

PowerPoint File available:

**[http://bl831.als.lbl.gov/~jamesh/powerpoint/
IGBMC_SvN_2016.pptx](http://bl831.als.lbl.gov/~jamesh/powerpoint/IGBMC_SvN_2016.pptx)**

Acknowledgements

Robert Stroud James Fraser John Spence
Chris Nielsen Clemens Schulze-Briese Aina Cohen Ana Gonzalez

UCSF LBNL SLAC

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UC Multicampus Research Programs and Initiatives (MRPI)

UCSF Program for Breakthrough Biomedical Research (PBBR)

one-time NIH-DOE Inter-agency agreement (IAA)

Integrated Diffraction Analysis Technologies (IDAT)

Plexxikon, Inc.

M D Anderson CRC

Synchrotron Radiation Structural Biology Resource (SLAC)

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Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

The “success rate” of structure determination

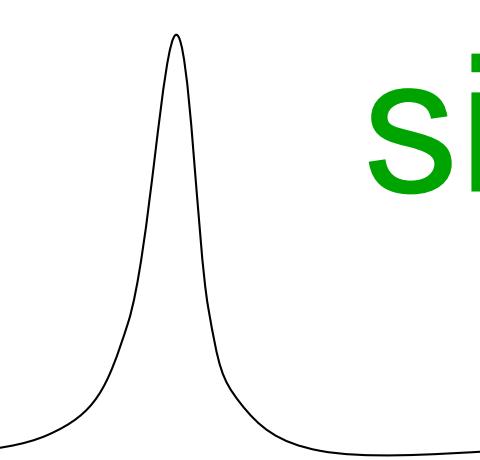
100 s/dataset

200 days/year

~150 beamlines

~26,000,000 datasets/year

9333 PDBs in 2015

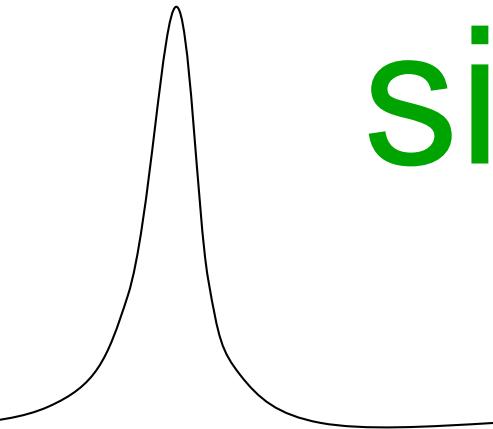


signal

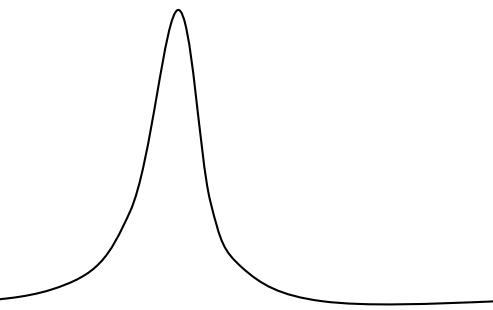
“If you don’t have
good data,
then you have
no data at all.”

-Sung-Hou Kim

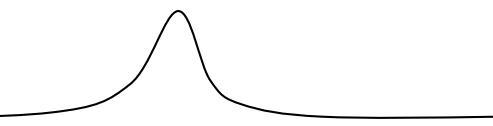
signal vs noise



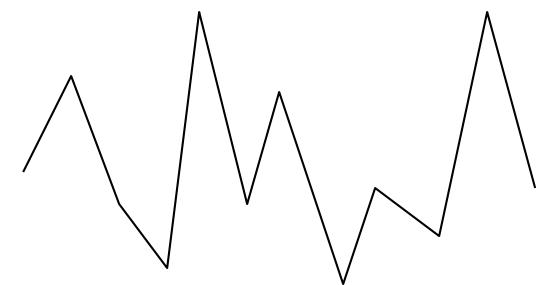
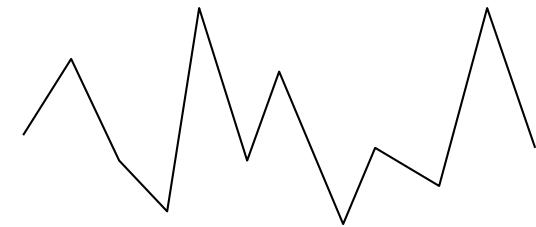
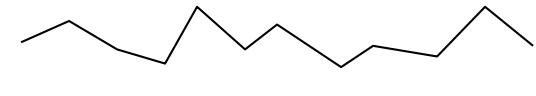
easy



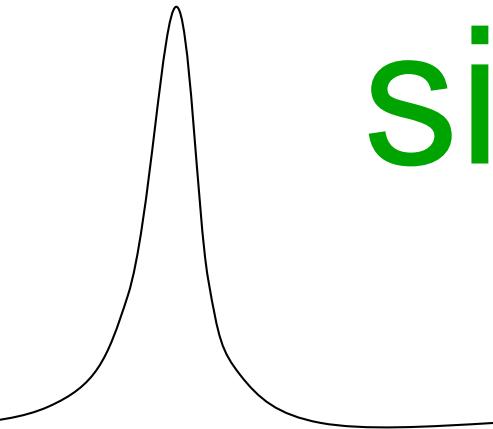
hard



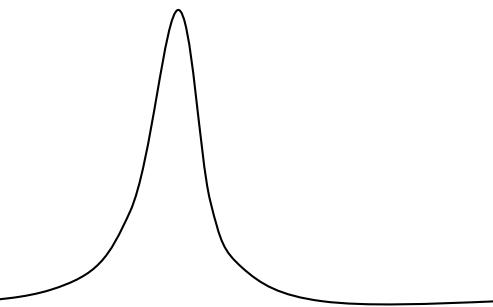
impossible



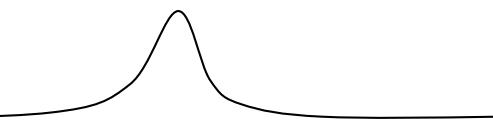
signal vs noise



easy

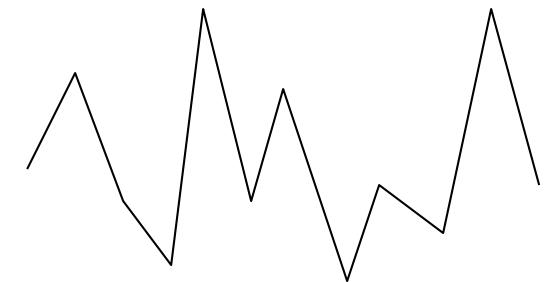
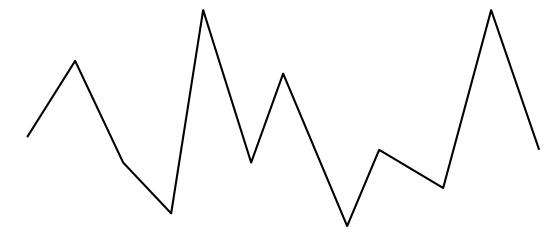
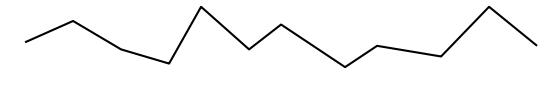


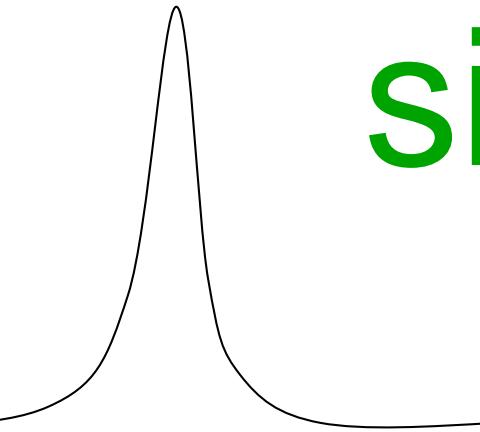
hard



threshold of “solvability”

impossible



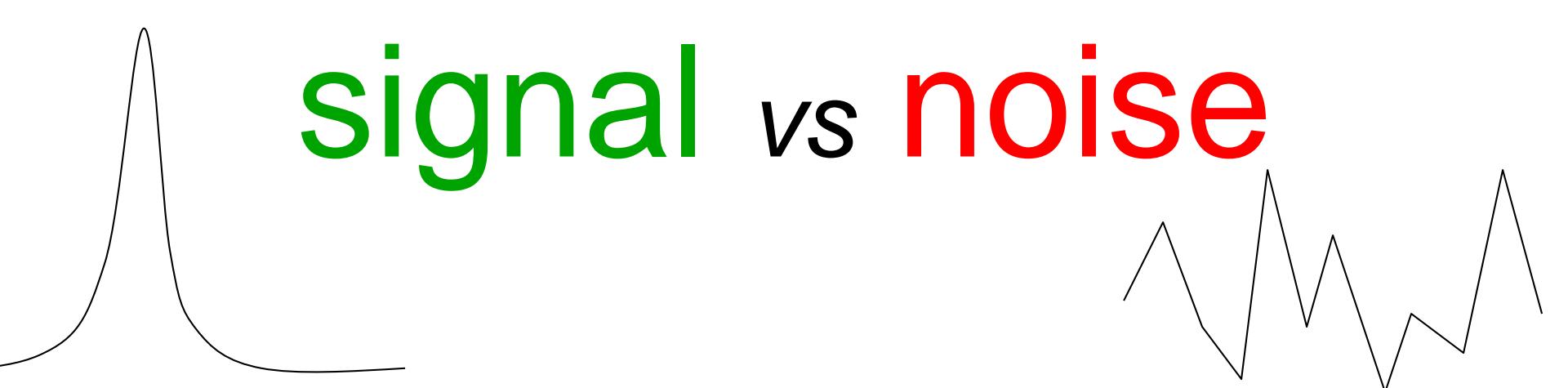


signal / noise =

$$\sqrt{\text{photons}} ?$$


	none	sqrt	proportional
none	CCD Read-out	Photon counting	Detector calibration attenuation partiality Non-isomorphism Radiation damage
1/sqrt			Beam flicker
1/prop.			Shutter jitter Sample vibration Pile-up

Time



signal vs noise

“If you don’t have
good data,
then you must
learn statistics.”

