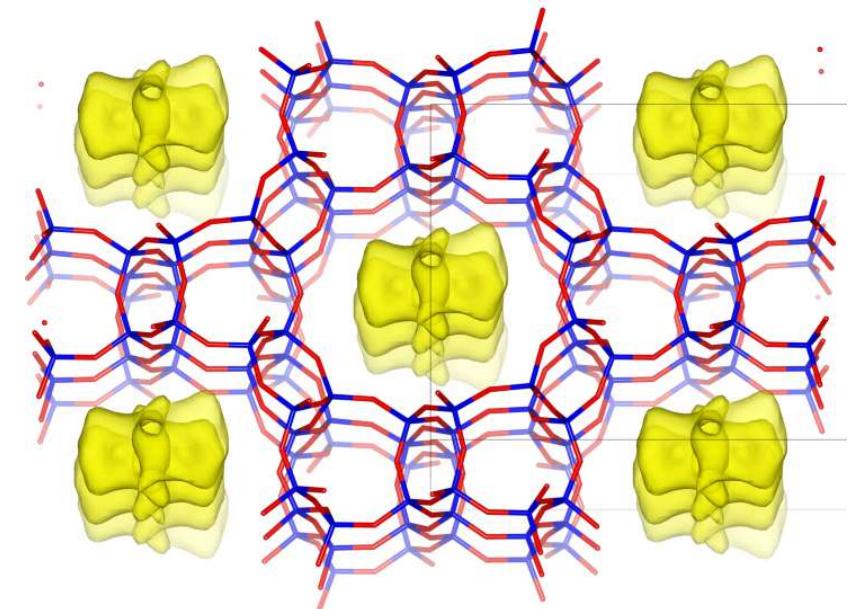
Rietveld refinement of SSZ-53



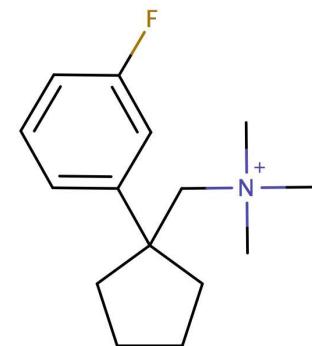
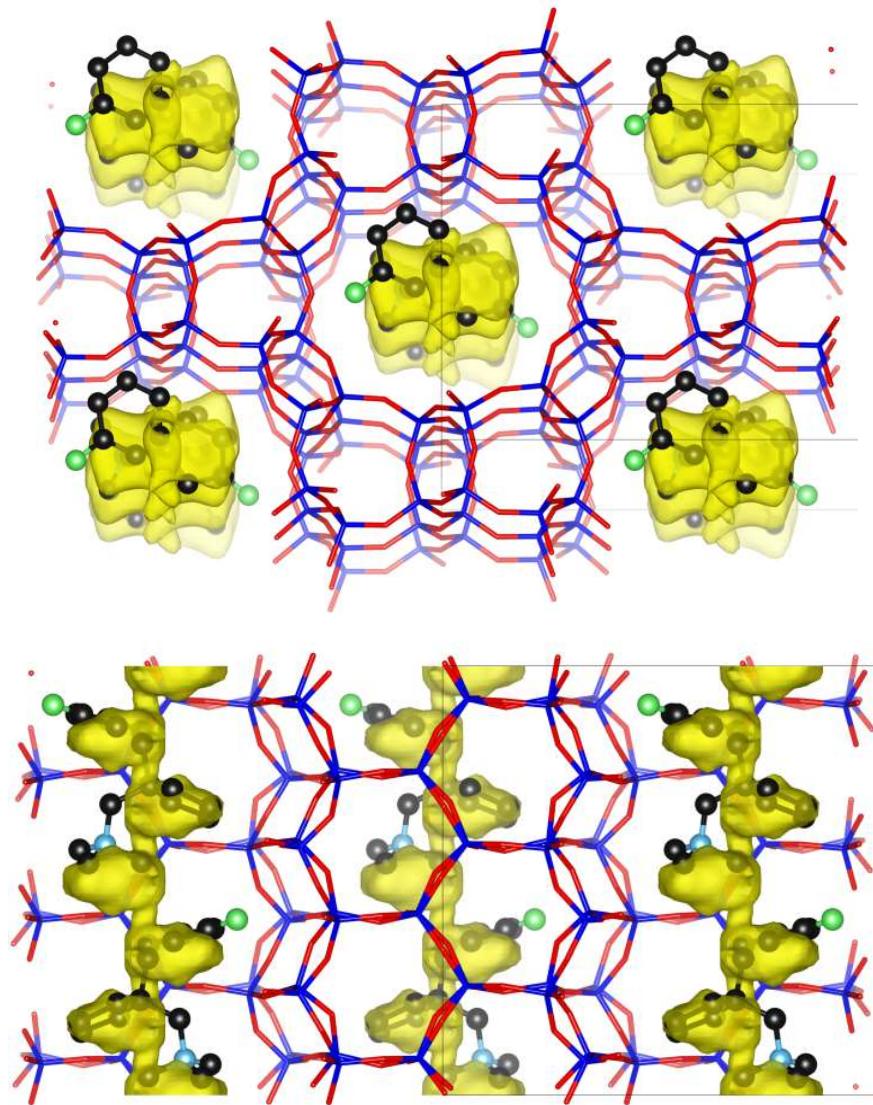
Comparison with molecular modeling



