

Quantifying and enforcing plane symmetry in images of 2D periodic objects

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Electron Microscopy Berkeley

Lord Kelvin (William Thompson), 1883:

“I often say that when you can measure what you are speaking about, and express it in numbers, you know something about it; but when you cannot measure it, when you cannot express it in numbers, your knowledge is of a meager and unsatisfactory kind; it may be the beginning of knowledge, but you have scarcely, in your thoughts, advanced to the state of science.”

Outline

1. Plane symmetry, a well defined mathematical concept, deviations from it are quantifiable
2. Experimental and simulated examples:
Scanning Tunneling Microscopy (STM), 2D periodic and highly symmetric calibration samples, symmetrizing a blunt STM tip,
Scanning Electron Microscopy (SEM), Transmission Electron Microscopy (TEM), better surface and nano-science, *but problems of deciding between plane groups without prior knowledge*, other limitations
3. Knowledge of 2D symmetry makes for more sophisticated art (M. C. Escher)
4. Summary and Conclusions

LATTICE UNITS WITH SYMMETRIES OF PERIODIC PLANE PATTERNS

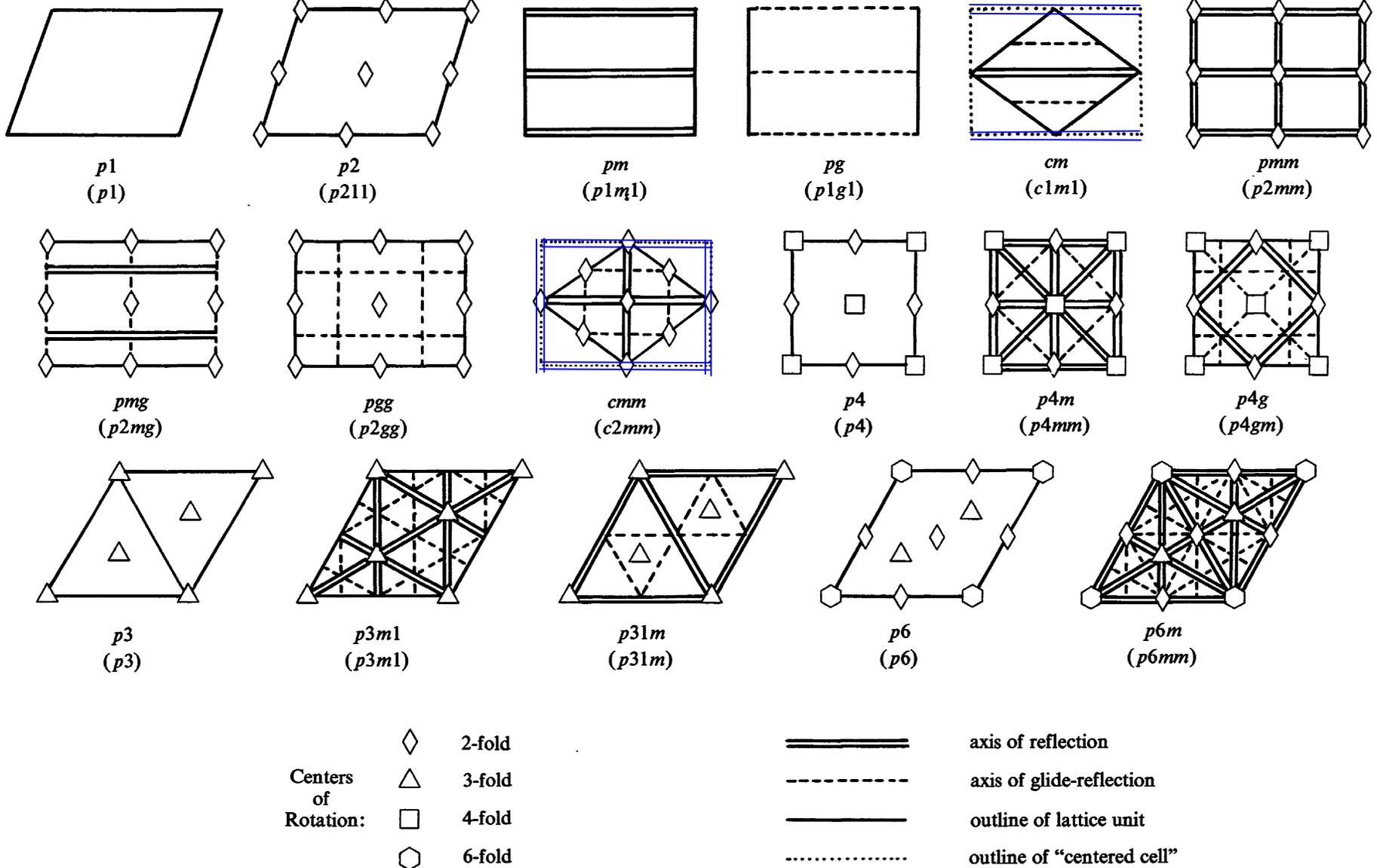
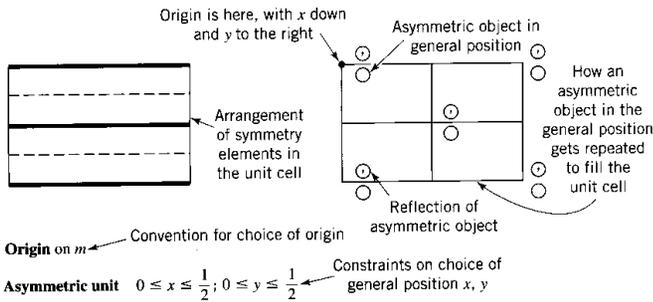


CHART 2. International notation identifies the seventeen two-dimensional crystallographic groups. The short form is given first, with the full notation in parentheses.

international notation, short form cm
 No. 5
 Point group symmetry m
 international notation, full form $c 1 m 1$
 Plane lattice type Rectangular
 Patterson symmetry $c 2 m m$



Positions	Coordinates
Multiplicity, Wyckoff letter, Site symmetry	$(0, 0) + (\frac{1}{2}, \frac{1}{2}) +$
4 b 1 (1) x, y (2) \bar{x}, y	Identifies this as a double cell; gives coordinates of lattice points
2 a .m. 0, y	

Designates general and special positions in the unit cell using fractional coordinates x, y . General position listed first, higher symmetry positions lower in the table.

Reflection conditions

General:
 $hk: h + k = 2n$
 $h0: h = 2n$
 $0k: k = 2n$

Special: no extra conditions

Diffraction information for this structure

Figure 3.32 Annotated copy of plane group No. 5, cm , from the *International Tables for Crystallography*, 1996.

T. Hahn (Ed.), Brief Teaching Edition of Volume A, Space-group symmetry, International Tables for Crystallography, 5th revised edition, International Union of Crystallography (IUCr), Chester 2005.

Robotics and machine vision community favors the direct space approach to plane symmetry detection and quantification, Y. Liu et al. IEEE Transactions on Pattern Analysis and Machine Intelligence 28 (2004) 354-374

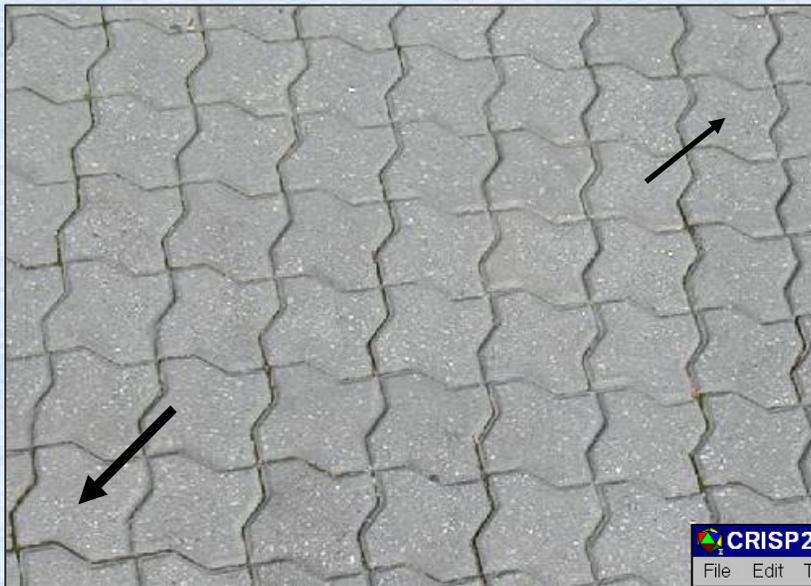
But crystallographic community prefers the reciprocal space, kinematic scattering, Fourier transform approach !

$$f(x, y) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} F(H, K) \exp\{-2\pi i(xH + yK)\} dH dK = FT^{-1}[F(H, K)]$$

The Fourier transform of a 2D periodic object is a discrete set of complex numbers, which contains exactly the same information!

$$f(x, y)^{periodic} = \sum_{-H}^H \sum_{-K}^K F(H, K)^{discrete} \exp\{-2\pi i(xH + yK)\}$$

Instead of analyzing the symmetry relations between *each pixel* of a 2D image, it is sufficient to analyze the symmetry relations of *a few Fourier coefficients*, in X-ray crystallography known as “structure factors”!!



Digital photograph of this street pavement was not recorded perpendicular to the pavement. Note the size difference between pavement bricks marked by arrows.

plane group: p4, point group of motif: 4

Crystallographic Image processing quantifies the plane symmetry correctly. Enforcing this symmetry leads to removal of the “imperfections of the imaging process” and “enhancement of surface details” of the pavement bricks.

CRISP2.1

File Edit Tools Area Fourier CIP ELD PhIDO Statistics Calculate Options Window Help

chatt1.jpg - 501x358x8 (1:1)

Origin refinement of chatt1.jpg

Symm RA% qRes Ao/Ae

p1	-	-	-	-
p2	-	-	32.8	-
pm	a\lx	37.2	19.0	-
pm	a\ly	37.2	15.8	-
pg	g\lx	38.1	15.7	1
pg	g\ly	40.0	19.1	1
cm	a\lx	41.0	12.3	1
cm	a\ly	41.0	14.1	1
pmm	-	37.2	43.3	-
pmg	a\lx	40.0	43.0	1
pmg	a\ly	38.1	37.6	1
p2gg	-	41.0	42.0	1
cmm	-	41.0	38.6	1
p4m	-	62.2	51.1	-
p4g	-	64.0	49.9	1
p3	-	-	-	-
p3m1	-	-	-	-

a=60.4Å b=60.4Å v=90.0° Shift h=104.8 Shift k=-44.6

FFT from ch...

p4 - density map from chatt...

Origin Refinement

Projection symmetry	Crystal symmetry	Real space glide parallel to	Systematic absences (n = any integer)
pg	P12 ₁	b	(0 k): k = 2n + 1
cm	C12	—	(h k): h + k = 2n + 1
p2mg	P222 ₁	a	(h 0): h = 2n + 1
p2gg	P22 ₁	a, b	(0 k): k = 2n + 1 (h 0): h = 2n + 1
c2mm	C222	—	(h k): h + k = 2n + 1
p4gm	P42 ₁	a, b	(0 k): k = 2n + 1 (h 0): h = 2n + 1

A* = 3.9Å Negate A

+1.905,+2.720, Den=288 Ready

Start Inb... http... 2 M... 2 W... CR... Mic... 10:43 PM

$$F_{res} = \frac{\sum_{HK} \left| |F_{obs}(H, K)| - |F_{sym}(H, K)| \right|}{\sum_{HK} |F_{obs}(H, K)|} \quad \alpha_{res} = \frac{\sum_{HK} w(H, K) \cdot |\alpha_{obs}(H, K) - \alpha_{sym}(H, K)|}{\sum_{HK} w(H, K)}$$

$$|F_{sym}(H, K)| = \frac{\sum_j^n |F_{obs}^j(H, K)|}{n}$$

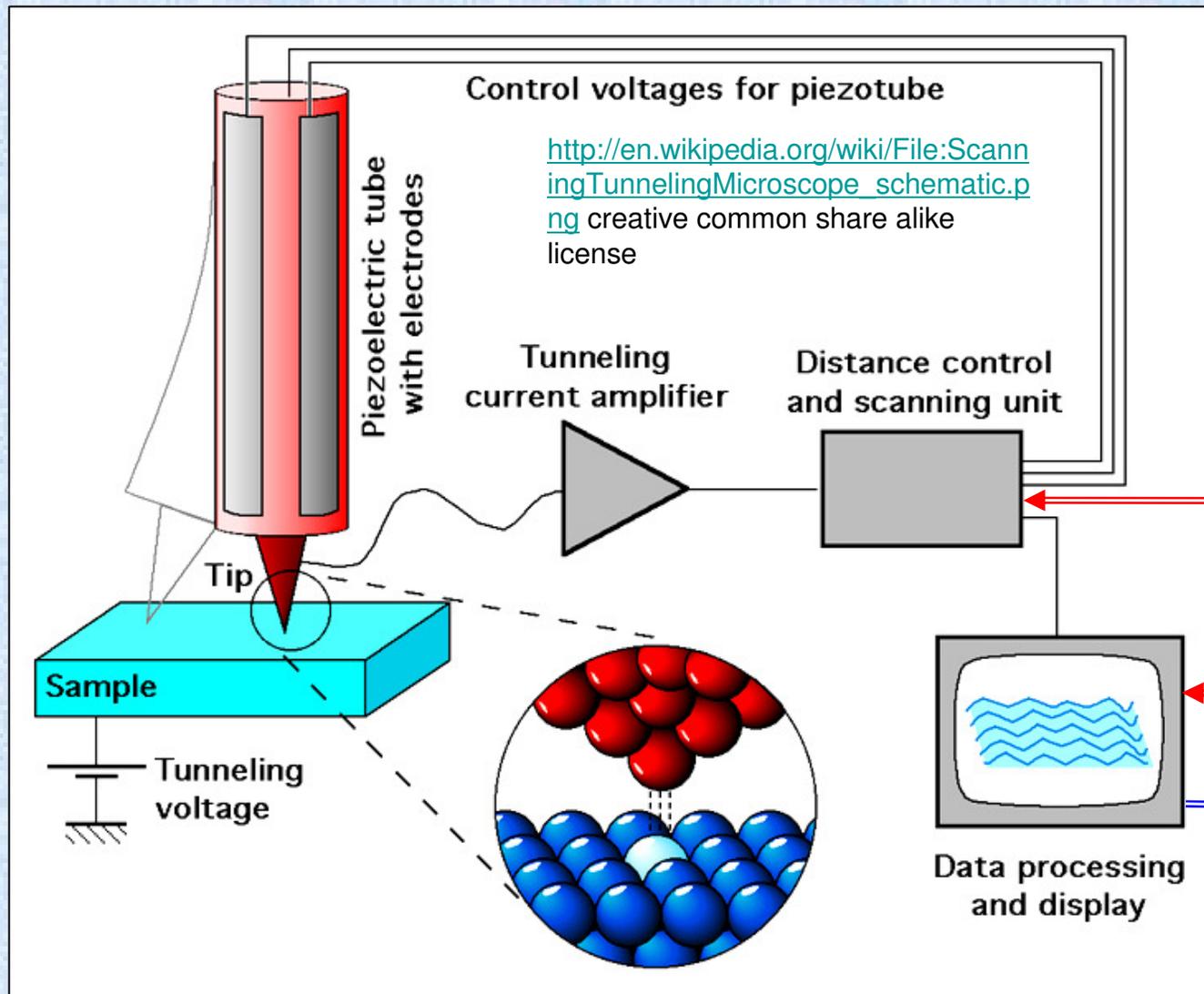
$$\alpha_{sym}(H, K) = \arctan \left[\frac{\sum_j w^j s^j \sin\{\alpha_{obs}^j(H, K)\}}{\sum_j w^j s^j \cos\{\alpha_{obs}^j(H, K)\}} \right] + \begin{cases} 0^\circ & [\text{if } \sum_j w^j s^j \cos\{\alpha_{obs}^j(H, K)\} > 0] \\ 180^\circ & [\text{if } \sum_j w^j s^j \cos\{\alpha_{obs}^j(H, K)\} < 0] \end{cases}$$

where w^j is a weighting factor, proportional to $|F_{obs}^j(H, K)|$; $s^j = 1$ in the absence of an approximately 180° flipped phase or $s^j = -1$. For centrosymmetric plane groups finally set to the nearest 0° or 180° .

$$\alpha_{sym}(H, K) = \alpha_{obs}(H, K) \text{ if not symmetry related except for Friedel's law } \alpha_{obs}(H, K) = -\alpha_{obs}(-H, -K)$$

$$|F_{sym}(H, K)| = |F_{obs}(H, K)| \text{ if not symmetry related except for Friedel's law } |F_{obs}(H, K)| = |F_{obs}(-H, -K)|$$

Note that known 2D symmetry information about a sample has not been utilized before !!



Crystallographic processing of an image from a known SPM calibration sample with 2D periodic (and highly symmetric) features in order to

1. Assess the performance of the microscope (determine prevalent point spread function) ;
2. Correct subsequently recorded images from unknowns with this point spread function ! (valid for one set of experimental conditions and scanning probe tip at a time)

Scanning Tunneling Microscopy: **left:** state of the art; **right:** our proposed enhancements, provisional patent application pending

Challenge: environmentally stable SPM standards with 2D periodicities at the atomic level, preferably high plane symmetry so that all subgroups can be used for quantification of the performance of the microscope and its proper calibration, HOPG should do !