-James Holton

Adding noise

Adding noise

$$1 + 1 = 1.4$$

Adding noise

$$1 + 1 = 1.4$$

$$\sigma_{\text{total}}^2 = \sigma_1^2 + \sigma_2^2$$

Adding noise

$$1^2 + 1^2 = 1.4^2$$

$$\sigma_{\text{total}}^2 = \sigma_1^2 + \sigma_2^2$$

Adding noise

$$1^2 + 1^2 = 1.4^2$$

$$3^2 + 1^2 = 3.2^2$$

$$\sigma_{\text{total}}^2 = \sigma_1^2 + \sigma_2^2$$

Adding noise

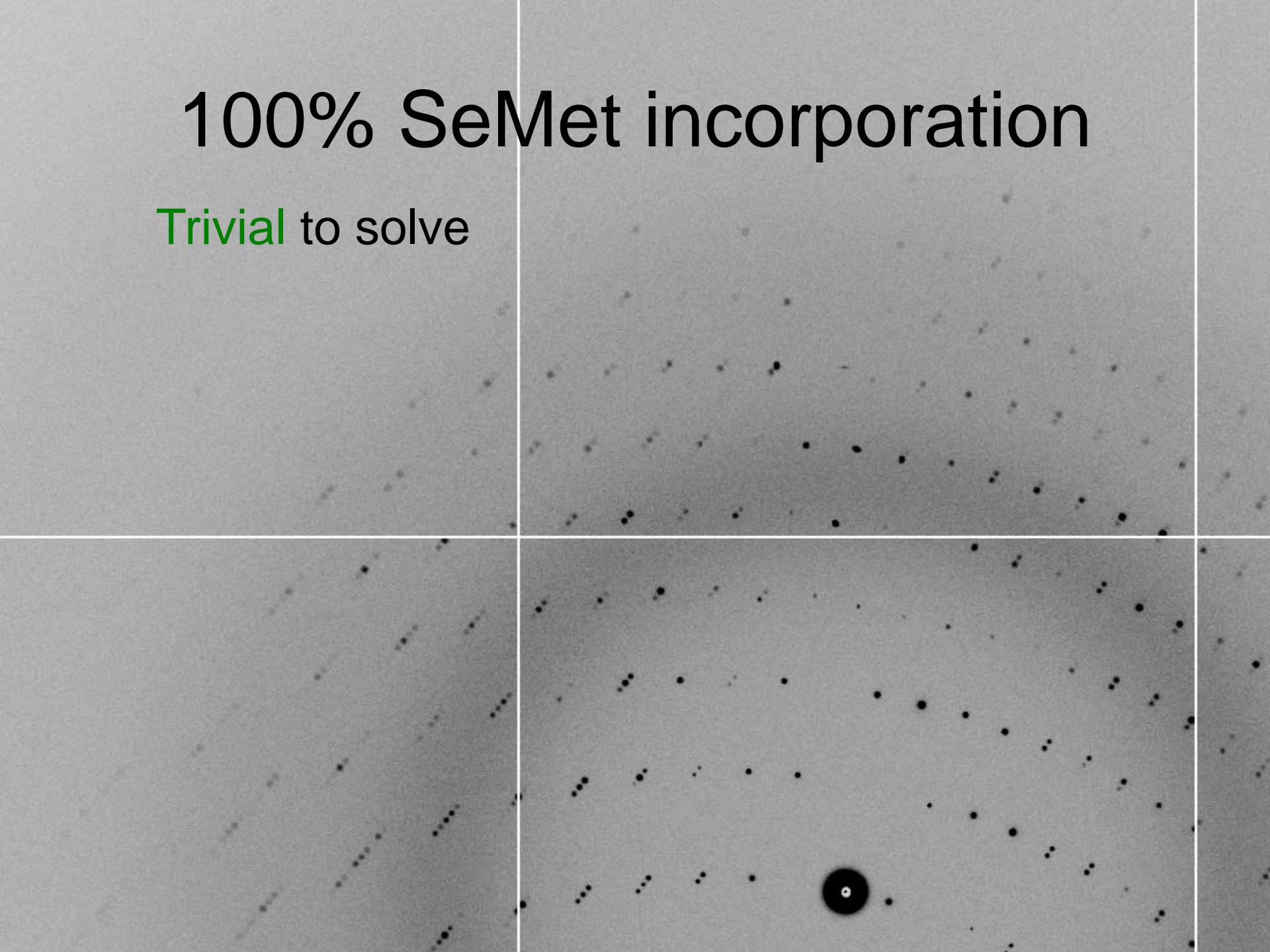
$$1^2 + 1^2 = 1.4^2$$

$$3^2 + 1^2 = 3.2^2$$

$$10^2 + 1^2 = 10.05^2$$

100% SeMet incorporation

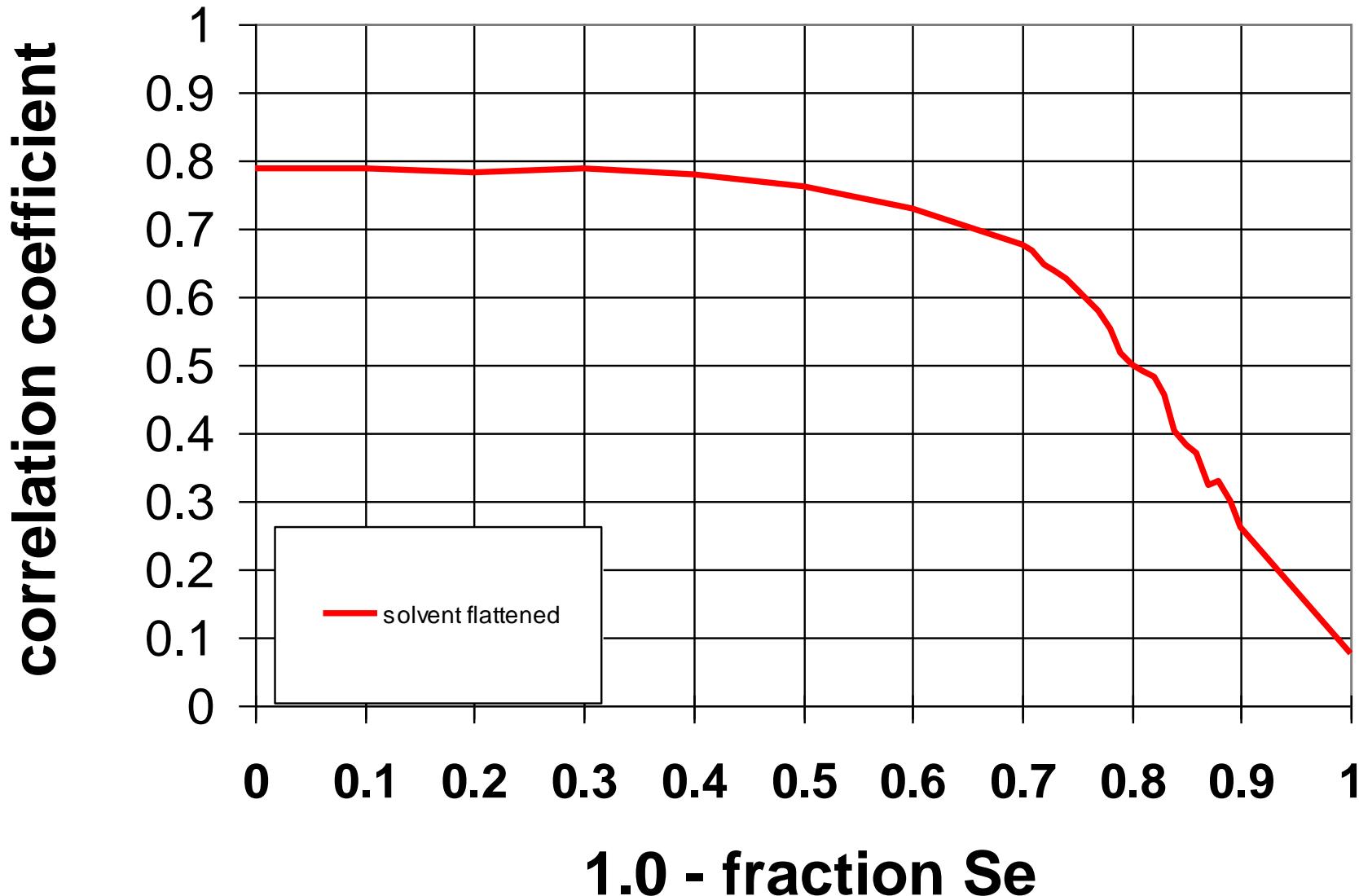
Trivial to solve



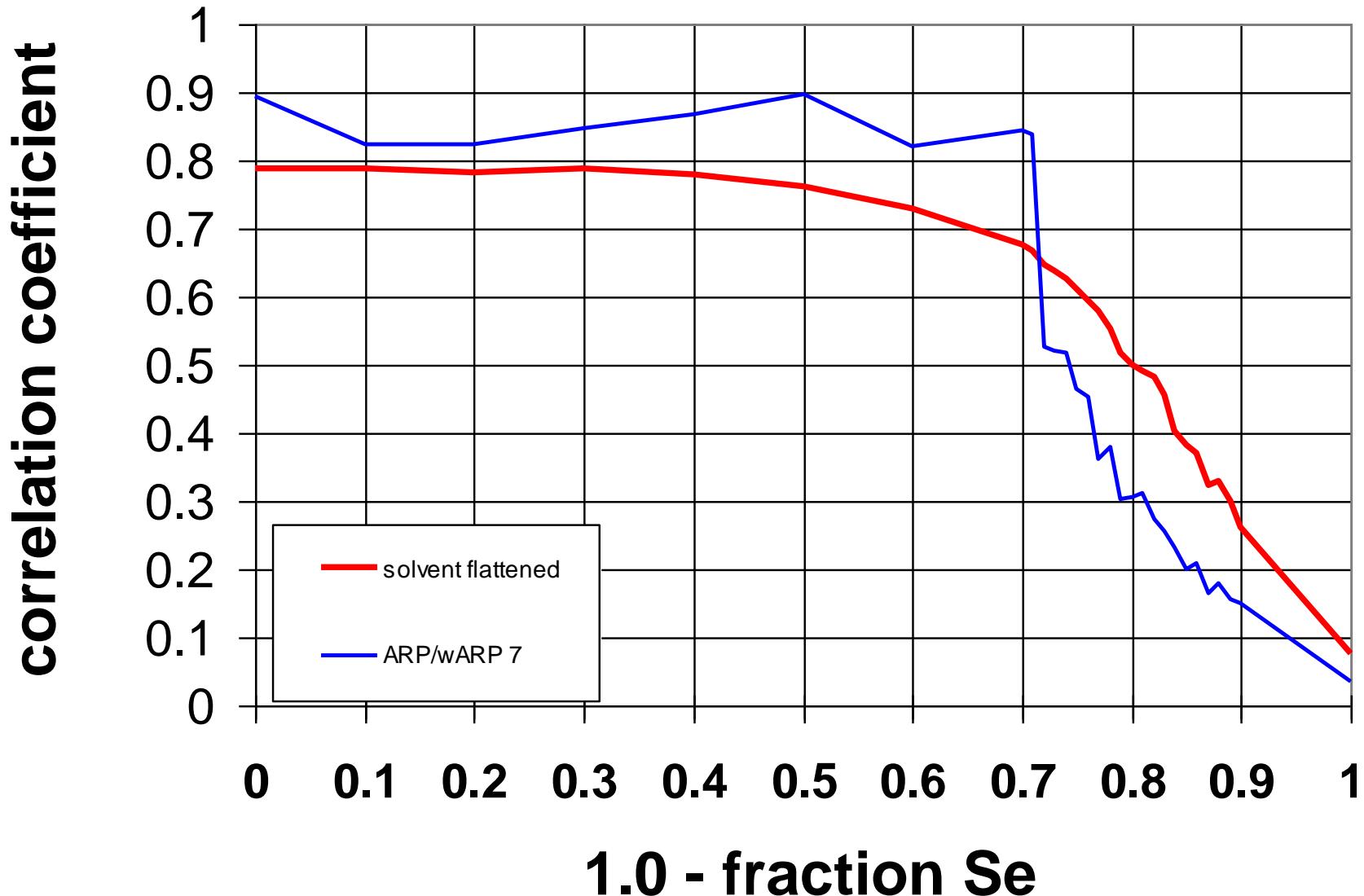
100% S -Met incorporation

Impossible to solve

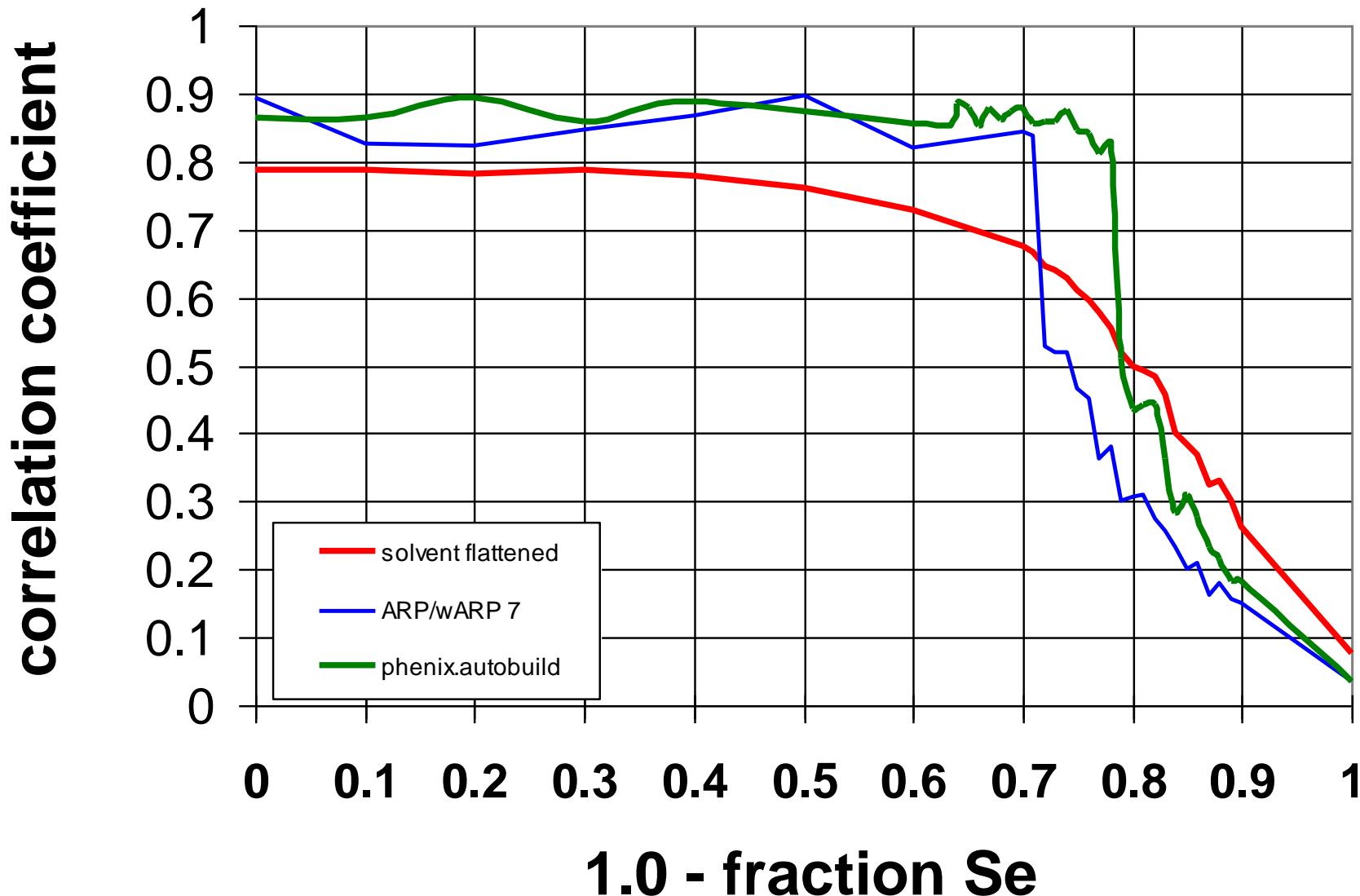
Phase Problem: Se vs S



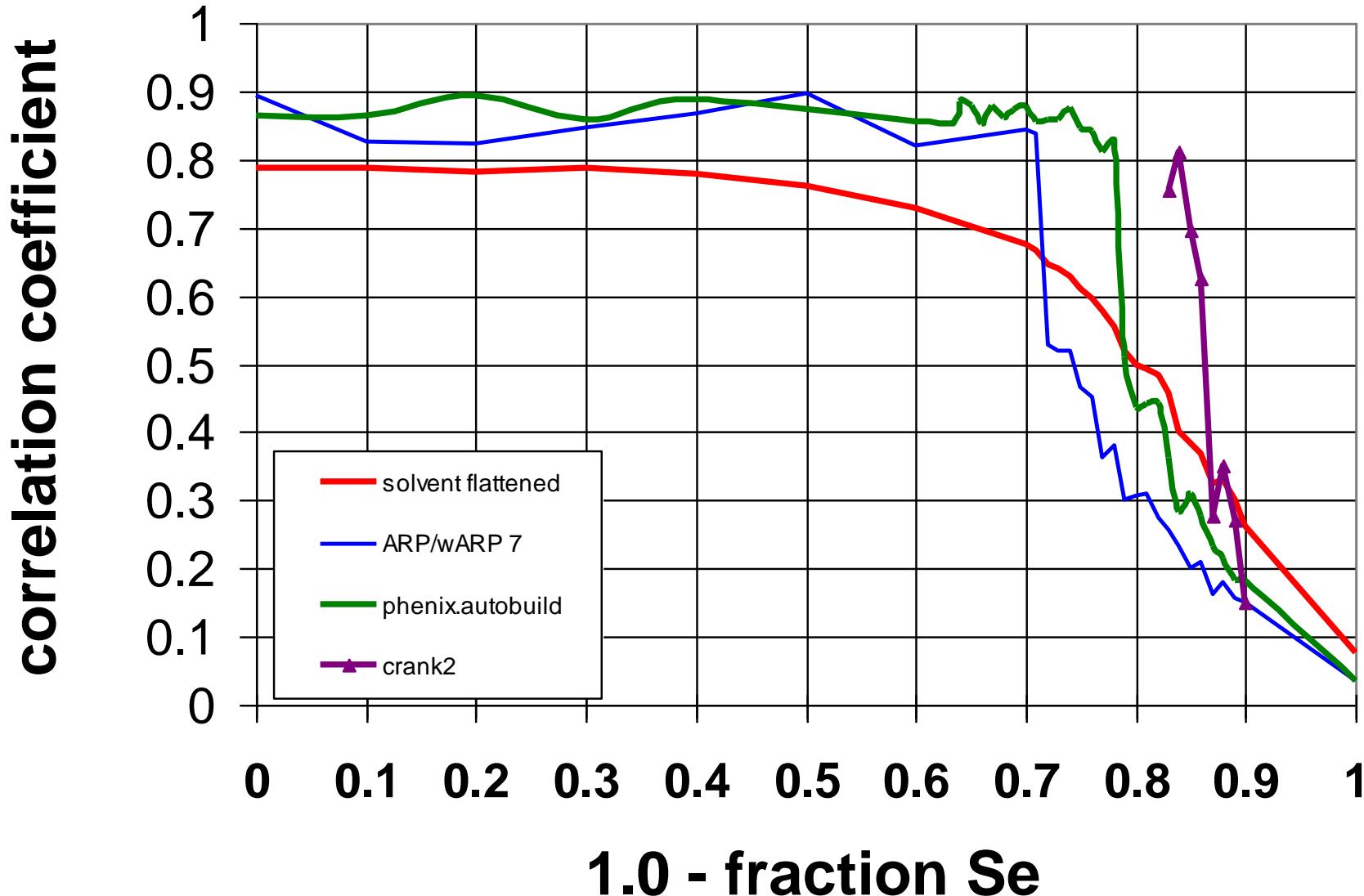
Phase Problem: Se vs S



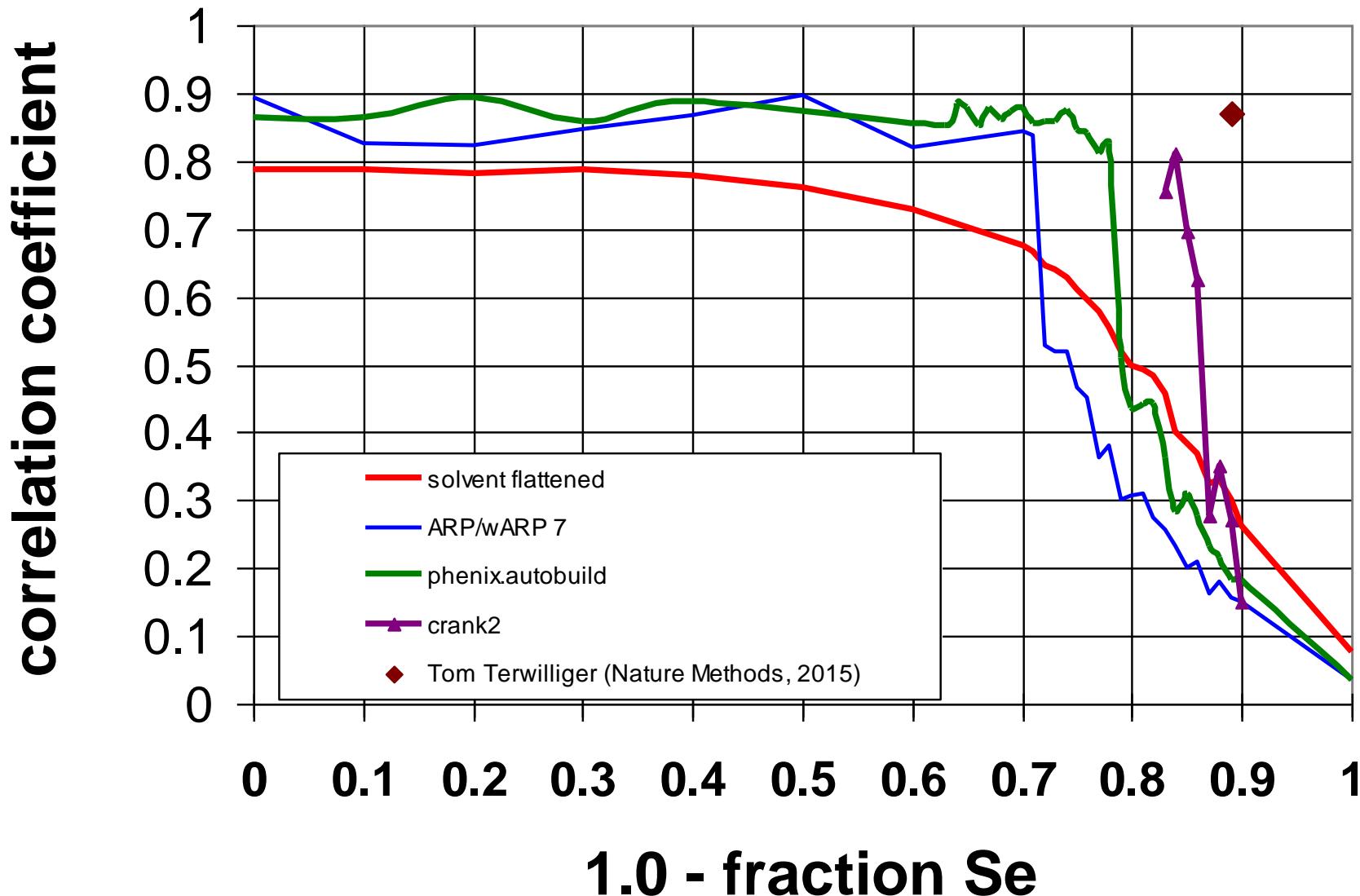
Phase Problem: Se vs S



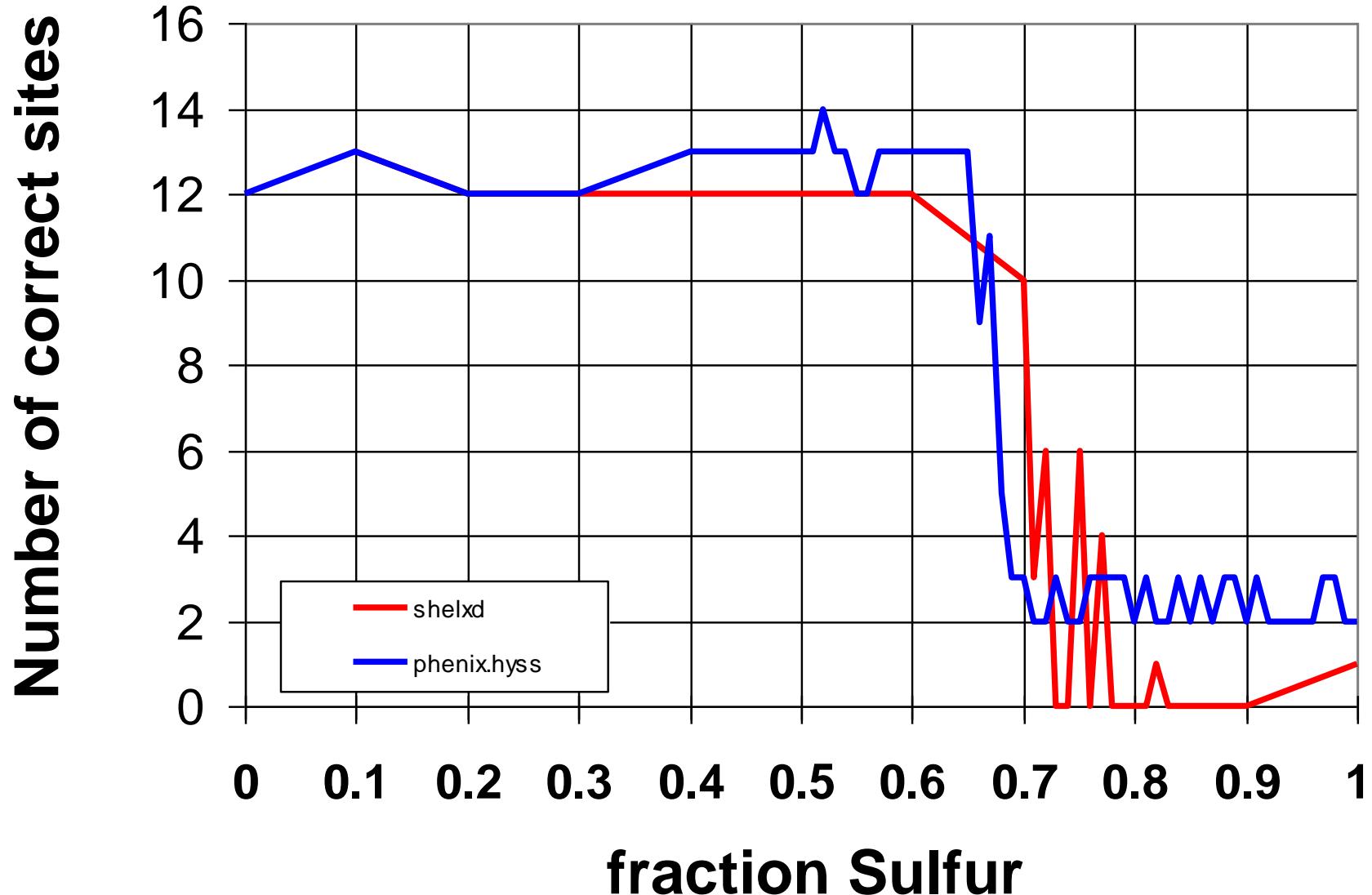
Phase Problem: Se vs S



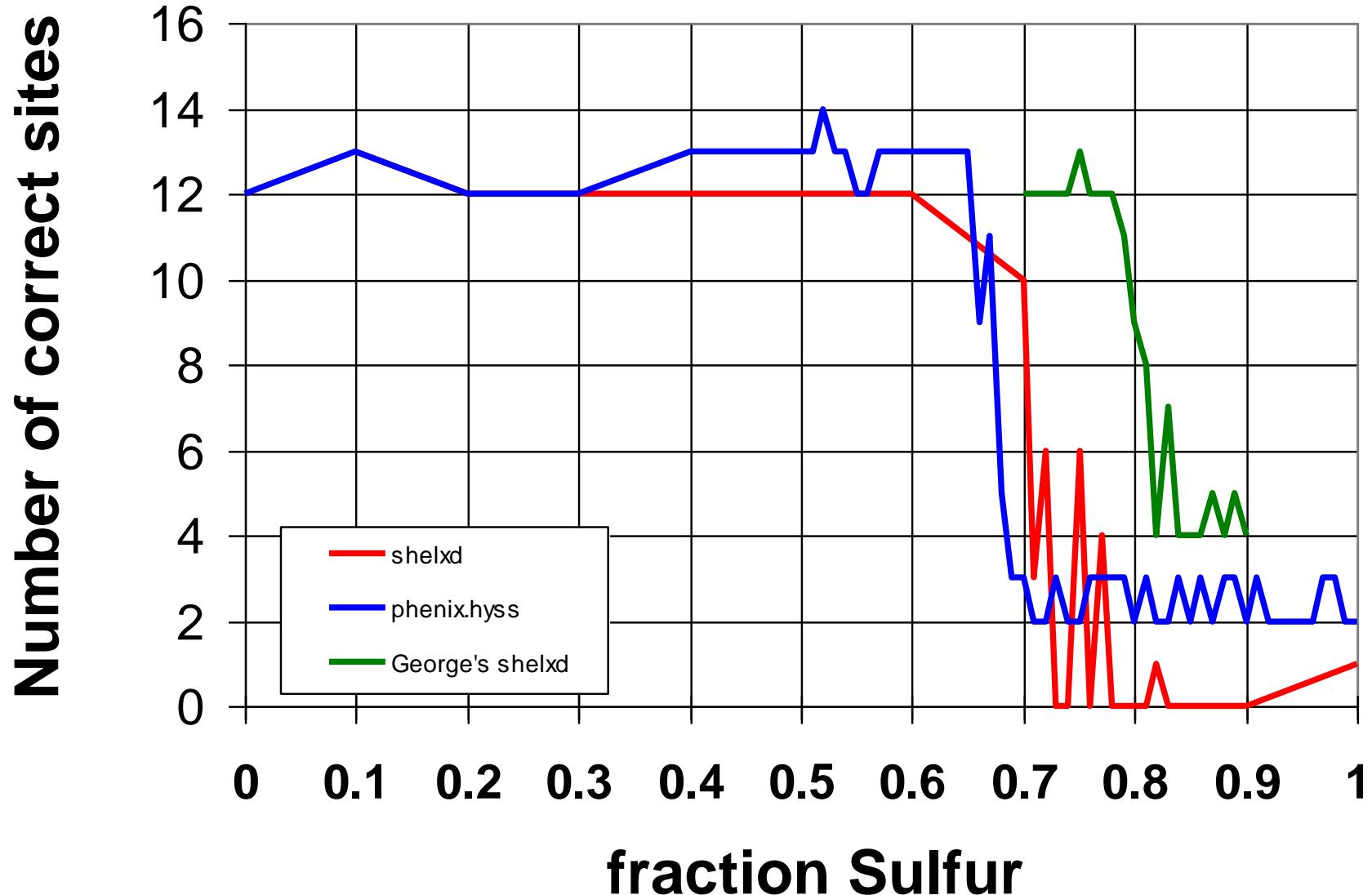
Phase Problem: Se vs S



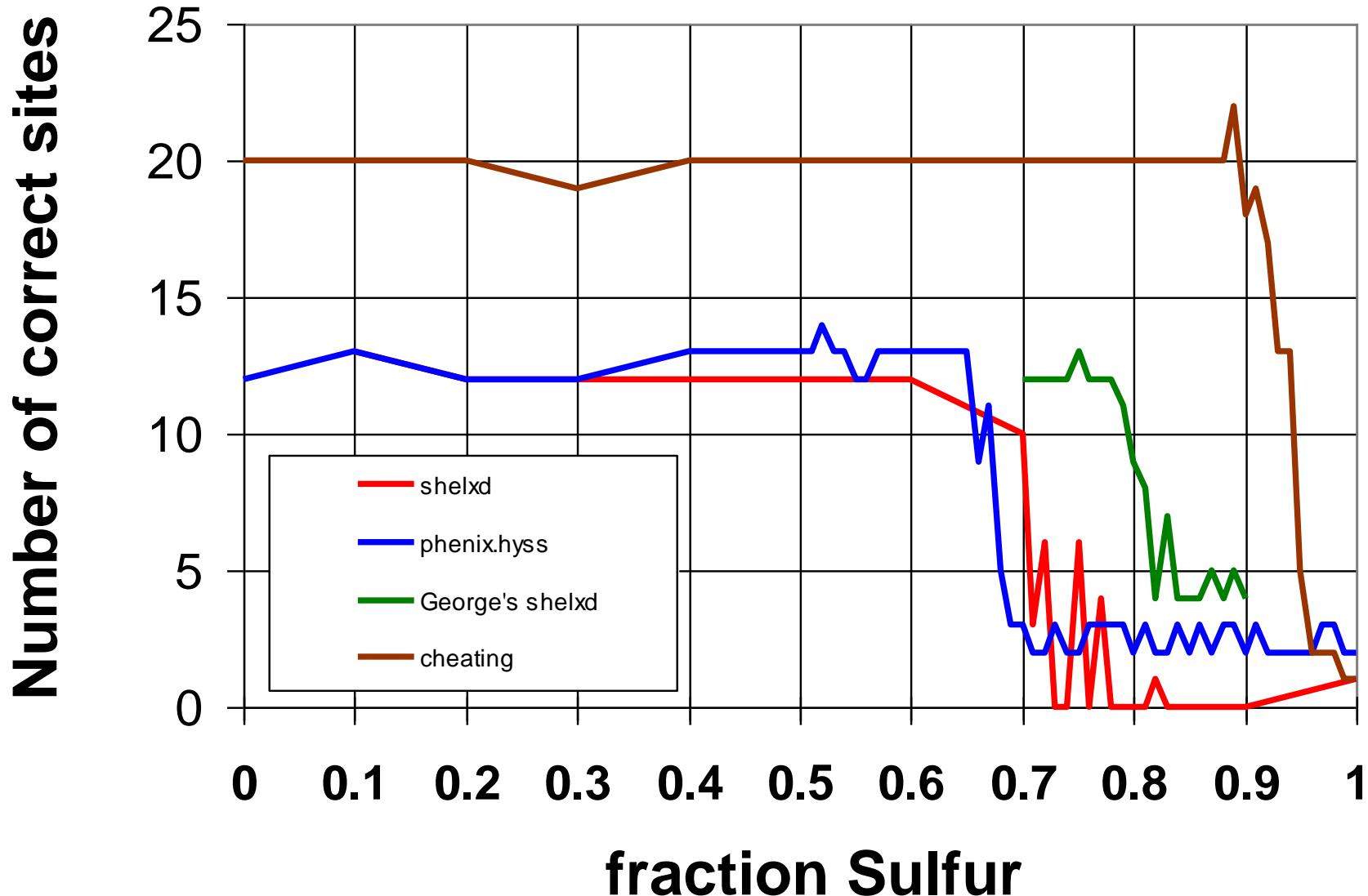
Finding sites



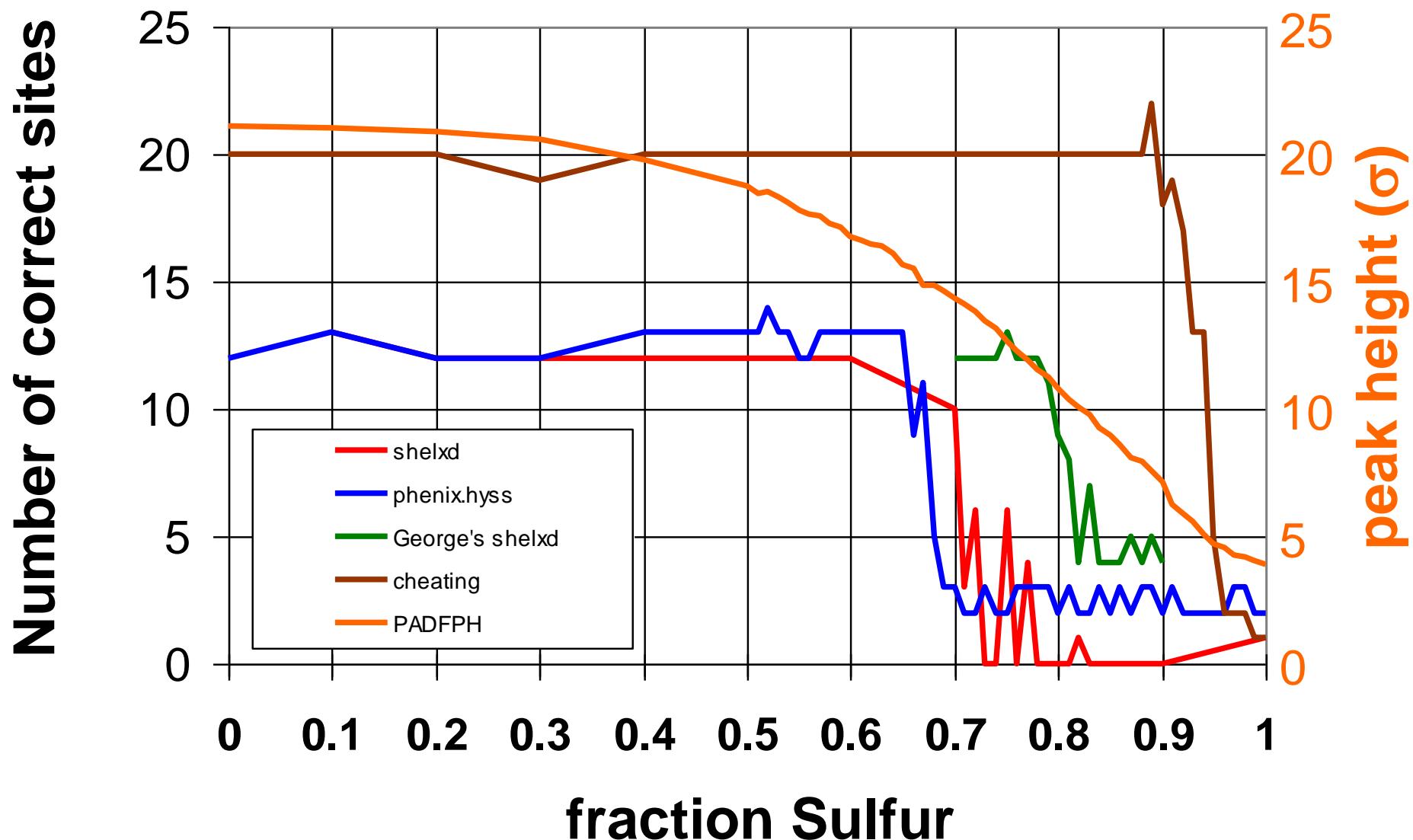
Finding sites



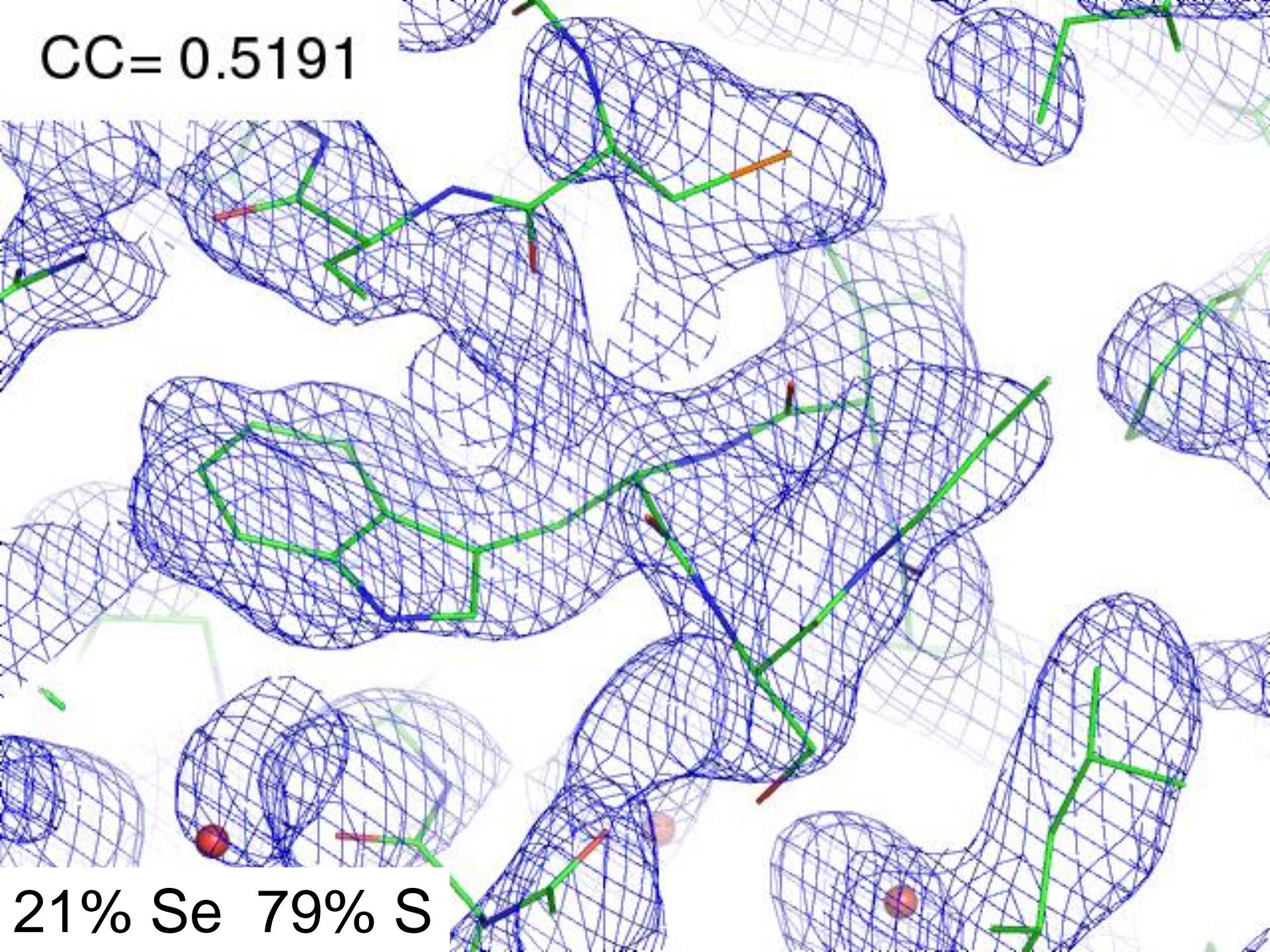
Finding sites



Finding sites

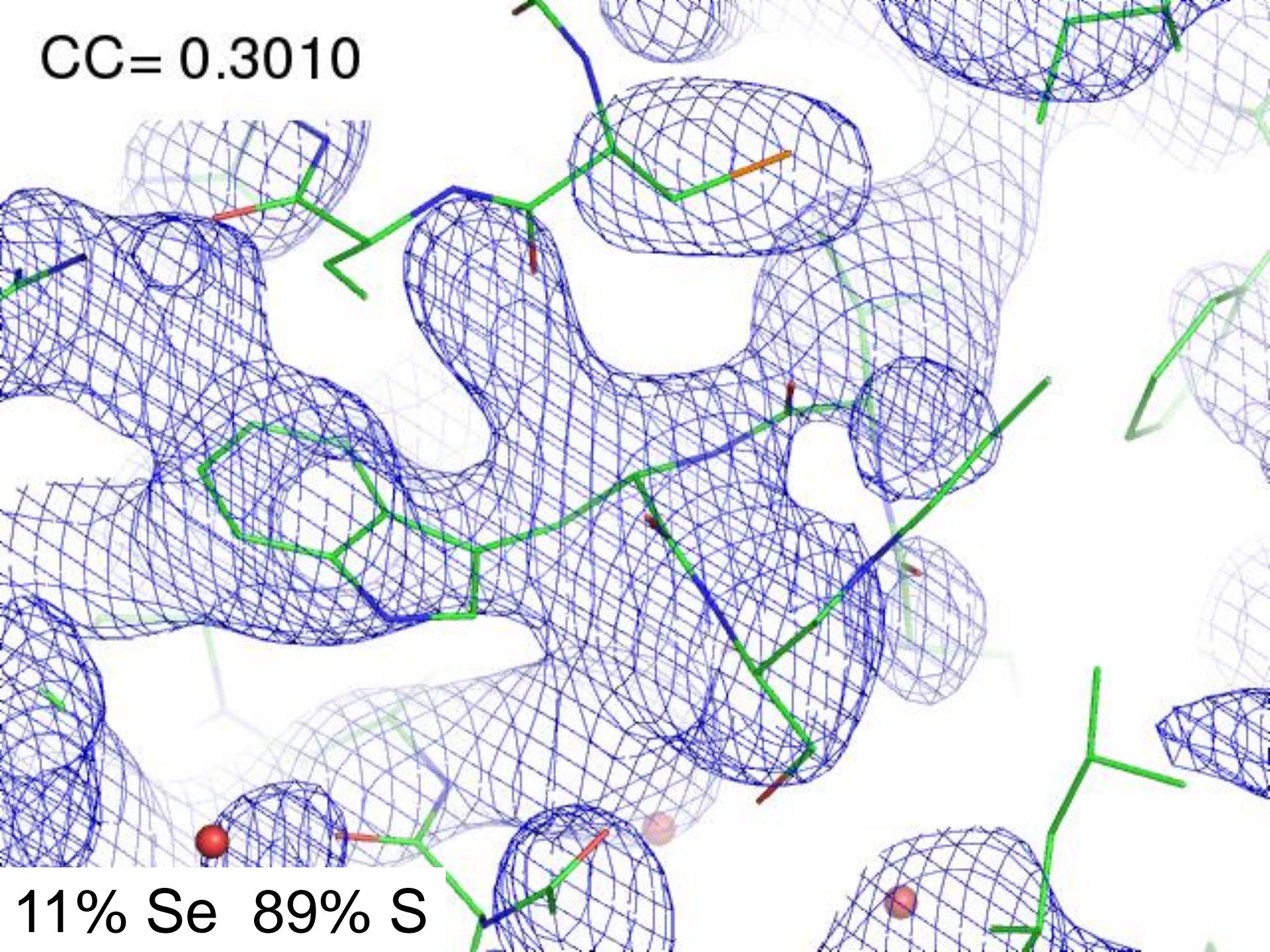


CC= 0.5191



21% Se 79% S

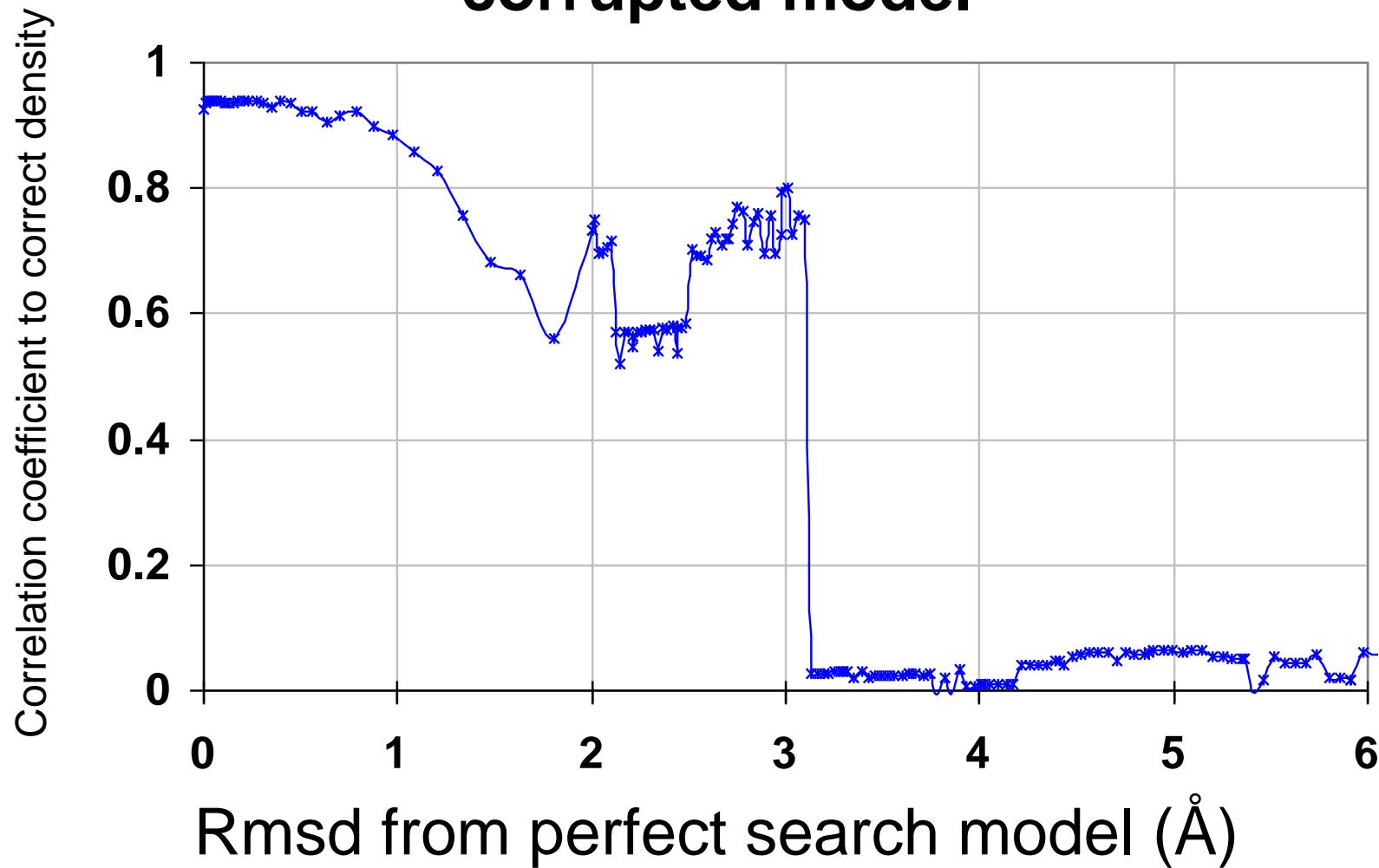
CC= 0.3010



11% Se 89% S

MR simulation

corrupted model



The transitions are sharp!

How can we predict success/failure?

Know Thy Experiment

Basic Principles

“Hell, there are **NO RULES** here - we're trying to accomplish something.”

Thomas A. Edison – inventor

“You've got to have an **ASSAY**.”

Arthur Kornberg – Nobel Laureate

“Control, control, you must learn **CONTROL!**”

Yoda – Jedi Master

Where do photons go?