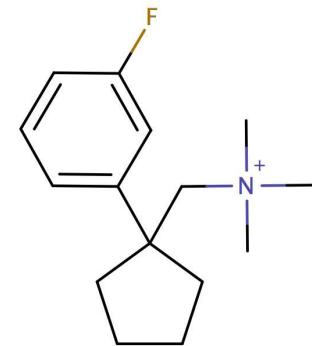
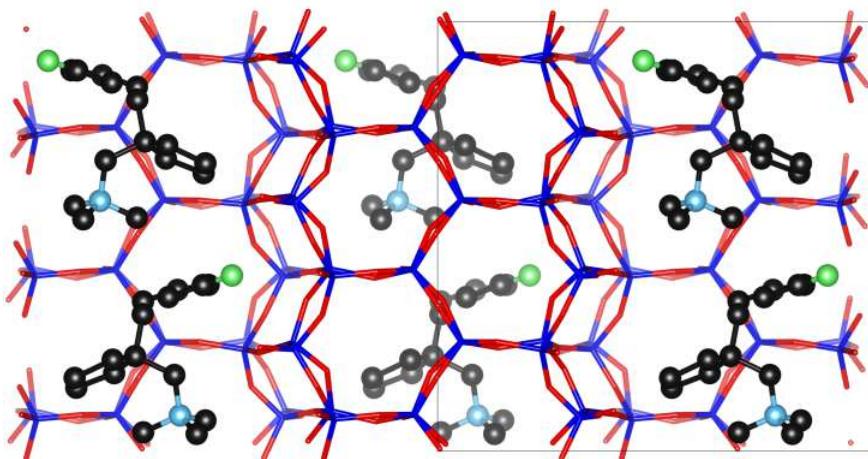
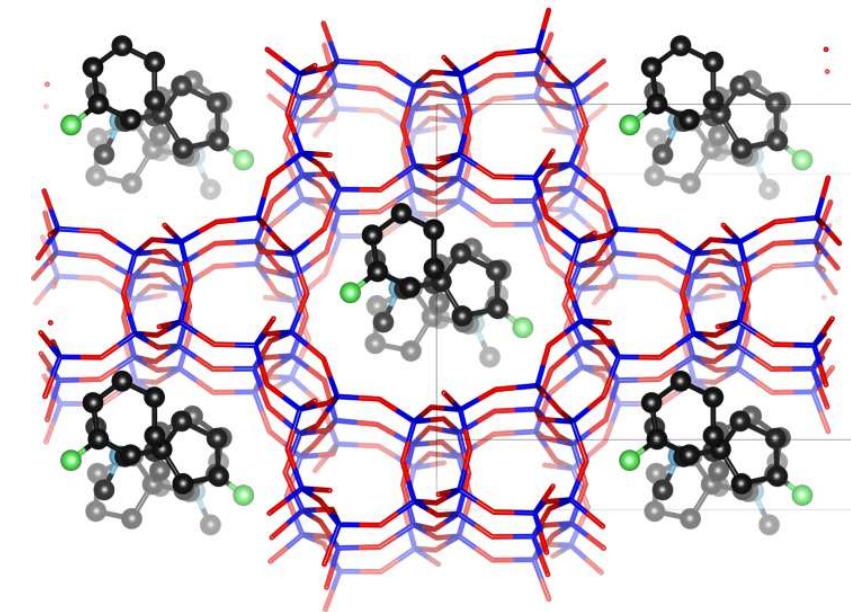
Difference map SSZ-55