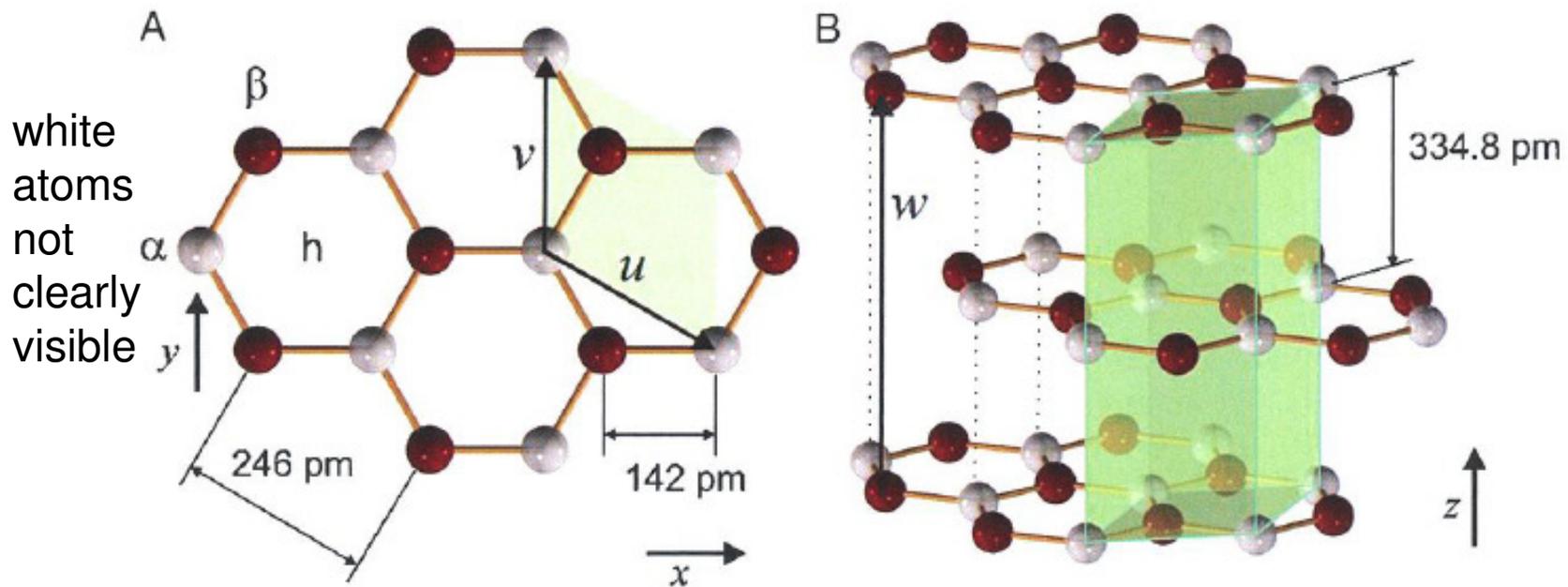
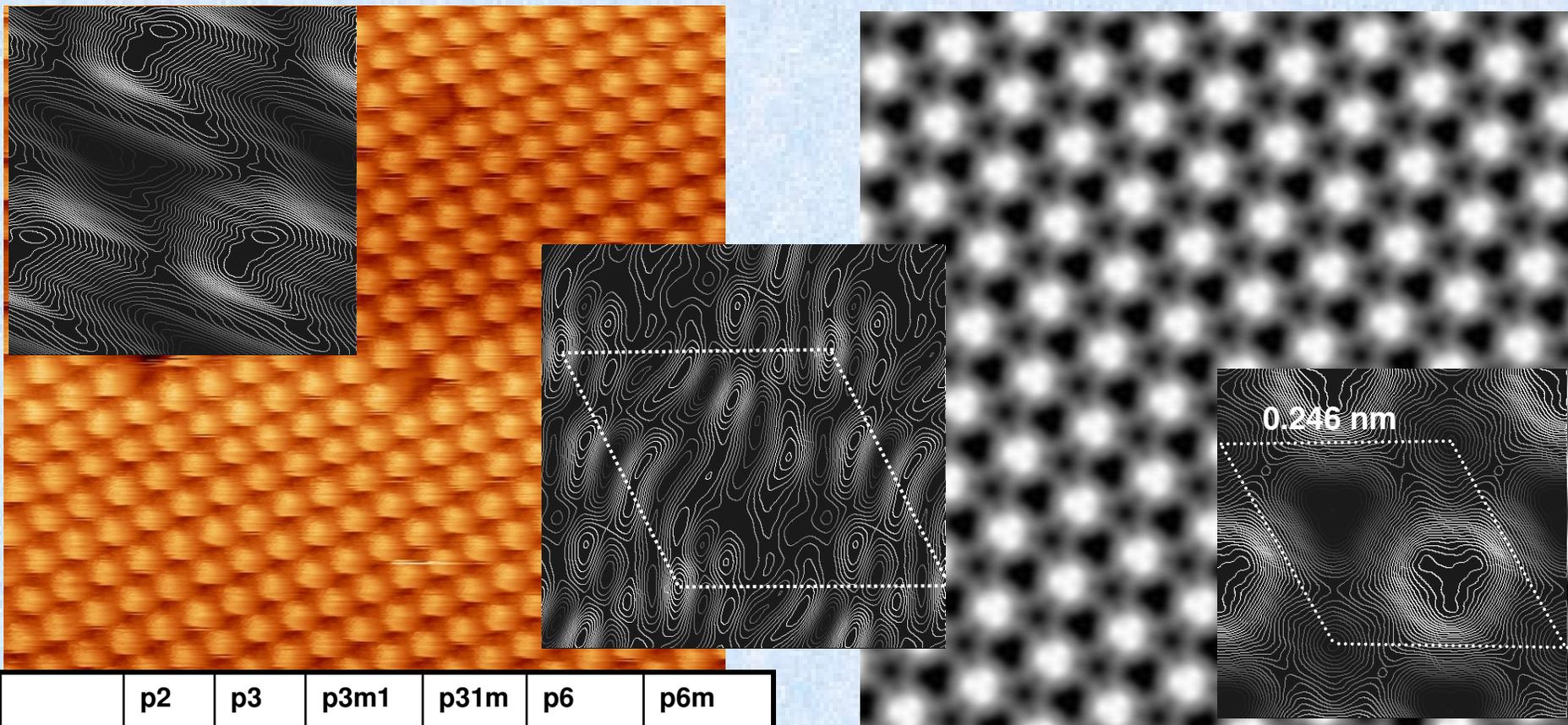


Fig. 1. Crystal structure of graphite. The unit cell is shaded in green. (A) Top view on the surface layer. The hexagonal surface lattice is defined by two unit vectors, u and v , in the xy plane with a length of 246 pm and an angle of 120° forming a honeycomb web of hexagonal rings. The basis of the lattice consists of two carbon atoms α (white) and β (red) with a distance of 142 pm. (B) Perspective view, showing the layered structure. The distance between layers is 2.36 times the next-neighbor distance of atoms within one layer, and the bond between layers is weak. The α atoms (white) are directly above an α atom in the layer directly underneath at a distance of 334.8 pm; the β atoms (red) are over a hollow sites (h). The unit vector w is parallel to the z -axis with a length of 669.6 pm.

S. Hembacher, et al., Revealing the Hidden Atom in Graphite by Low-Temperature Atomic Force Microscopy, Proceedings of the National Academy of Sciences of the United States of America, Vol. 100, No. 22 (Oct. 28, 2003), pp. 12539-12542

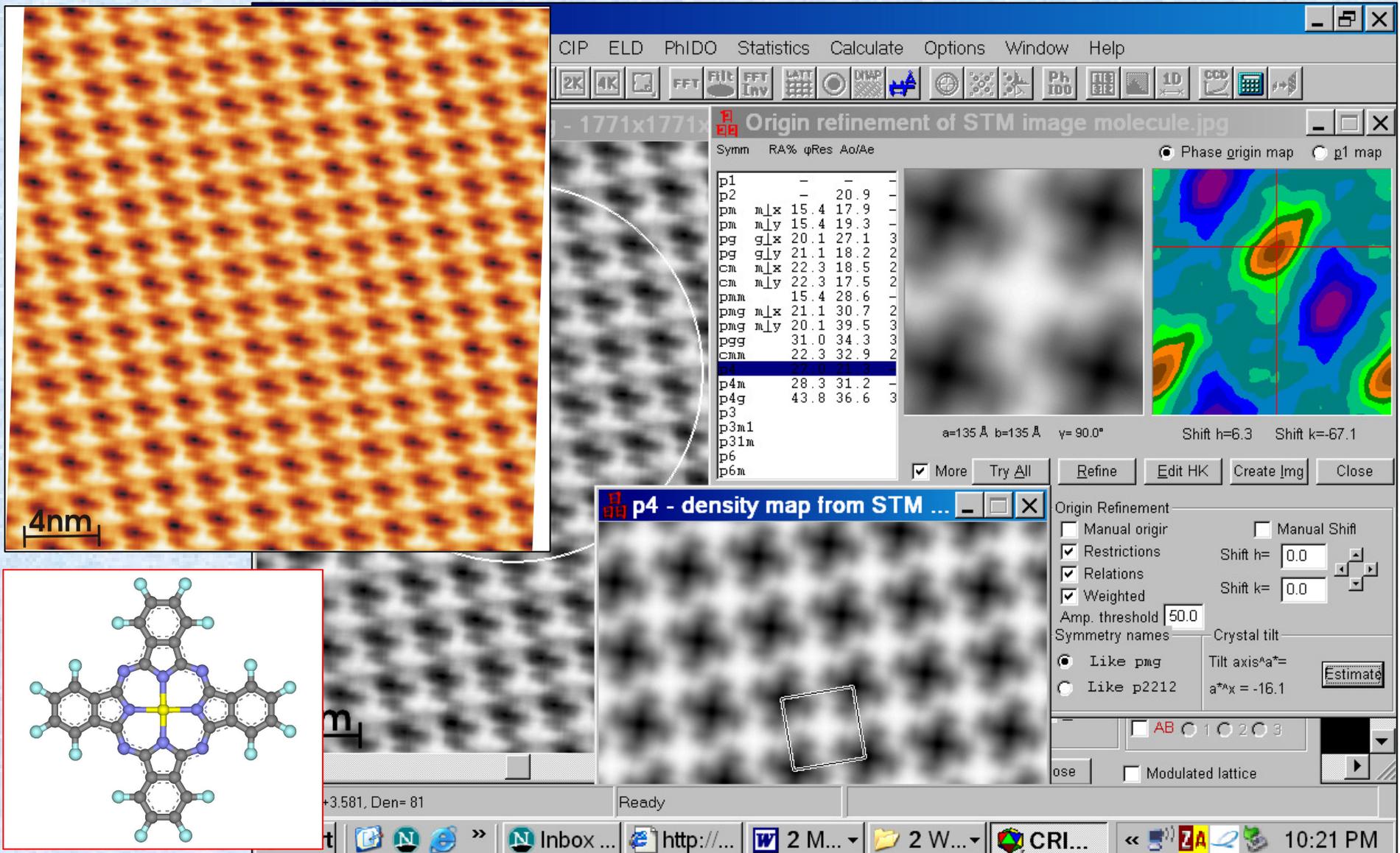


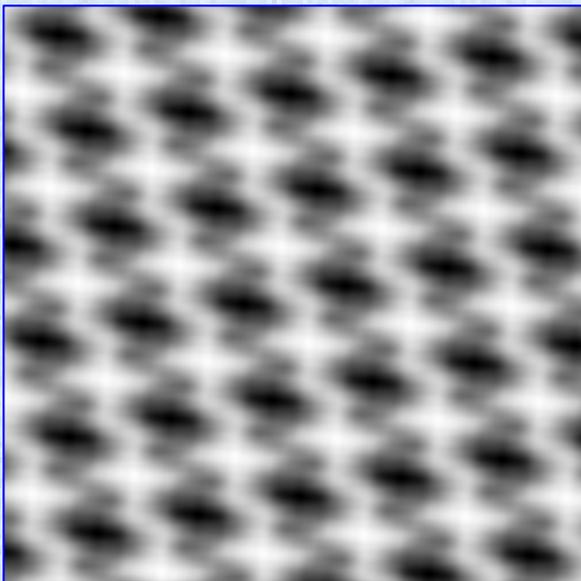
	p2	p3	p3m1	p31m	p6	p6m
F_{res} [%]		50.4	51.3	51.3	50.4	51.3
α_{res} [°]	19.9	14.2	17.1	29.2	31.3	32.2

The relative high amplitude residuals are due to a unit cell angle of 63.7° in the raw image.

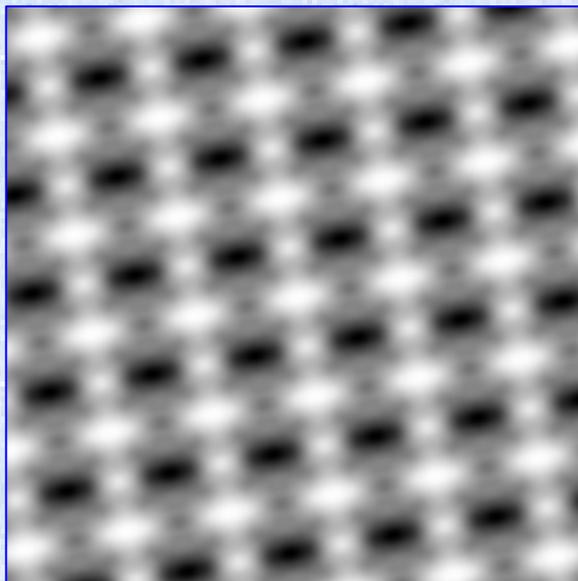
STM images of HOPG **(a)** raw data with 512 by 512 pixels, constant current mode, tungsten tip, 1 V bias, 0.15 nA (left) and **(b)** p3m1 enforced version of raw data (right). Note that there are atomic vacancies in the raw data. While the carbon atom at position $(\frac{2}{3}, \frac{1}{3})$ is clearly revealed, its counterpart at position (0,0) is rather faint, but at a higher intensity than the “empty spot” at position $(\frac{1}{3}, \frac{2}{3})$.

Scanning Tunneling Microscopy Image of a mono-layer of fluorinated cobalt phthalocyanine ($F_{16}CoPc$) on graphite, UHV and 20 K, tungsten tip bias 1 V, constant 0.1 nA tunneling current !

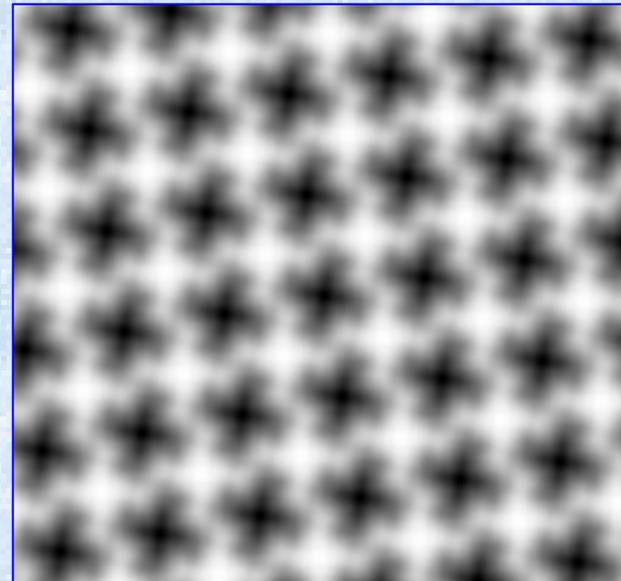




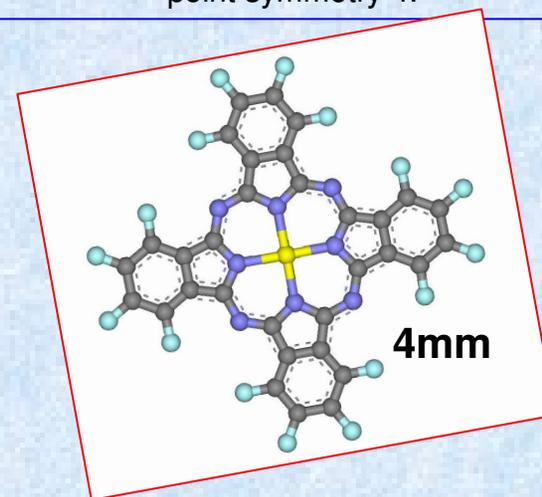
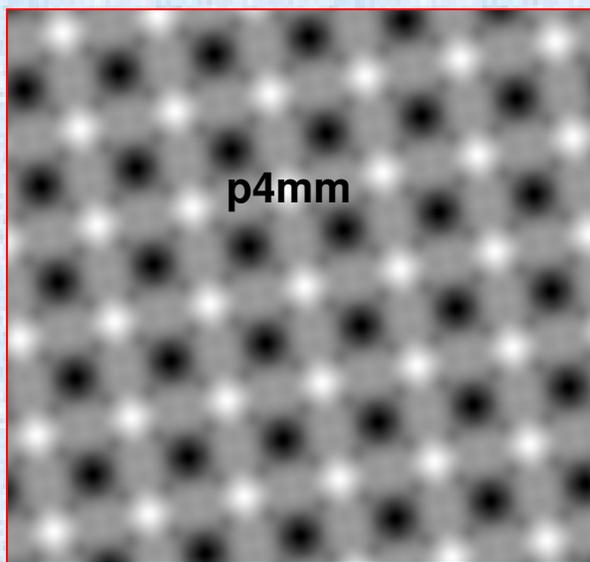
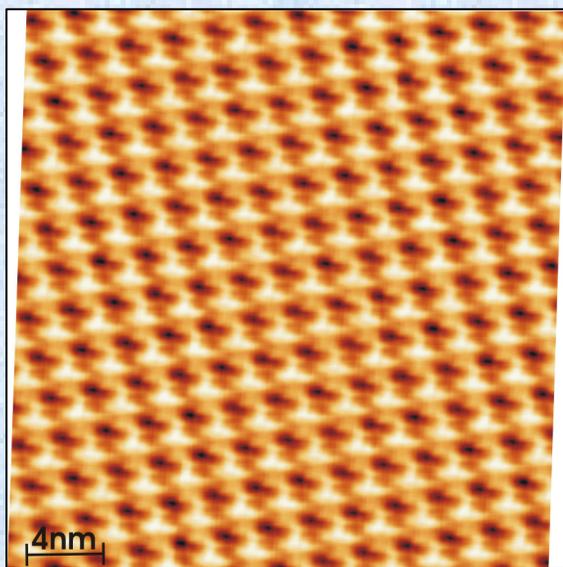
p2 symmetry enforced STM data. A two-fold axis is imposed. The lattice remains oblique. The motif acquires point symmetry 2 and the peculiar "overall image twist" is removed.



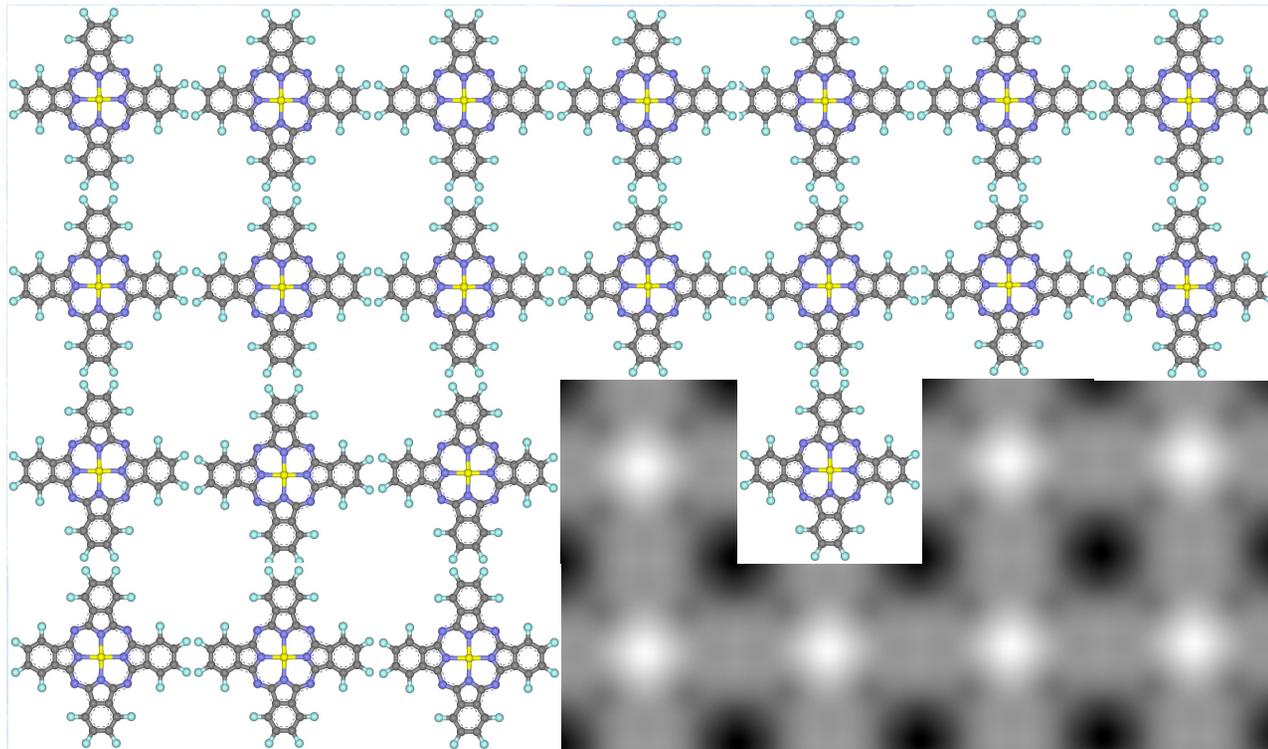
p2mm symmetry enforced STM data. A two-fold axis with a set of perpendicular mirrors is imposed. The lattice becomes rectangular. The motif acquires point symmetry 2mm



p4 symmetry enforced STM data. A four-fold axis is imposed. The lattice becomes quadratic and the motif acquires the point symmetry 4.