Protein
1A x-rays



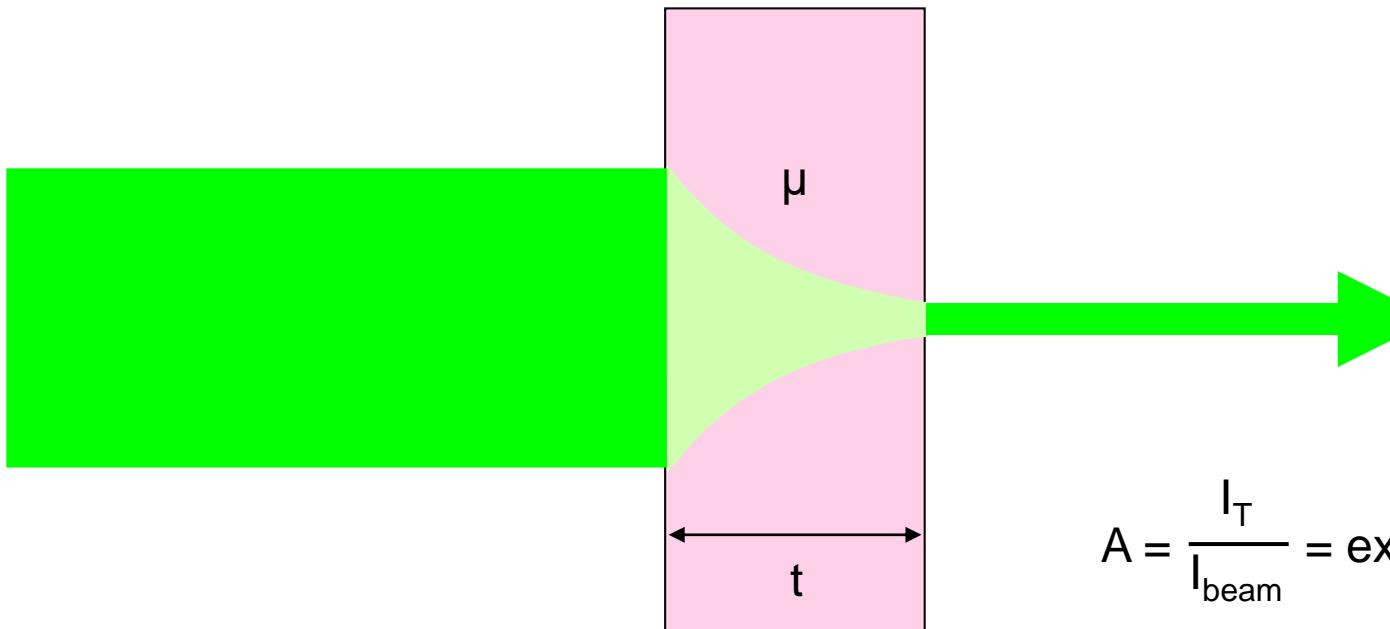
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1Å x-rays



2% total → error < 2%

attenuation factor

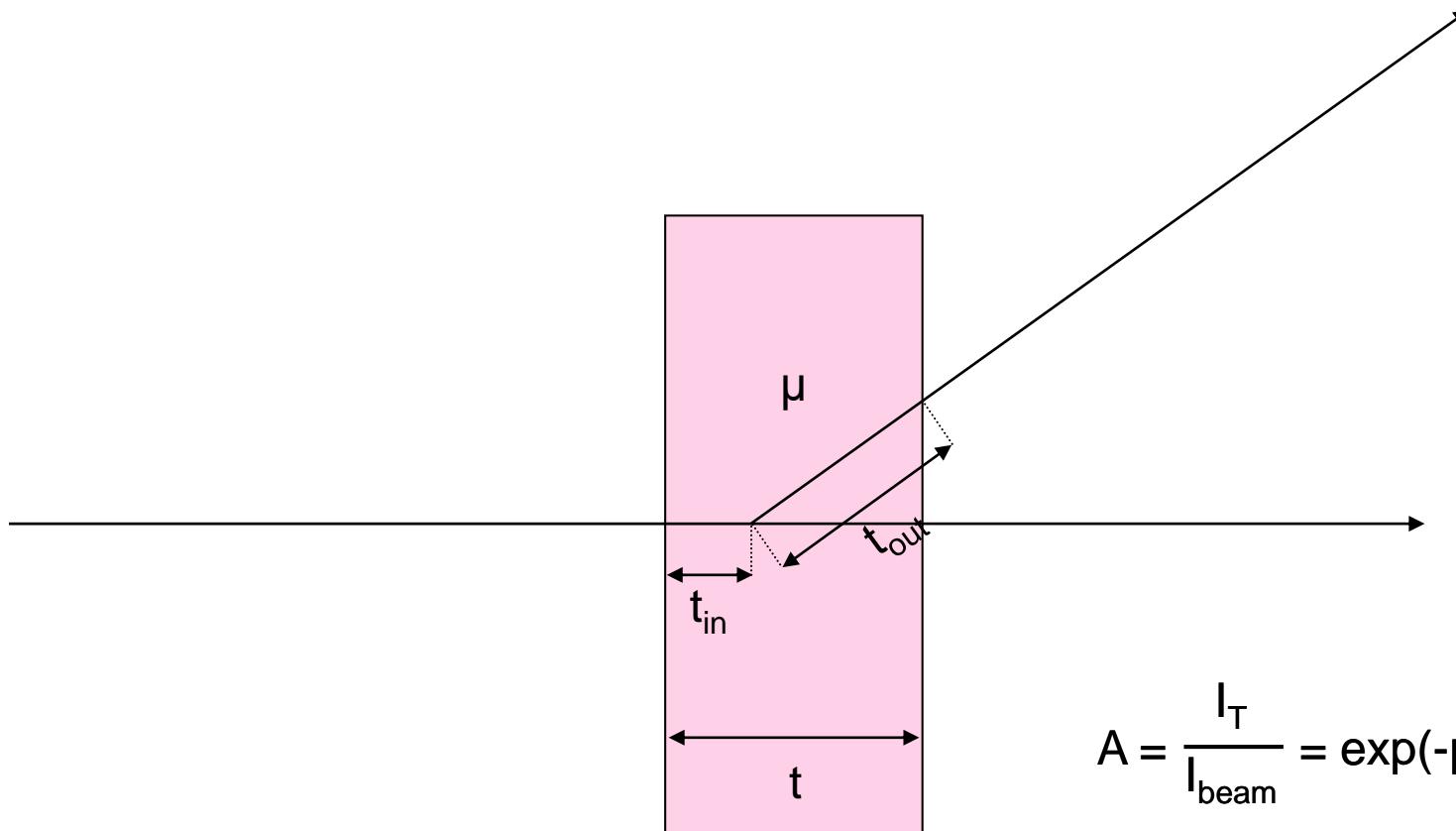


Bouguer, P. (1729). *Essai d'optique sur la gradation de la lumière*.

Lambert, J. H. (1760). *Photometria: sive De mensura et gradibus luminis, colorum et umbrae*. E. Klett.

Beer, A. (1852). "Bestimmung der Absorption des rothen Lichts in farbigen Flüssigkeiten", *Ann. Phys. Chem* **86**, 78-90.

attenuation factor

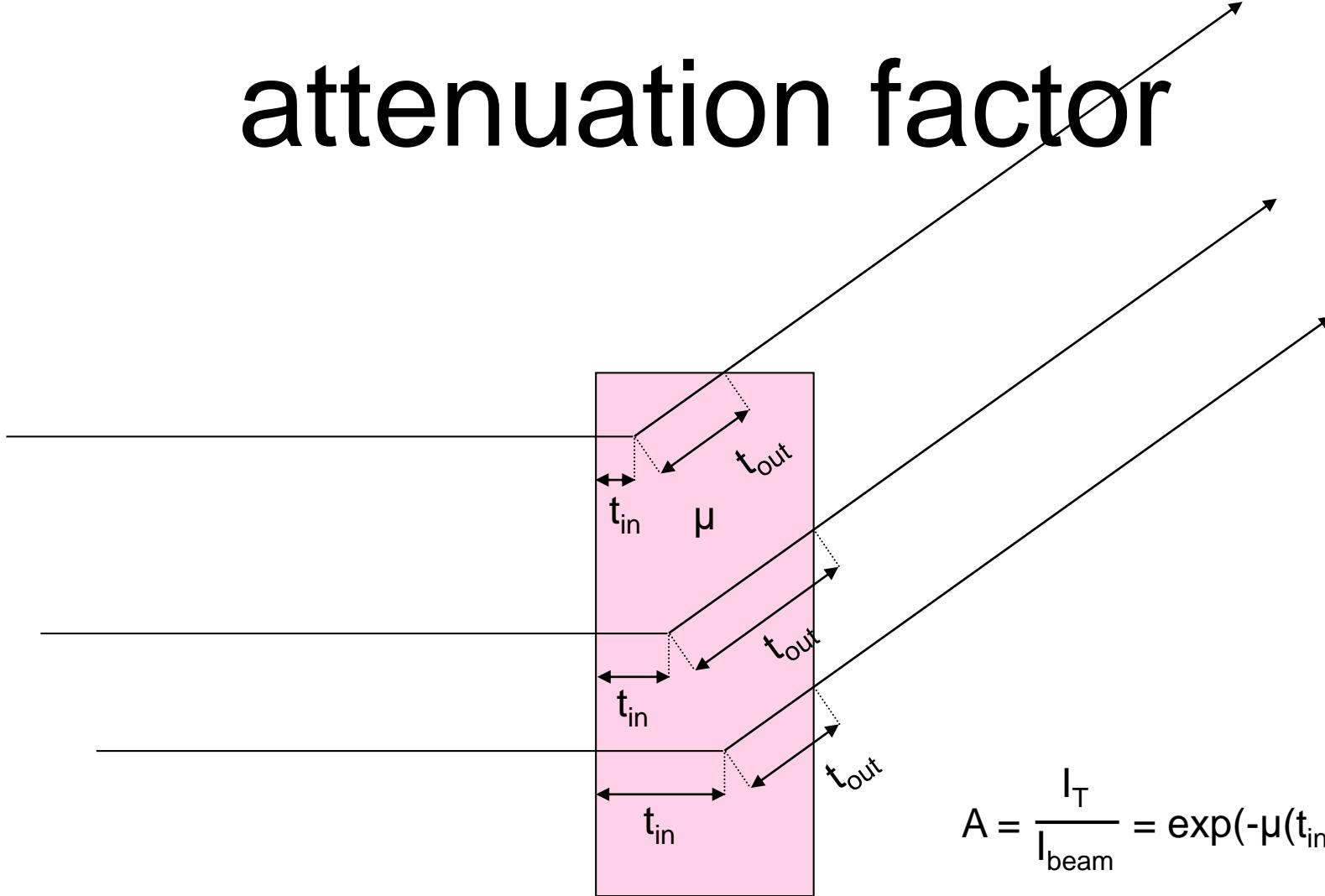


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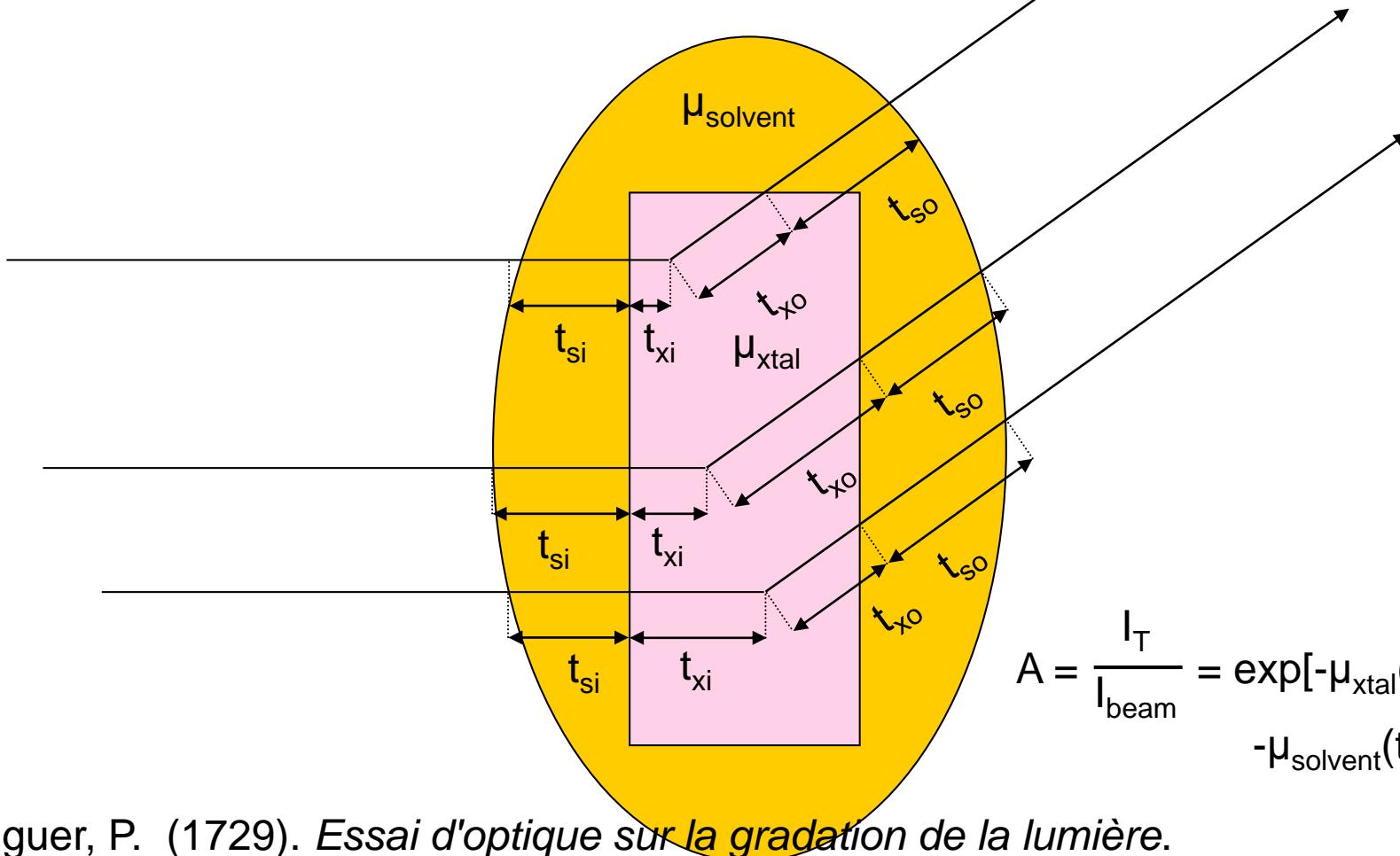


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attenuation factor



$$A = \frac{I_T}{I_{\text{beam}}} = \exp[-\mu_{\text{xtal}}(t_{\text{xi}} + t_{\text{xo}}) - \mu_{\text{solvent}}(t_{\text{si}} + t_{\text{so}})]$$

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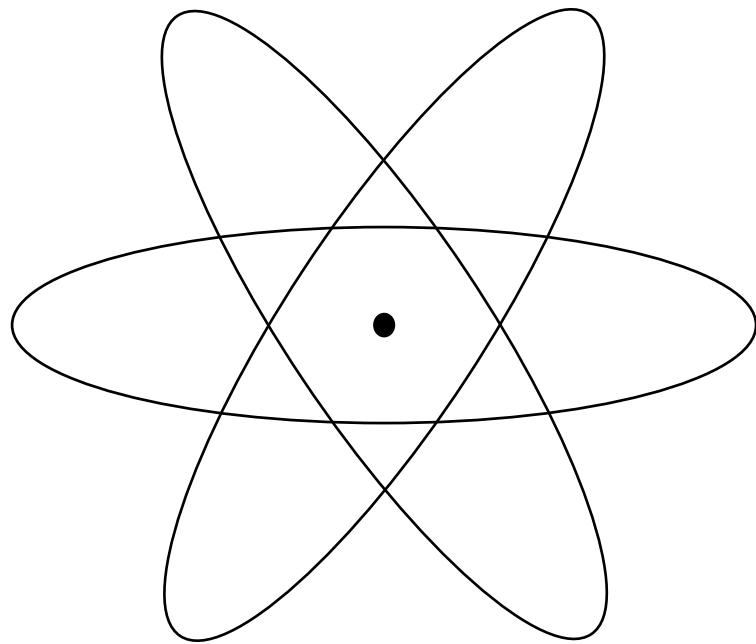
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Decisions, Decisions, Decisions

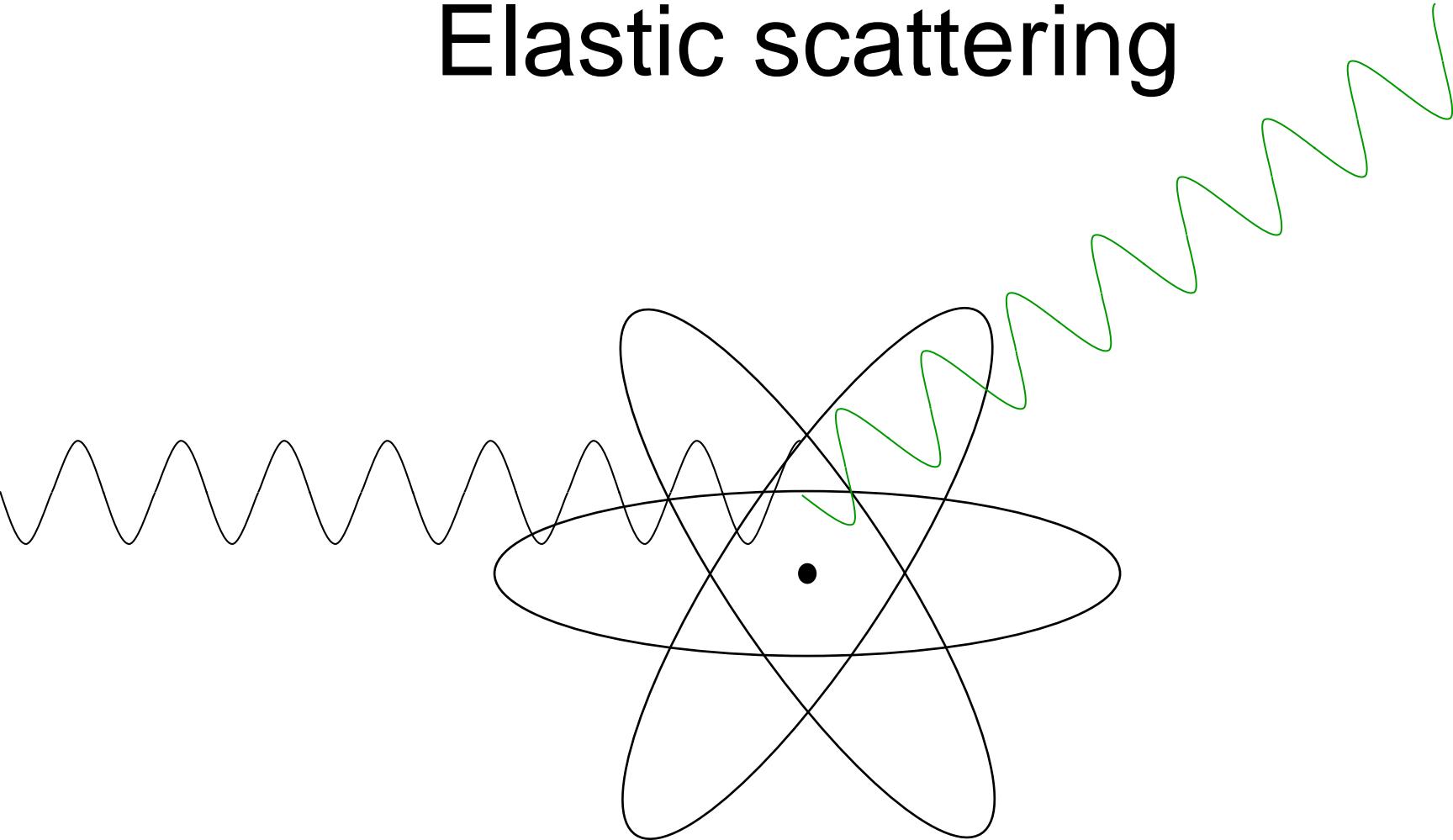
- Exposure time
- Number of images
- **Wavelengths**
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
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Run Strategy

Elastic scattering

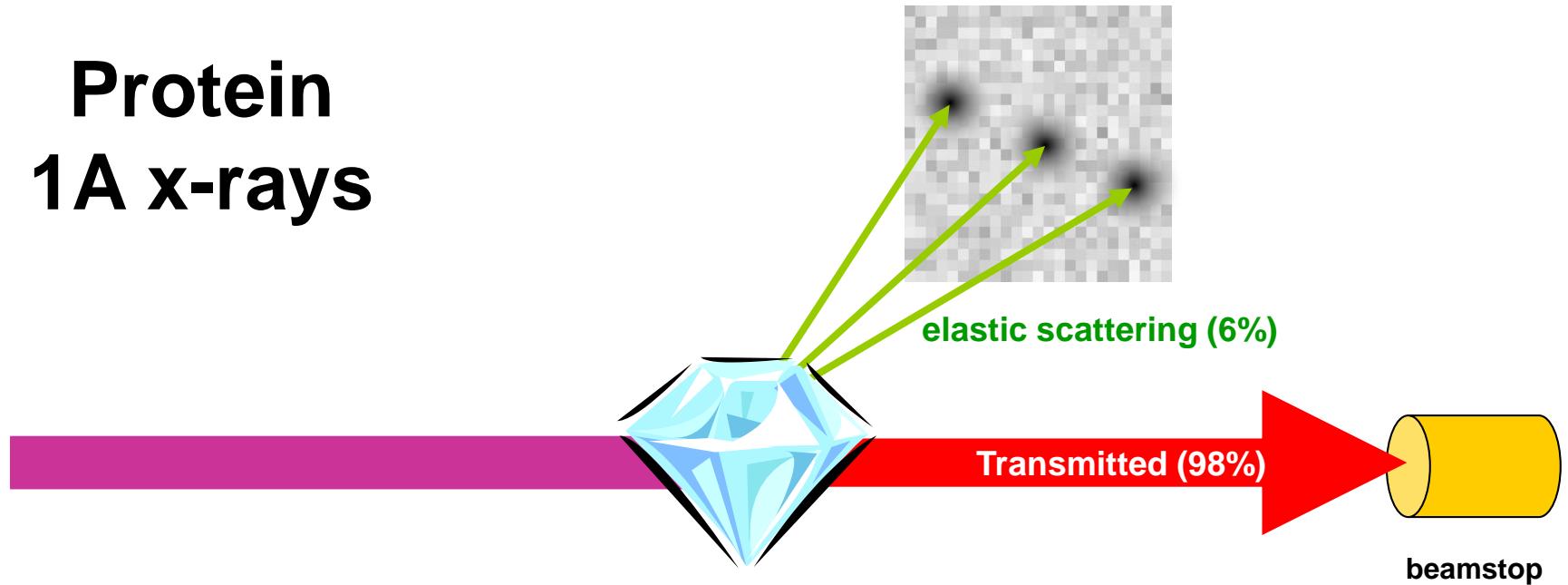


Elastic scattering

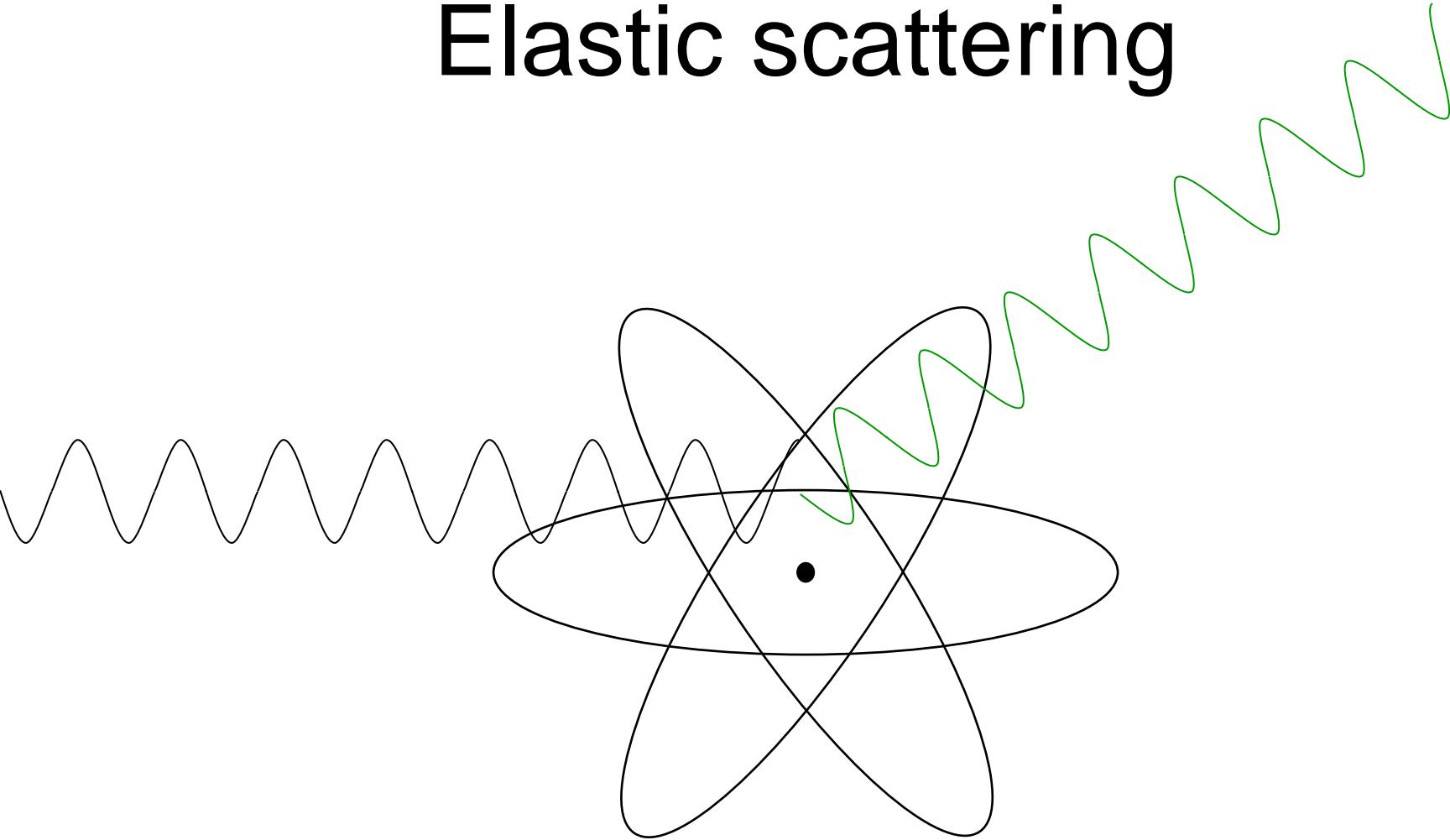


Where do photons go?

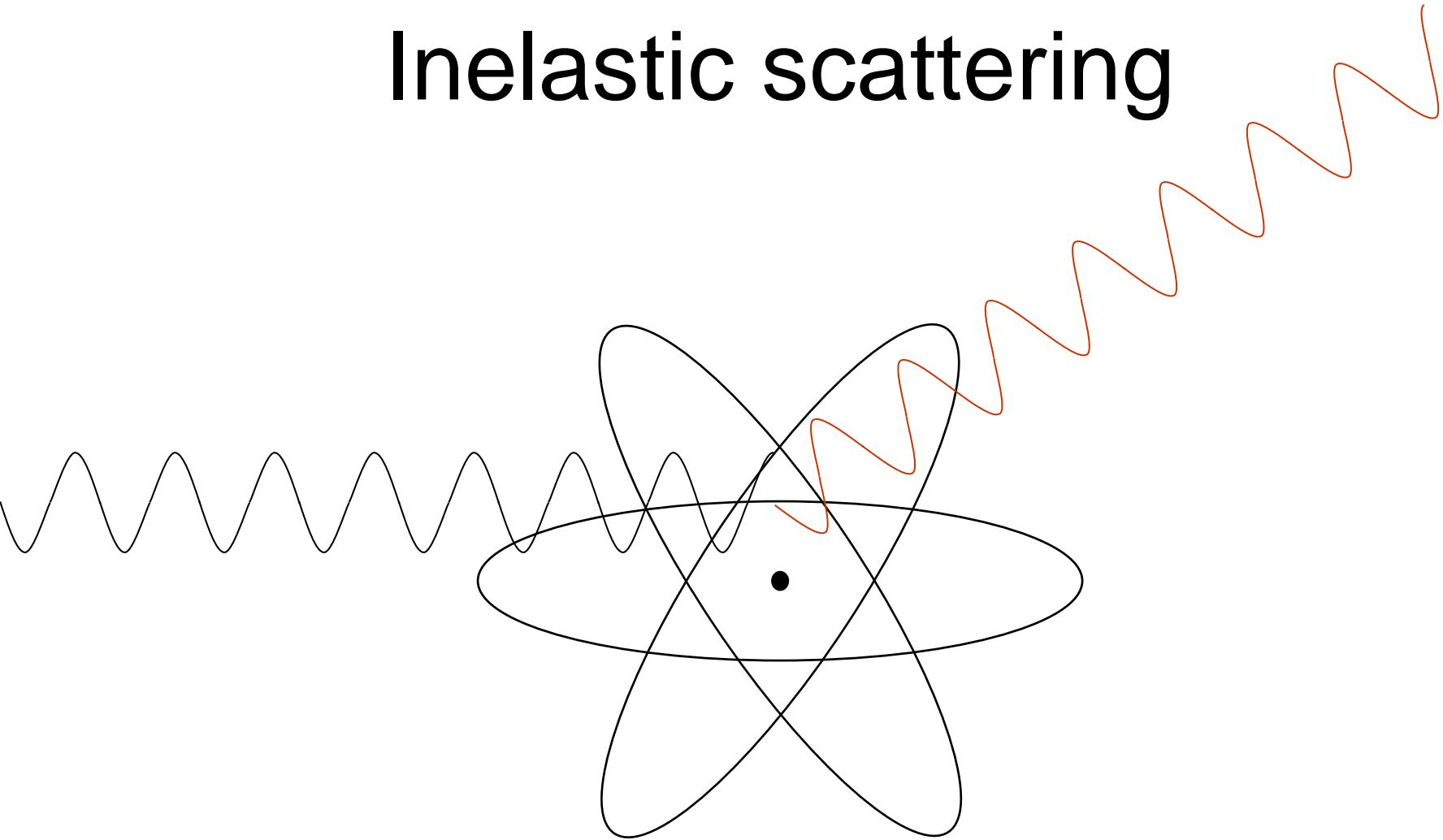
Protein
1A x-rays



Elastic scattering

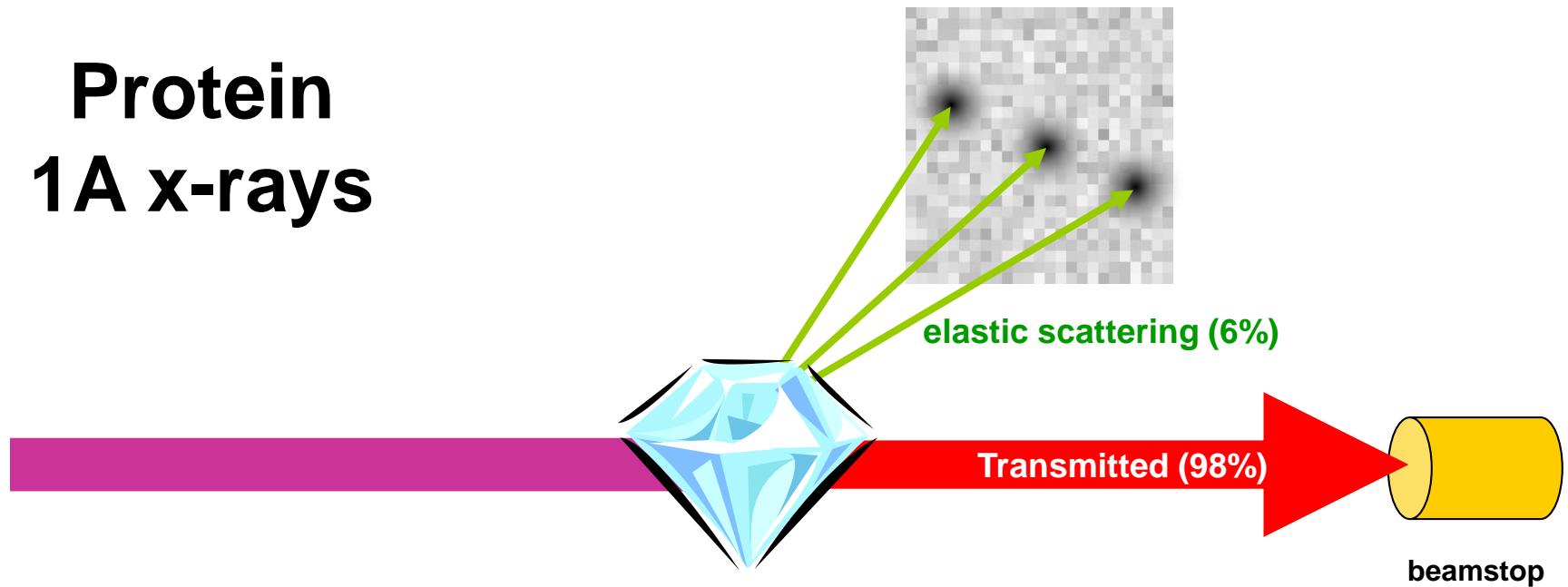


Inelastic scattering



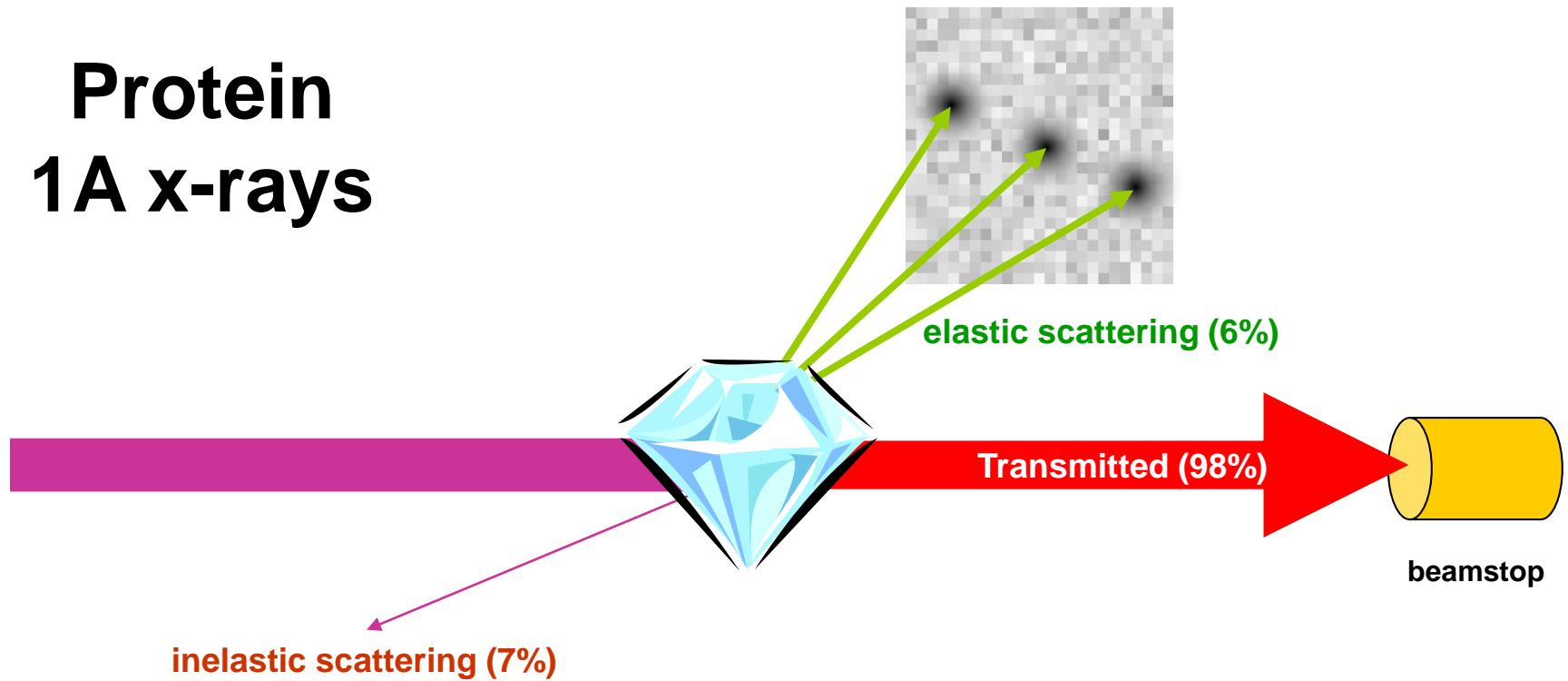
Where do photons go?