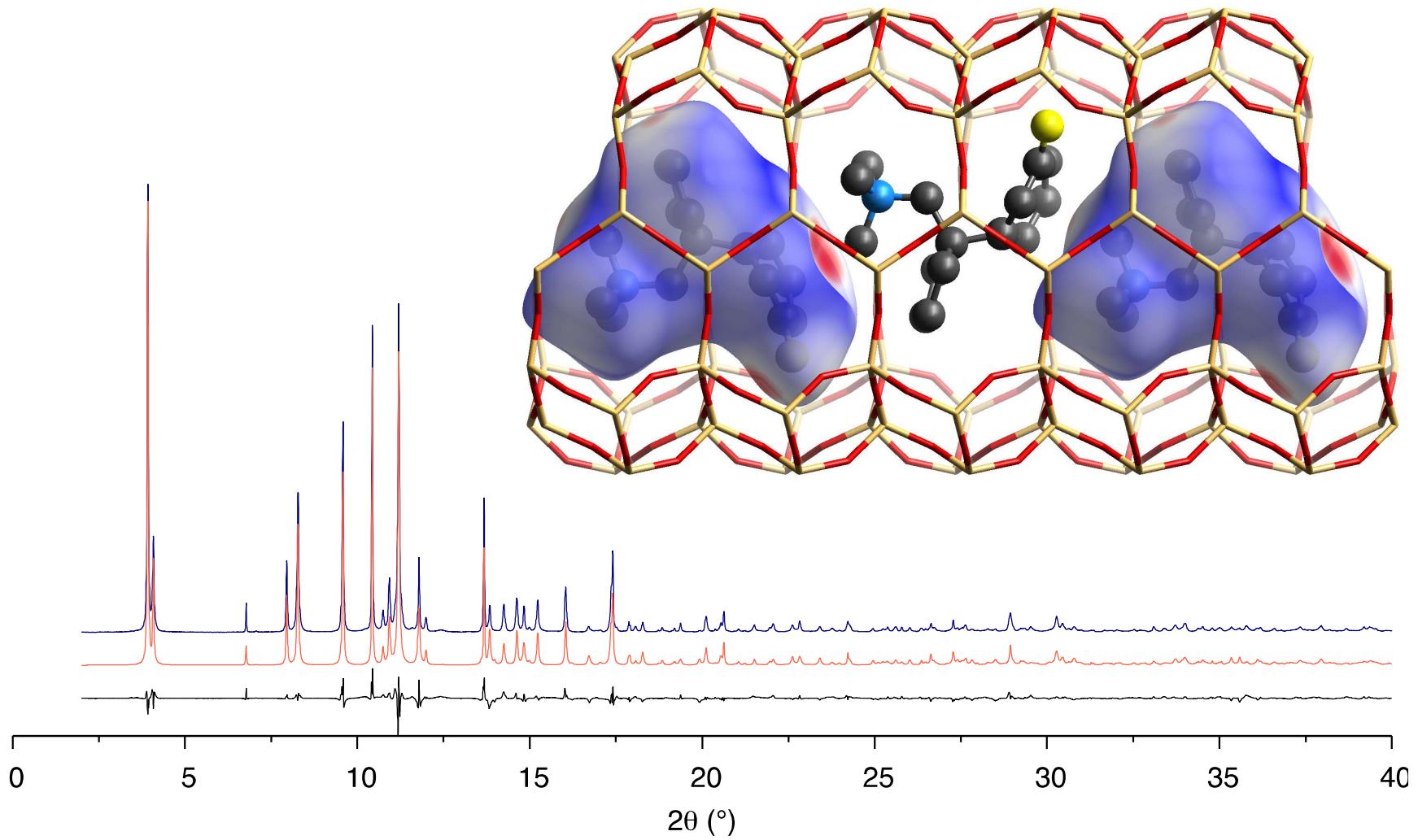
Difference map SSZ-55



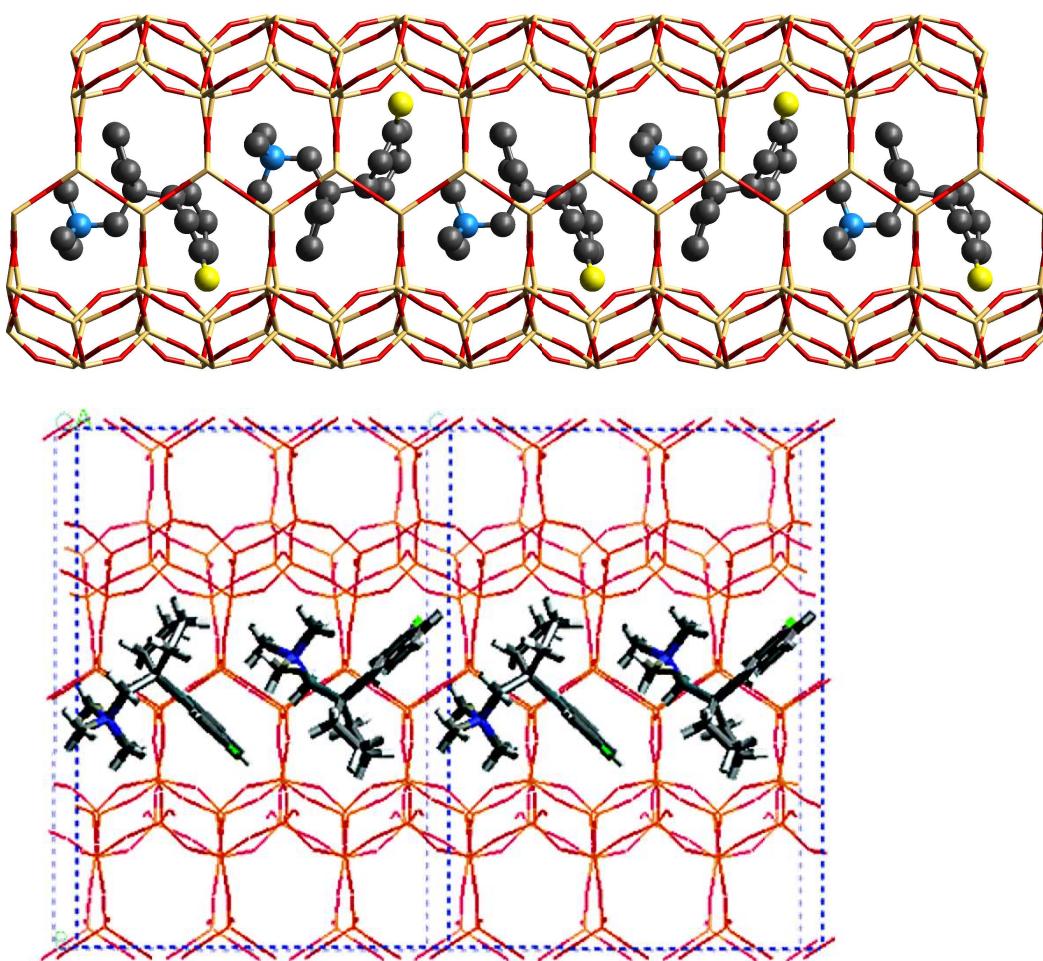
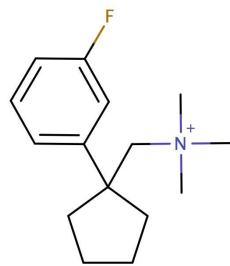
Difference map SSZ-55