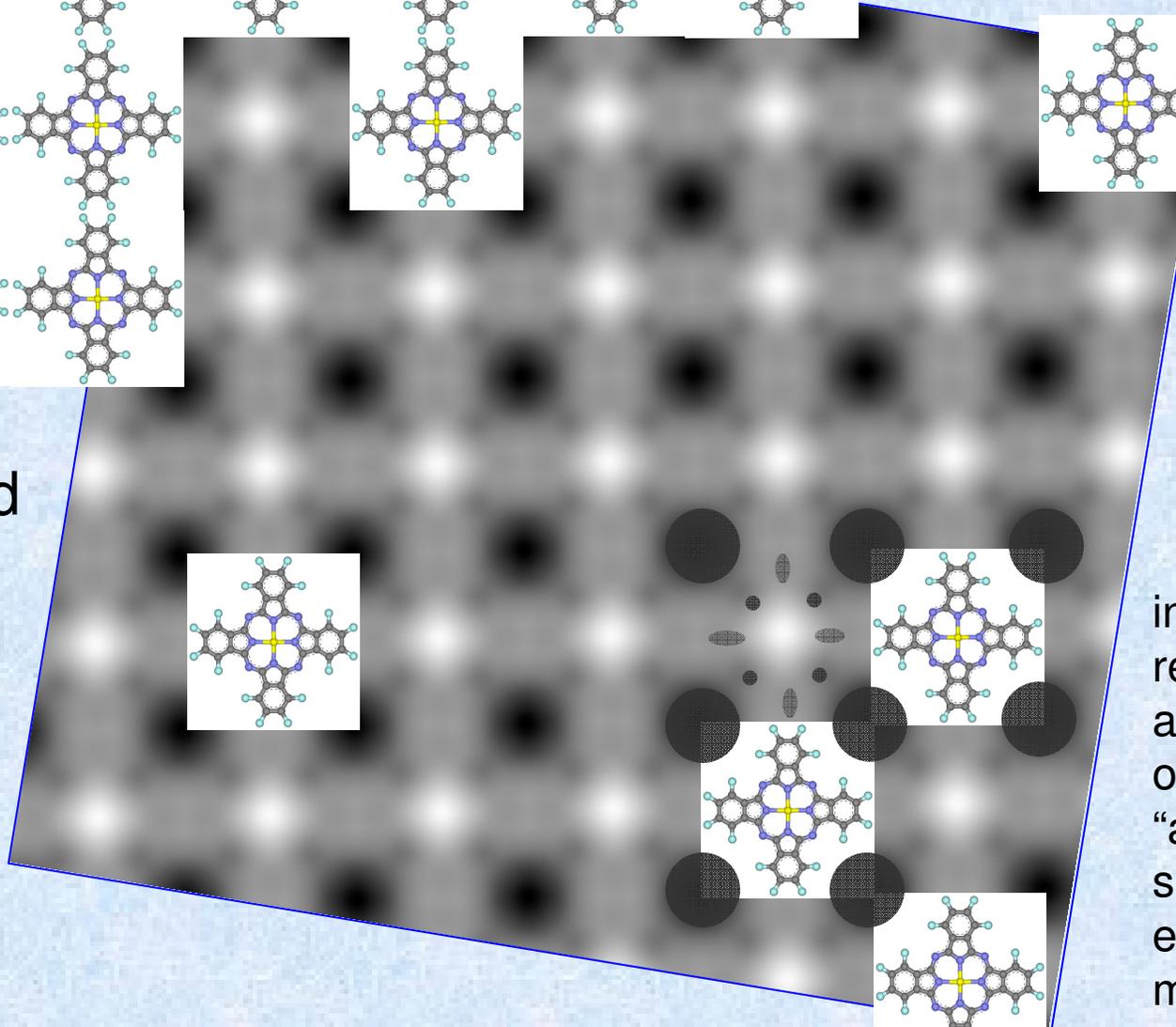


"Size" of molecule and basic periodicities in STM images approximately 1.5 – 2 nm



“Size” of molecule and
basic 2D periodicities
in STM images
approximately 1.5 - 2
nm

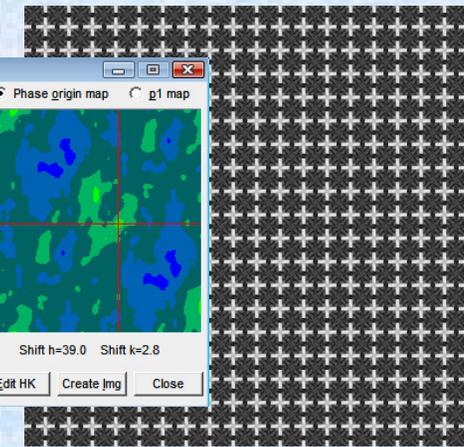
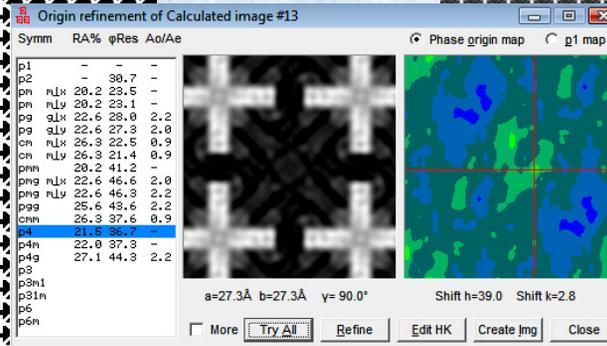
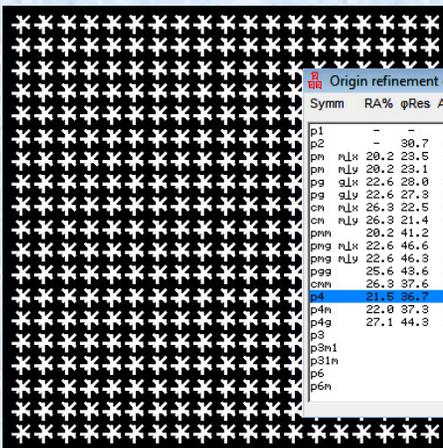
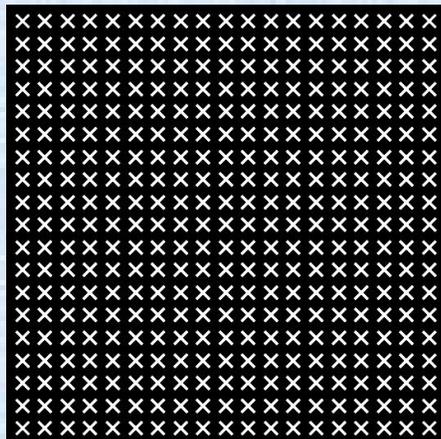
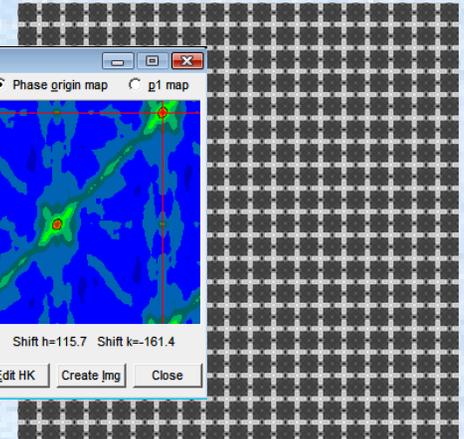
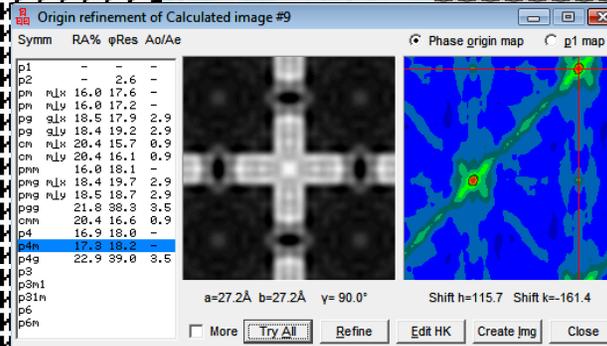
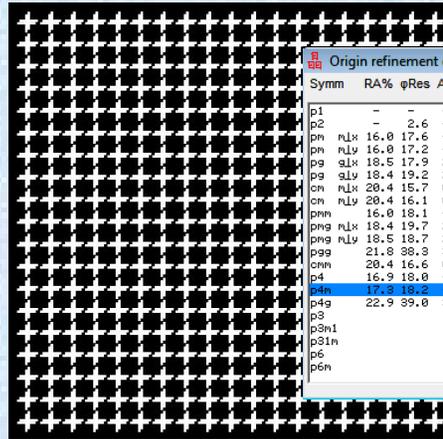
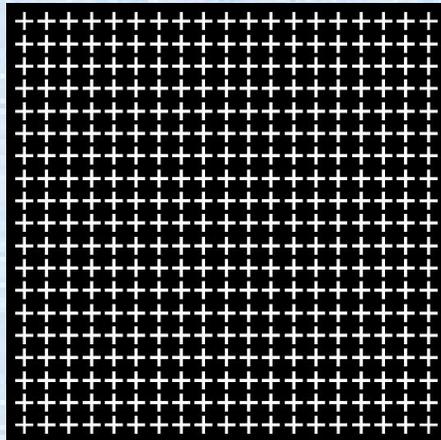
p4mm
symmetrized
version of
STM data in
comparison
to a model
array of
 $F_{16}CoPc$
molecules



in essence
regularly
arrayed copies
of the
“average” and
symmetry
enforced
molecule

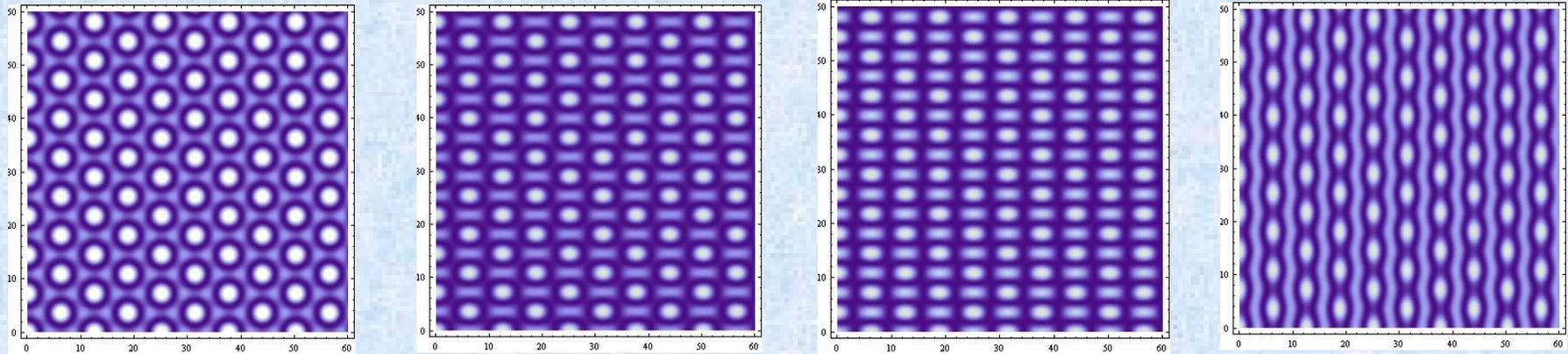
recovering one
image of the
original 2D array

two images of the “same kind” from the same 2D array
superimposed with translation

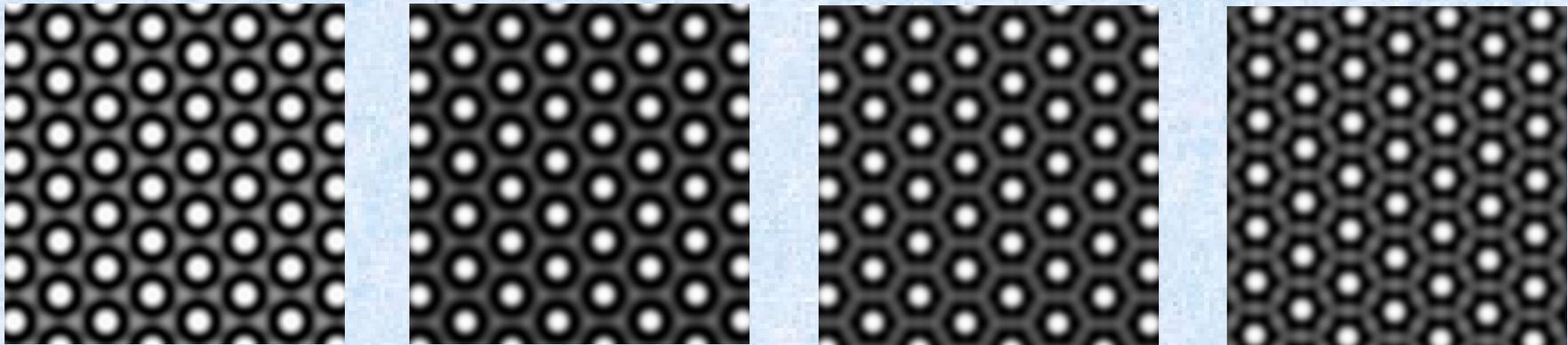


both p4mm

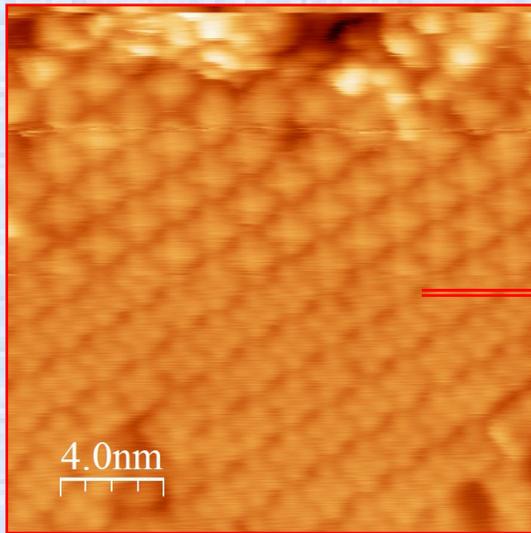
two images of “each kind” from the same 2D array
superimposed with translation



Simulated series of STM images of a sample surface with plane symmetry $p6mm$. The first image was simulated for a single tip. Two mini-tips (as a simple model of a blunt STM tip) were scanning from top to bottom in the other three simulations. The distance between the two mini-tips increases from left to right. **Electron interference between the two tips was included in the simulations.**

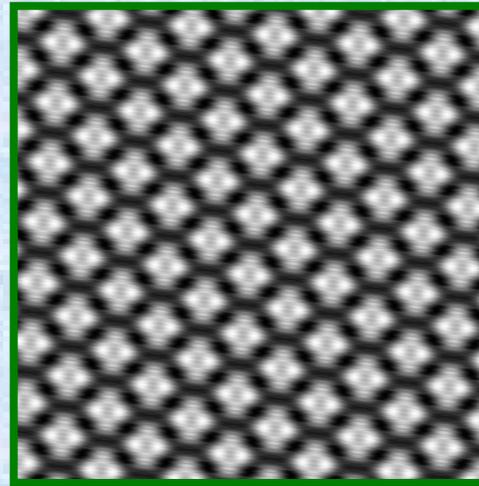


CIP processed STM images (of a sample surface with plane symmetry $p6mm$) as simulated in the figures above. **Despite the two mini-tips and electron interference effects between these two tips in three of the simulated image, crystallographic image processing recovers in all cases the underlying plane symmetry of the sample surface reasonably well !**

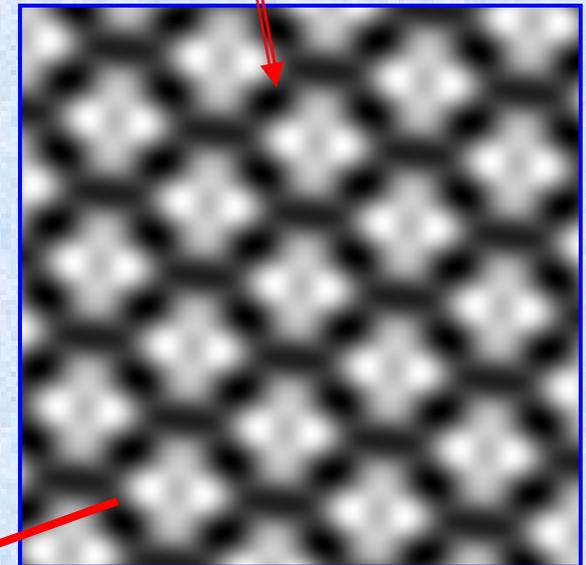
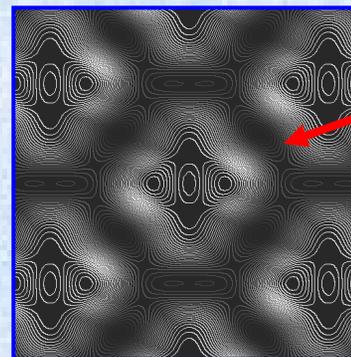
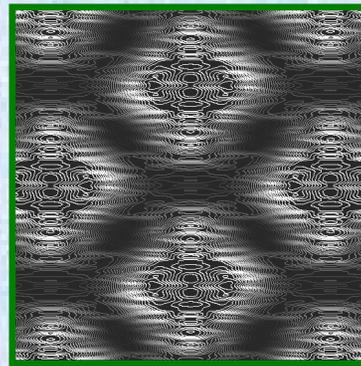
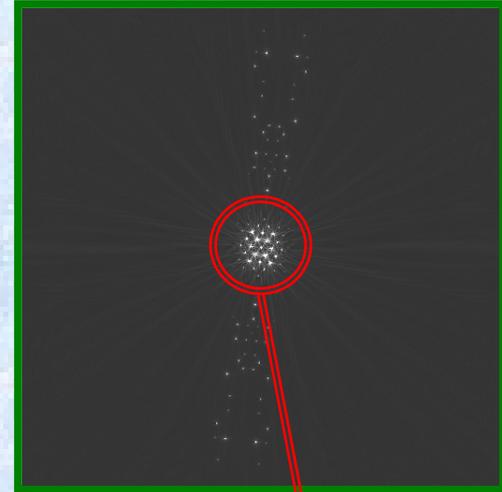


F₁₆CoPc on silver (110), UHV and 20 K, tungsten tip bias 1 V, constant 0.1 nA tunneling current

Working hypotheses: since the substrate has plane group *c2mm*, it is no coincidence that the monolayer also has this plane group (Curie's principle for strong epitaxial interaction), also directly above every 5th Ag atom, there is the central Co atom of every F₁₆CoPc molecule



c2mm enforced ↓



Fourier filtered and contour map

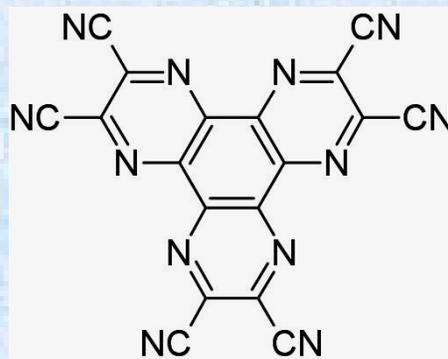
“Soft” Metallic Contact to Isolated C Molecules

Hendrik Glowatzki, Benjamin Broßker, Ralf-Peter Blum, Oliver T. Hofmann, Antje Vollmer, Ralph Rieger, Klaus Müllen, Egbert Zojer, Jürgen P. Rabe, and Norbert Koch

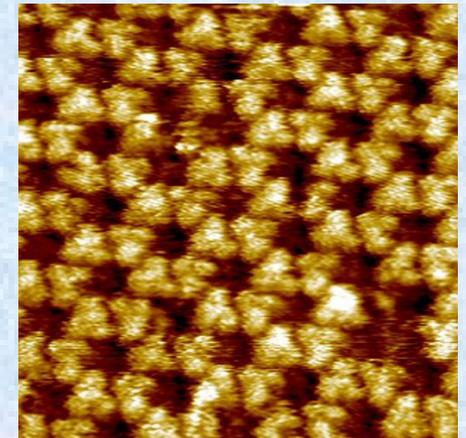
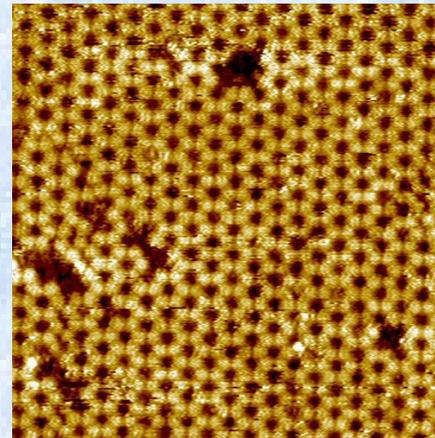
Nano Lett., 2008, 8 (11), 3825-3829 • DOI: 10.1021/nl8021797 • Publication Date (Web): 28 October 2008