Protein
1A x-rays



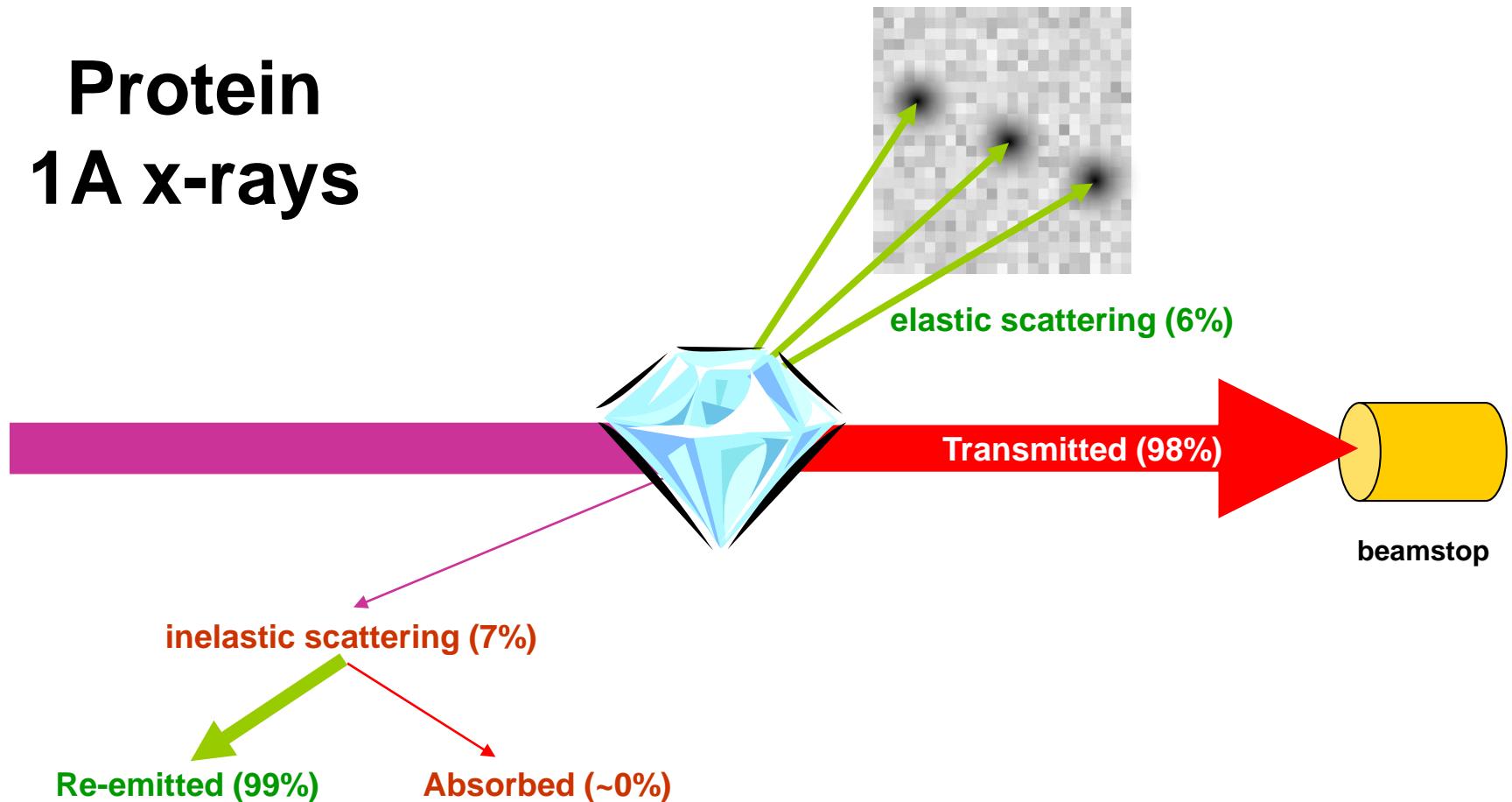
Where do photons go?

Protein
1A x-rays

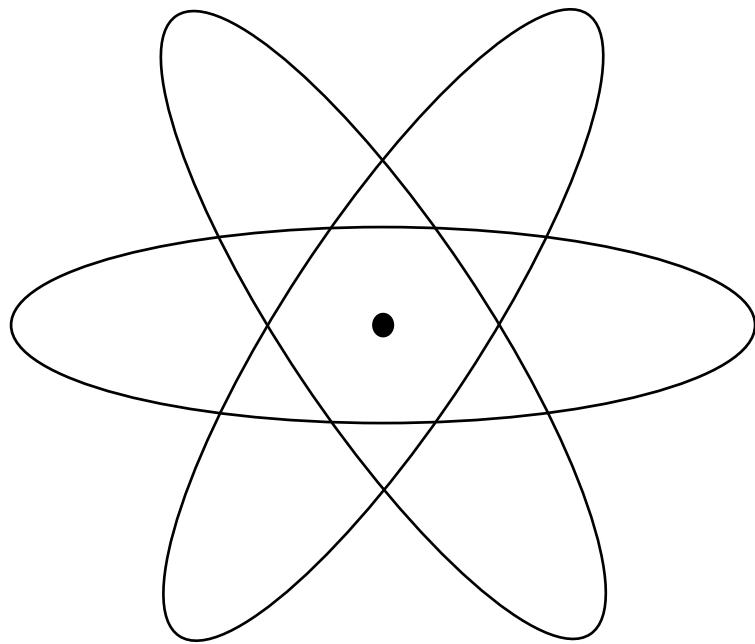


Where do photons go?

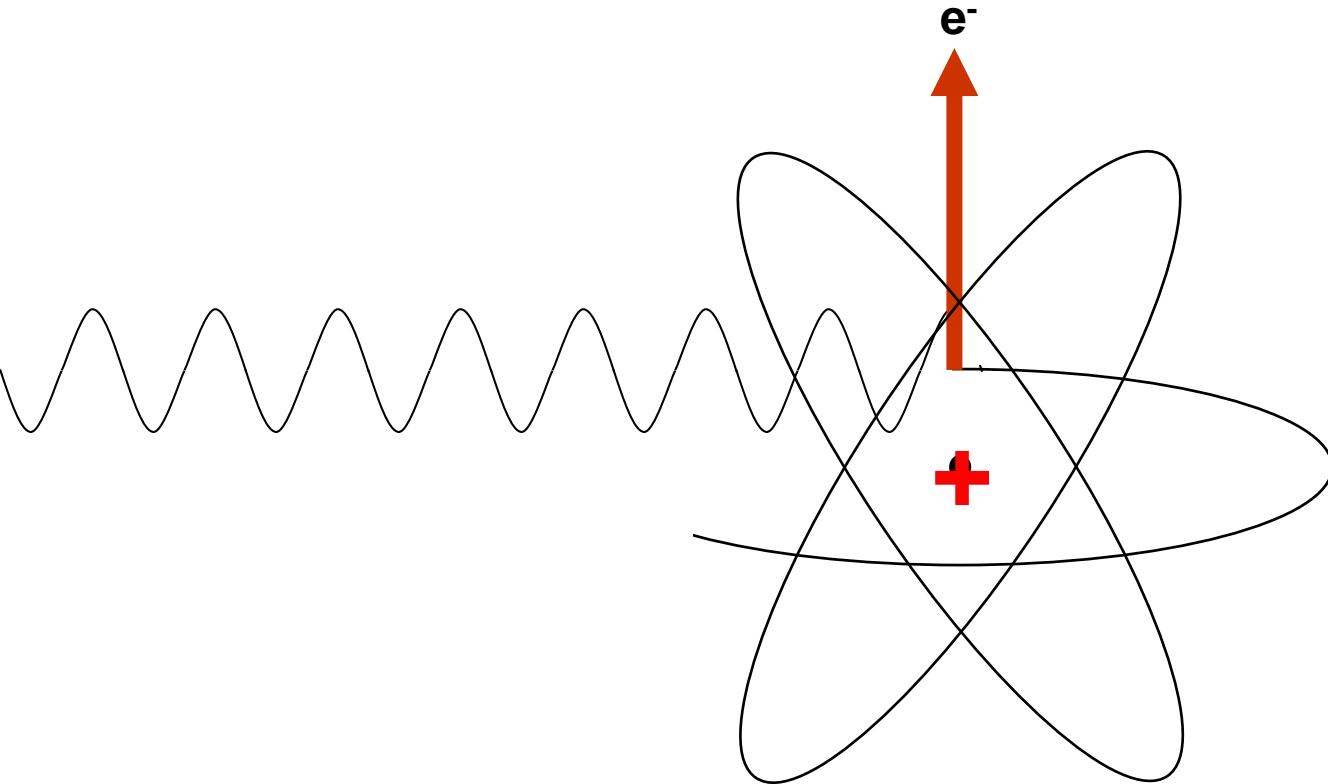
Protein
1A x-rays



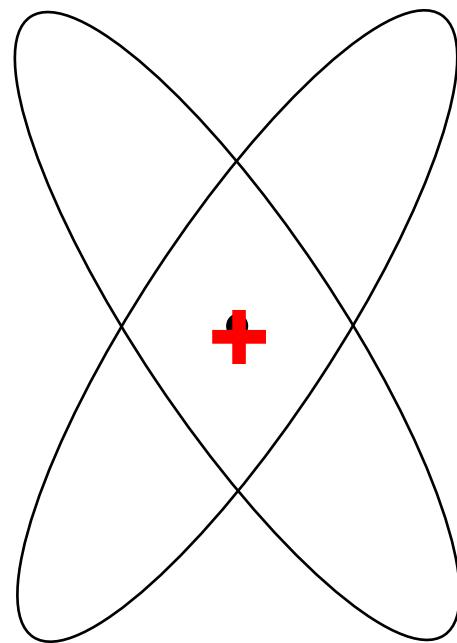
Photoelectric absorption



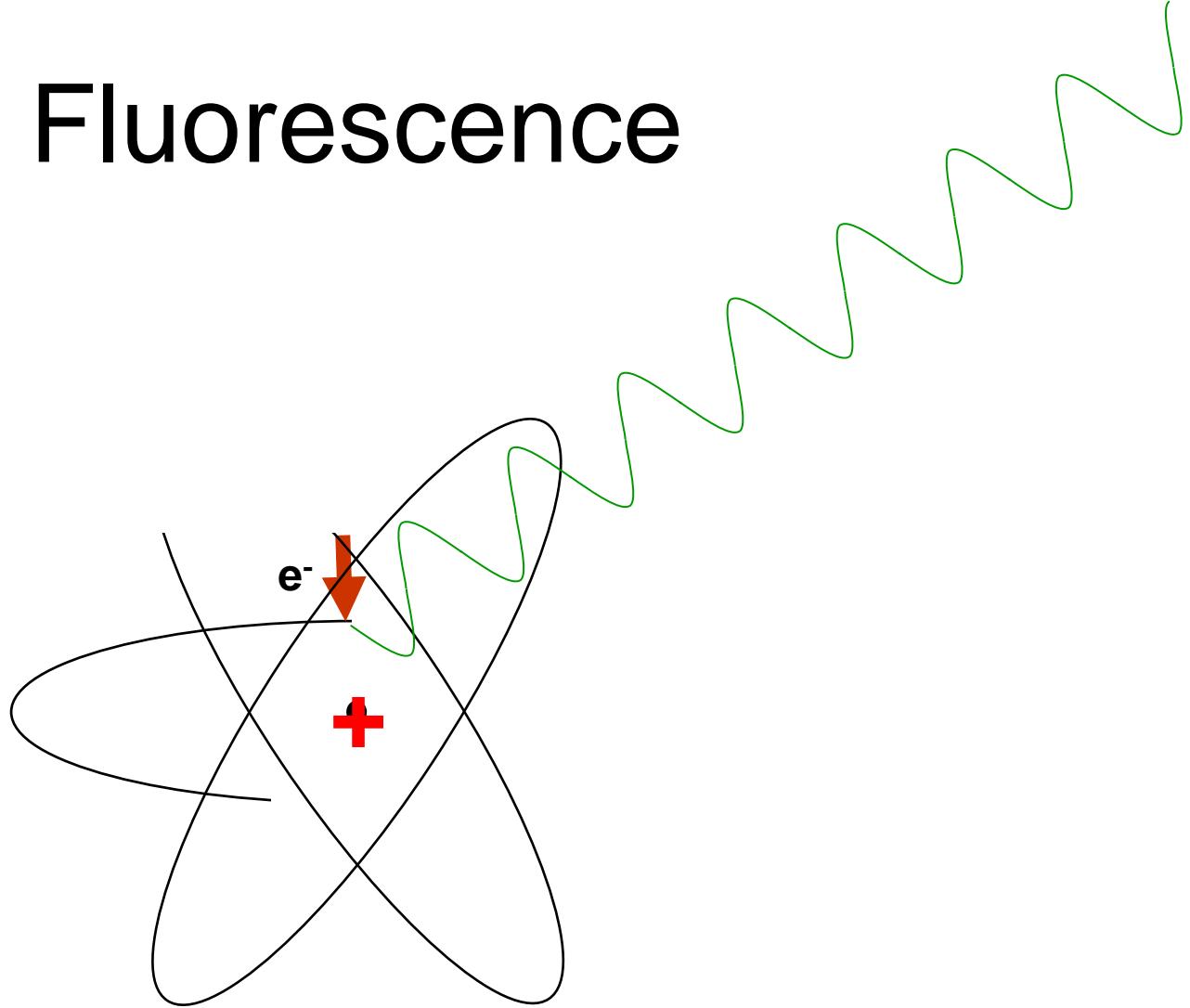
Photoelectric absorption



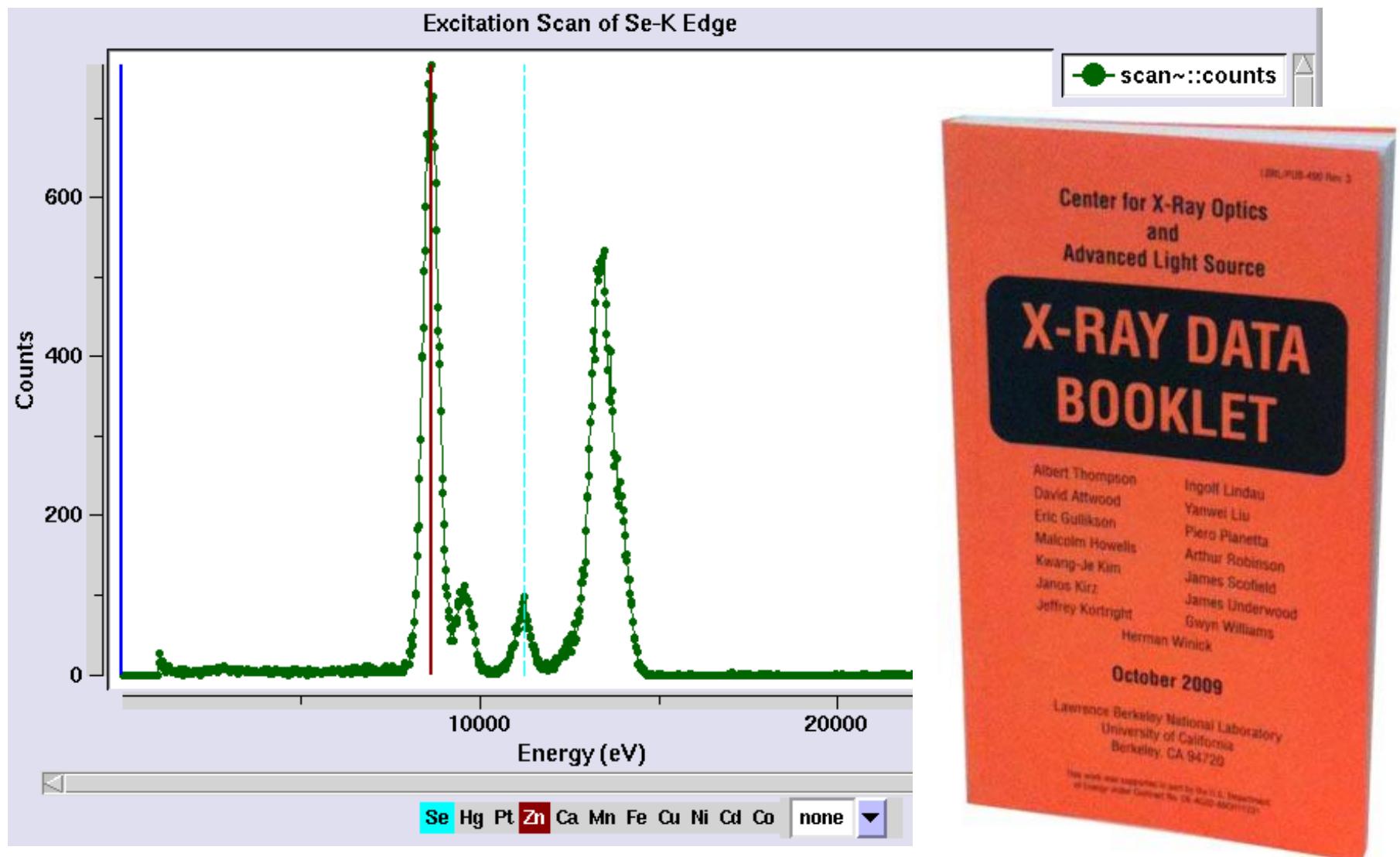
Fluorescence



Fluorescence

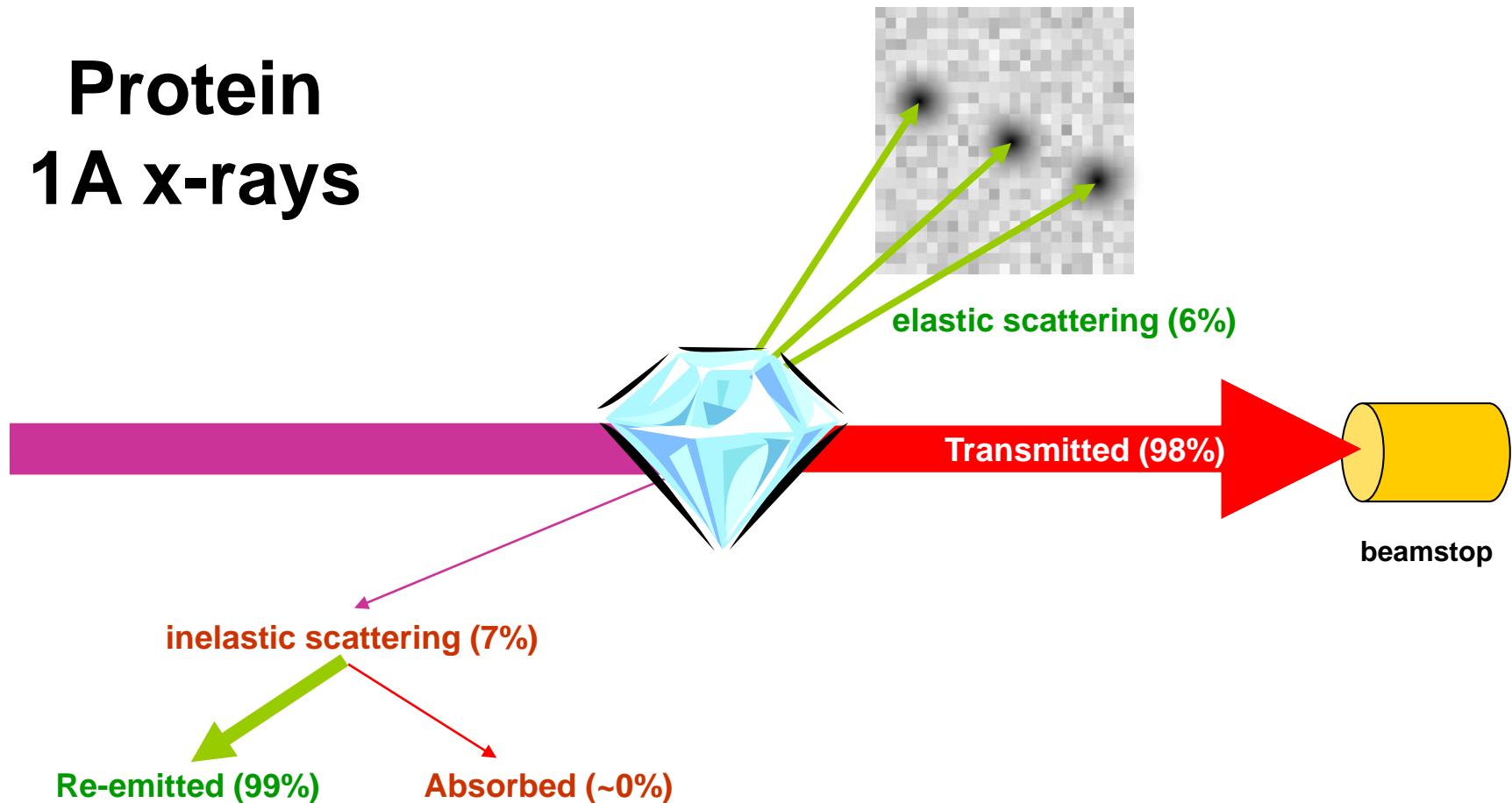


Metal identification



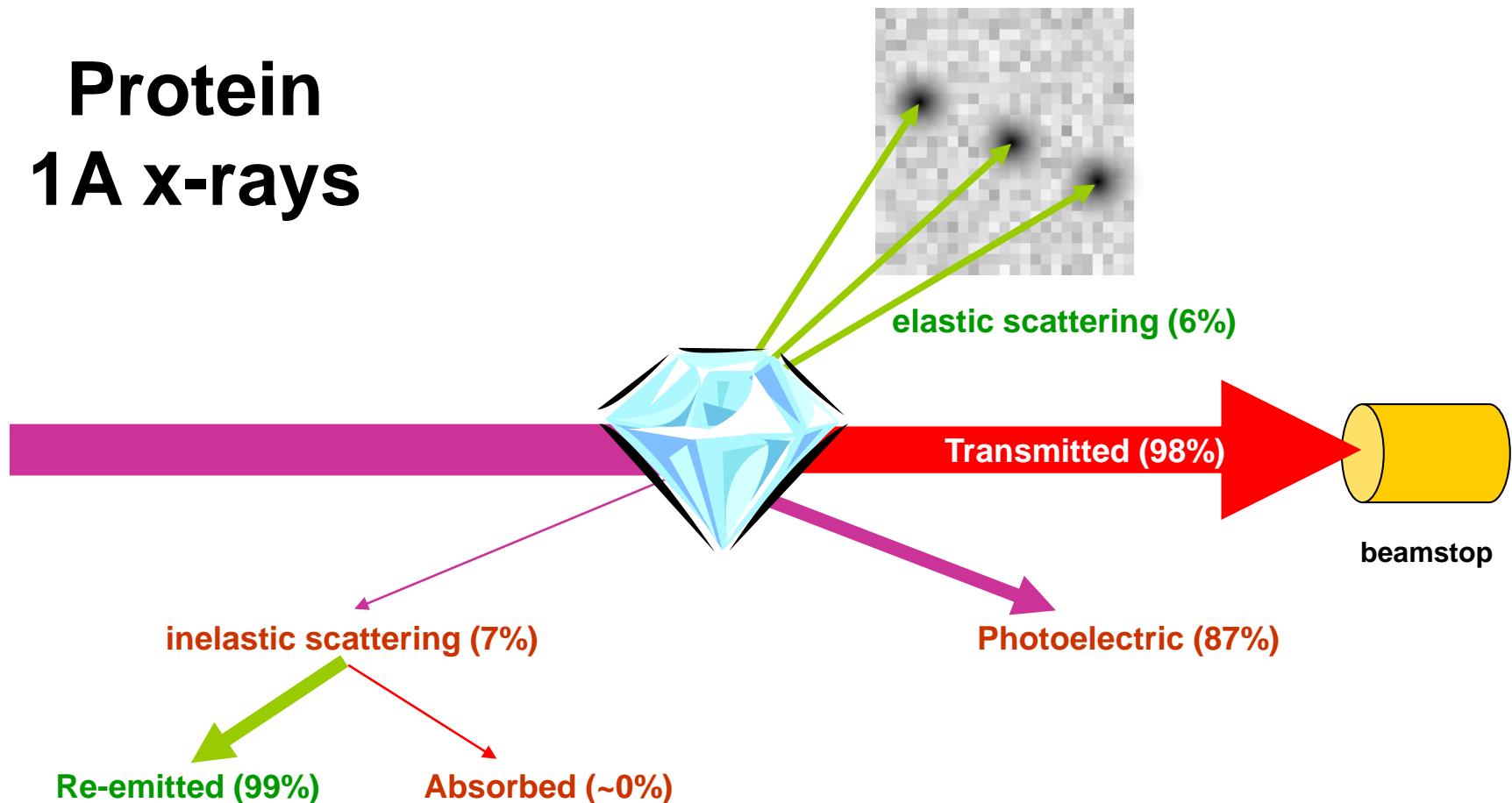
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Protein
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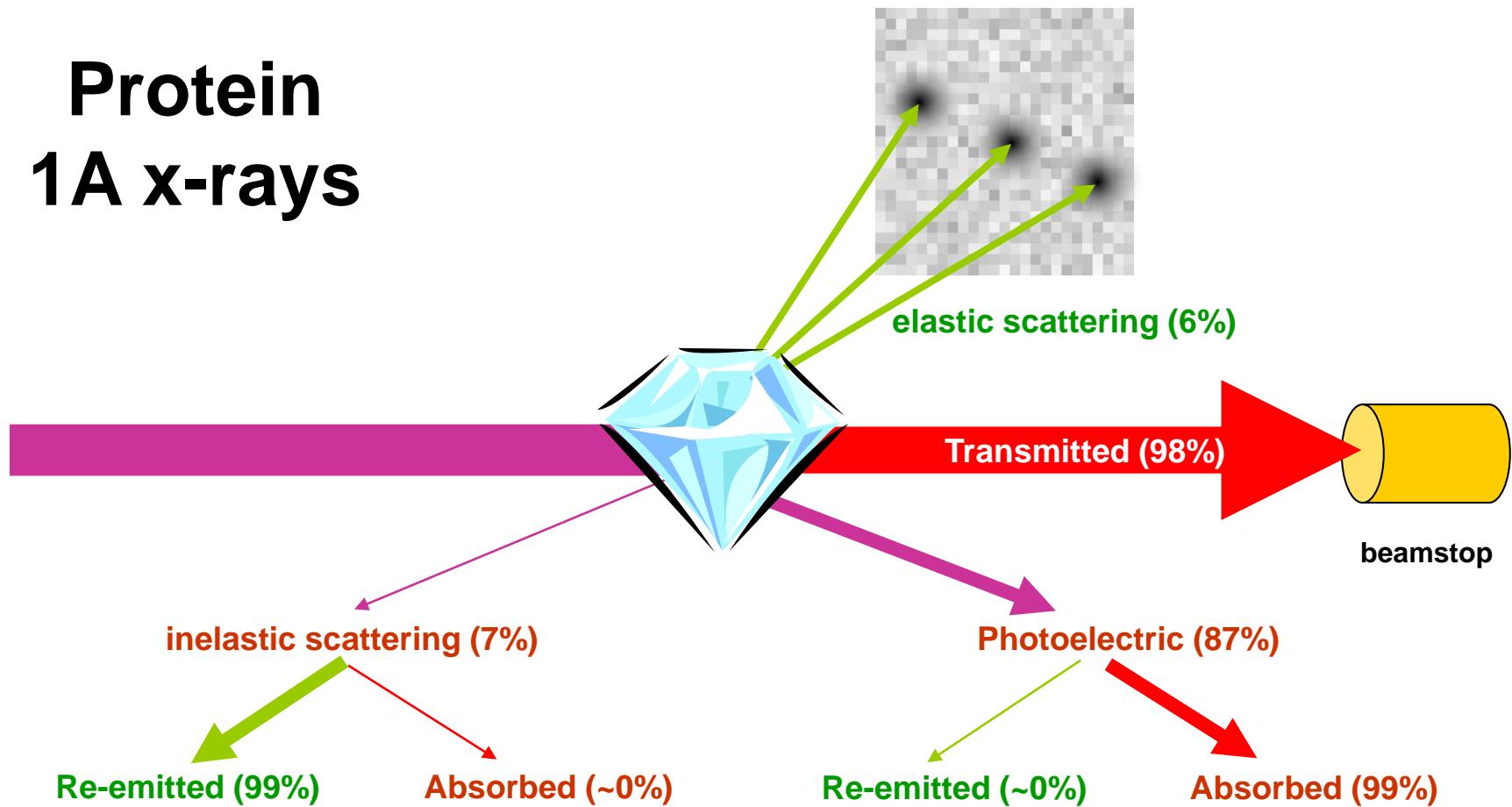
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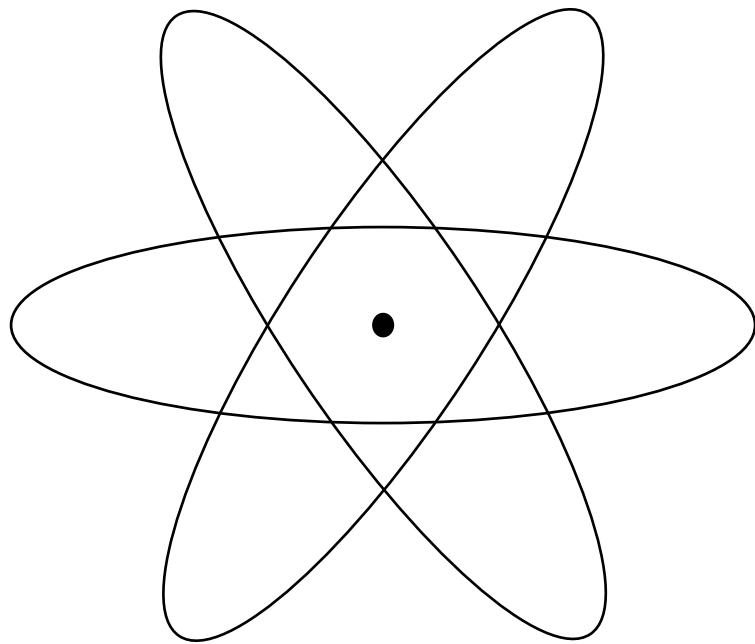


Where do photons go?