Rietveld refinement of SSZ-55

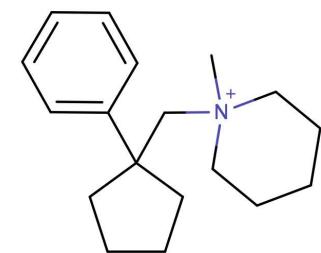
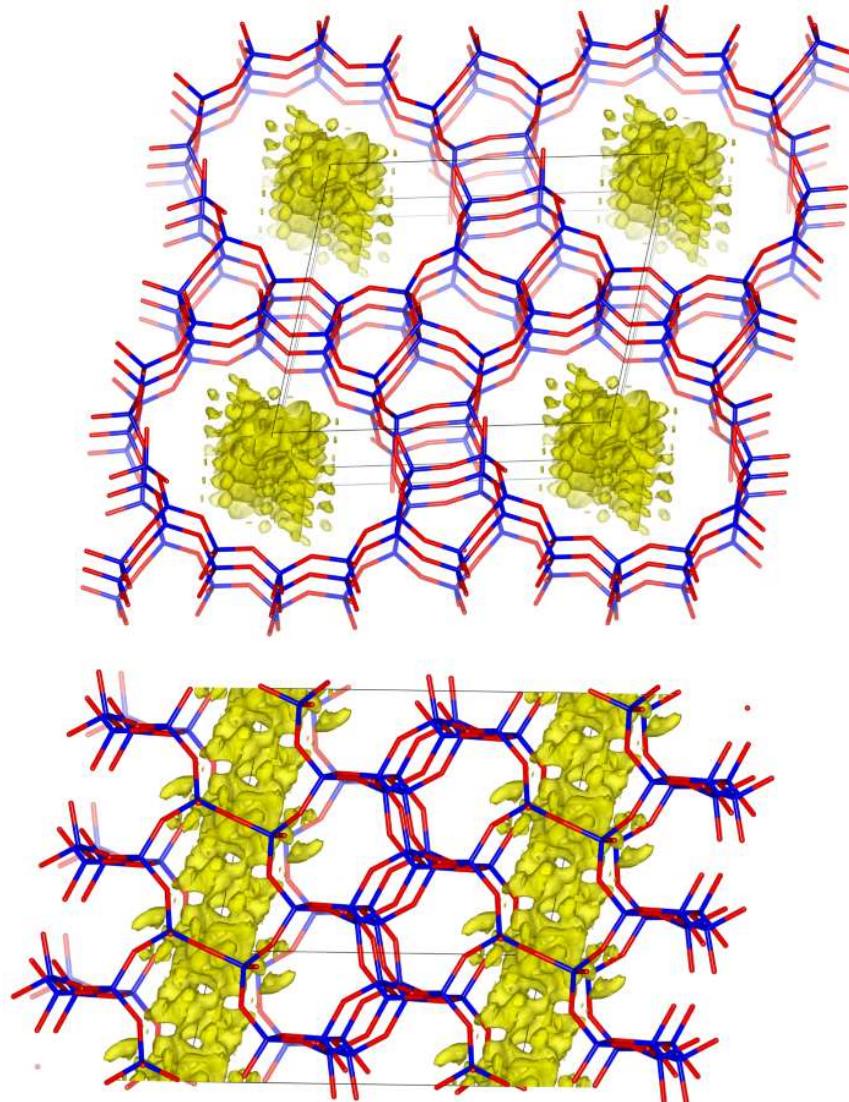


Comparison with molecular modeling

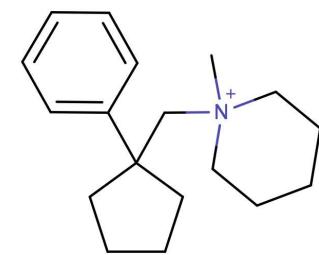
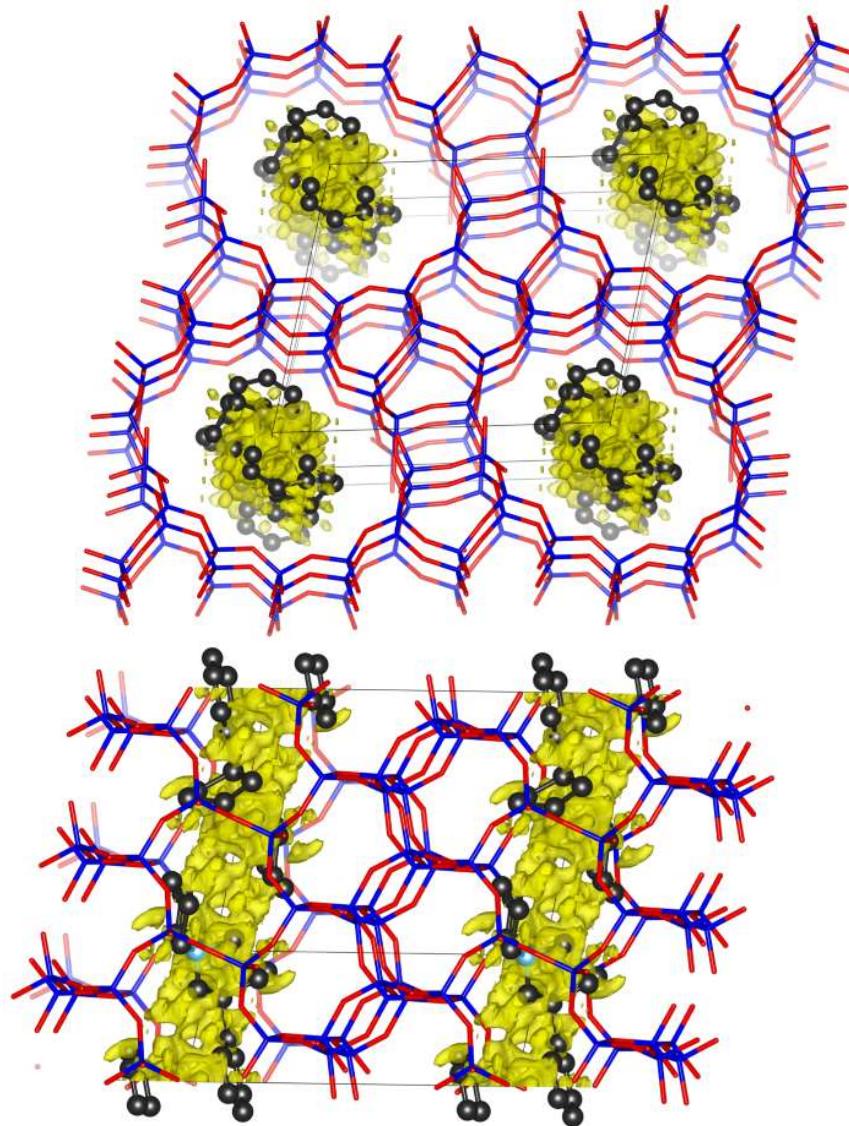