STM, UHV, 300 K, a genuine breakthrough

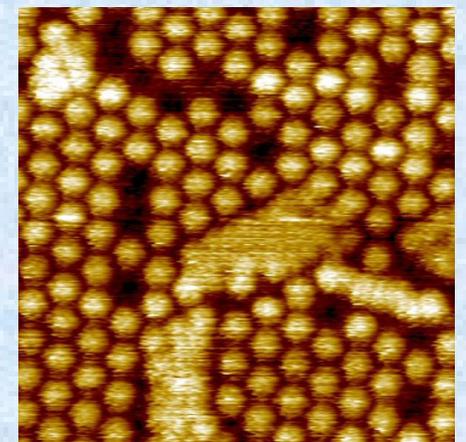
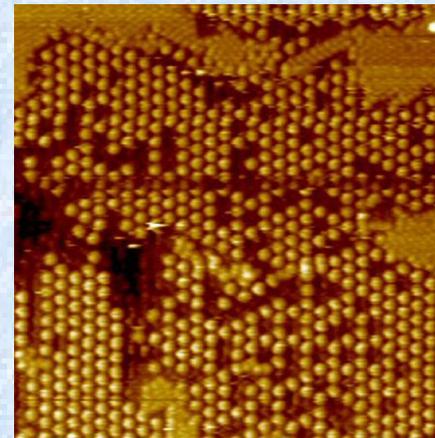
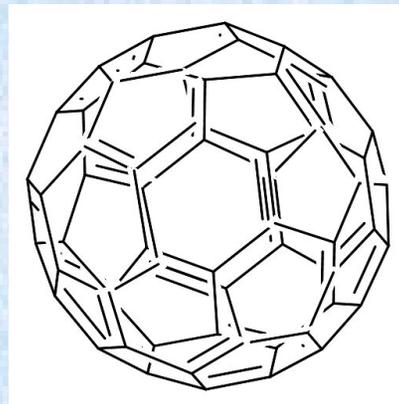
Ag (111), plane
group $p6mm$ +
hexaazatriphenylene
-hexanitrile
HATCN



metallic when on Ag (111)



+ buckyballs
in the centers
of the
molecules



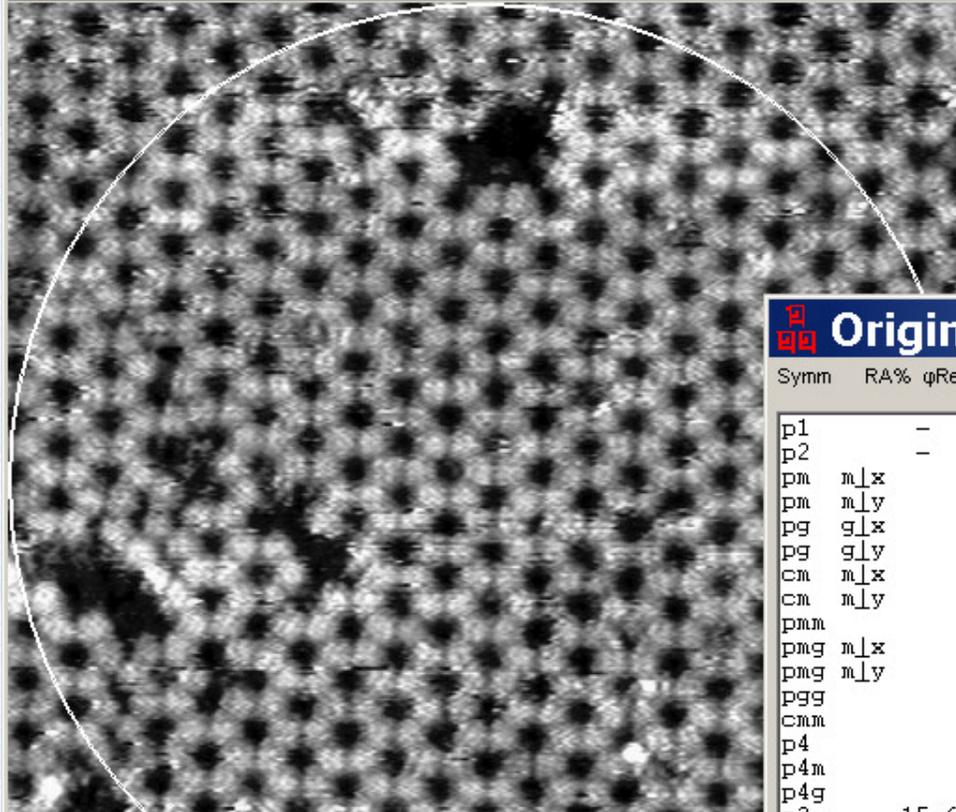
electronic structure not modified by the soft metallic contact !

CRISP2.1

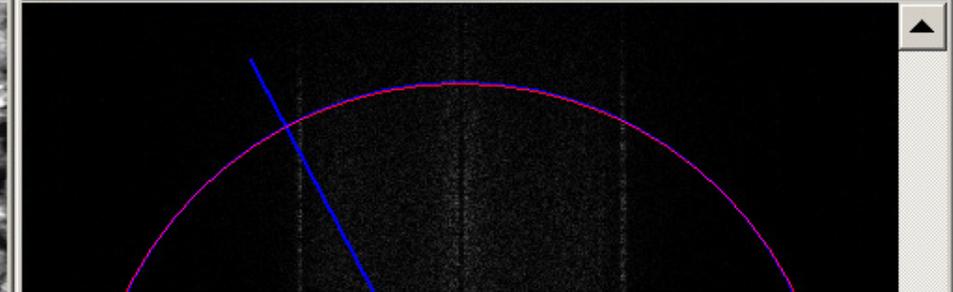
File Edit Tools Area Fourier CIP ELD PhIDO Statistics Calculate Options Window Help

128 256 512 1K 2K 4K FFT File FFT Inv LATT DMAP Ph IDO 1D CCD

HATCN-Ag(111).jpg - 512x512



FFT from HATCN-Ag(111).jp...



Origin refinement of HATCN-Ag(111).jpg

Symm RA% qRes Ao/Ae

Phase origin map p1 map

p1	-	-	-
p2	-	11.9	-
pm	m x	-	-
pm	m y	-	-
pg	g x	-	-
pg	g y	-	-
cm	m x	-	-
cm	m y	-	-
pmm	-	-	-
pmg	m x	-	-
pmg	m y	-	-
pgg	-	-	-
cm	-	-	-
p4	-	-	-
p4m	-	-	-
p4g	-	-	-
p3	15.6	7.6	-
p3m1	16.7	8.5	-
p31m	16.7	11.9	-
p6	15.6	12.3	-
p6m	16.7	12.7	-

a=28.8Å b=28.8Å γ=120.0° Shift h=-169.1 Shift k=-5.6

More

Lattice refinement ...

Detect Lattice 'AB'

Refl	H	K
1	1	0
2	0	1

Modify Lattice 'AB'

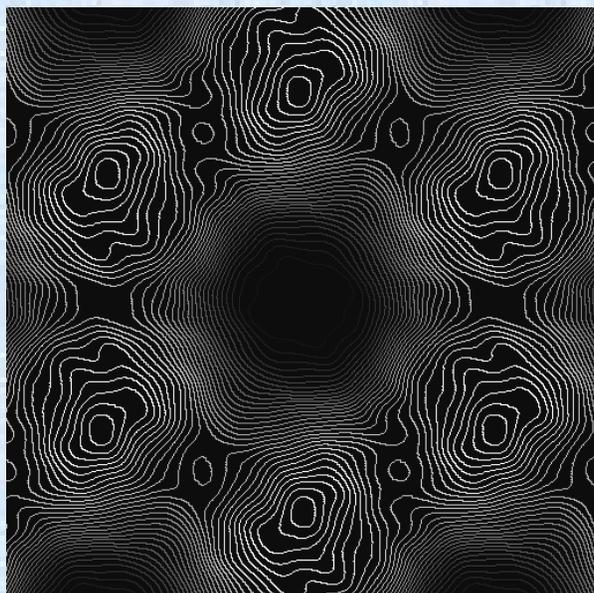
A*=20.5Å Negate A

B*=20.6Å Negate B

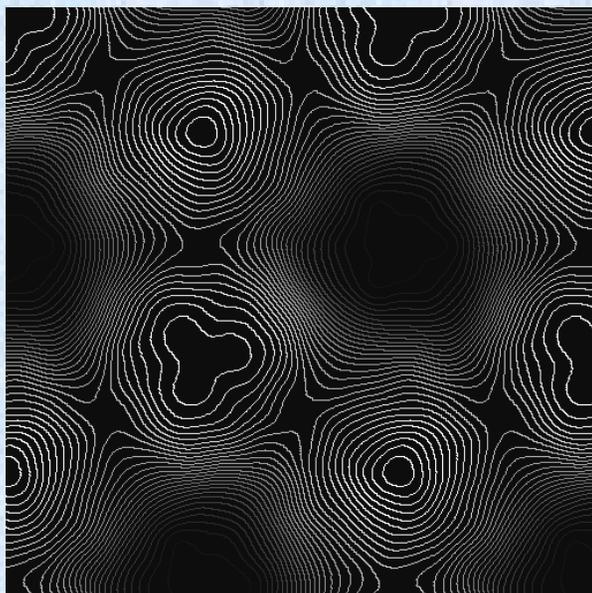
γ*=59.7°

+0.587,-0.371, Den=237 Ready Microsoft PowerPoint - [HUB 1.ppt]

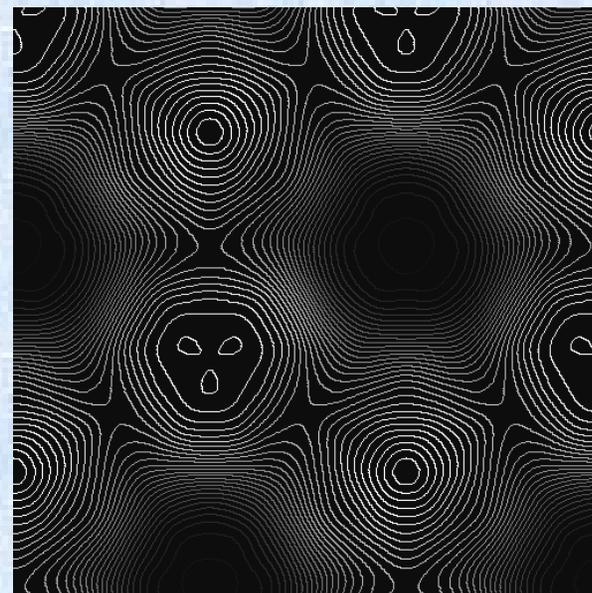
Start 2 N 3 I Wi... 2 W Sof... CR... Mic... 1:24 PM



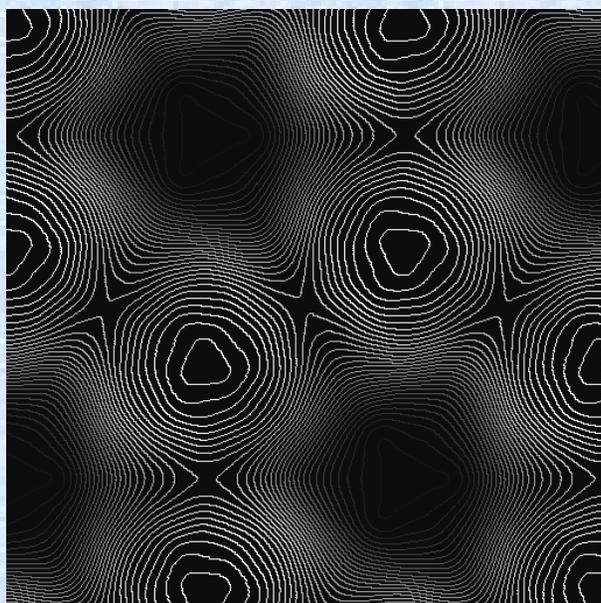
p2, $\alpha_{\text{res}}:11.9^\circ$



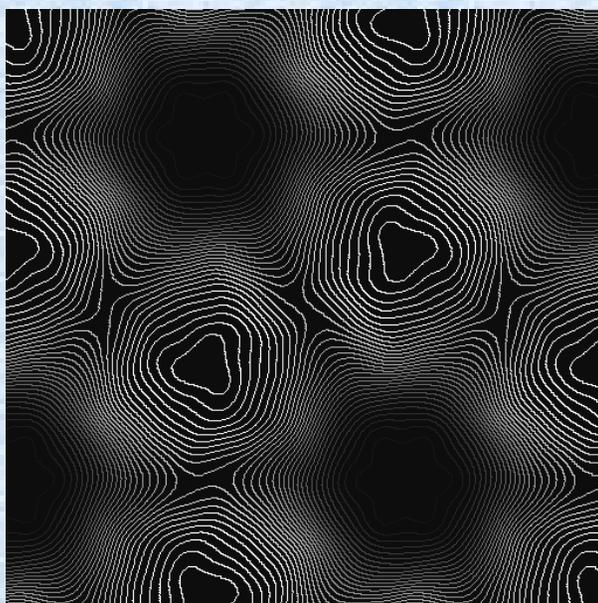
p3, $F_{\text{res}}:15.6\%$, $\alpha_{\text{res}}:7.6^\circ$



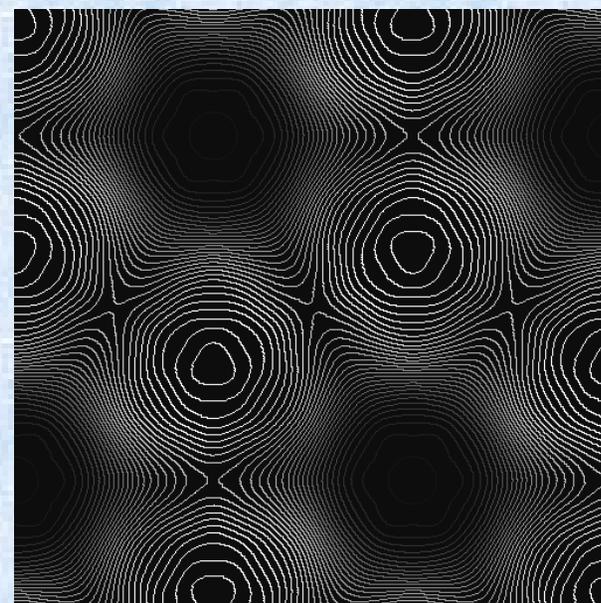
p3m1, $F_{\text{res}}:16.7\%$, $\alpha_{\text{res}}:8.5^\circ$



p31m, $F_{\text{res}}:16.7\%$, $\alpha_{\text{res}}:11.9^\circ$

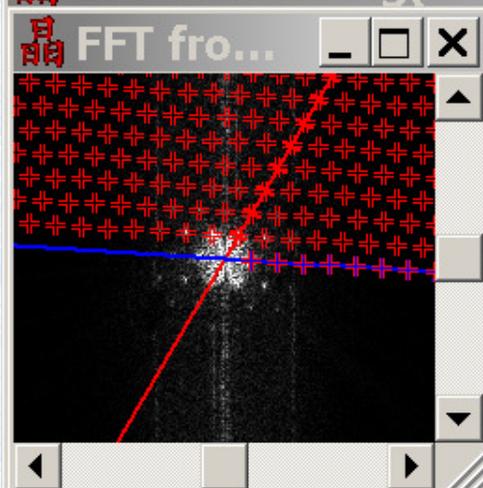


p6, $F_{\text{res}}:15.6\%$, $\alpha_{\text{res}}:12.3^\circ$



p6mm, $F_{\text{res}}:16.7\%$, $\alpha_{\text{res}}:12.7^\circ$

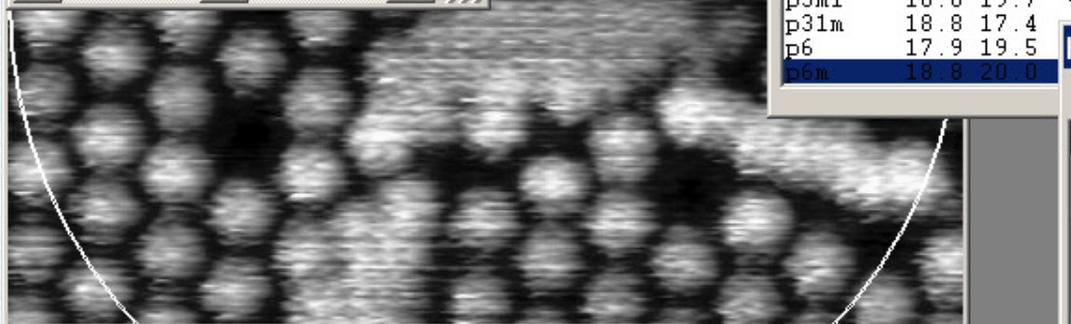
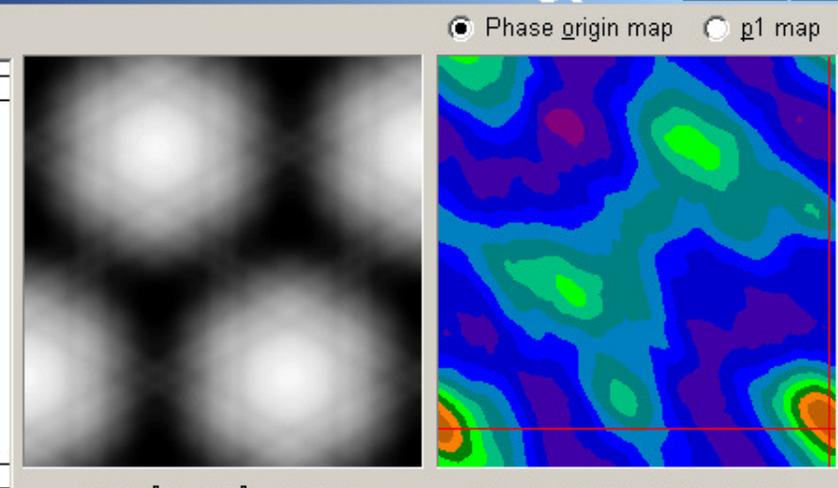
C60-HATCN-Ag(111)_zoom.jp... Origin refinement of C60-HATCN-Ag(1...



Symm RA% qRes Ao/Ae

```

p1 - -
p2 - 17.5 -
pm m|x
pm m|y
pg g|x
pg g|y
cm m|x
cm m|y
pmm
pmg m|x
pmg m|y
pgg
cmm
p4
p4m
p4g
p3 17.9 15.0 -
p3m1 18.8 19.7 -
p31m 18.8 17.4 -
p6 17.9 19.5 -
p6m 18.8 20.0 -
    
```



Lattice refinement ...