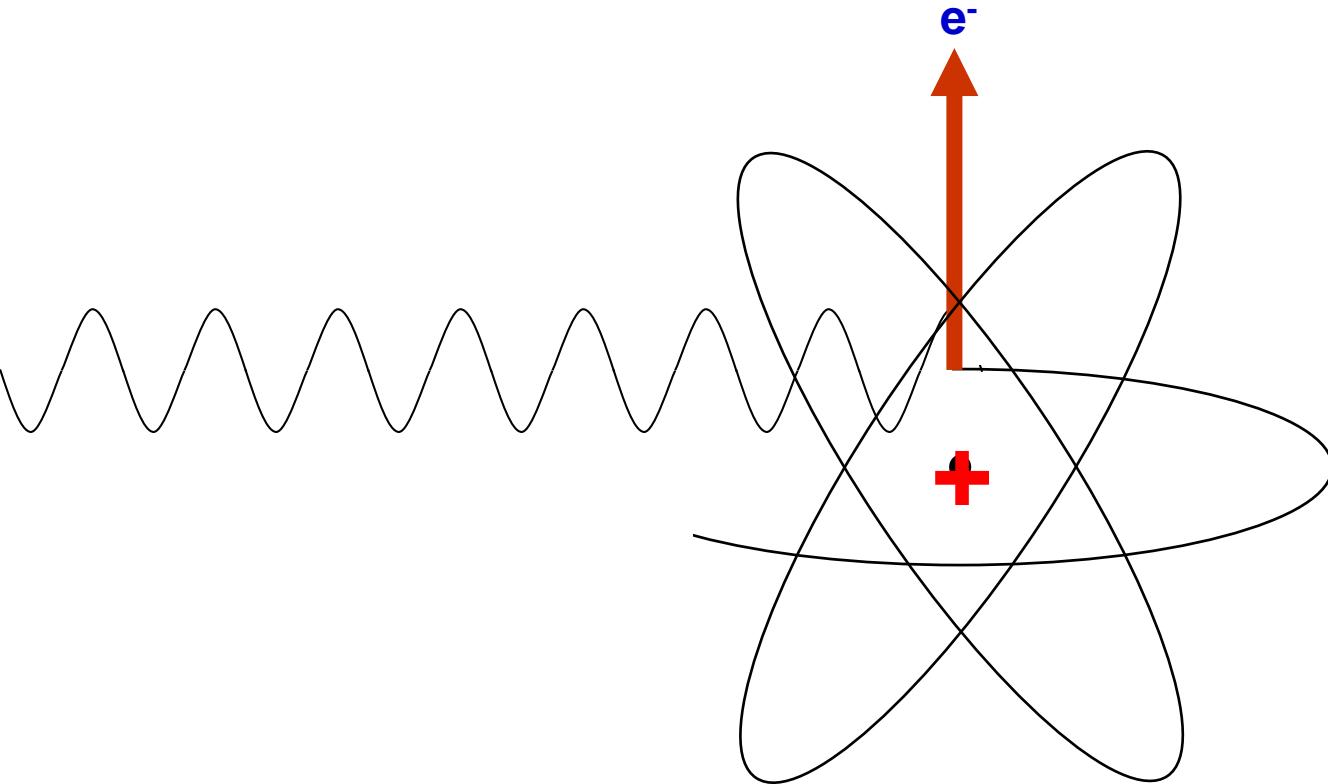
Protein
1A x-rays



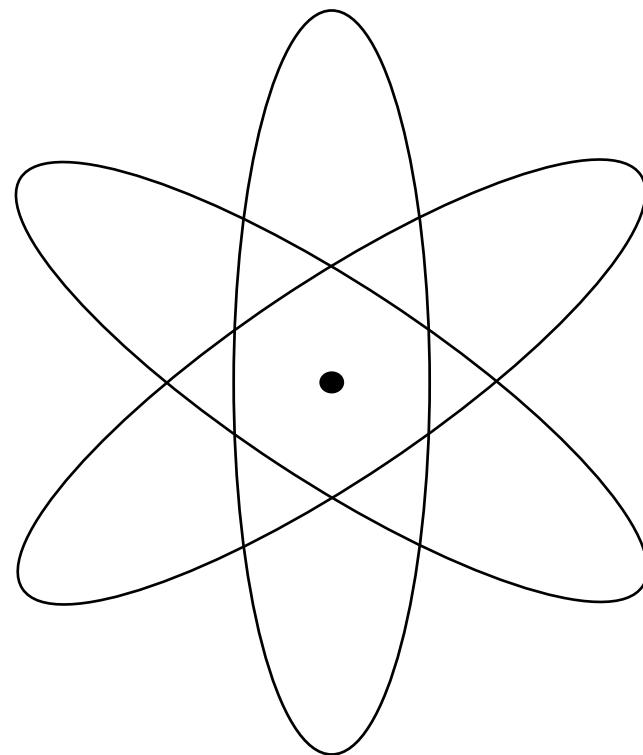
Photoelectric absorption



Photoelectric absorption

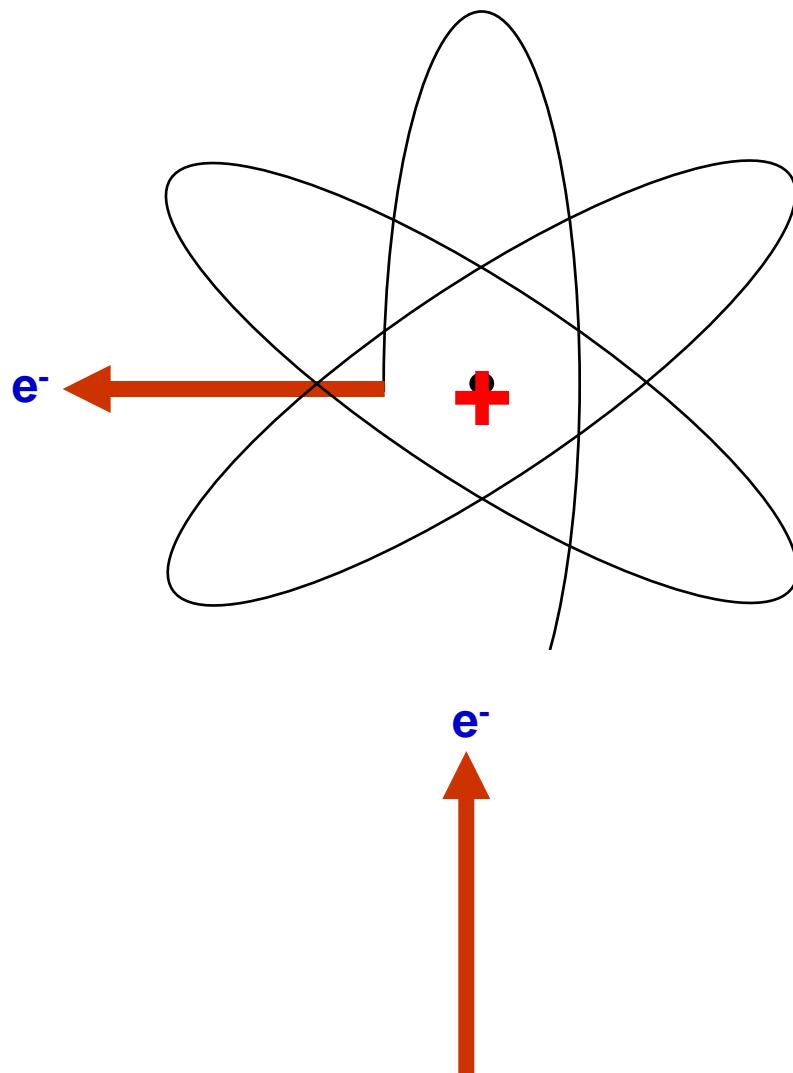


Secondary ionization

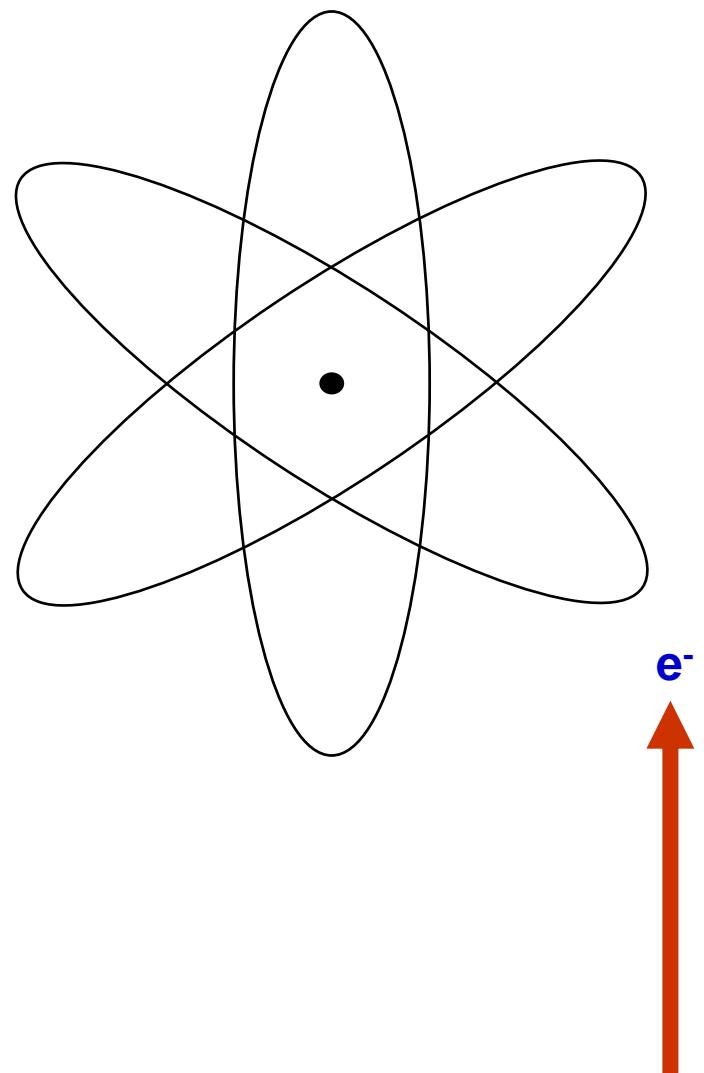


e^-

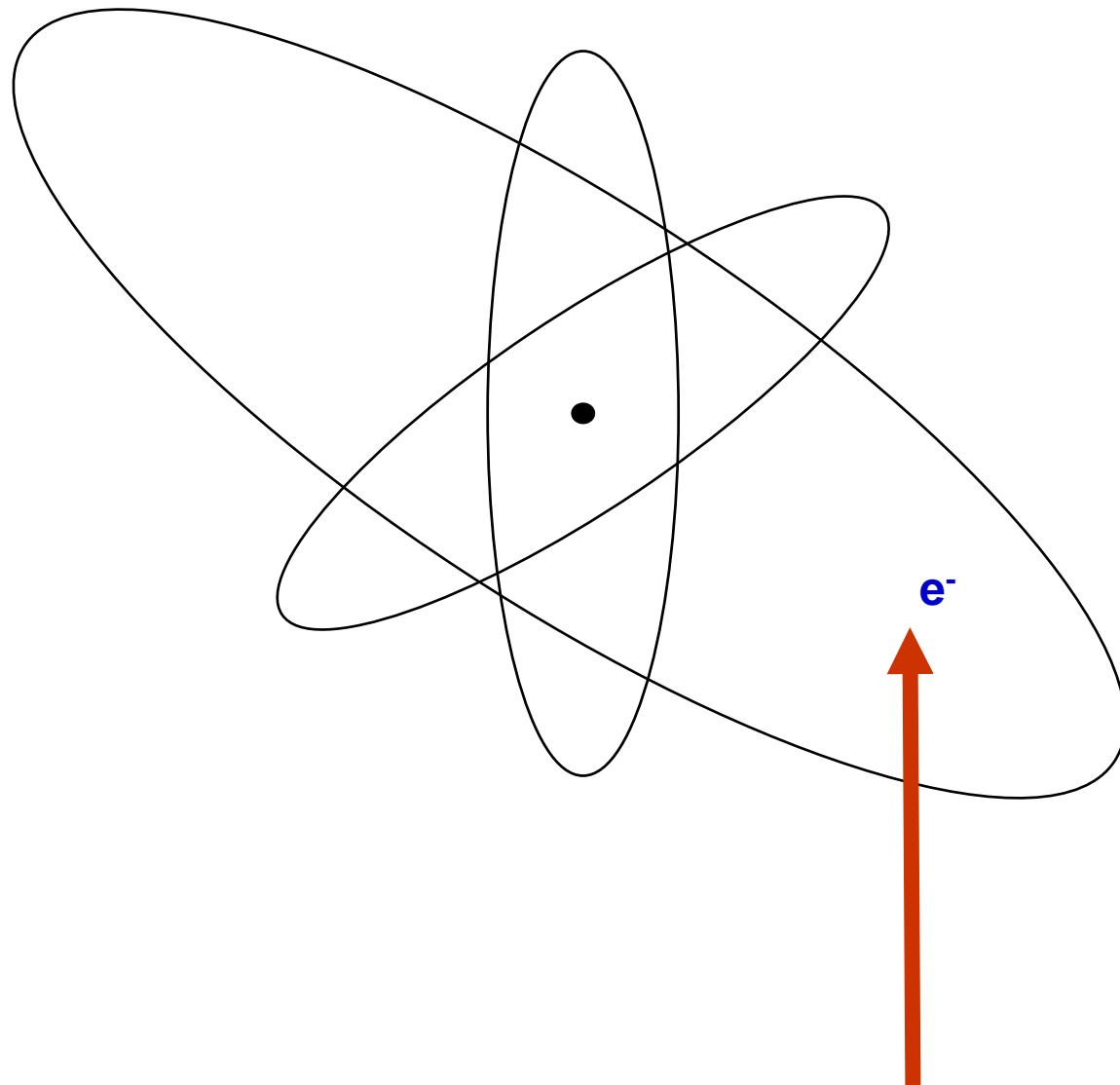
Secondary ionization



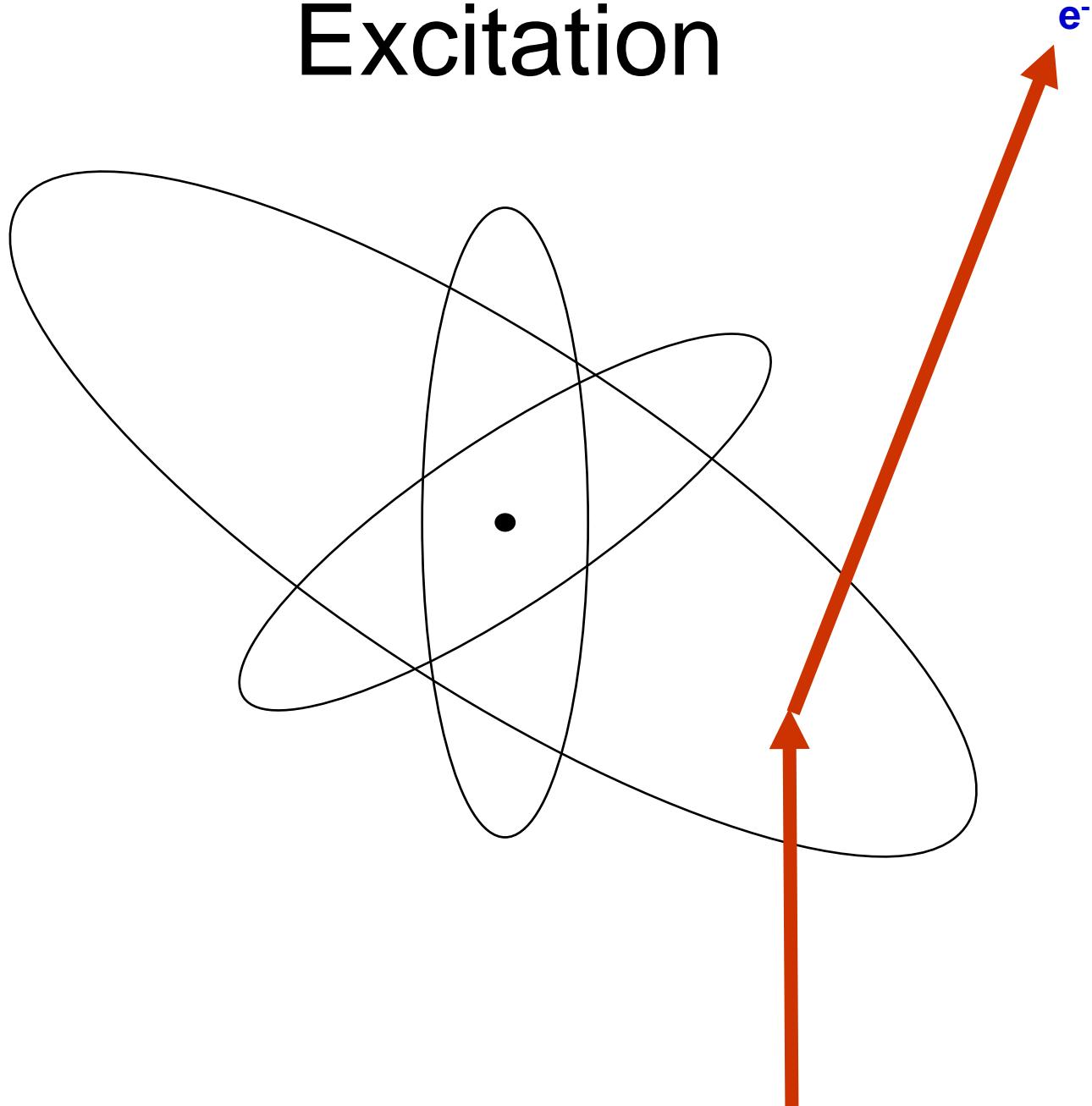
Excitation



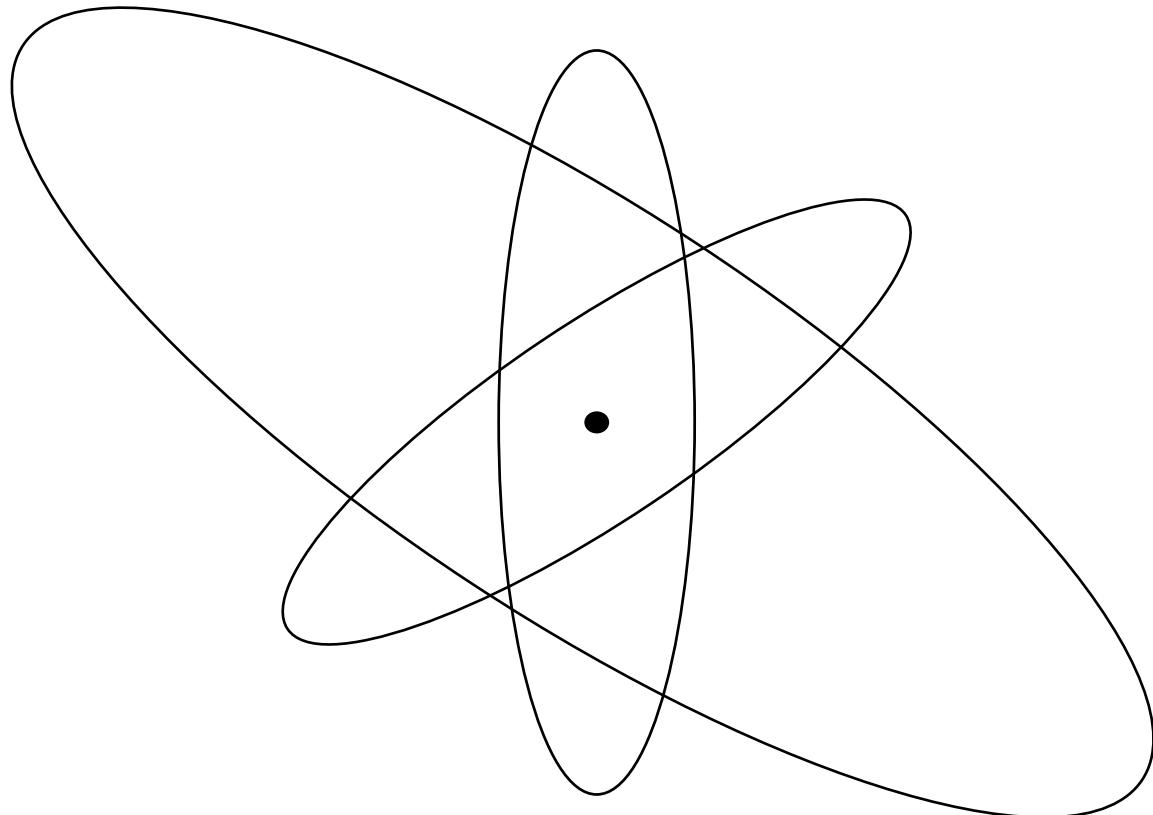
Excitation



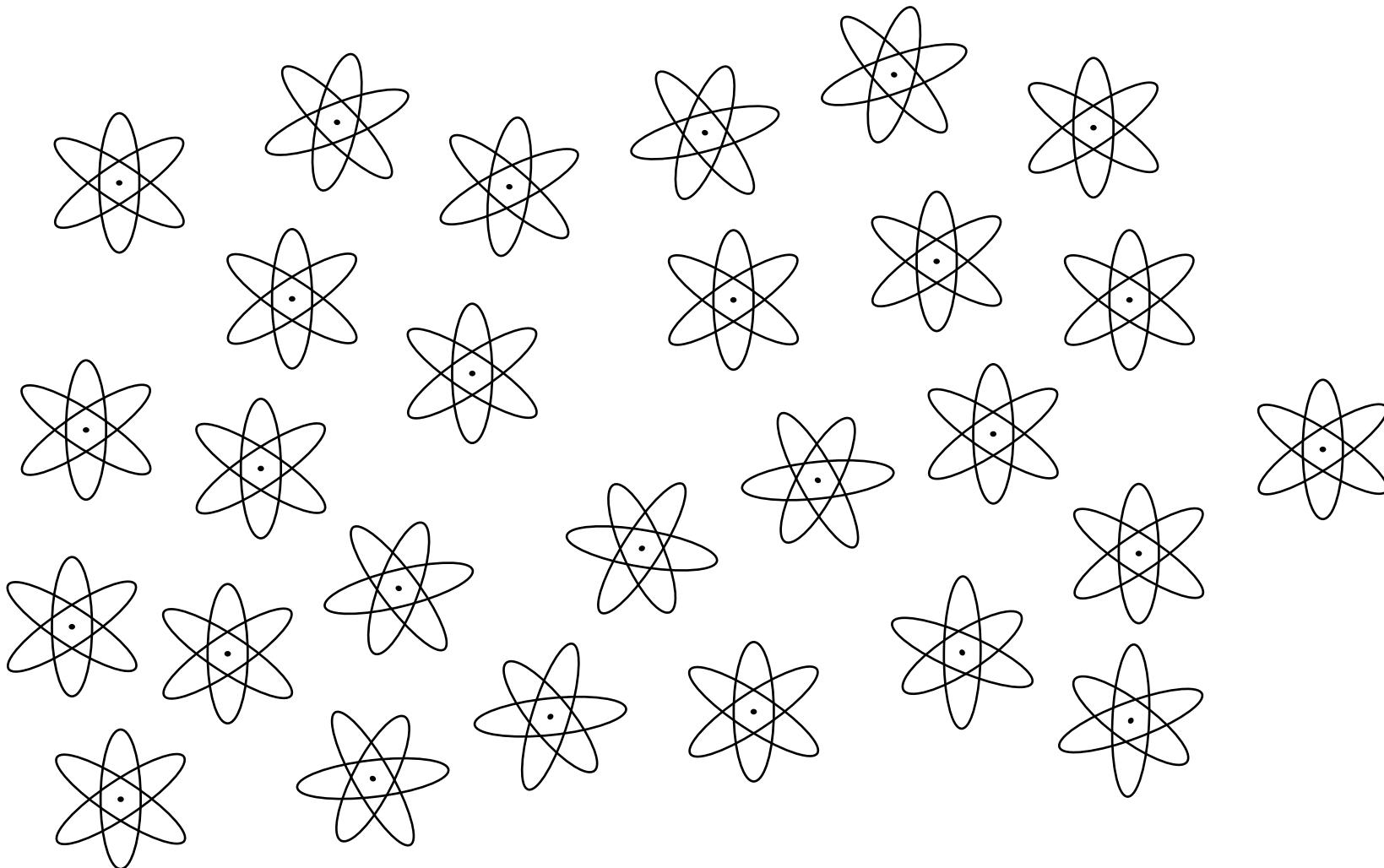
Excitation



Excitation

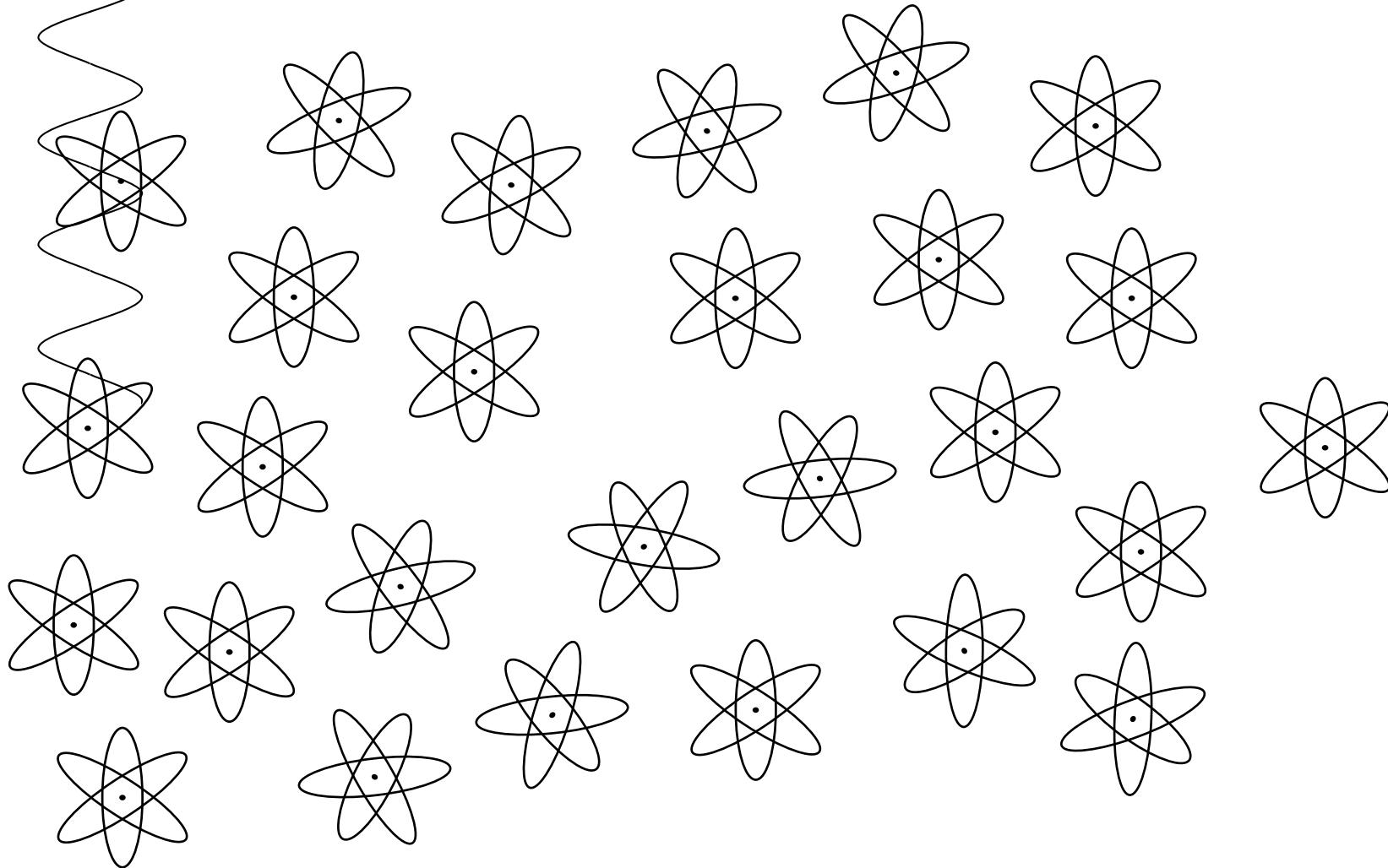