Burton *et al.*, J. Phys. Chem. B, 110, 5273, 2006

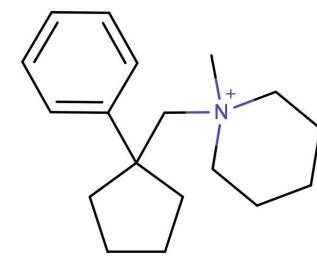
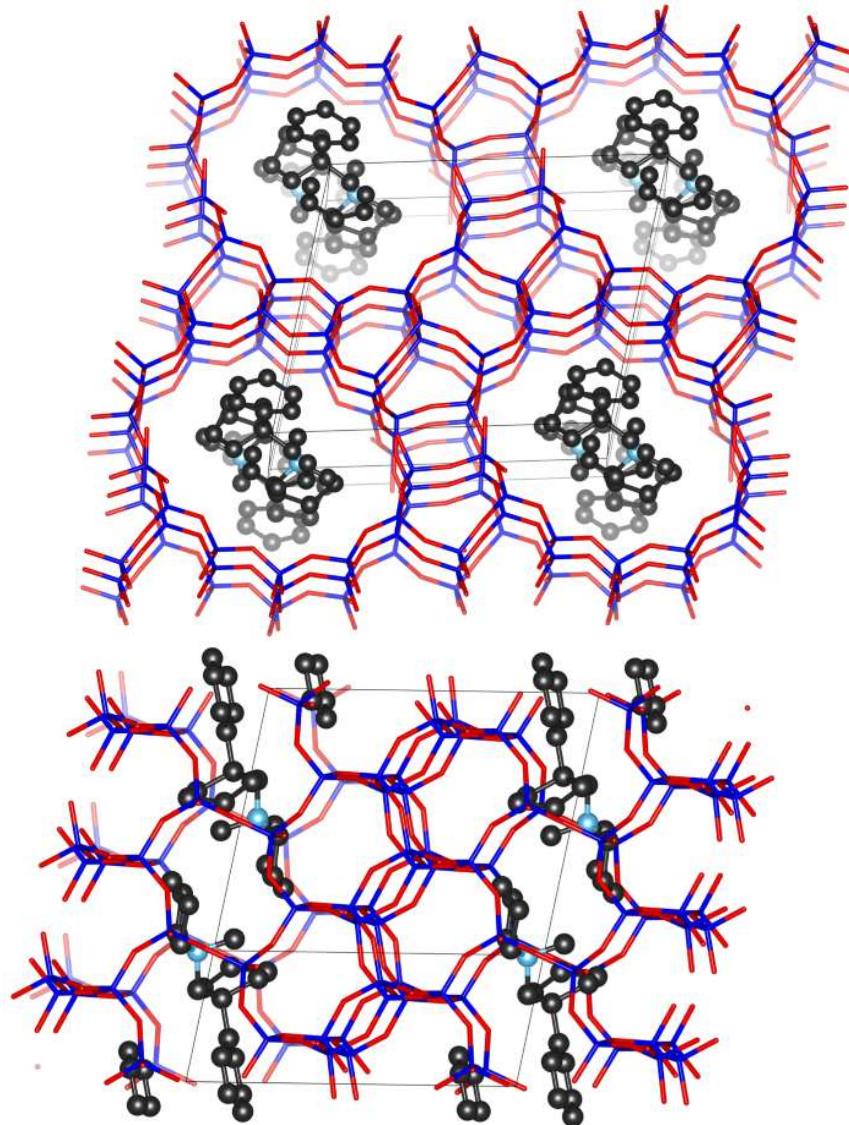
Difference map SSZ-59