Detect Lattice 'AB'

Refl	H	K
1	1	0
2	0	1

Modify Lattice 'AB'

A*=14.1Å Negate A

B*=14.1Å Negate B

γ*=63.1° Swap AB

HK Edit

h	k	Amp	AmpS	Pha	PhaS	Err
0	1	9931	9766	-2	0	
1	0	10000	9766	-16	0	
1	-1	9367	9766	0	0	
0	2	1531	1253	-143	180	
2	0	1150	1253	-158	180	
2	-2	1077	1253	179	180	
0	3	169	84	56	0	P
3	0	0	84	32	0	A
3	-3	84	84	-68	0	P
0	5	14	92	-134	180	PU
5	0	135	92	-19	180	PU
5	-5	127	92	-161	180	U
1	1	2161	1745	-149	180	
1	-2	2167	1745	150	180	
2	-1	908	1745	-58	180	P
1	2	776	589	67	0	P

Edit

h = 0

k = 1

Amp = 9766

Pha = 0

Change

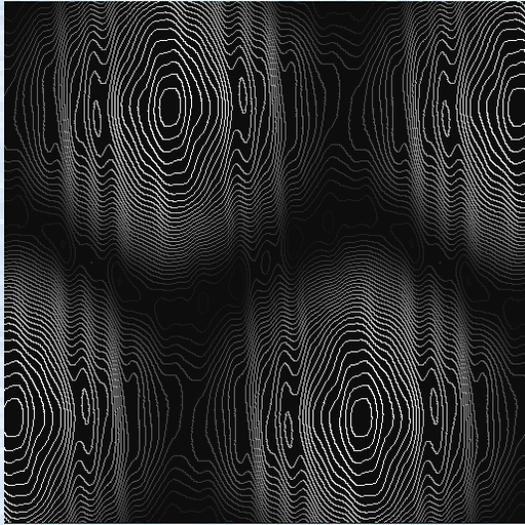
Add

Delete

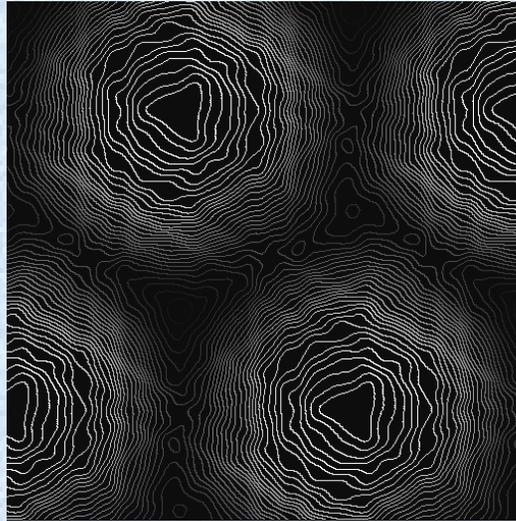
Inverse

Amp Threshold

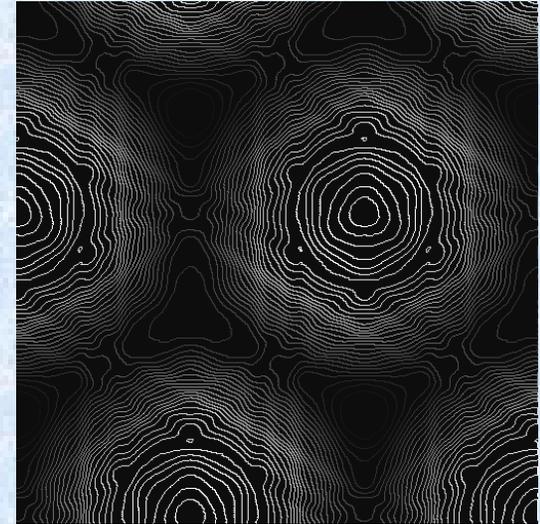
50 Apply



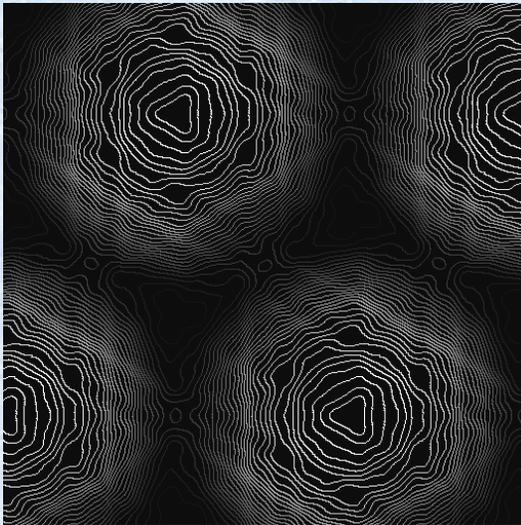
p2, α_{res} :17.5°



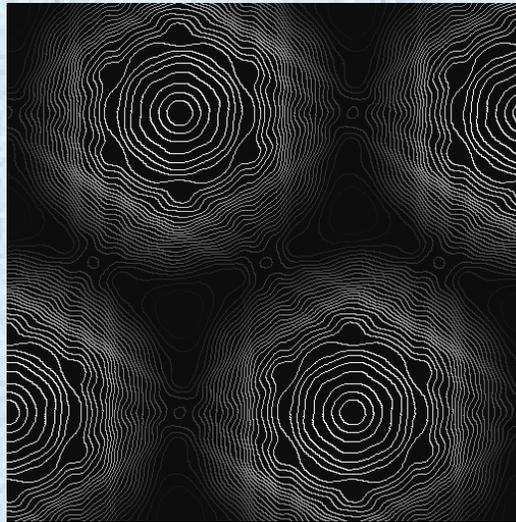
p3, F_{res} :17.9%, α_{res} :15.0°



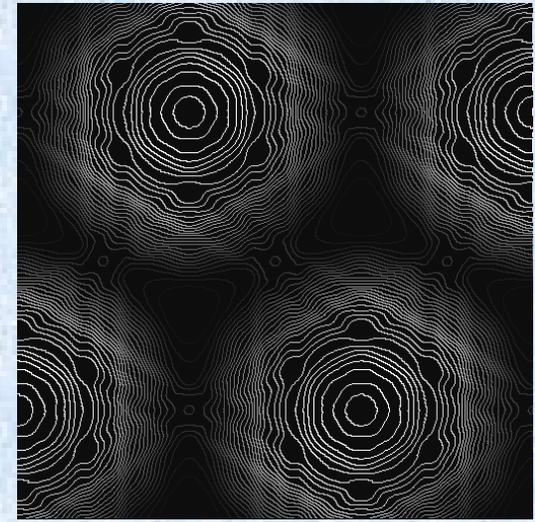
p3m1, F_{res} :18.8%, α_{res} :19.7°



p31m, F_{res} :18.8%, α_{res} :17.4°

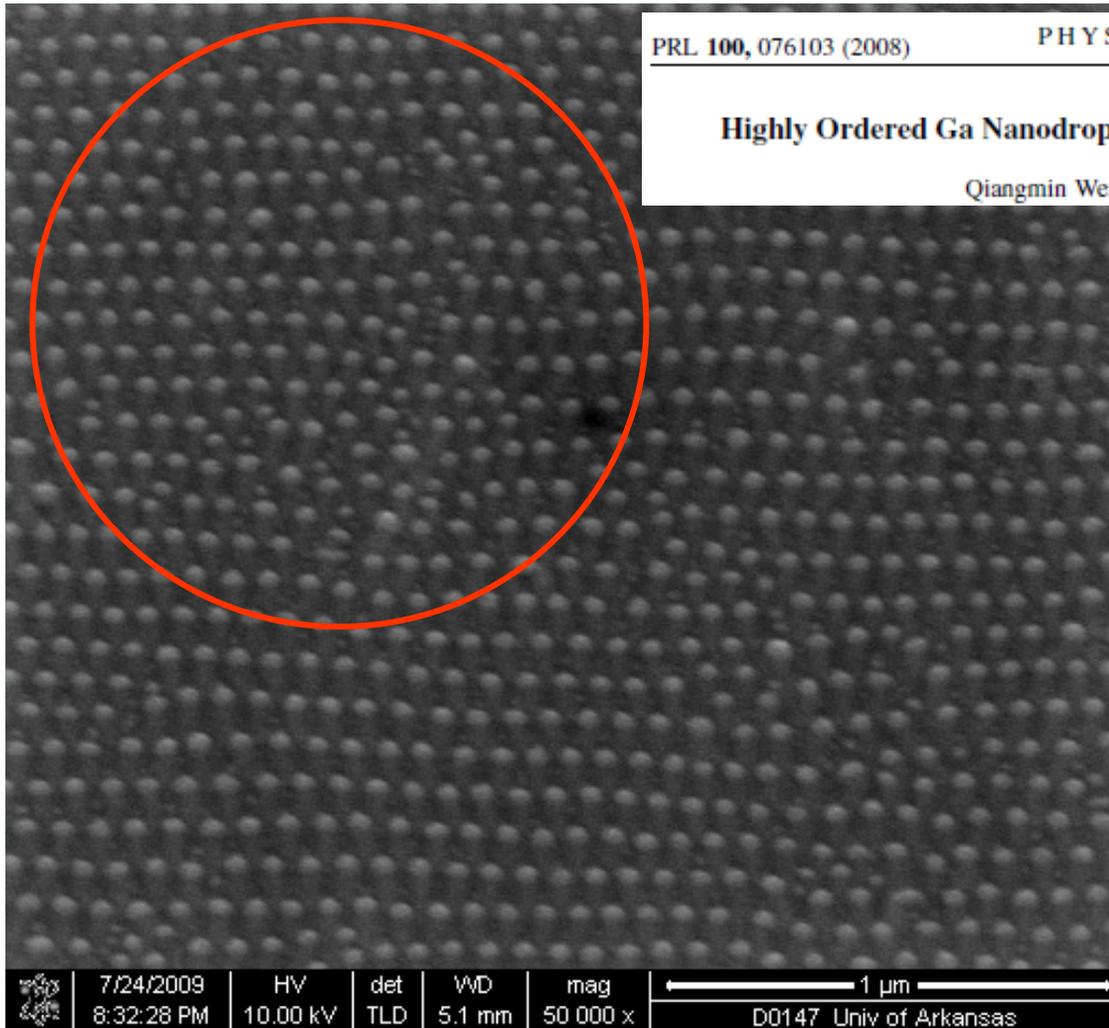


p6, F_{res} :17.9%, α_{res} :19.5°



p6mm, F_{res} :18.8%, α_{res} :20.0°

Highly Ordered Ga Nanodroplets on a GaAs Surface Formed by a Focused Ion Beam

Qiangmin Wei,¹ Jie Lian,^{2,3} Wei Lu,⁴ and Lumin Wang^{1,5,*}

SEM of a “uniform” array of Ga droplets on crystalline GaAs obtained by simultaneous sputtering and low energy Ga-ion milling at a large inclination angle, residuals for p6mm, **amplitude: 37.2%**, **phase angle 23.6°**, i.e. the ideal closed packed hexagonal array of identical entities with a point symmetry higher than 6mm

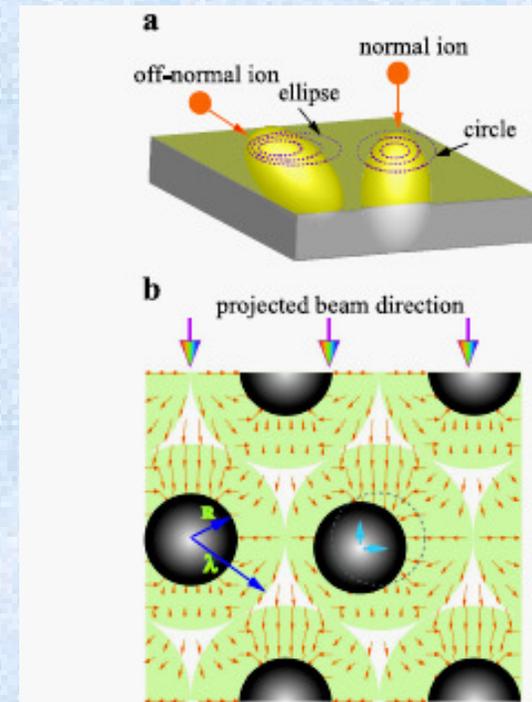
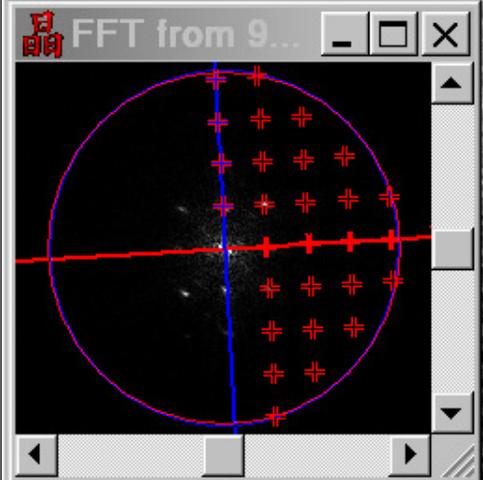


FIG. 4 (color online). Schematic illustration of a model for the formation of ordered droplet patterns. (a) Average energy distribution for ion bombardment. For normal bombardment, the deposited energy on the surface is circular, while for off-normal bombardment, it is elliptical where the energy contour along the projected beam direction is longer than other directions. The dashed curves represent the equal energy contours. (b) Atom supply and movement directions that cause an off-center particle to move to the center of nanoparticle lattice. Small arrows represent the direction and magnitude of local Ga atom migration induced by the ion beam on the substrate surface, radial dark shaded circle denotes the droplet, the light circle with radius λ is the exclusive zone for Ga, arrows inside droplet indicate the movement direction of droplets, and dashed circle shows the final position of partially aligned droplets.

911_FeO_Am6_04_B3_19_M70kx.tif

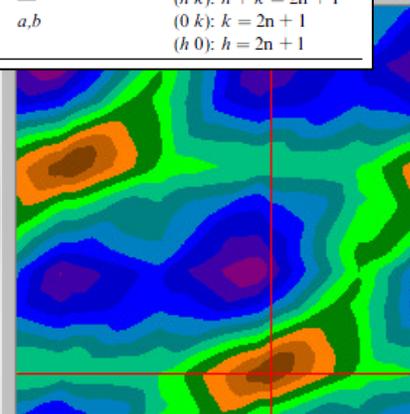
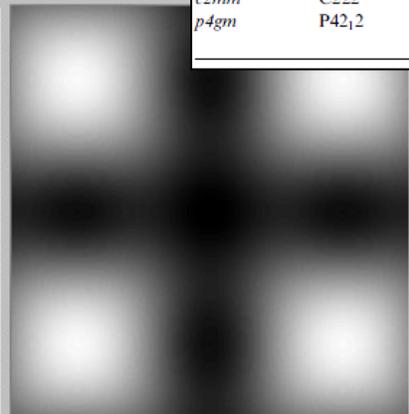


Origin refinement of 911

Symm	RA%	qRes	Ao/Ae
p1	-	-	-
p2	-	6.0	-
pm	m x	11.1	10.2
pm	m y	11.1	19.7
pg	g x	21.2	5.0
pg	g y	12.6	4.7
cm	m x	25.9	0.6
cm	m y	25.9	6.9
pmm		11.1	22.0
pmg	m x	12.6	5.2
pmg	m y	21.2	10.3
pgg		27.5	12.7
cm		25.9	8.2
p4		55.4	22.0
p4m		55.8	22.0
p4g		49.9	13.2
p3			
p3m1			
p31m			
p6			
p6m			

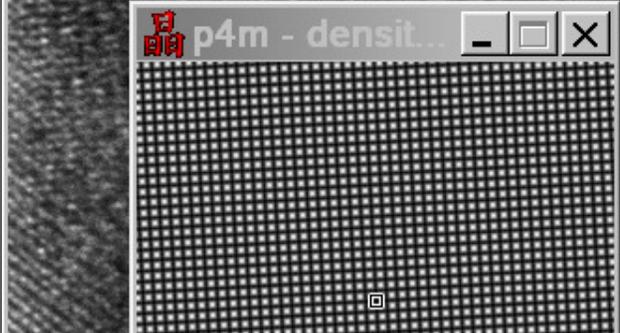
Symmetry-forbidden reflections in 2D space groups

Projection symmetry	Crystal symmetry	Real space glide parallel to	Systematic absences (n = any integer)
pg	P12 ₁	b	(0 k): k = 2n + 1
cm	C12	—	(h k): h + k = 2n + 1
p2mg	P222 ₁	a	(h 0): h = 2n + 1
p2gg	P22 ₁ 2 ₁	a,b	(0 k): k = 2n + 1
			(h 0): h = 2n + 1
c2mm	C222	—	(h k): h + k = 2n + 1
p4gm	P42 ₁ 2	a,b	(0 k): k = 2n + 1
			(h 0): h = 2n + 1



a=22.7Å b=22.7Å γ=90.0° Shift h=50.6 Shift k=142.4

More Try All Refine Edit HK Create Img Close



Lattice refinement of 911...

Detect Lattice 'AB'

Ref	H	K
1	1	0
2	0	1

Modify Lattice 'AB'

A*=22.3Å Negate A

B*=22.8Å Negate B

γ*=89.7° Swap AB

HK Edit

h	k	Amp	AmpS	Pha	PhaS	Err
0	1	10000	6259	-0	0	
1	0	2518	6259	0	0	
0	2	1957	1062	-149	180	
2	0	167	1062	176	180	
1	1	3853	2853	14	0	
1	-1	1854	2853	-170	0	P
1	2	371	148	20	0	
1	-2	138	148	117	0	P
2	1	83	148	50	0	P
2	-1	0	148	-156	0	AP

Edit

h = 0

k = 1

Amp = 6259

Pha = 0

Change

Add

Delete

Inverse

(1976,1860) Y=169 Ready

Residuals for c2mm: amplitude: 18.3%, phase angle: 3.4°, all subgroups c1m1, c11m, p2 (*translationsgleiche*), p2gg, p2gm, p2mg, p2mm (*klassengleiche*) have similarly low residuals

The screenshot displays the CRISP2.1 software interface. The main window shows a TEM image of a material with a grid pattern. A white circle is drawn on the image. To the right, the FFT (Fast Fourier Transform) of the image is shown, with a red arrow pointing to a specific spot and a blue line indicating a direction. The FFT window has a title bar that reads "FFT from TEM01full.jpg, 512x...".

Below the main window, there is a panel titled "Origin refinement of TEM01full.jpg". This panel contains a table of symmetry data and a color-coded phase map. The table lists various symmetry groups and their corresponding parameters. The phase map shows a periodic pattern of colors (red, yellow, green, blue) representing phase variations.

At the bottom of the origin refinement panel, there is a table with the following data:

h	k	Amp	AmpS	Pha	PhaS	Err
0	2	314	314	-24	0	
1	1	5876	7938	0	0	
1	-1	10000	7938	-0	0	
2	0	7556	7556	-9	0	
2	2	38	94	-33	0	
2	-2	150	94	25	0	
3	1	163	316	12	0	
3	-1	470	316	3	0	
4	0	57	57	-15	0	

Below the table, there are several buttons: "More", "Try All", "Refine", "Edit HK", "Create Img", and "Close". The status bar at the bottom of the origin refinement panel shows "(-208,60) A=4 P=-168" and "Ready".

The Windows taskbar at the bottom shows the Start button, the CRISP2.1 application icon, and the Microsoft... icon. The system tray on the right shows the time as 11:27 PM and the date as Z.A.

A scale bar in the bottom right corner indicates 10 nm.