Sample atoms

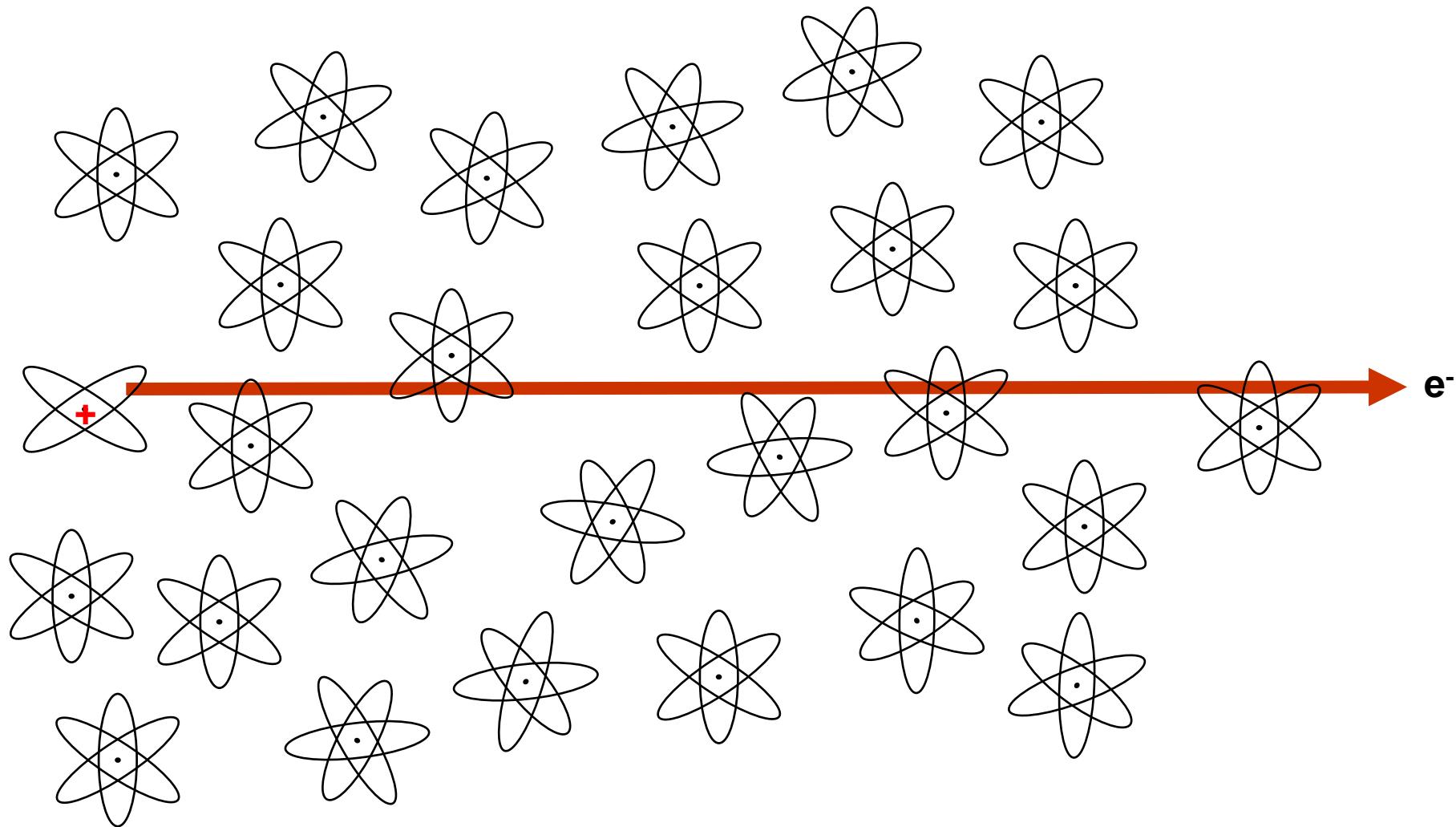




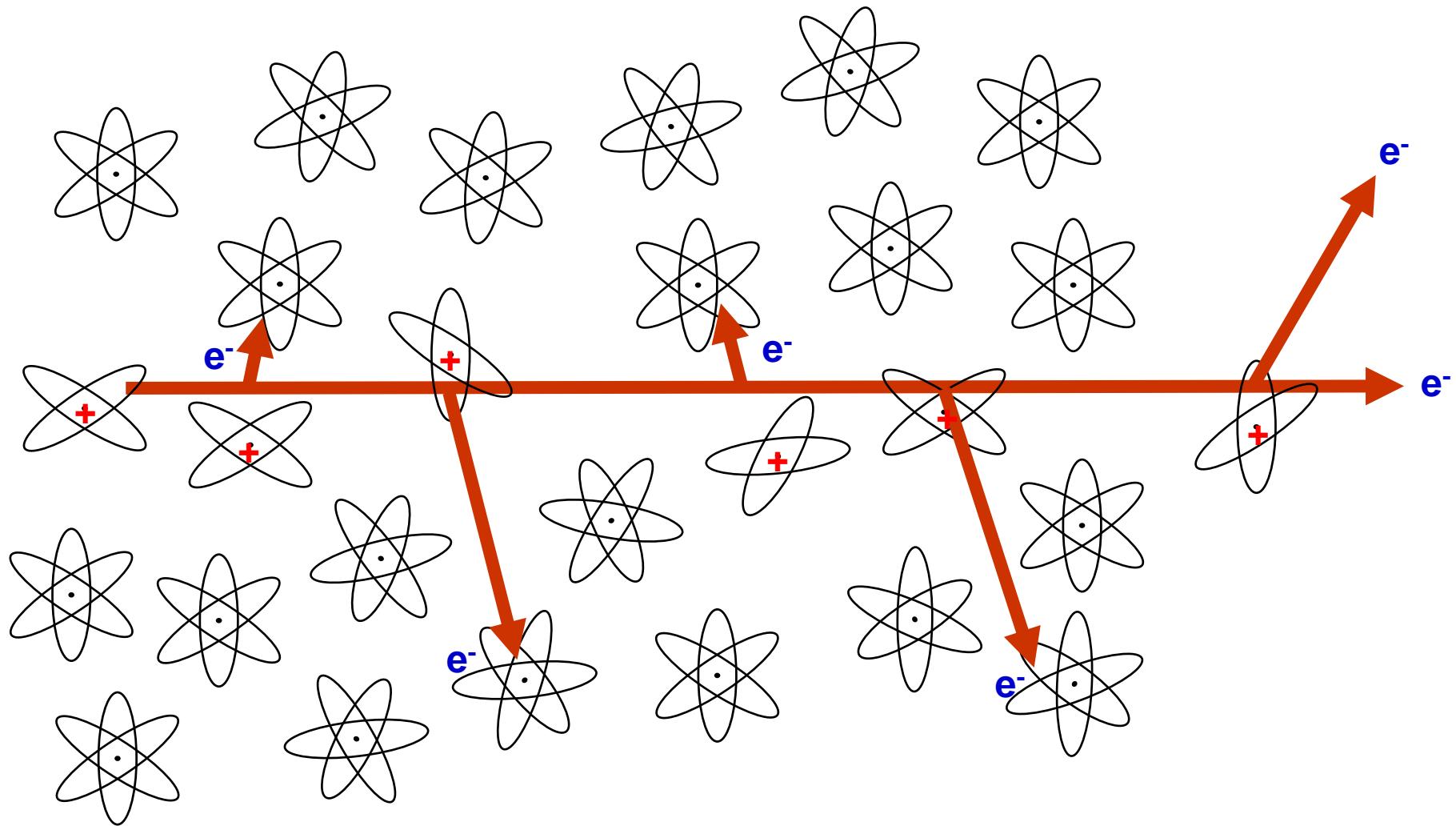
Sample atoms



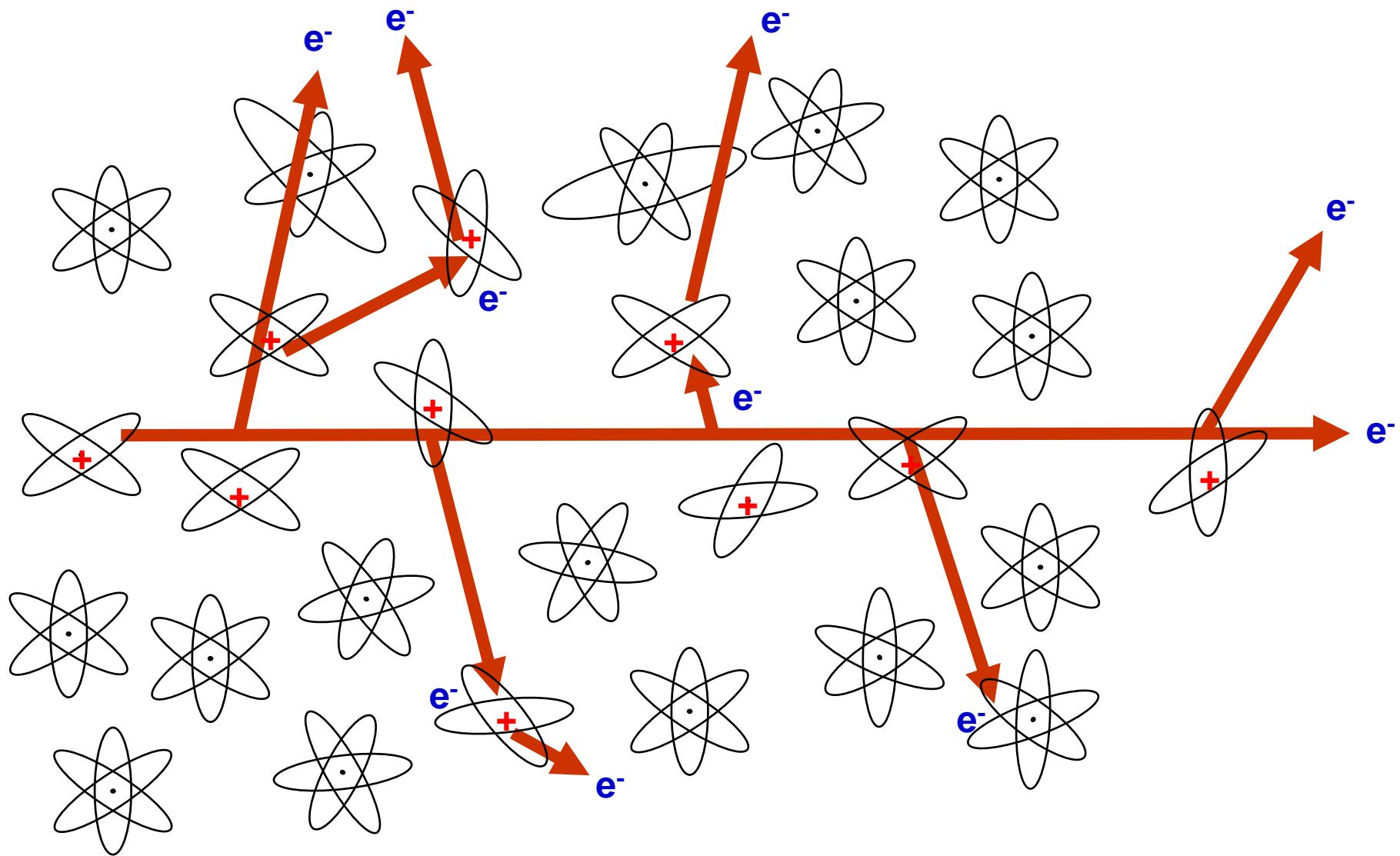
Primary ionization



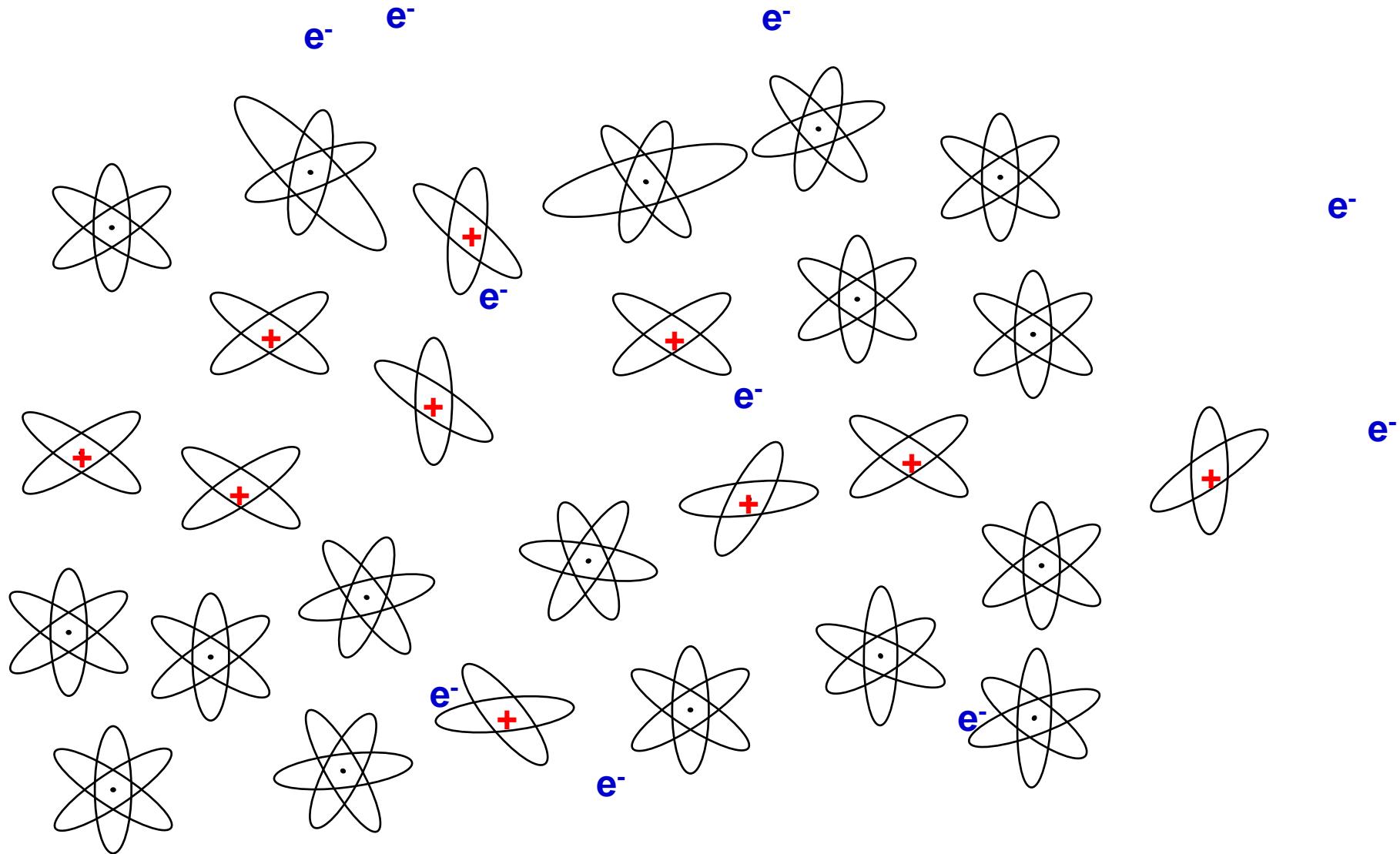
Secondary ionizations



Ionization track



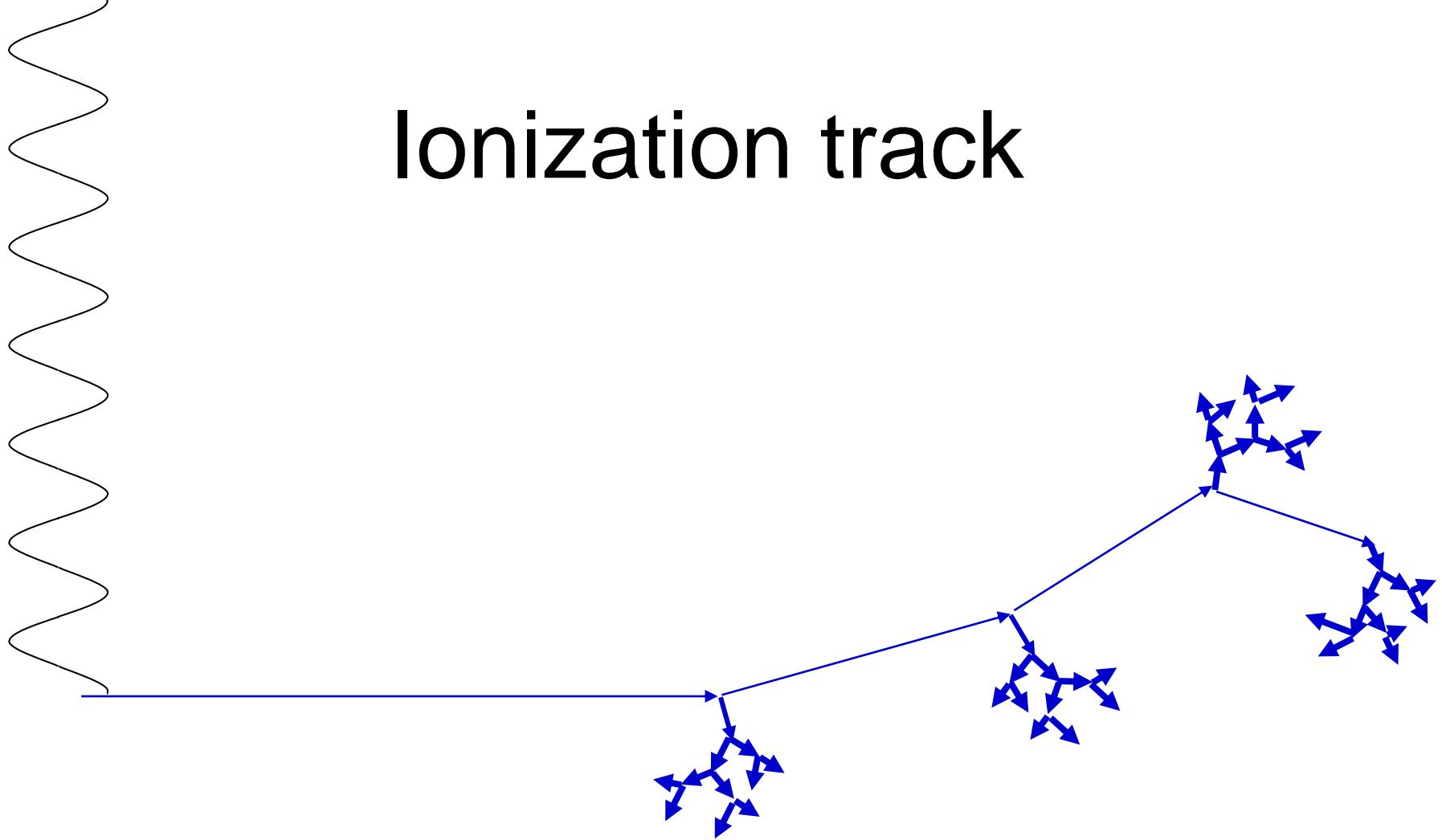
Ionization track



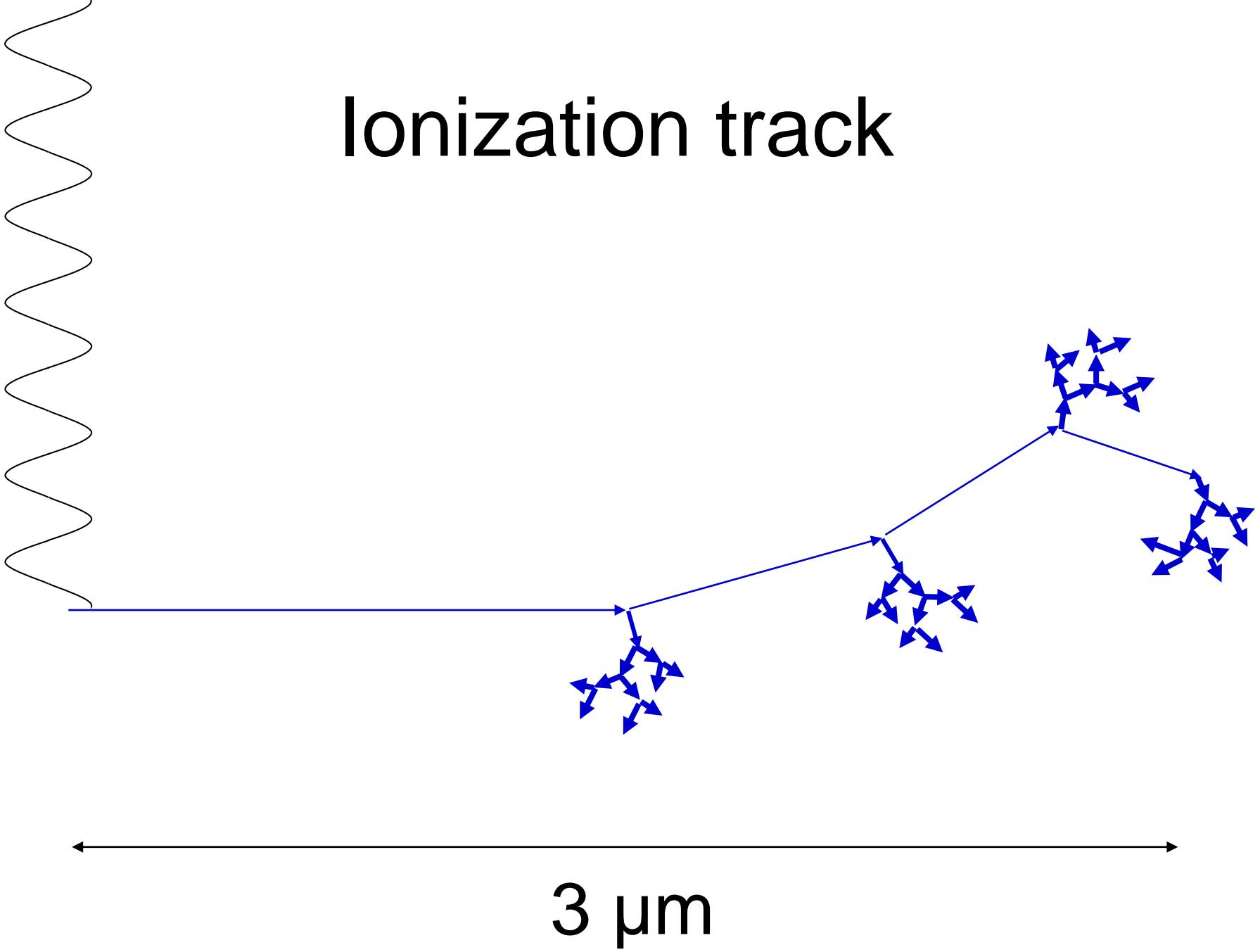


Ionization track

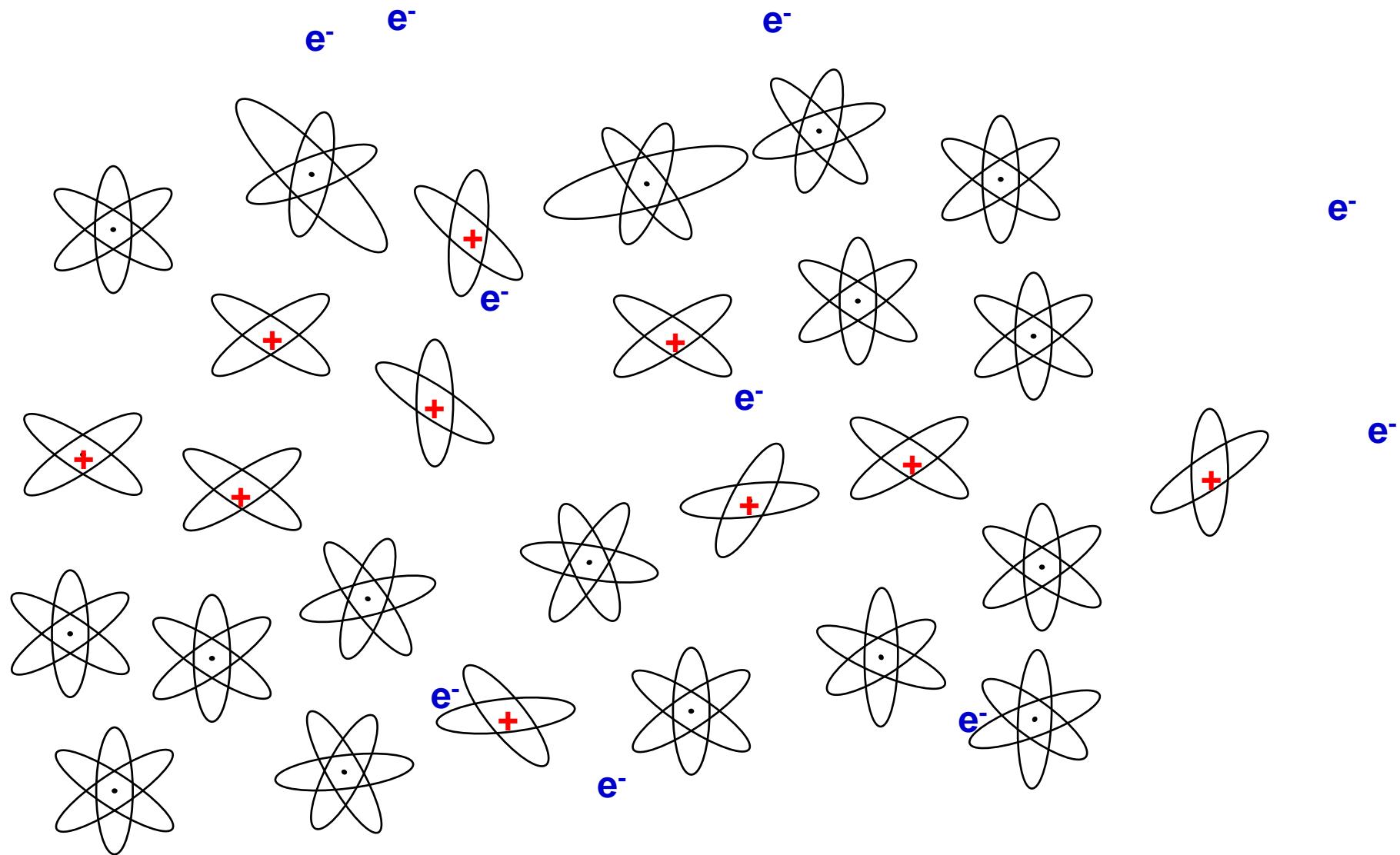
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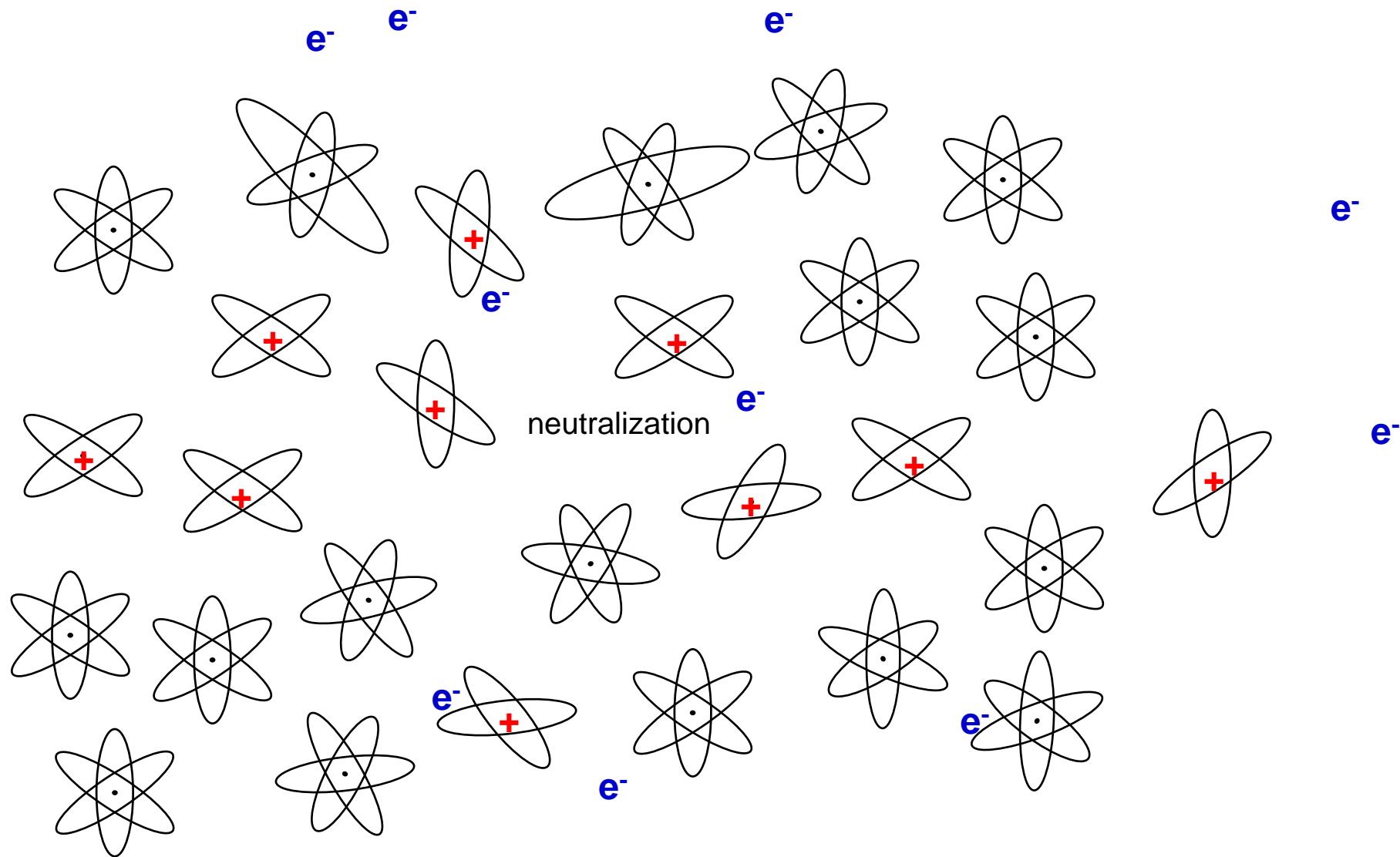
Ionization track