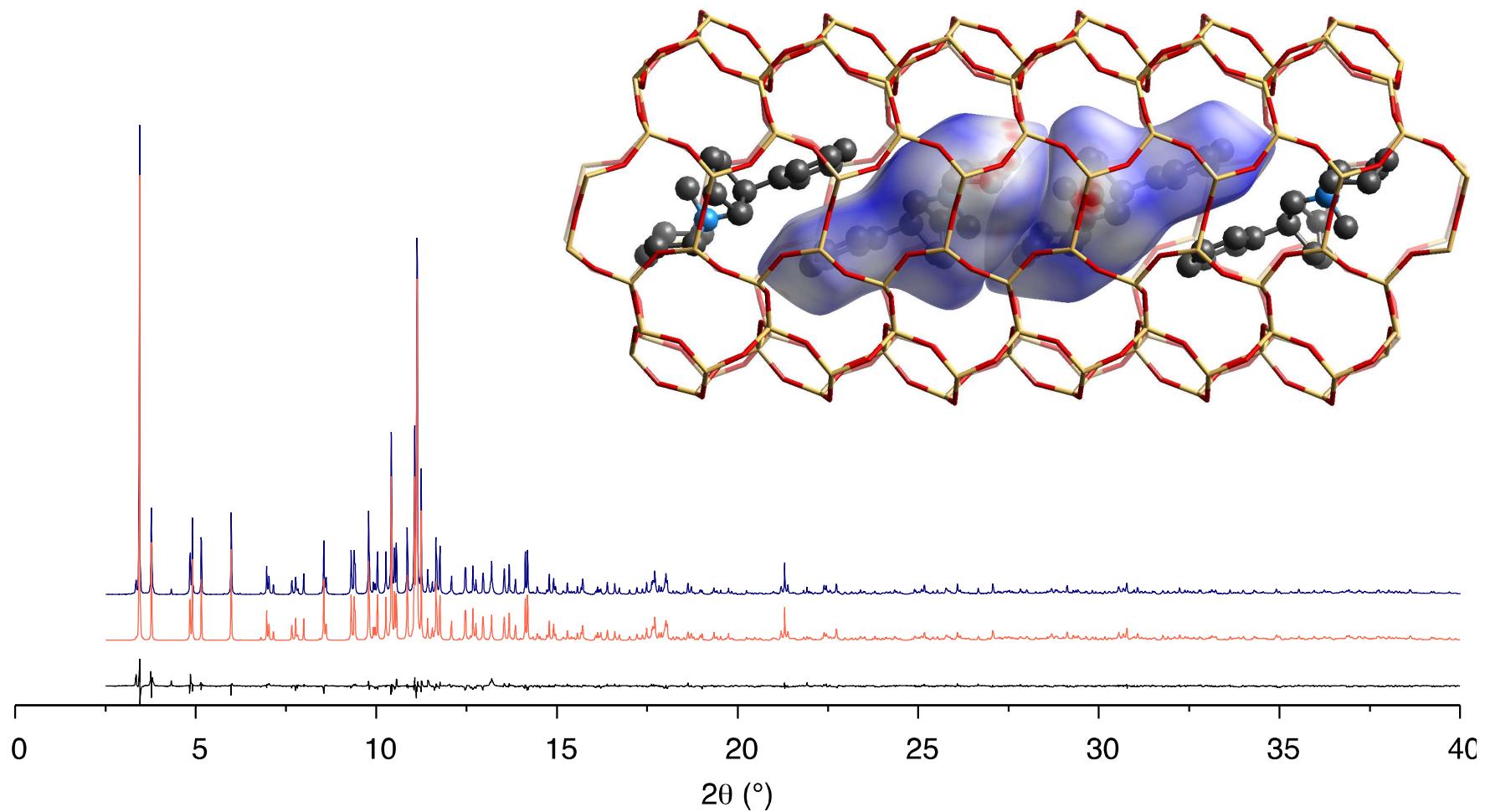
Difference map SSZ-59



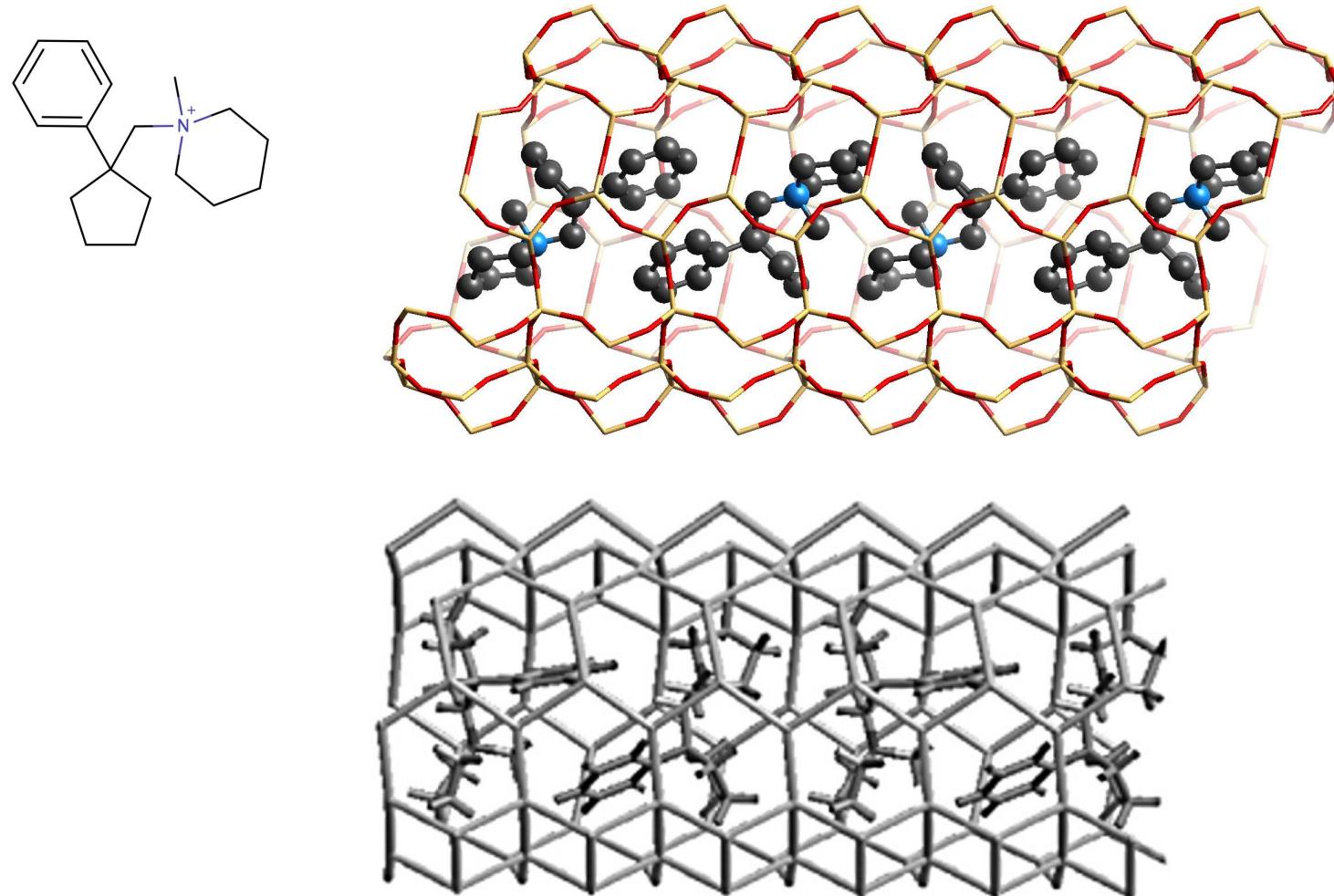
Difference map SSZ-59



Rietveld refinement of SSZ-59



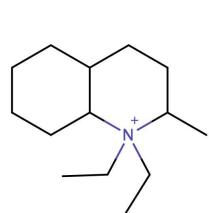
Comparison with molecular modeling



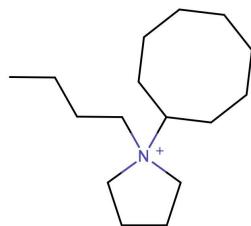
Burton *et al.*, Chem.-Eur. J., 9, 5737, 2003

SSZ-60

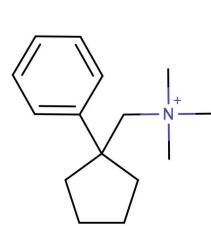