CRISP2.1

File Edit Tools Area Fourier CIP ELD PhIDO Statistics Calculate Options Window Help

128 256 512 1K 2K 4K FFT FILE FFT Inv LATT DMAP Ph IDO CCD

1.jpg - 1799x1799x8 (1:3)

Origin refinement of 1.jpg

Symm RA% qRes Ao/Ae

Phase origin map p1 map

Lattice refinement of 1.jpg

Detect Lattice 'AB'

Ref	H	K
1	1	0
2	0	1

Modify Lattice 'AB'

A* = 6.3Å Negate A

B* = 8.2Å Negate B

$\gamma^* = 90.9^\circ$ Swap AB

Radius = 256 Auto

Centred Lattice 'AB'

AB 1 2 3

Modulated lattice

Refine Close

```

p1 - - -
p2 - - 8.5 -
pm m_lx 11.1 4.3 -
pm m_ly 11.1 0.0 -
pg g_lx 11.1 4.3 -
pg g_ly 41.8 0.1 10
cm m_lx 35.4 0.5 4
cm m_ly 35.4 0.0 4
pmm 11.1 8.5 -
pmg m_lx 41.8 14.3 10
pmg m_ly 11.1 8.5 -
pgg 41.8 14.3 10
cmm 35.4 9.7 4
p4
p4m
p4g
p3
  
```

Skewed plane group relations: special group - supergroups

2	-1	0	154	-38	180	AP
2	2	0	65	46	0	AP
2	-2	131	65	-12	0	
3	0	416	416	-80	0	PU
3	1	585	292	1	0	
3	-1	0	292	25	0	A
4	0	134	134	15	0	
4	2	160	80	-47	0	PU
4	2	0	80	176	0	APPT

(1547,1193) Y=159 Ready

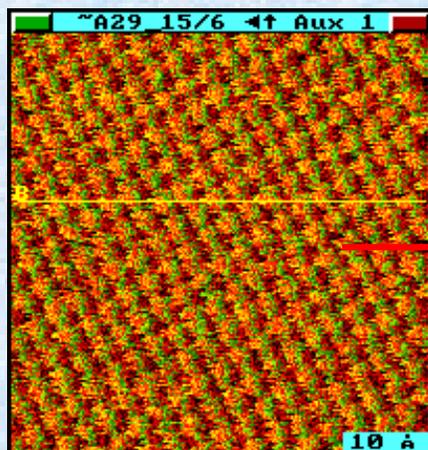
Start Desktop CRISP2.1 Microsoft P... 4:49 PM

there are limits to detecting symmetries in STM images, but 2-fold axis and set of intersecting mirror lines, i.e. subgroup pmm at least identified correctly, skewed symmetry, due to affine transformation, different step sizes, non-orthogonality of fast and slow scan directions

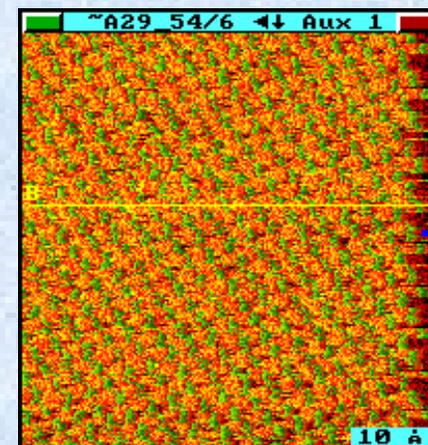
Limits of raw data quality for CIP procedure to work properly

Raw data courtesy Dr. Rachel Cannara, NIST

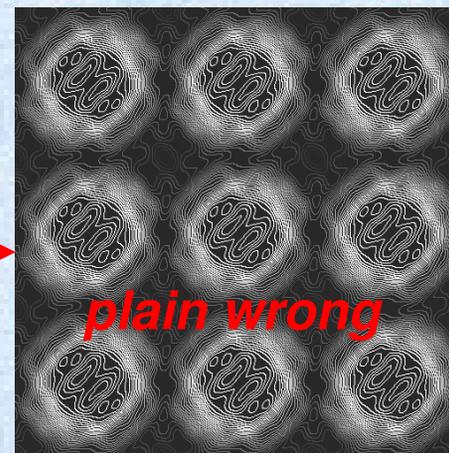
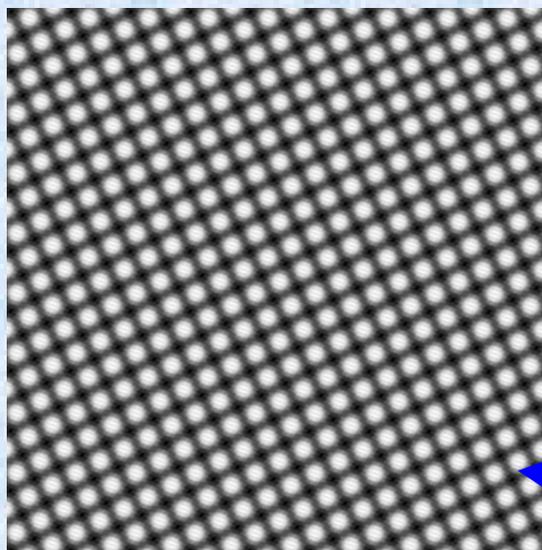
AFM images on LiF, only about 200 by 200 pixels, so “128 pixel masks” are necessary for FFT calculations



128 pixel diameter circular mask

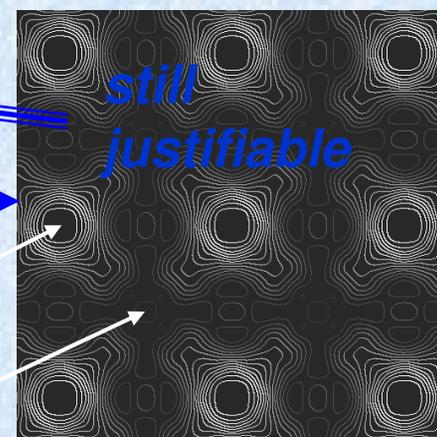


128² pixel square mask

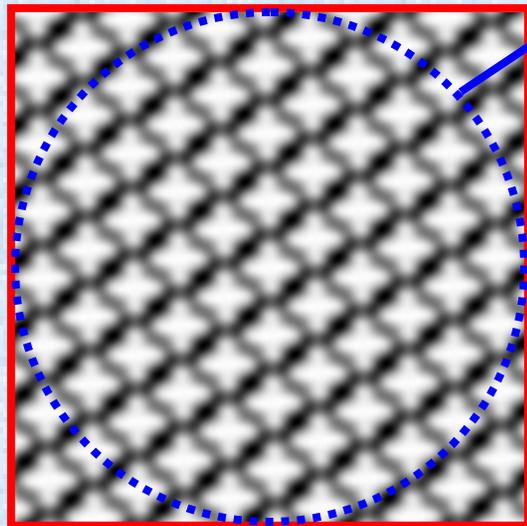


only large F anions are clearly visible, H. Heinzlmann et al., Zeits. Physik B Condensed Matter 79 (1990) 3-4

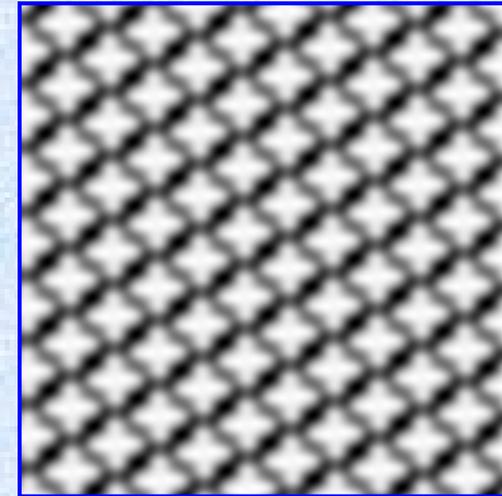
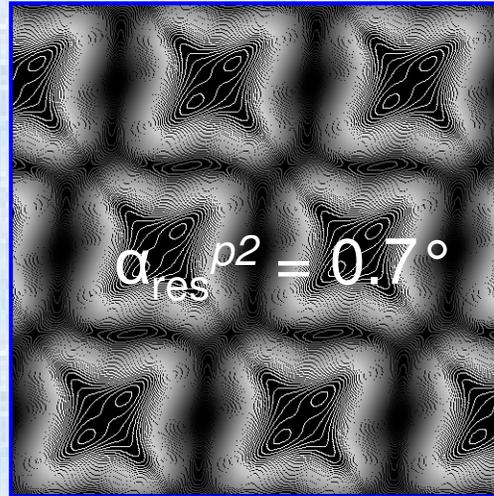
p4mm is detected and enforced for both raw images, but only for the square mask is 4mm enforced as site symmetry of (0,0) and (1/2,1/2) in the detailed counter map



Circular mask for calculation of fast Fourier transform, 256 pixel diameter

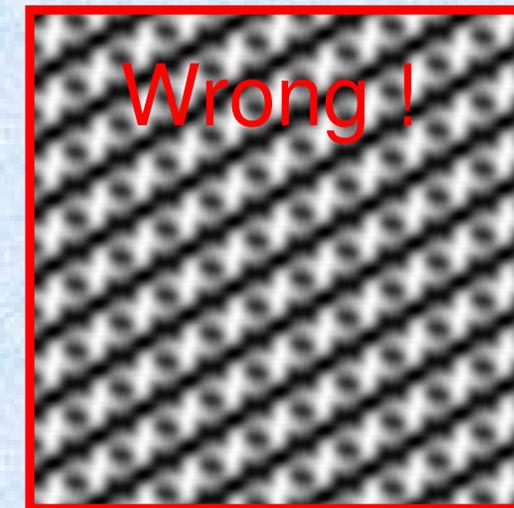
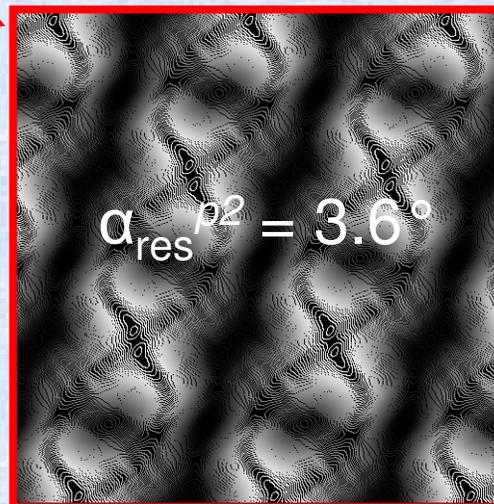


256 pixel square mask for calculation of fast Fourier transform



Avoid inclusion of image edges, example 256 by 256 pixel image

lattices are almost the same, but phase angles of a few "medium strong" Fourier coefficients flipped by about 180° due to convolution of image & mask



Wrong!

“Placebo Test”: Point symmetry 1 combined with oblique lattice = plane symmetry p1, note that all amplitude and phase residuals are very high, crystallographic image processing works well

CRISP2.1

File Edit Tools Area Fourier CIP ELD PhID0 Statistics Calculate Options Window Help

128 256 512 1K 2K 4K FFT PhID0

pegasus.jpg - 804x804x8 (1...)

Origin refinement of pegasus.jpg

Symm RA% qRes Ao/Ae

p2	-	28.5	-
pm	m_lx	53.3	9.6
pm	m_ly	53.3	10.9
pg	g_lx	59.2	10.5
pg	g_ly	53.3	9.4
cm	m_lx	60.2	10.4
cm	m_ly	60.2	8.1
pmm		53.3	32.2
pmg	m_lx	53.3	34.0
pmg	m_ly	59.2	33.6
pgg		59.2	35.6
cmm		60.2	31.0
p4		63.7	32.3
p4m		91.2	36.9
p4g		90.6	36.5
p3			
p3m1			
p31m			

a=77.7Å b=76.0Å γ=88.0° Shift h=0.0 Shift k=0.0

HK Edit

h	k	Amp	AmpS	Pha	PhaS	Err
0	1	4605	4605	-99	-99	
0	3	766	766	-138	-138	
0	6	719	719	94	94	
0	8	67	67	-166	-166	
0	9	132	132	124	124	
0	10	238	238	-125	-125	
1	1	10000	10000	-25	-25	
1	-1	217	217	6	6	
1	3	1676	1676	-64	-64	
1	5	253	253	-14	-14	
1	-5	1092	1092	-116	-116	
1	8	187	187	-140	-140	
1	-8	645	645	136	136	
1	-9	535	535	-24	-24	
1	-10	160	160	14	14	
2	0	1706	1706	173	173	

Edit

h = 0

k = 1

Amp = 4605

Pha = -99

Change

Add

Delete

Inverse

Amp Threshold

50 Apply

Manual Shift

Shift h = 0.0

Shift k = 0.0

Crystal tilt

Axis 'a' =

k = -87.9 Estimate

(748,750) Y=92

Ready

Start

Inbox ...

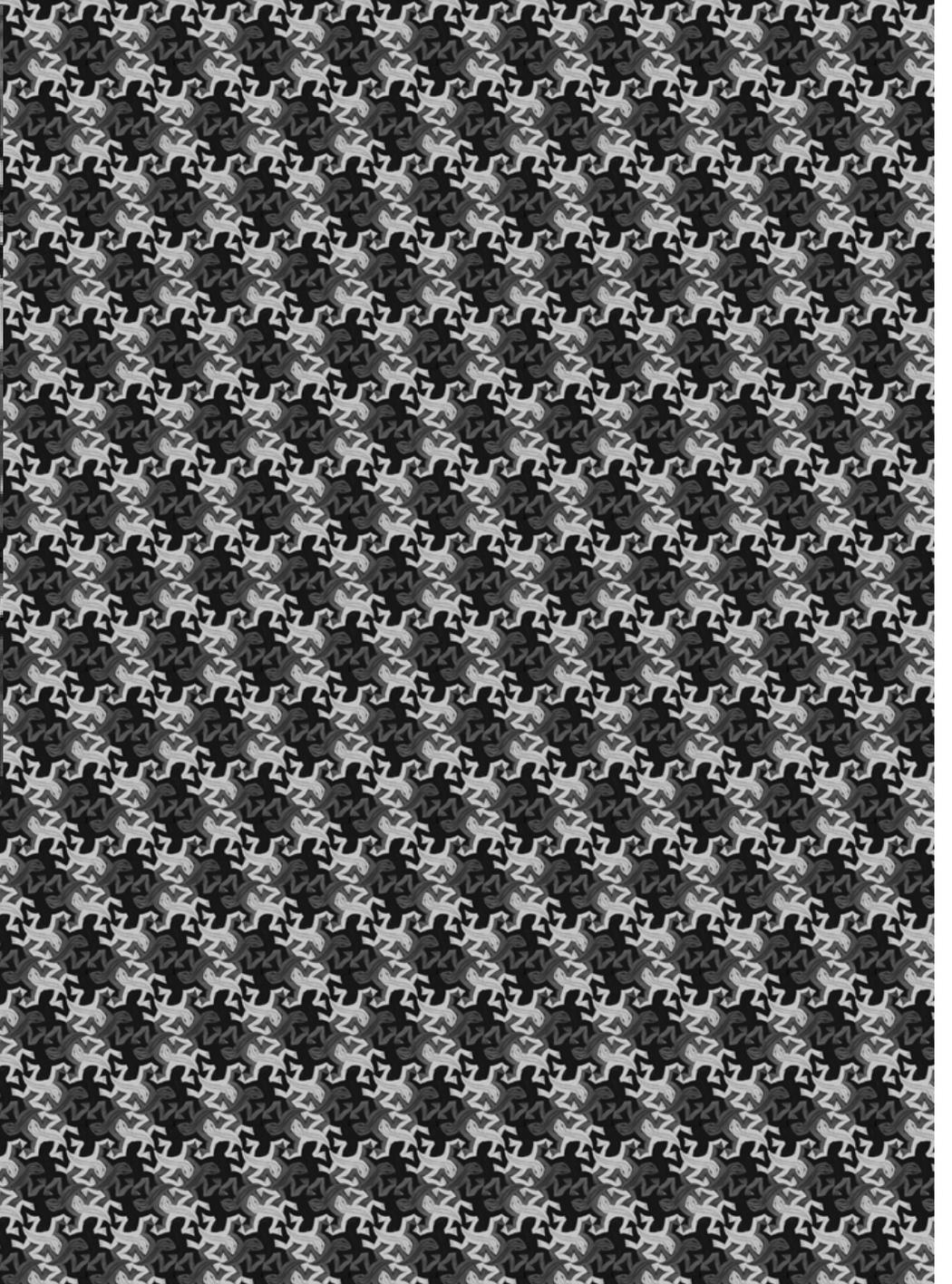
4 Mi...

2 Int...

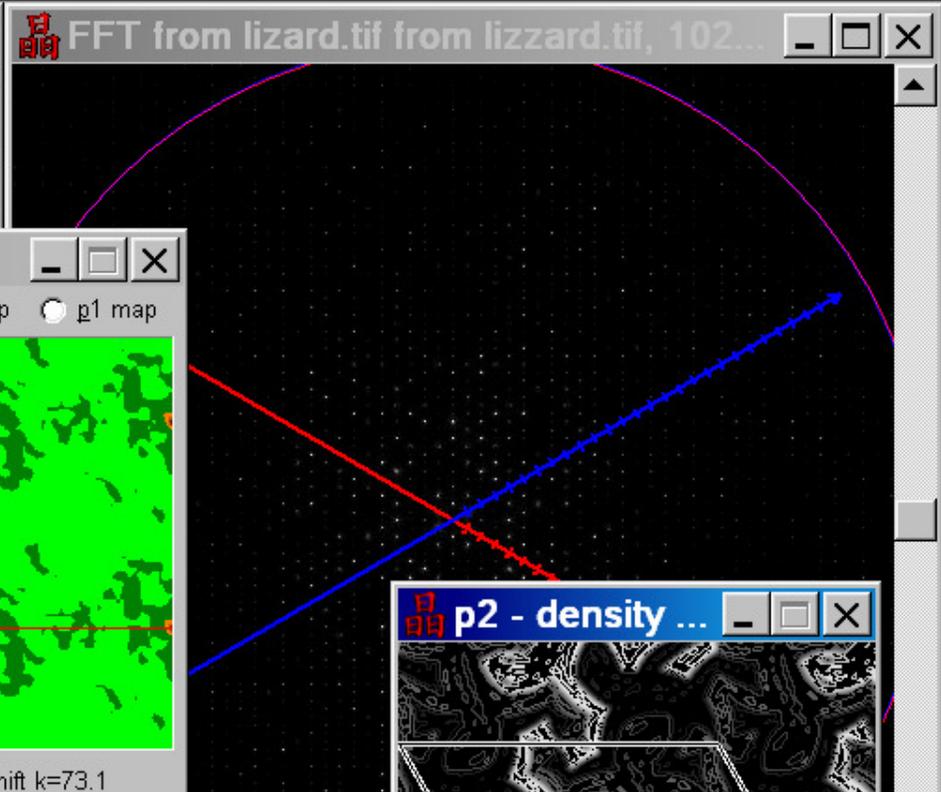
C:\Do...

CRIS...

7:43 PM



Escher's "E 56", only $p2$, four sets of two-fold axes, although the lattice seems to be close to hexagonal, if gray levels in motif were discarded (incorrectly) one would obtain $p6$



Origin refinement of lizzard.tif

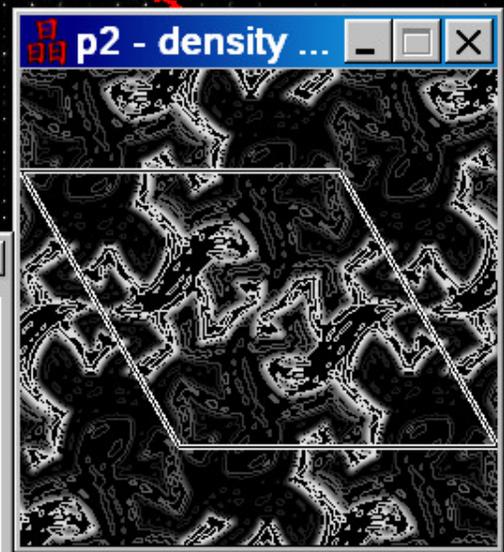
Symm RA% σ Res Ao/Ae

Phase origin map p1 map

p1	-	-	-
p2	-	7.8	-
pm	m x	-	-
pm	m y	-	-
pg	g x	-	-
pg	g y	-	-
cm	m x	-	-
cm	m y	-	-
pmm	-	-	-
pmg	m x	-	-
pmg	m y	-	-
pgg	-	-	-
cm	-	-	-
p4	-	-	-
p4m	-	-	-
p4g	-	-	-
p3	42.1	38.9	-
p3m1	46.2	50.0	-
p31m	46.2	49.7	-
p6	42.1	40.6	-
p6m	46.2	50.7	-

a=67.3Å b=67.3Å γ =119.8° Shift h=0.0 Shift k=73.1

More



Lattice refinement of lizz...