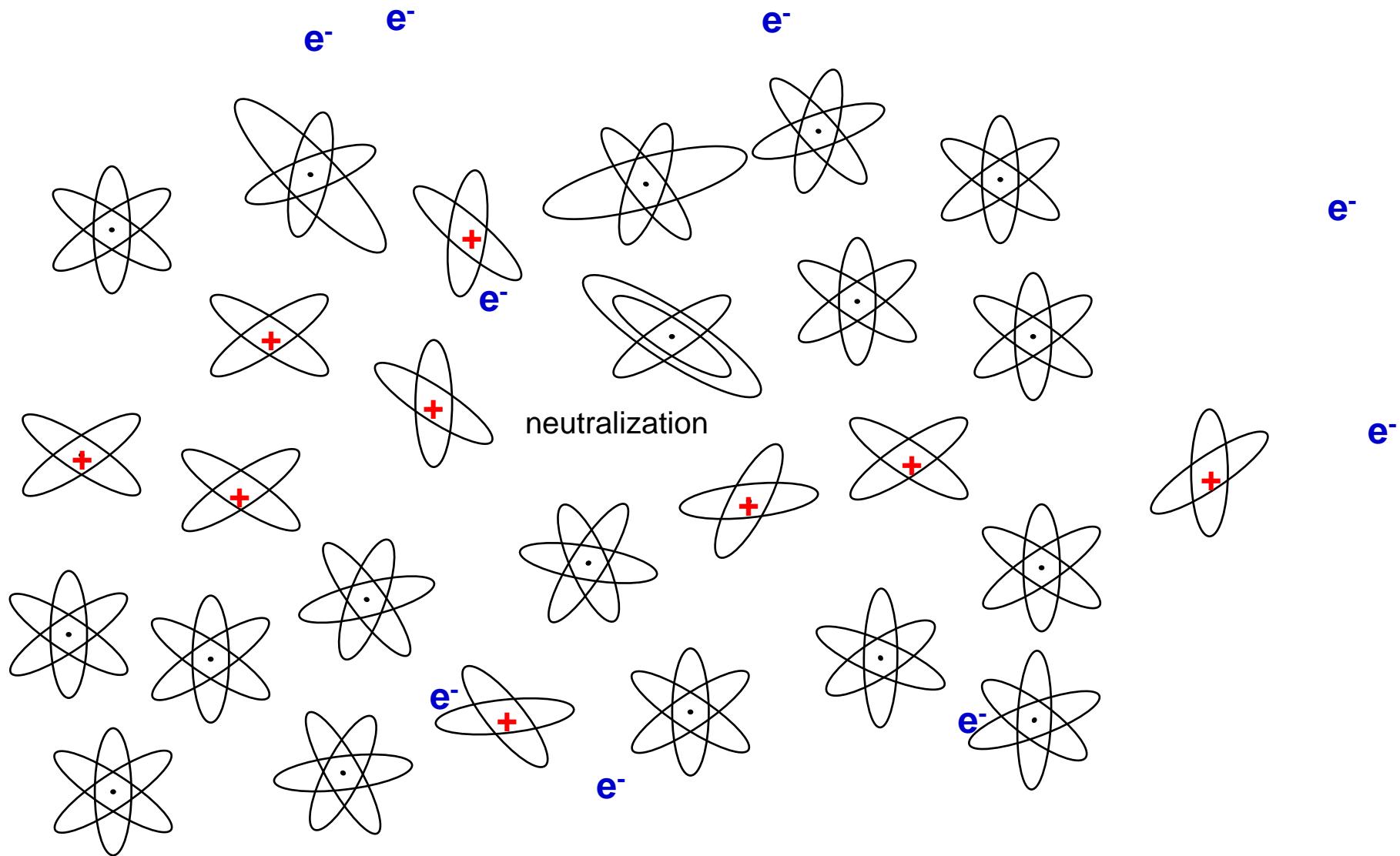
initial effects



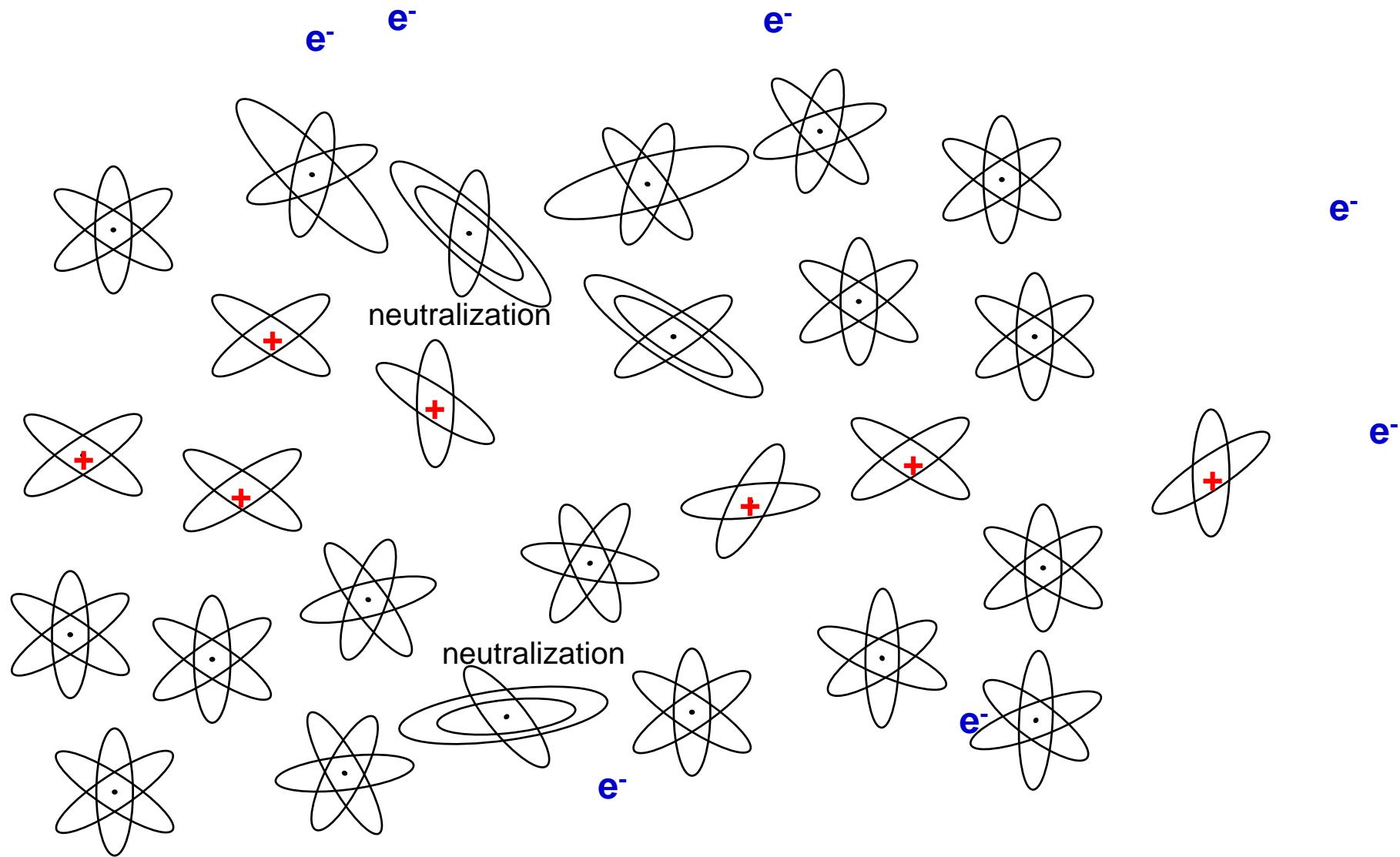
initial effects



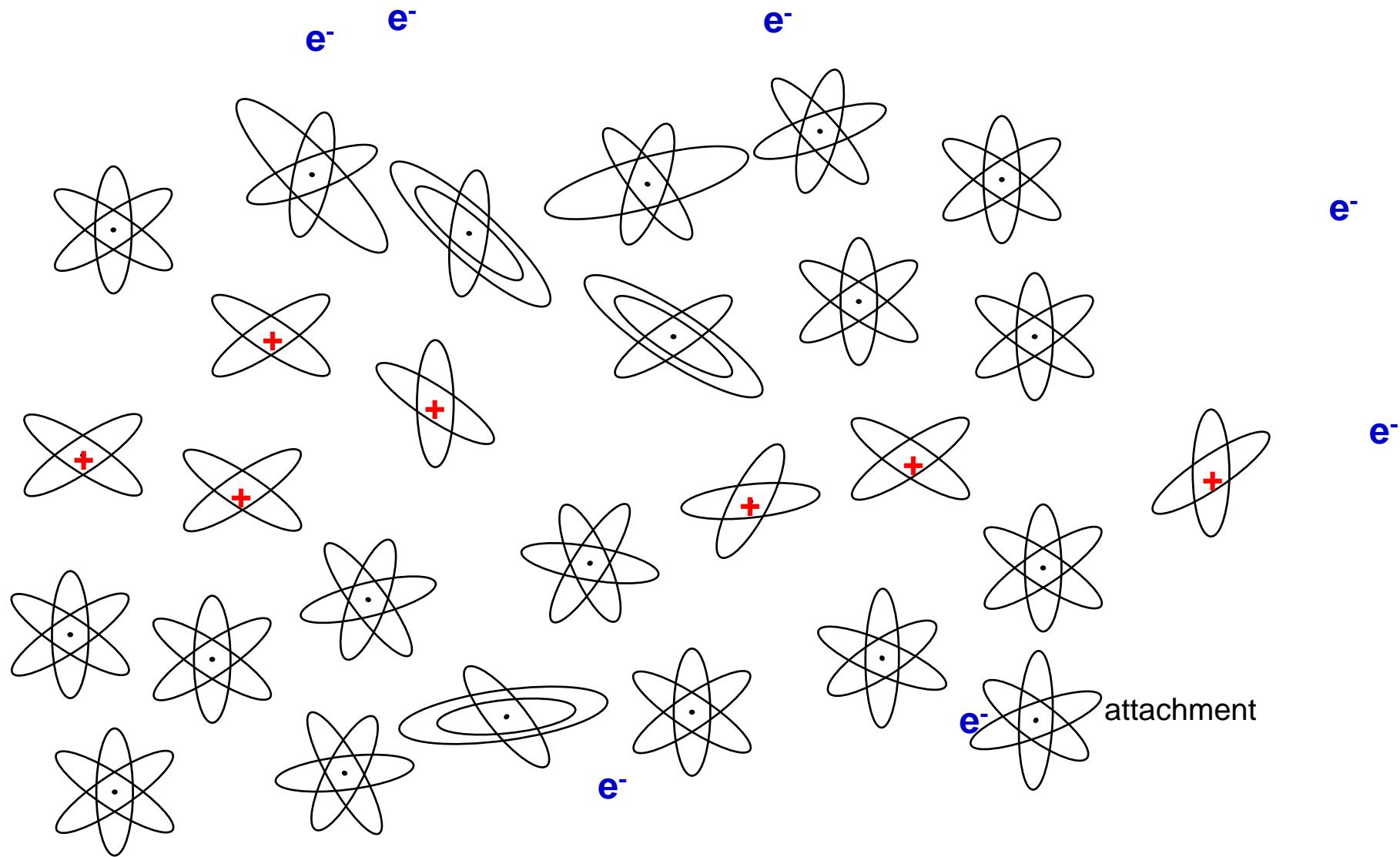
initial effects



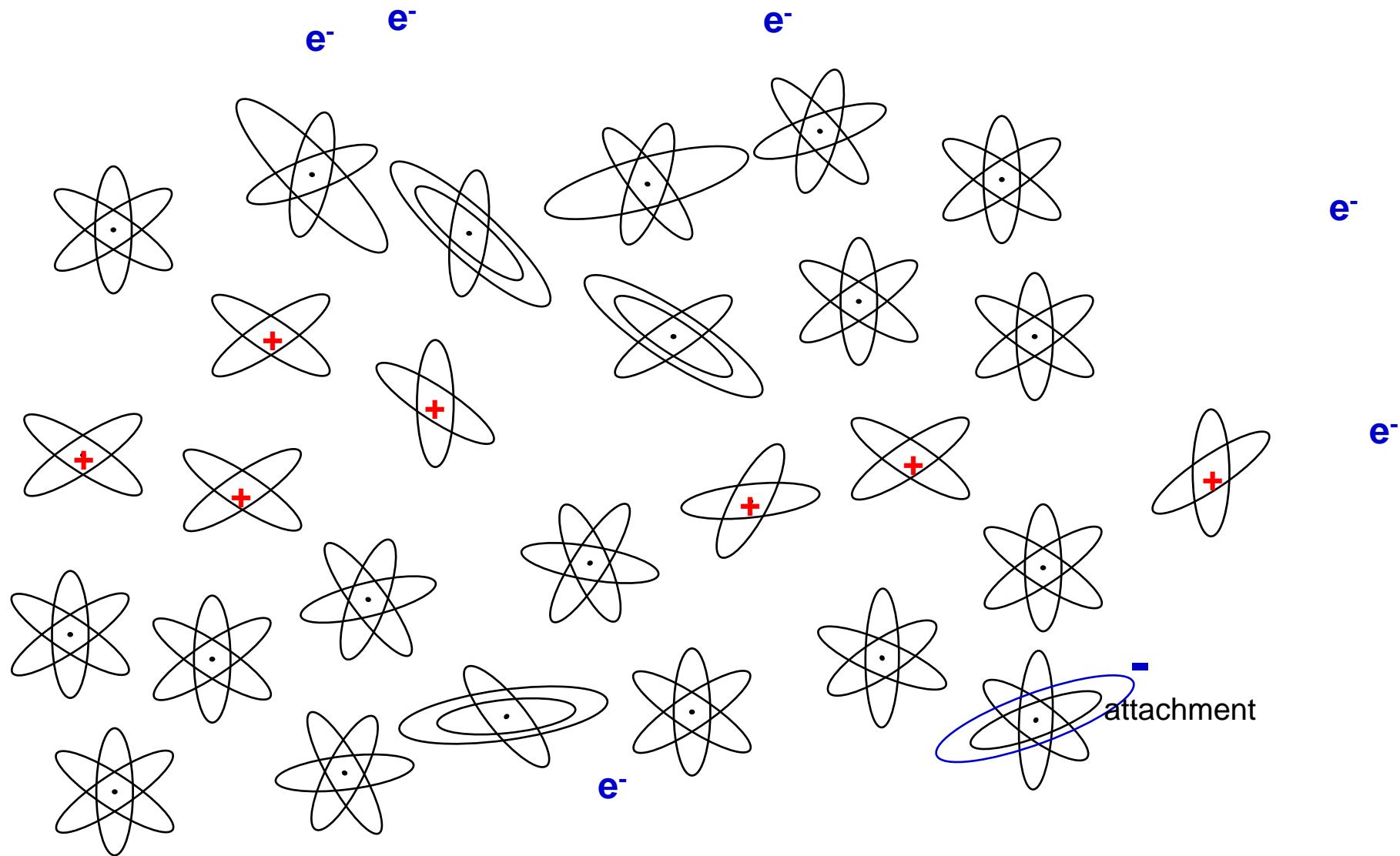
initial effects



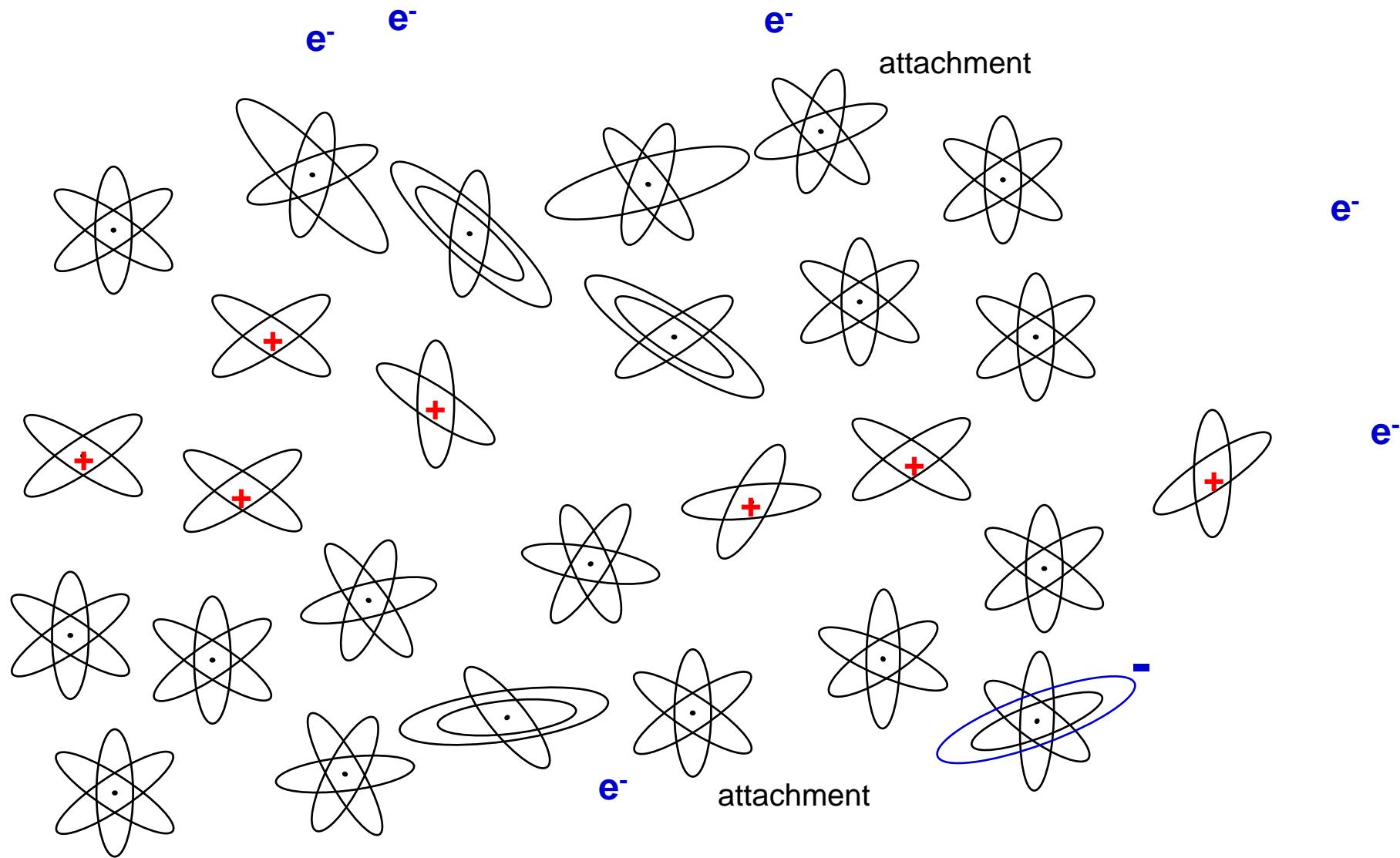
initial effects



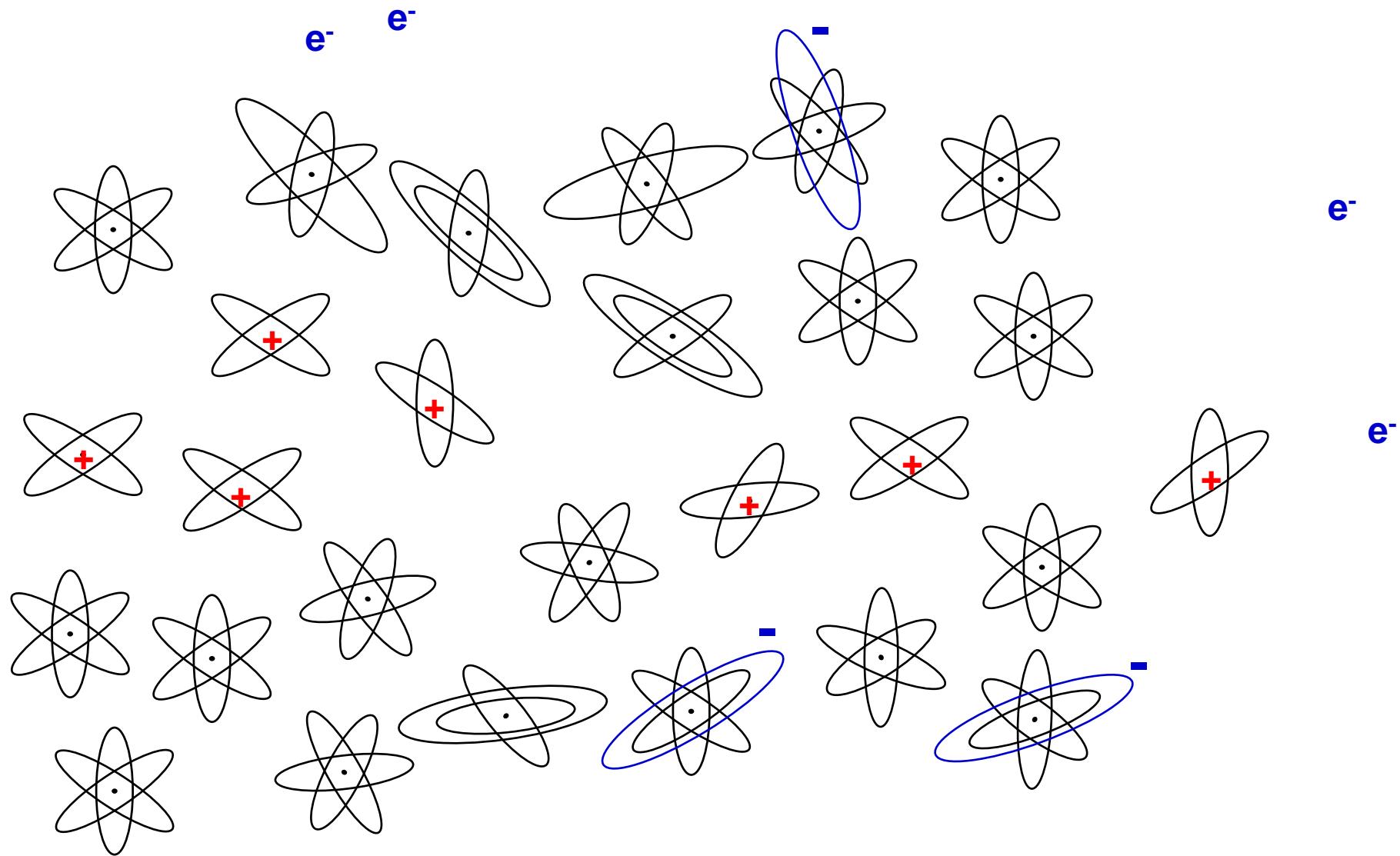
initial effects



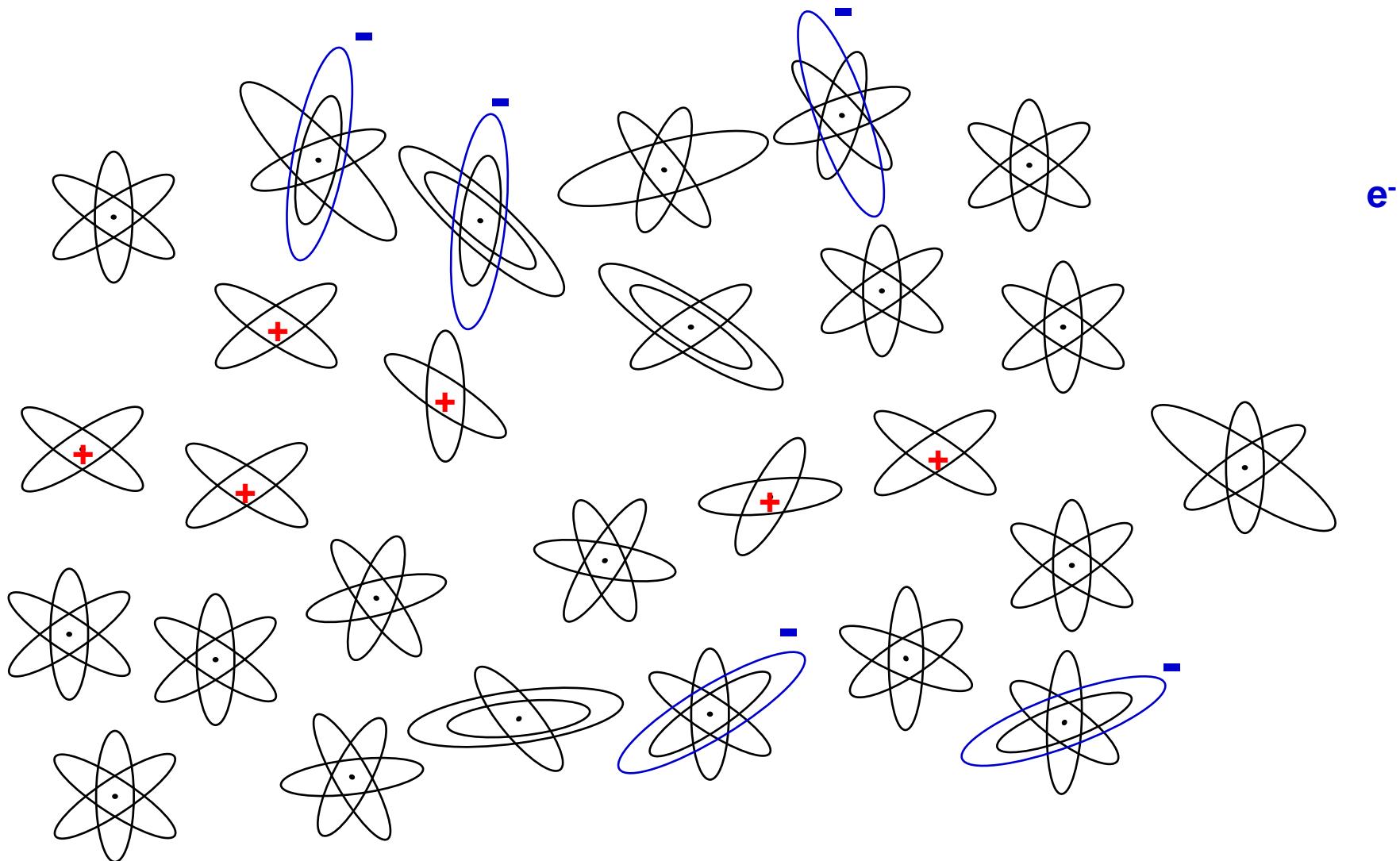
initial effects



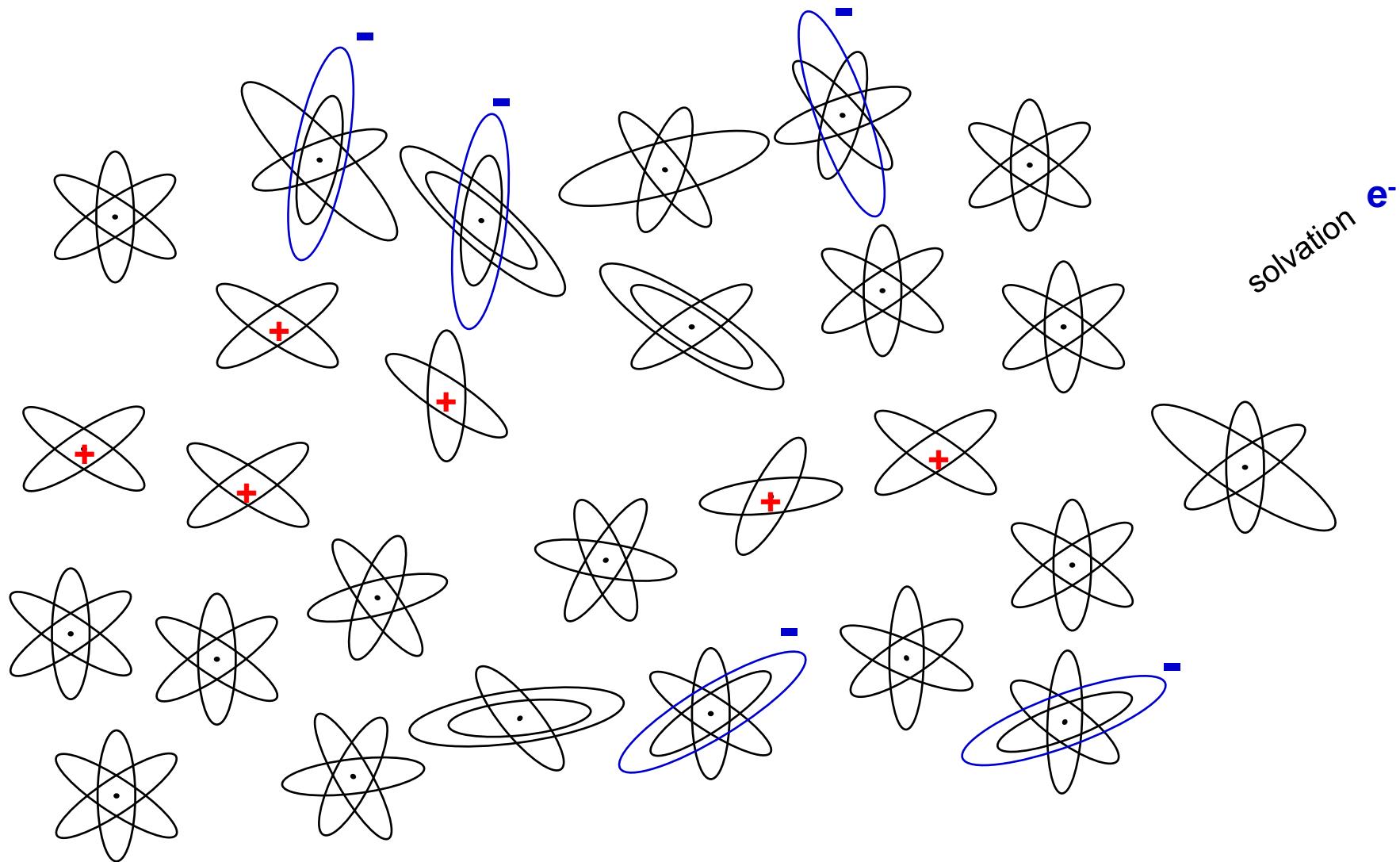
initial effects



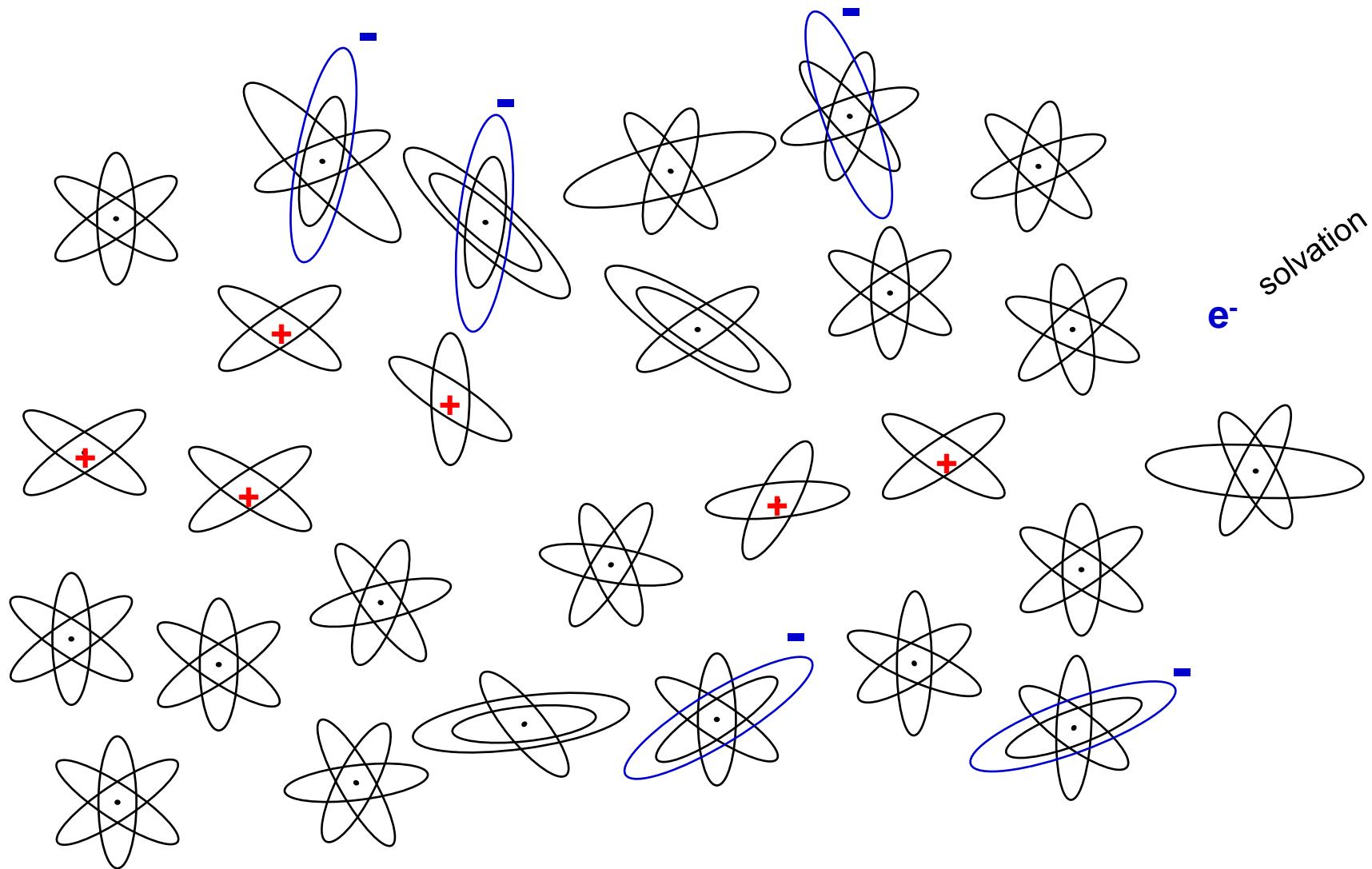
initial effects



initial effects

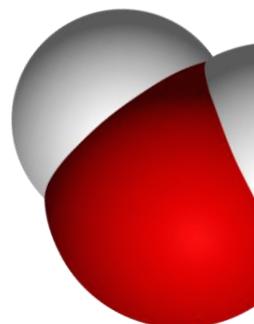
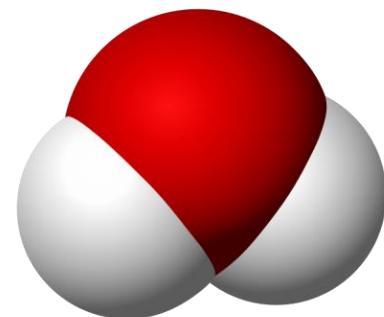
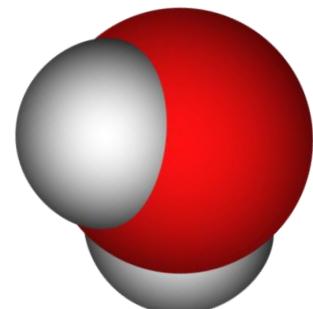
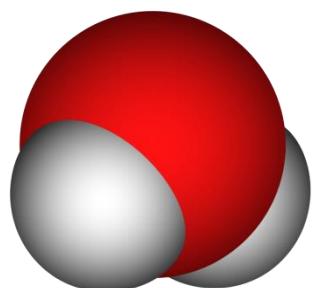


initial effects



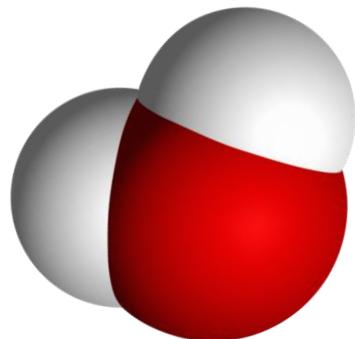
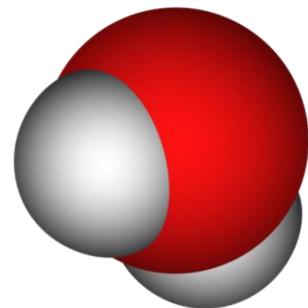
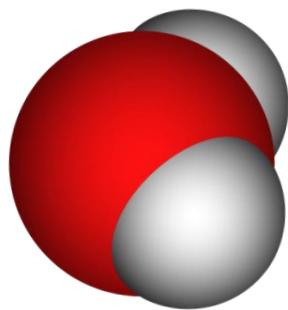
Solvated electron