	ftc	<i>a</i>	<i>b</i>	<i>c</i>	α	β	γ	Spgr.
SSZ-60	SSY	21.95	13.70	5.01				<i>P</i> 2 ₁



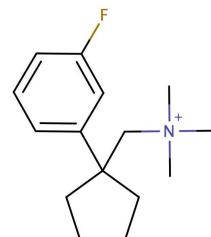
SSZ-56



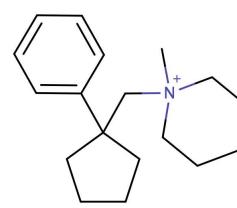
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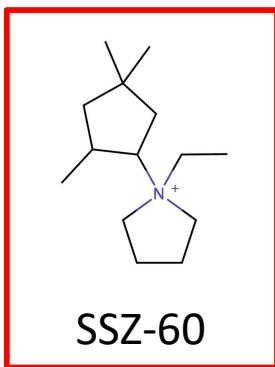
SSZ-53



SSZ-55

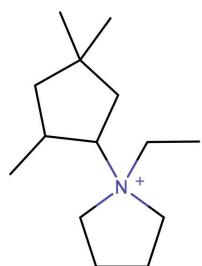
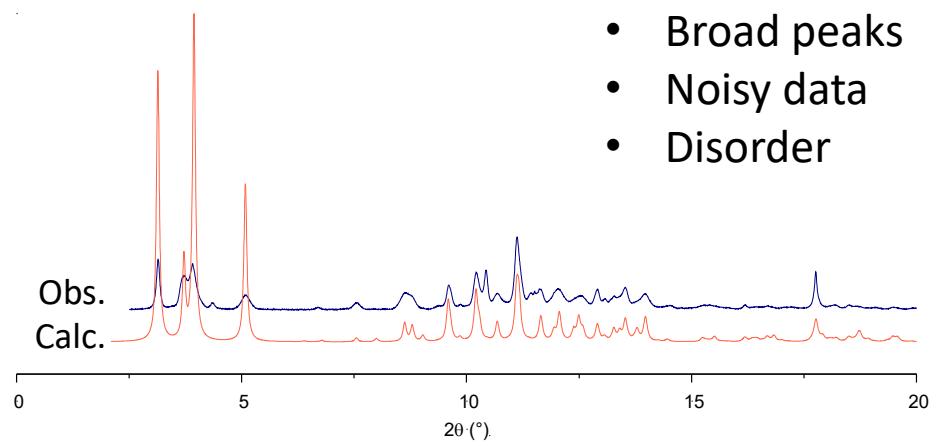
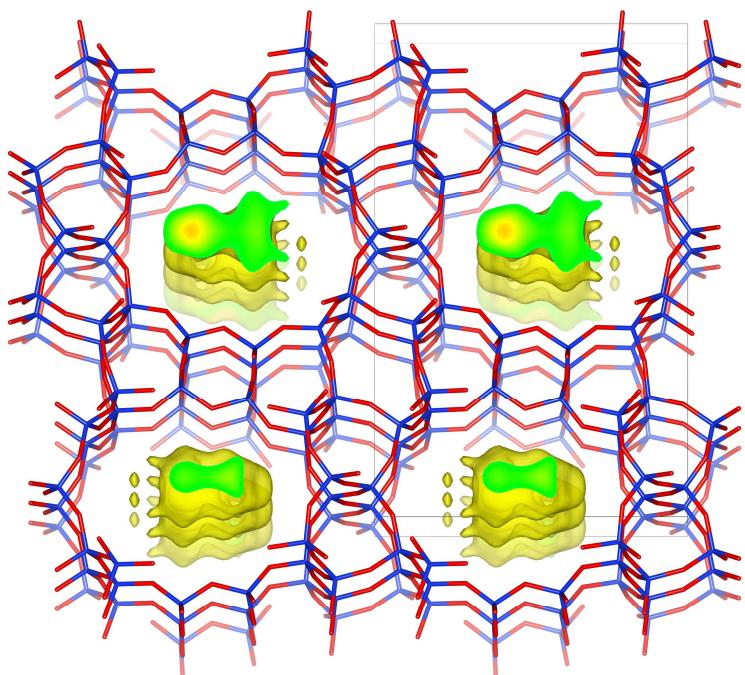