Detect Lattice 'AB'

Ref	H	K
1	1	0
2	0	1

Modify Lattice 'AB'

A*=17.5Å Negate A

B*=17.5Å Negate B

γ *=60.2° Swap AB

+0.965,+1.365, Den=158 Ready

Tune Density Map

Visualization

a offset b offset

Cells Rotation

Negative

Cell edge

Contour map Steps

Cell

a= 67.32Å

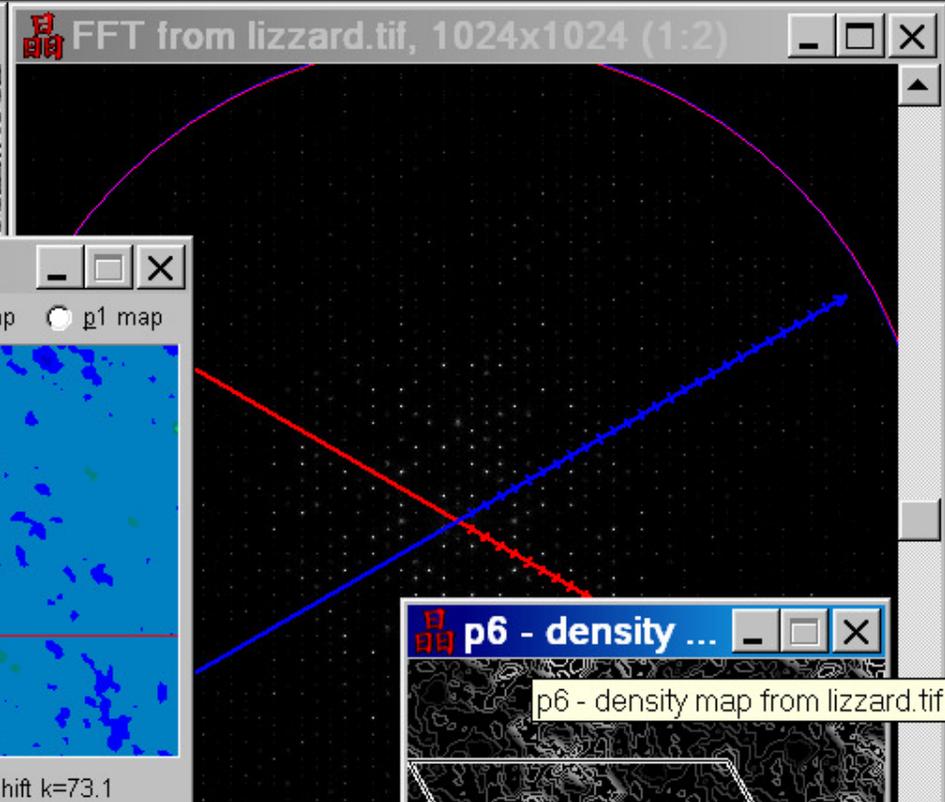
b= 67.32Å

γ = 119.8°

a-x angle=-120.1°

Size

X



Origin refinement of lizzard.tif

Symm RA% ϕ Res Ao/Ae

Phase origin map p1 map

p1	-	-	-
p2	-	7.8	-
pm	m x		
pm	m y		
pg	g x		
pg	g y		
cm	m x		
cm	m y		
pmm			
pmg	m x		
pmg	m y		
pgg			
cmm			
p4			
p4m			
p4g			
p3	42.1	38.9	-
p3m1	46.2	50.0	-
p31m	46.2	49.7	-
p6	42.1	40.6	-
p6m	46.2	50.7	-

a=67.3Å b=67.3Å γ =120.0° Shift h=-0.4 Shift k=73.1

More

p6 - density ...

p6 - density map from lizzard.tif

Tune Density Map

Visualization

a offset b offset

Cells Rotation

Negative

Cell edge Contour map Steps

Cell

a= 67.32Å
b= 67.32Å
 γ = 120.0°
a-x angle=-120.1°

Size X

Lattice refinement of lizzard.tif

Detect Lattice 'AB'

Ref	H	K
1	1	0
2	0	1

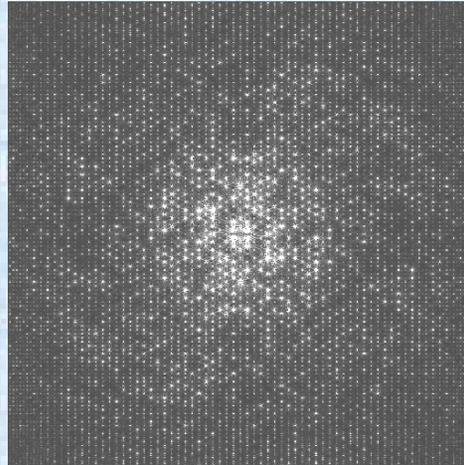
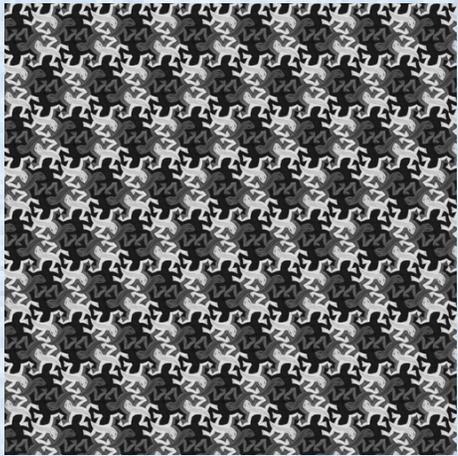
Modify Lattice 'AB'

A*=17.5Å Negate A

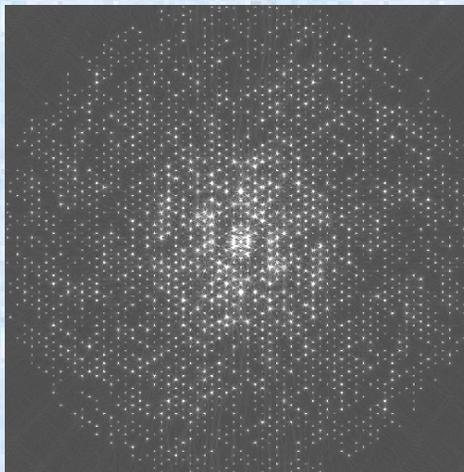
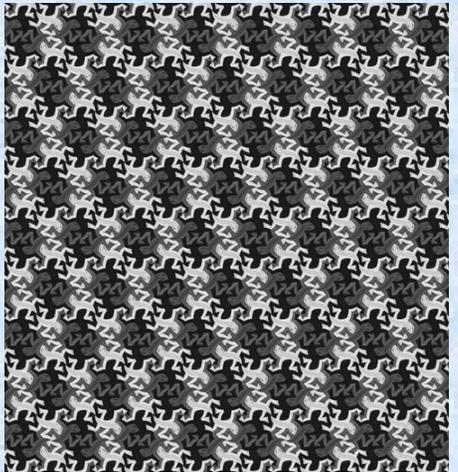
B*=17.5Å Negate B

γ *=60.2° Swap AB

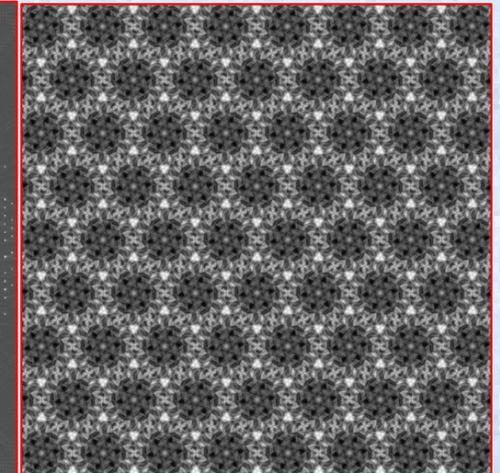
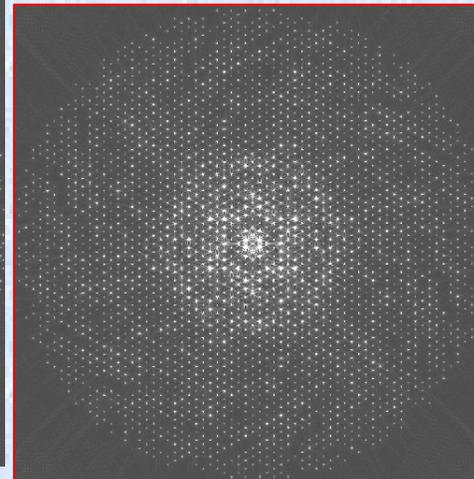
+0.886,+1.366, Den=559



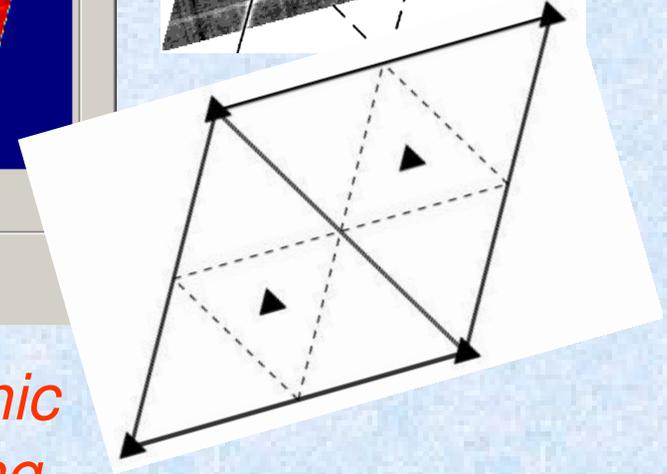
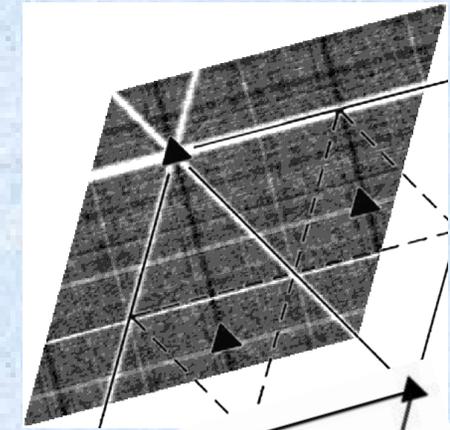
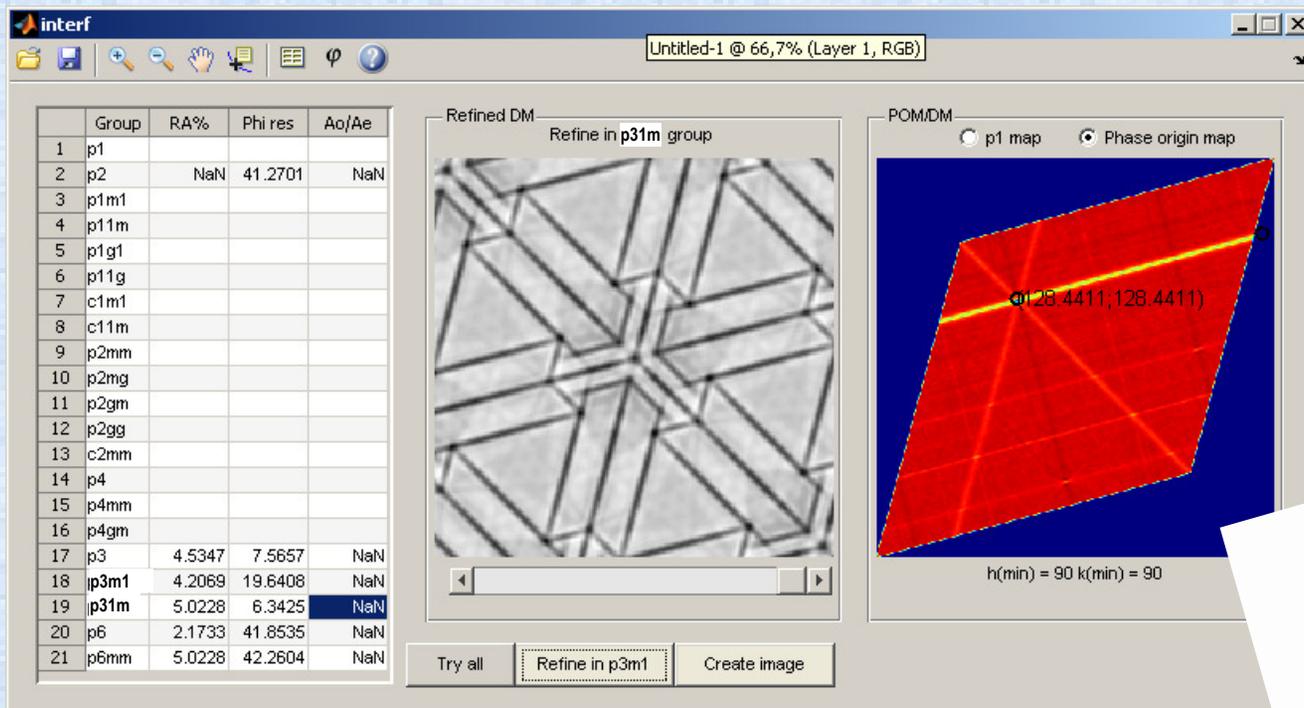
p1 version, no point symmetry elements besides 1, only translation symmetry of an oblique lattice, two-fold axis in reciprocal space for maps of amplitudes as property of Fourier transform



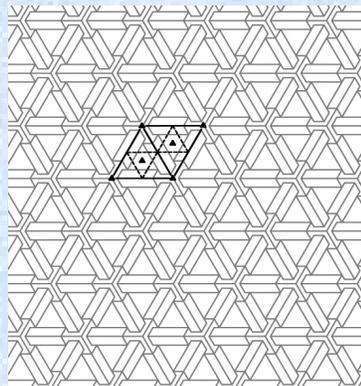
p2 enforced version, all phase angles 0 or 180 degrees, origin at the position of one set of these two-fold axes, one two-fold axis in maps of amplitudes, due to combined effect of symmetry in the *p2* image and Fourier transform



p6 enforced version, all phases 0 or 180 degrees, origin at the position of the six fold axis, almost same phase shift as for *p2*, three two-fold axes and two three-fold axes are also produced. Six-fold axis in map of Fourier coefficients amplitudes



development of a dedicated crystallographic image processing program for the scanning probe microscopy community

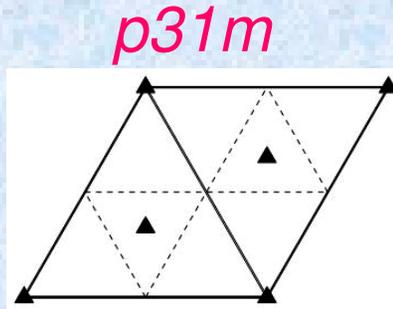
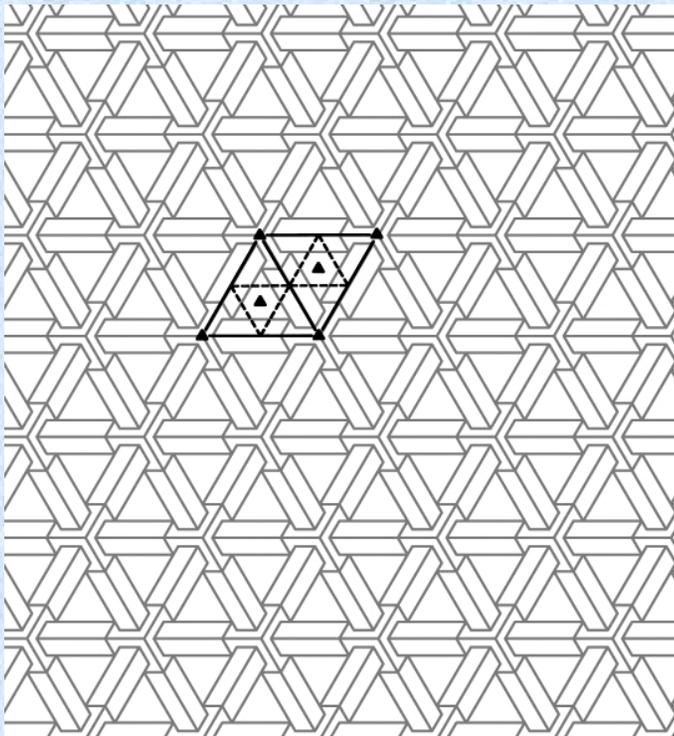


For “sufficiently” symmetric images without much distortion and noise, e.g. simulations or works of art, the map of the phase residuals indicates not only the correct origin, but also the location of the other symmetry elements,

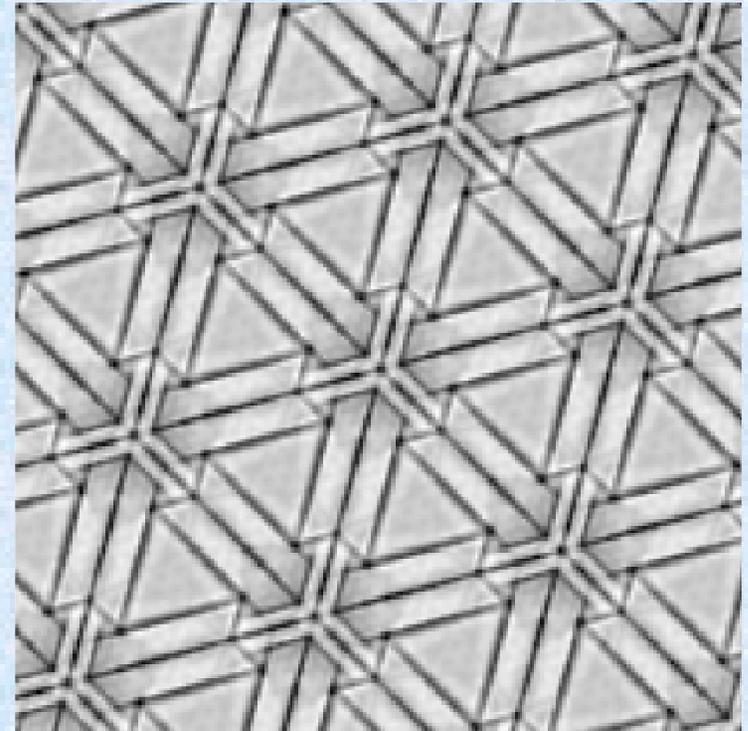
in case of $p31m$, the mirror lines, glide lines, and position of mirror equivalent three-fold rotation points at $(\frac{1}{3}, \frac{2}{3})$ and $(\frac{2}{3}, \frac{1}{3})$

Cross correlation of a 2D periodic image with the symmetrized tile

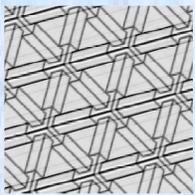
(that is Fourier back transformed after symmetrization of the Fourier coefficients of the image)



Idea: relative height of the central peak of CCF quantifies “similarity”, *p1* value serves as a “baseline” of the FFT procedures

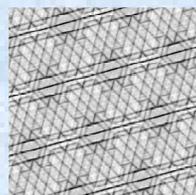


$$CCF(x, y) = \frac{\sum_{x'} \sum_{y'} [i(x', y') - \langle i(x', y') \rangle] \cdot [t(x'-x, y'-y) - \langle t \rangle]}{(\sum_{x'} \sum_{y'} [i(x', y') - \langle i(x', y') \rangle]^2 \cdot \sum_{x'} \sum_{y'} [t(x'-x, y'-y) - \langle t \rangle]^2)^{0.5}}$$



p1, 1.14

all site symmetries broken for p2



p2, 3.68



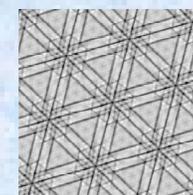
p3, 4.21



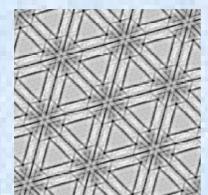
p3m1,
4.31



p31m,
4.48

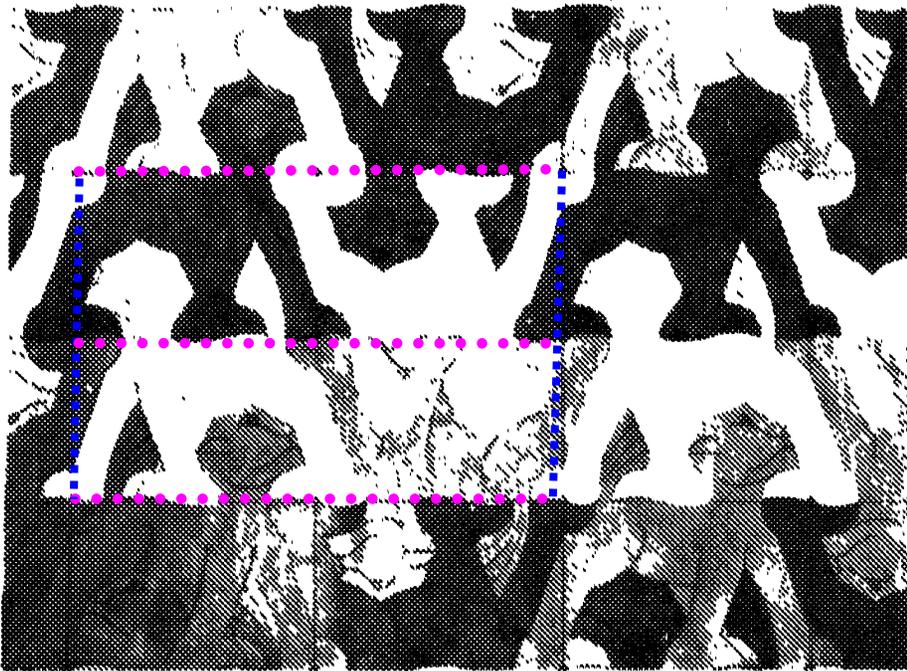


p6, 4.10



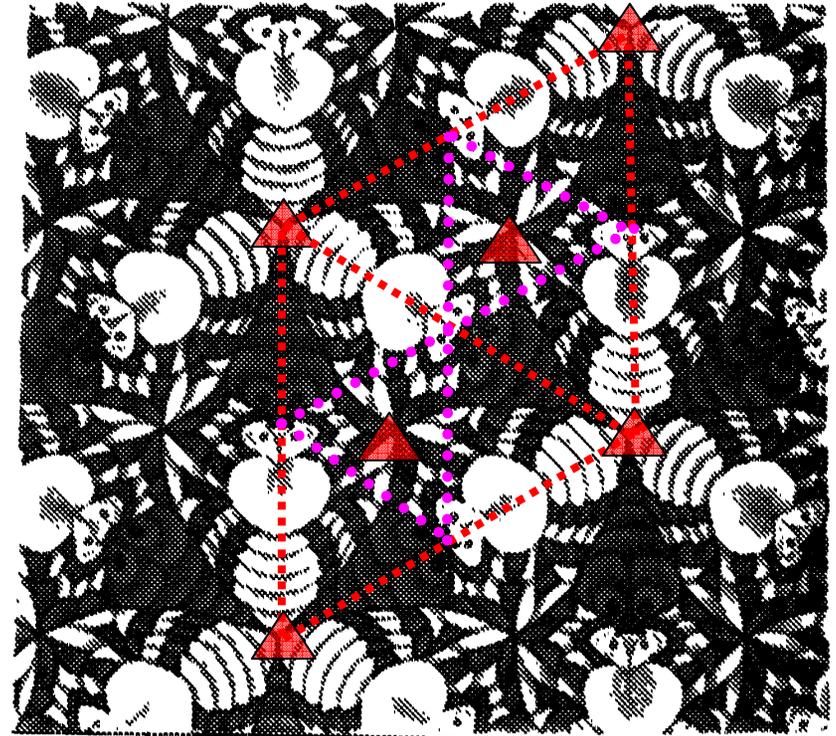
p6mm,
4.29

Exercise in mirror and glide lines plus three axes



pg unfinished

Escher Foundation, Haags Gemeentemuseum, The Hague.



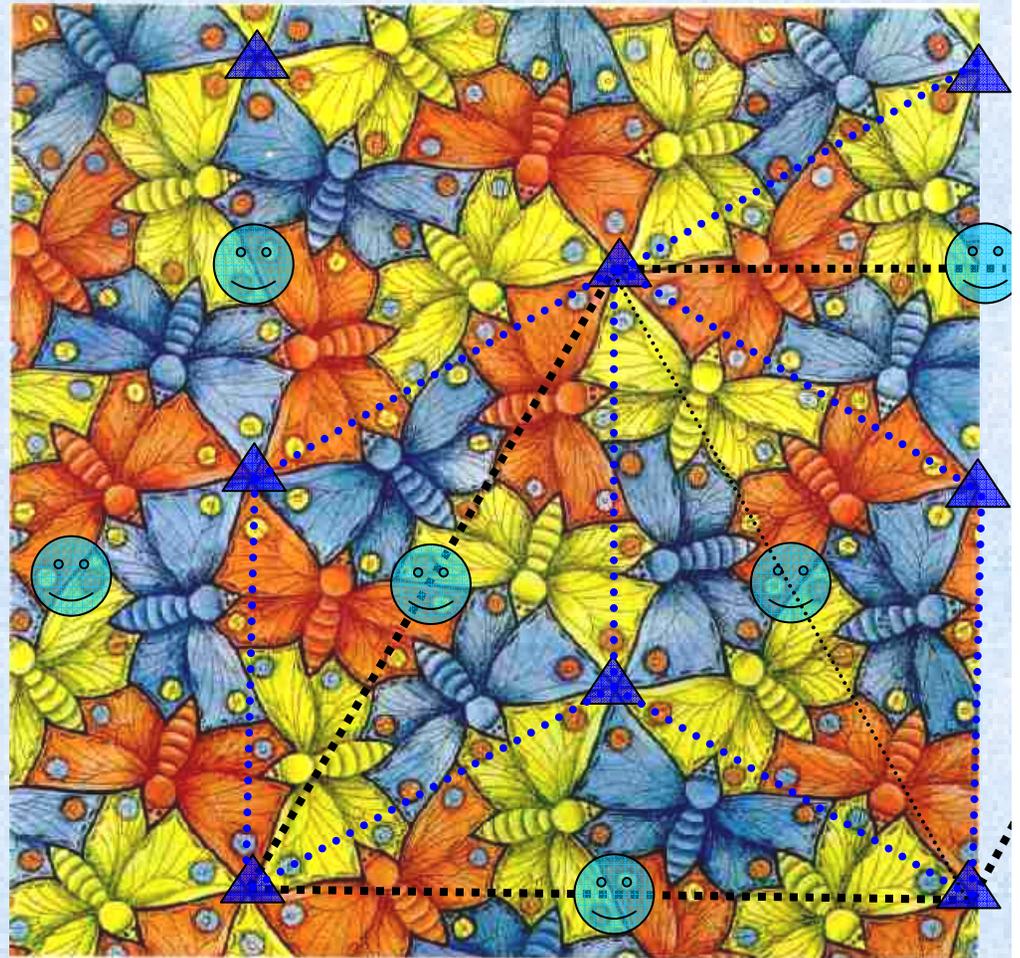
p31m

Two periodic drawings by M. C. Escher contrast his early effort at repeating design with his later masterful skill. The pattern of lions, dated “1926 or 1927,” was done before he developed a system which grew out of his study of ~~mathematical~~ articles and periodic designs on the Alhambra. The pattern of bugs is dated 1942, one year after Escher recorded his codified system in notebooks.

G. Pólya, Über die Analogie der Kristallsymmetrie in der Ebene, *Zeits. Krist.* 60 (1924) 278-282;
P. Niggli, Die Flächensymmetrien homogener Diskontinuen, *Zeits. Krist.* 60 (1924) 283-298.

D. Schattschneider, *The American Mathematical Monthly* 85 (1978) 439-450

almost square lattice, but no point symmetries at all, very special case of p1



*p6 deliberately broken (removing two fold axes ☺) by coloring so that drawing becomes **special case of p3**,*

18 moths in total per unit cell, half of a unit cell – a triangle - clearly visible, so let's account for 9 moths, three are blue the, other six are red and yellow, asymmetric unit – dotted blue lines - contains two moth of each color

Summary and conclusions

All periodic 2D images can be classified to belong to one of only 17 plane symmetry groups

Deviation from each of these groups can be quantified, correct plane group can often be identified on basis of quantitative measures (plus some prior knowledge), i.e. the amplitude and phase angle residuals of the Fourier coefficients of the image intensity

HOPG is a good calibration sample, one can determine the point spread function of the microscope on its basis, then the same function can be utilized to correct images from unknowns, allows for better measurements and surface science

A blunt STM tip can be symmetrized

There are problems on how to decide on the correct plane group if there is no prior knowledge, some clever geometric AIC is needed

Knowledge of 2D symmetry made Maurits Cornelis Escher (17 June 1898 – 27 March 1972) a more sophisticated artists, it may also result in better surface- and nano-science

Crystallographic Image Processing works even for noisy Fourier transforms

CRISP2.1

File Edit Tools Area Fourier CIP ELD PhIDO Statistics Calculate Options Window Help

128 256 512 1K 2K 4K FFT FILE FFT Inv LATTI UNMAP Ph IDO 1D CCD

escher4.jpg - 847x654x8 (1:2)

Origin refinement of escher4.jpg

Symm RA% qRes Ao/Ae

Phase origin map
 p1 map

p1				
p2				
pm	m _↓ x			
pn	m _↓ y			
pg	g _↓ x			
pn	g _↓ y			
cm	m _↓ x			
cn	m _↓ y			
pmn				
png	m _↓ x			
pn	m _↓ y			
pgg				
cmn				
p4				
p4m				
p4g				
p3		58.2	18.8	-
p3m		76.5	35.1	-
p31m		76.5	36.8	-
p6		59.2	51.6	-
p6m		83.7	56.1	-

a=121 Å b=121 Å γ=120.0° Shift h=-140.6 Shift k=28.1

More

FFT from escher4...

Lattice refinement of esc...

Detect Lattice 'AB'

Refl	H	K
1	1	0
2	0	1

Radius=256 Auto

Modify Lattice 'AB'

A*=4.0Å Negate A

B*=8.0Å Negate B

γ*=52.8° Swap AB

Centred Lattice 'AB'

AB 1 2 3

Origin Refinement

Manual origin Manual Shift

Restrictions Shift h= 0.0

Relations Shift k= 0.0

Weighted Amp. threshold 50.0

Symmetry names

Like png

Like p2212

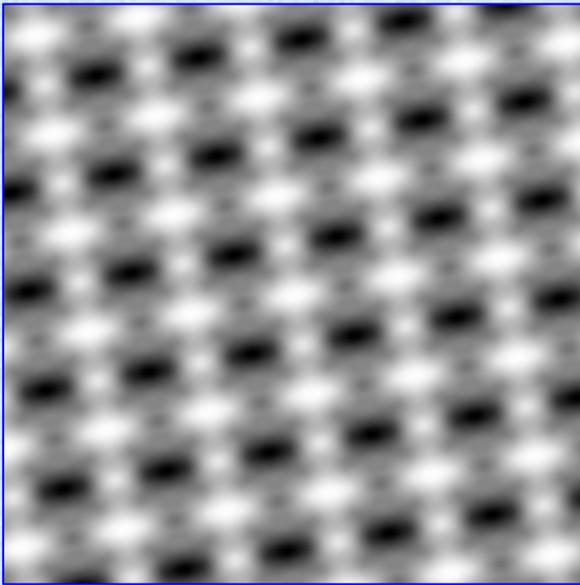
Crystal tilt

Tilt axis^a*=

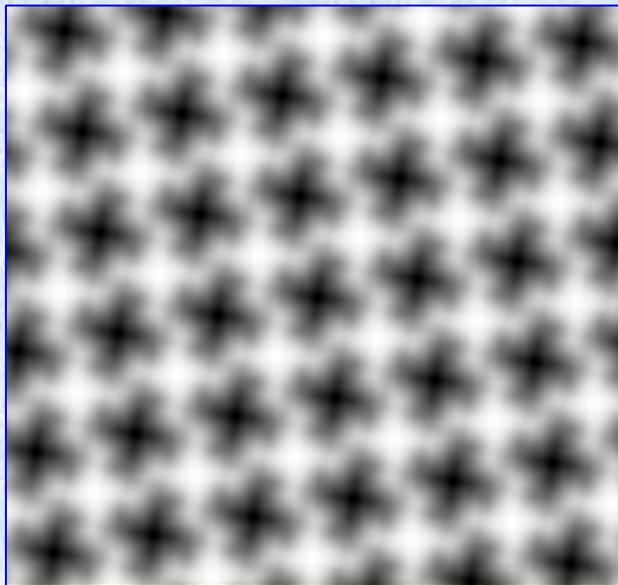
a^ax = -0.2

(130,-87) A=7 P=-153 Ready

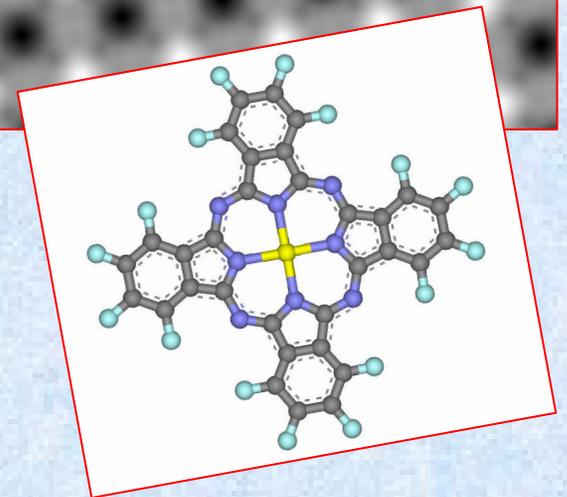
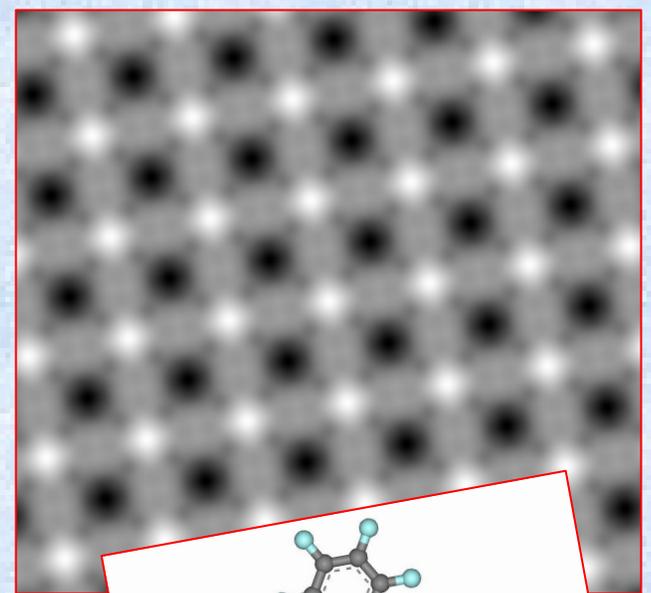
Start 3 Na... An im... 2 Wi... Micros... CRIS... 8:37 PM



p2mm symmetry enforced STM data. A two-fold axis with a set of perpendicular mirrors is imposed. The lattice becomes rectangular. There is no 4, so the molecules ‘line up’ at two of the 2mm positions.

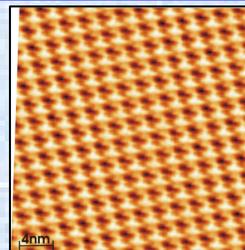


p4 symmetry enforced STM data. A four-fold axis is imposed. The lattice becomes quadratic. Because the “azimuthal orientations” of the molecules are not restricted, there is molecular misalignment and “washed out detail”



“Size” of molecule and basic periodicities in STM images approximately 1.5 – 2 nm

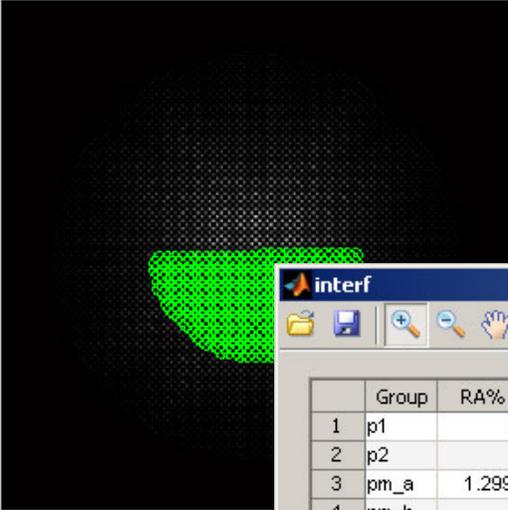
In effect, the “imperfections” of the scanning probe imaging process by a real microscope are removed from the data by imposing the most likely symmetry ! This is possible because all molecules are quantum mechanically identical and we may know the point group of a molecule in advance from quantum mechanics!



p4mm symmetry enforced STM data. Two additional sets of perpendicular mirror lines are imposed. The motif “rotates” counter clockwise and acquires the full point symmetry of the molecule.

FFT_import

128 256 512 1024



Display mode

- Oblique
- Orthorhombic
- Centered/Hexagonal

Lattice parameters

	h	k	F	phi
1	2		-2 1.3560e+06	-2.9560
2	2		2 1.3550e+06	-2.9559
3	4		0 9.4551e+05	-0.0494
4	0		4 9.2498e+05	1.4724e-05
5	8		0 6.5967e+05	-2.9176
6	0		8 6.5674e+05	-3.1415
7	-4		6 5.0794e+05	2.8633
8	4		6 5.0668e+05	0.2783
9	8		4 5.0468e+05	2.8656

FFT options

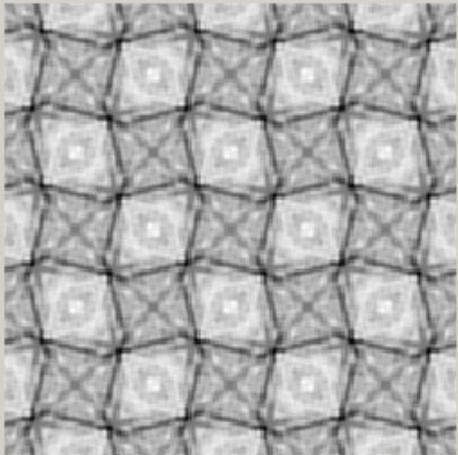
Open >>

interf

	Group	RA%	Phi res	Ao/Ae
1	p1	0	0	0
2	p2	0	5.5581	NaN
3	pm_a	1.2990	26.6806	NaN
4	pm_b	0	0	0
5	pg_a	0	0	0
6	pg_b	0	0	0
7	cm_a	0	0	0
8	cm_b	0	0	0
9	pmm	1.2990	45.8867	NaN
10	pmga	0	0	0
11	pmgb	0	0	0
12	pgg	0	0	0
13	cmm	0	0	0
14	p4	3.1565	8.5711	NaN
15	p4m	5.1251	30.0690	NaN
16	p4g	5.1251	12.6005	0.1612
17	p3	0	0	0
18	p31m	0	0	0
19	p3m1	0	0	0
20	p6	0	0	0
21	p6mm	0	0	0

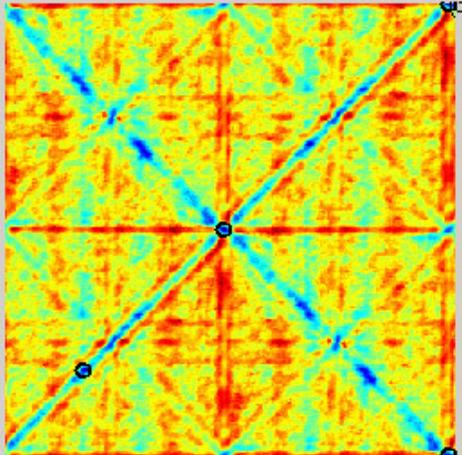
Refined DM

Refine in p4g group



POMDM

p1 map Phase origin map



h(min) = -180 k(min) = 175

Try all Refine in p4g Create image

Import Reset stat Save