SSZ-59

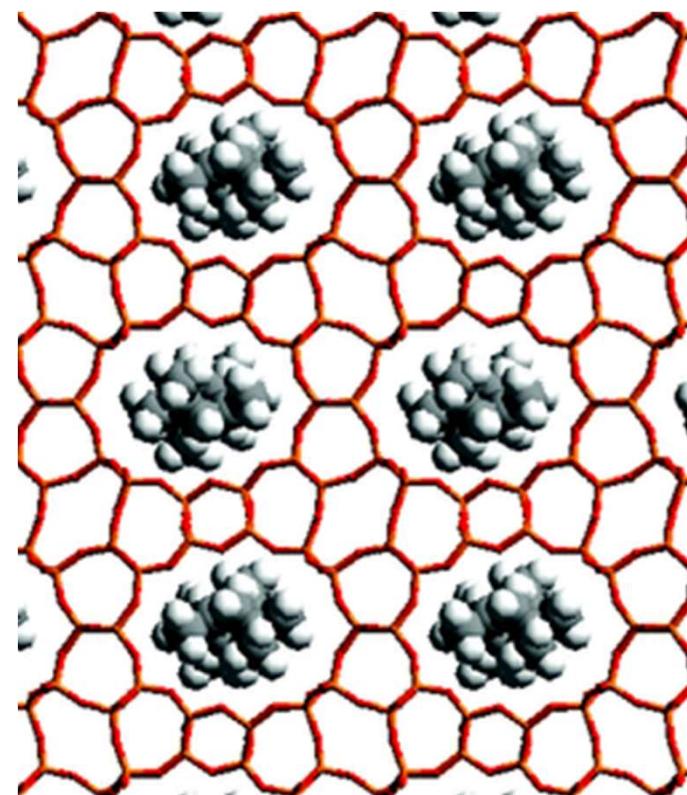
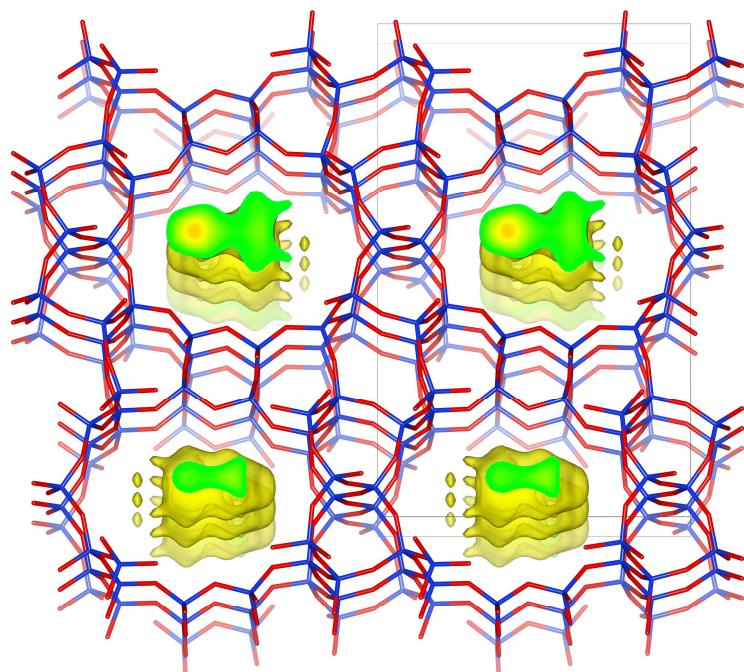


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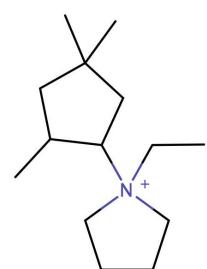
SSZ-60



SSZ-60



Burton & Elomari, Chem. Commun., 2618, 2004



Conclusions

- Can the location of the SDA be determined routinely?
 - Yes...
 - Flexibility of simulated annealing is ideal for structure completion
- How well do molecular modeling and XPD data agree?
 - Effect of Boron?
 - Difference map is very informative
 - Combine modeling and XPD data?