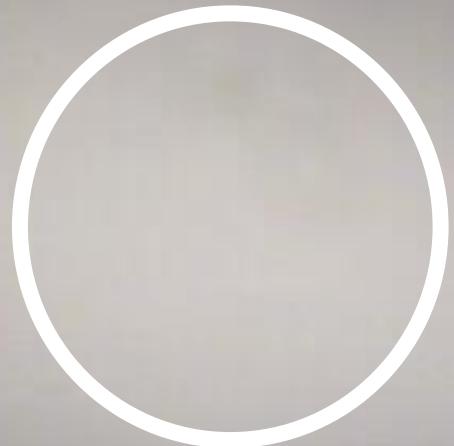
e^-

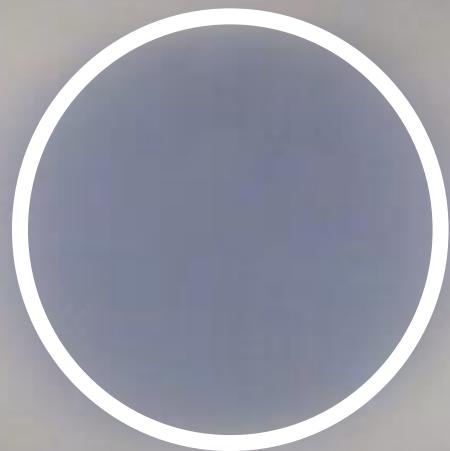


Solvated electron

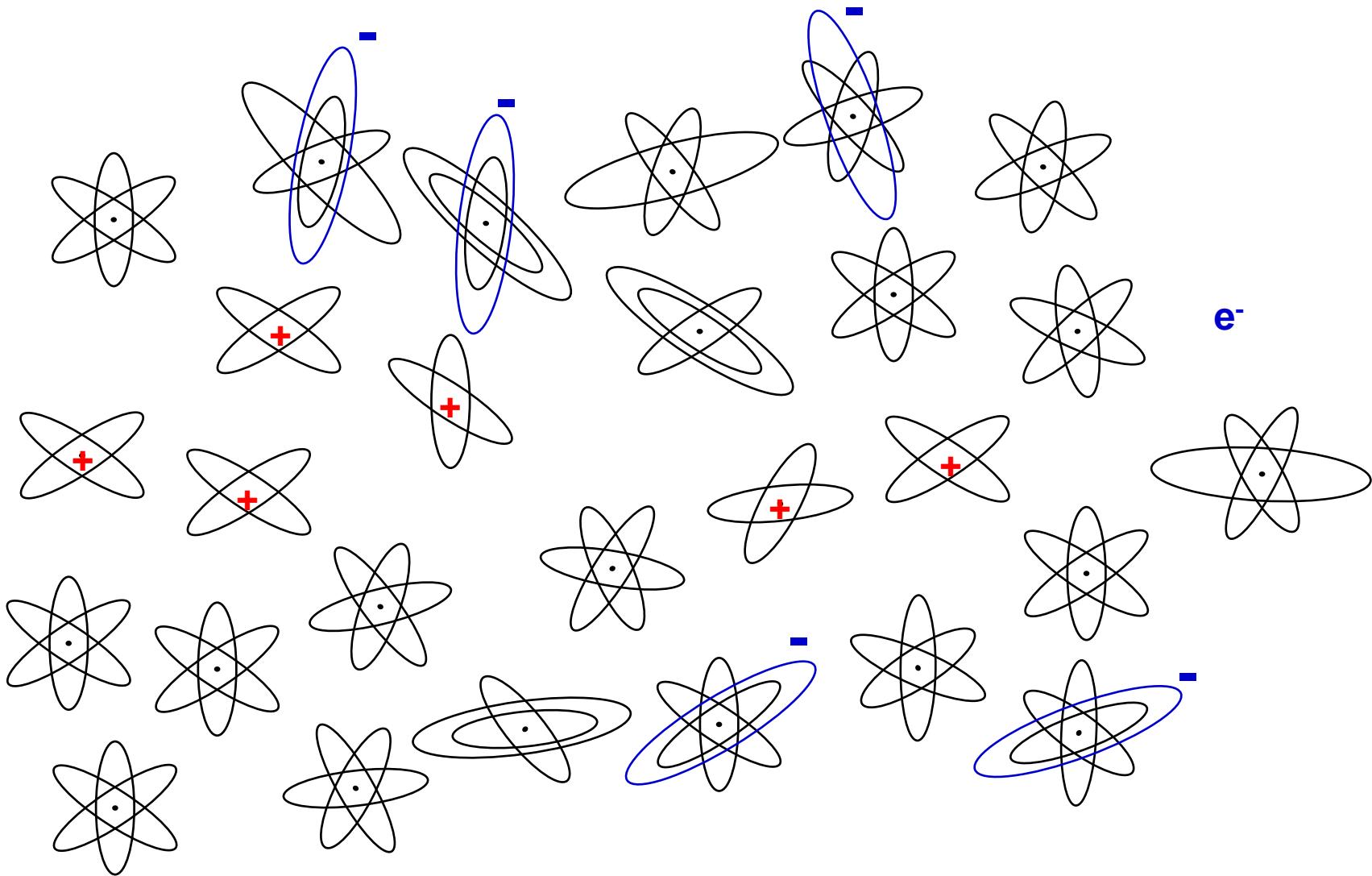
e^-



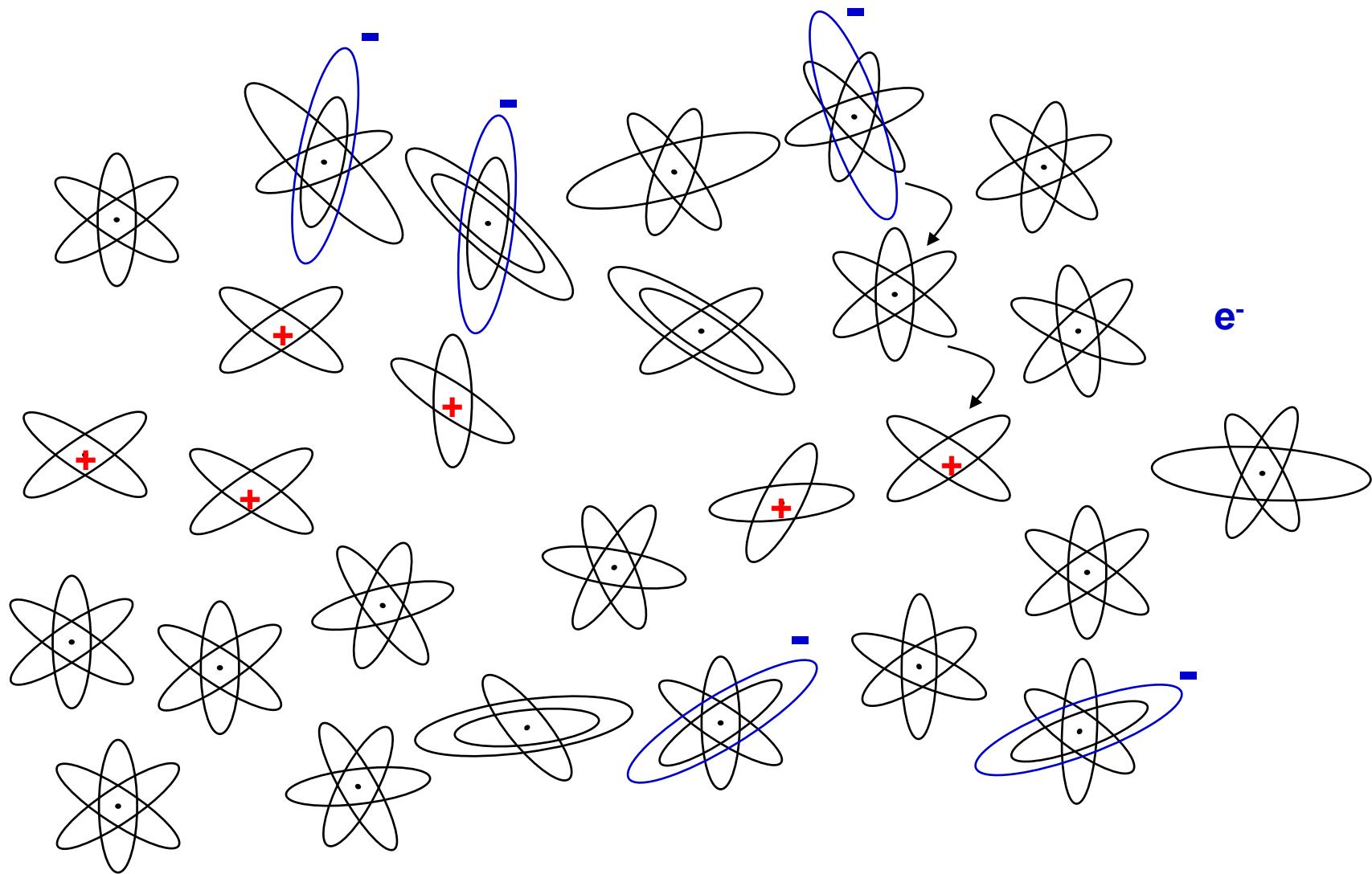




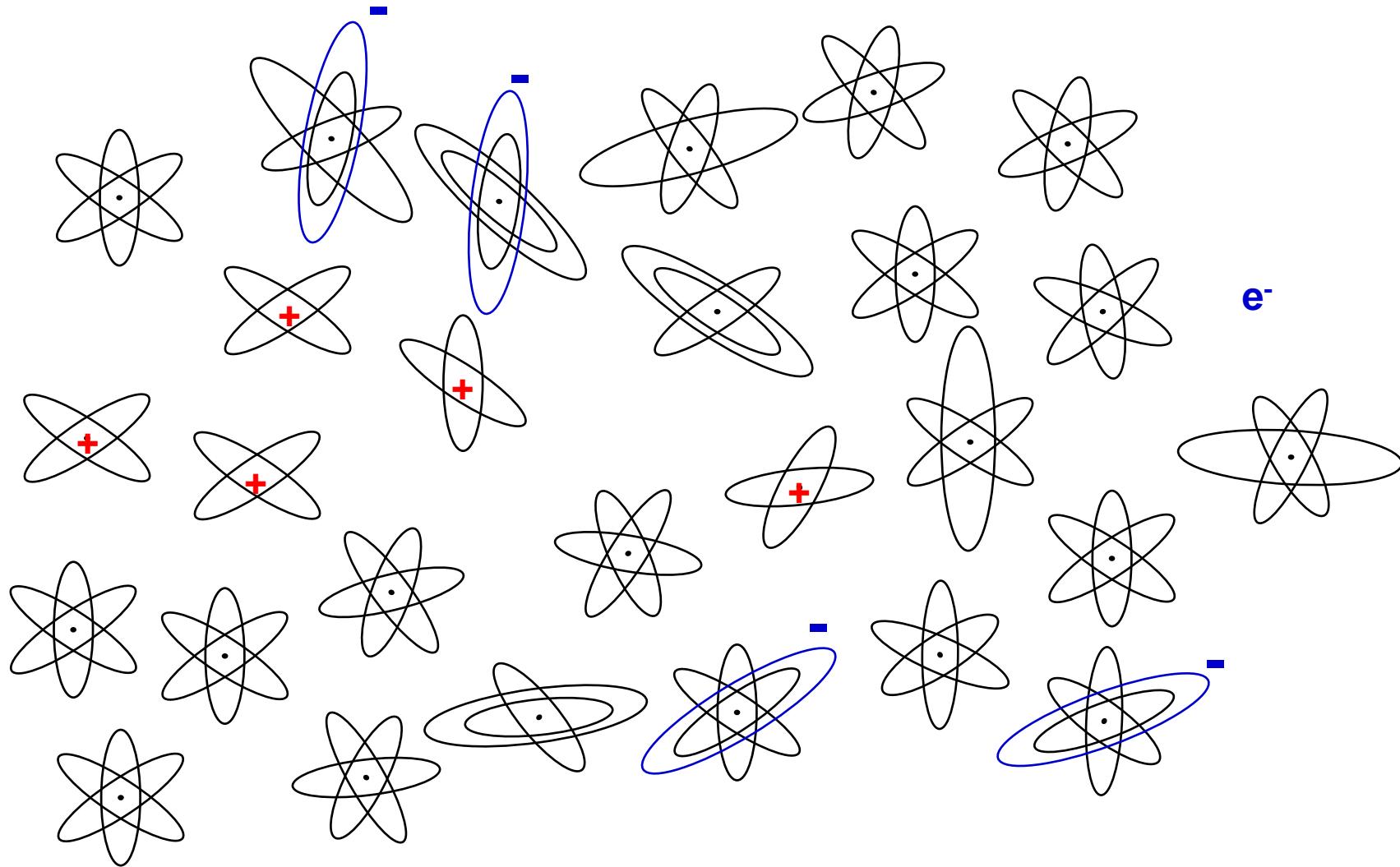
initial effects



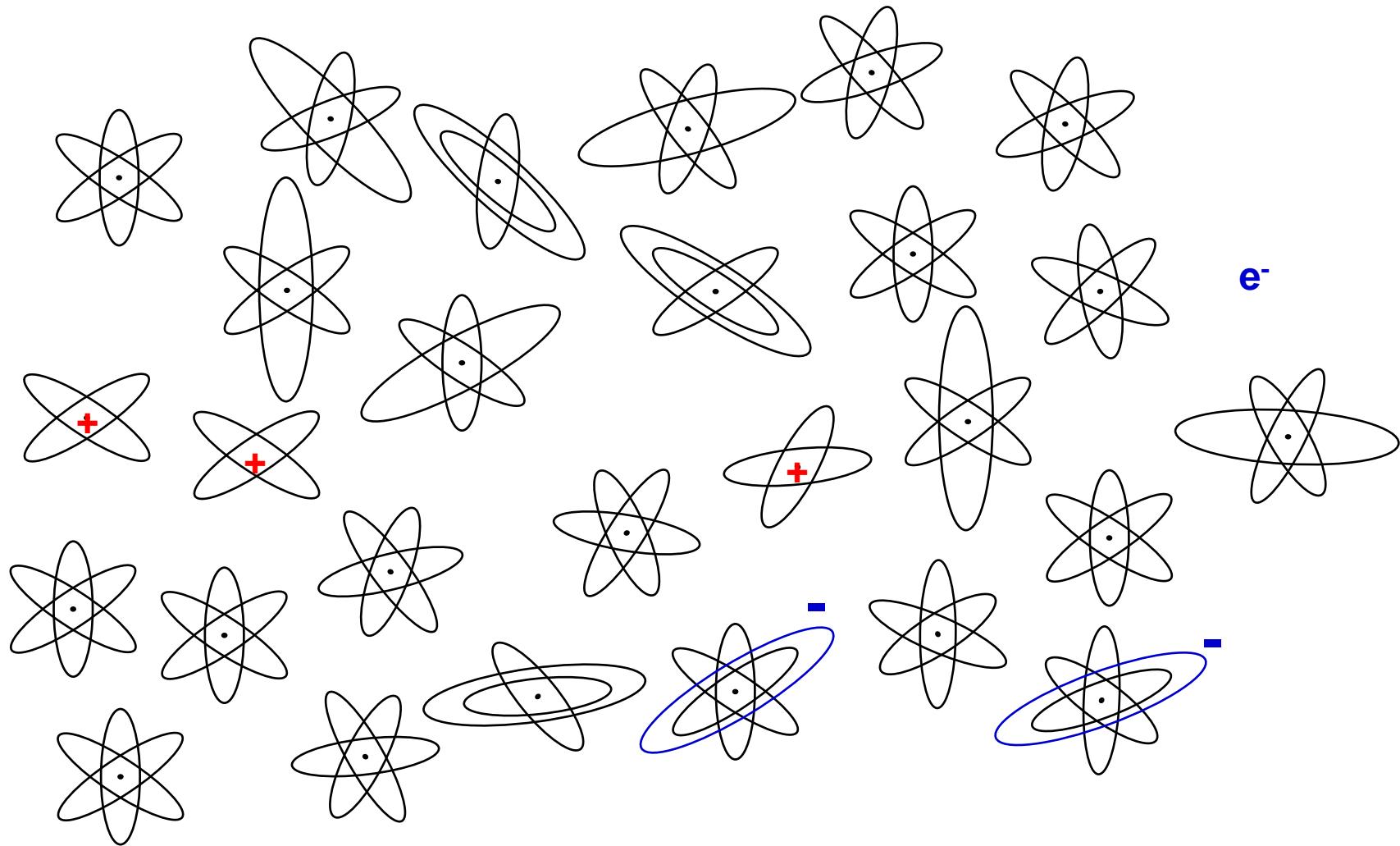
initial effects



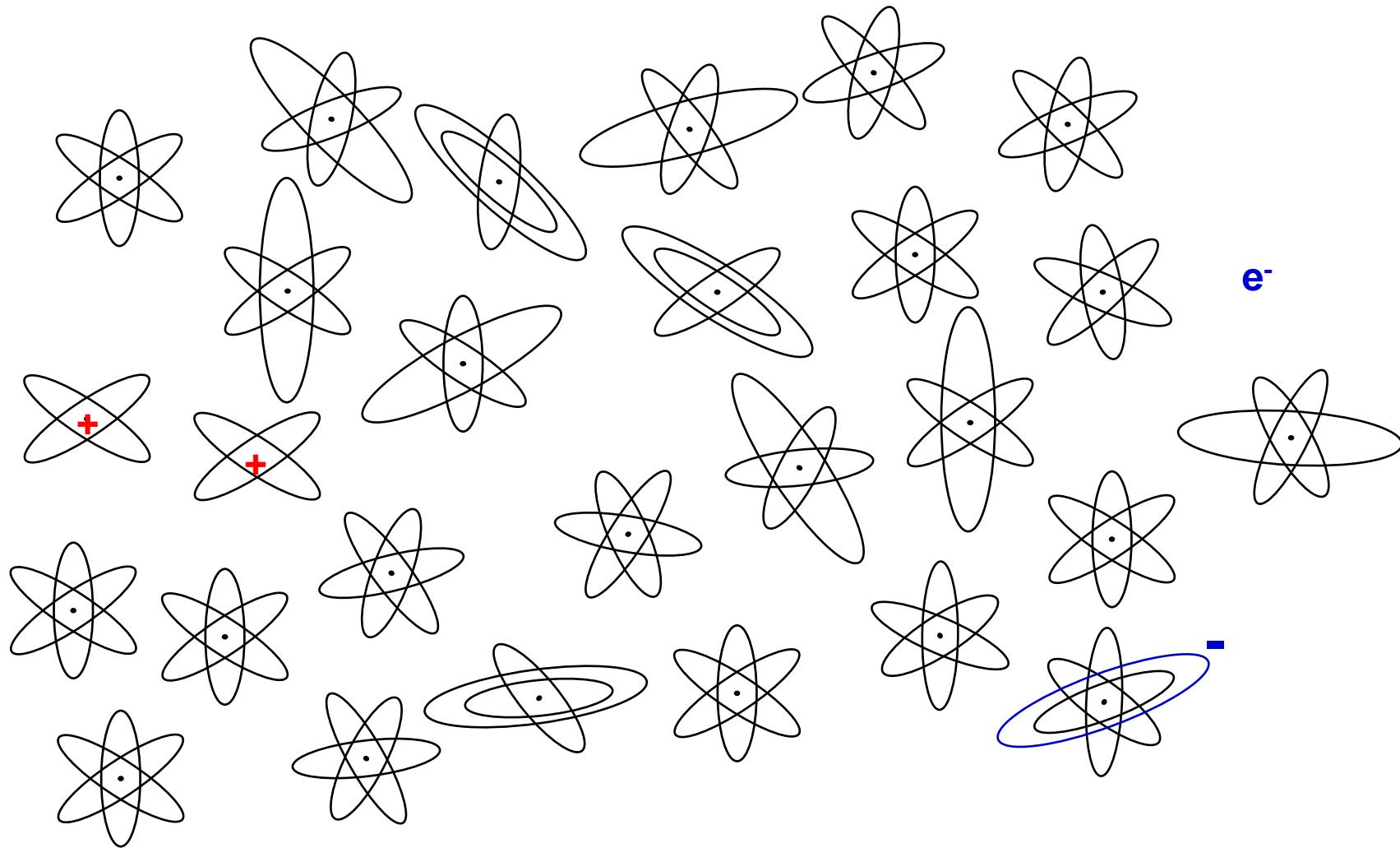
initial effects



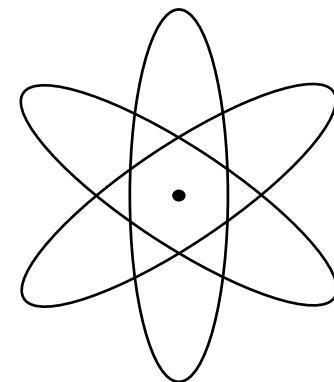
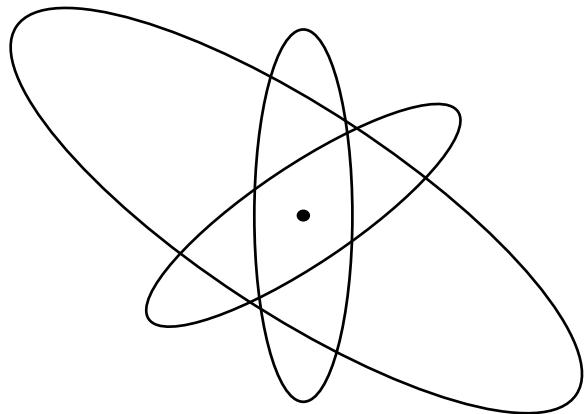
initial effects



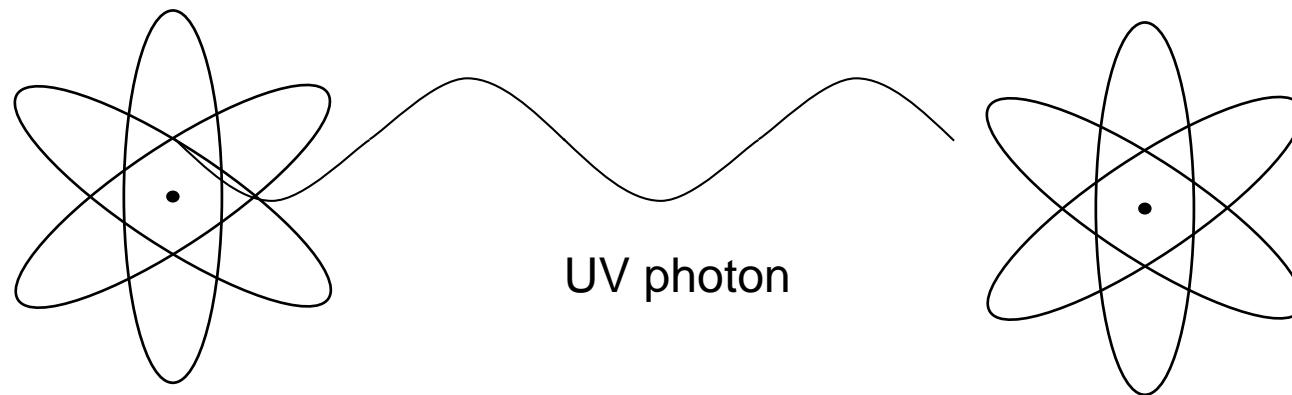
initial effects



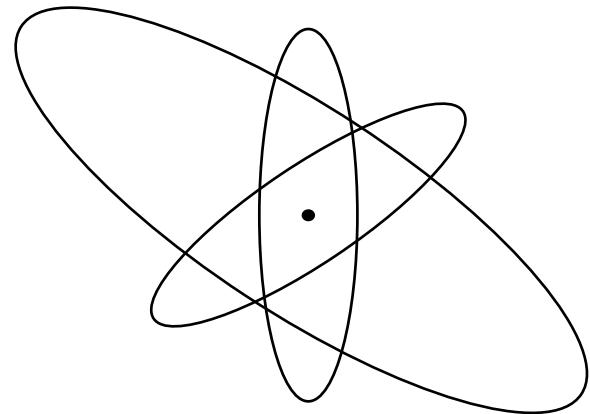
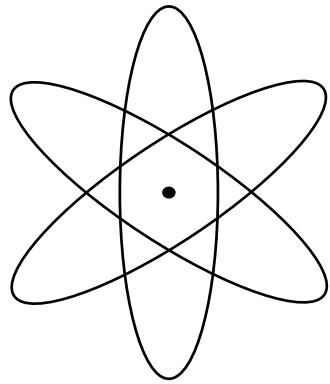
Excitation transfer



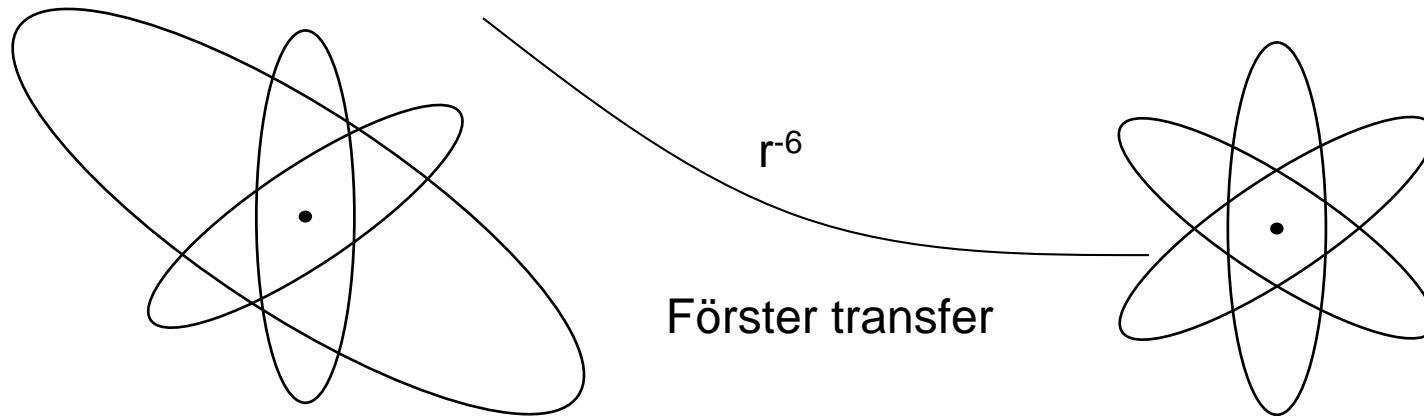
Excitation transfer



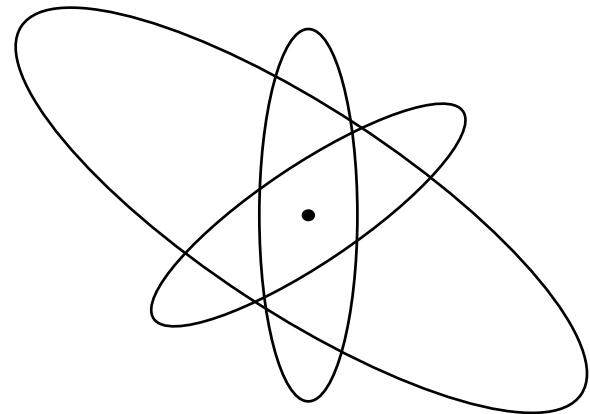
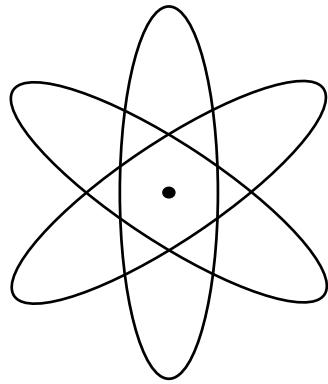
Excitation transfer



Excitation transfer

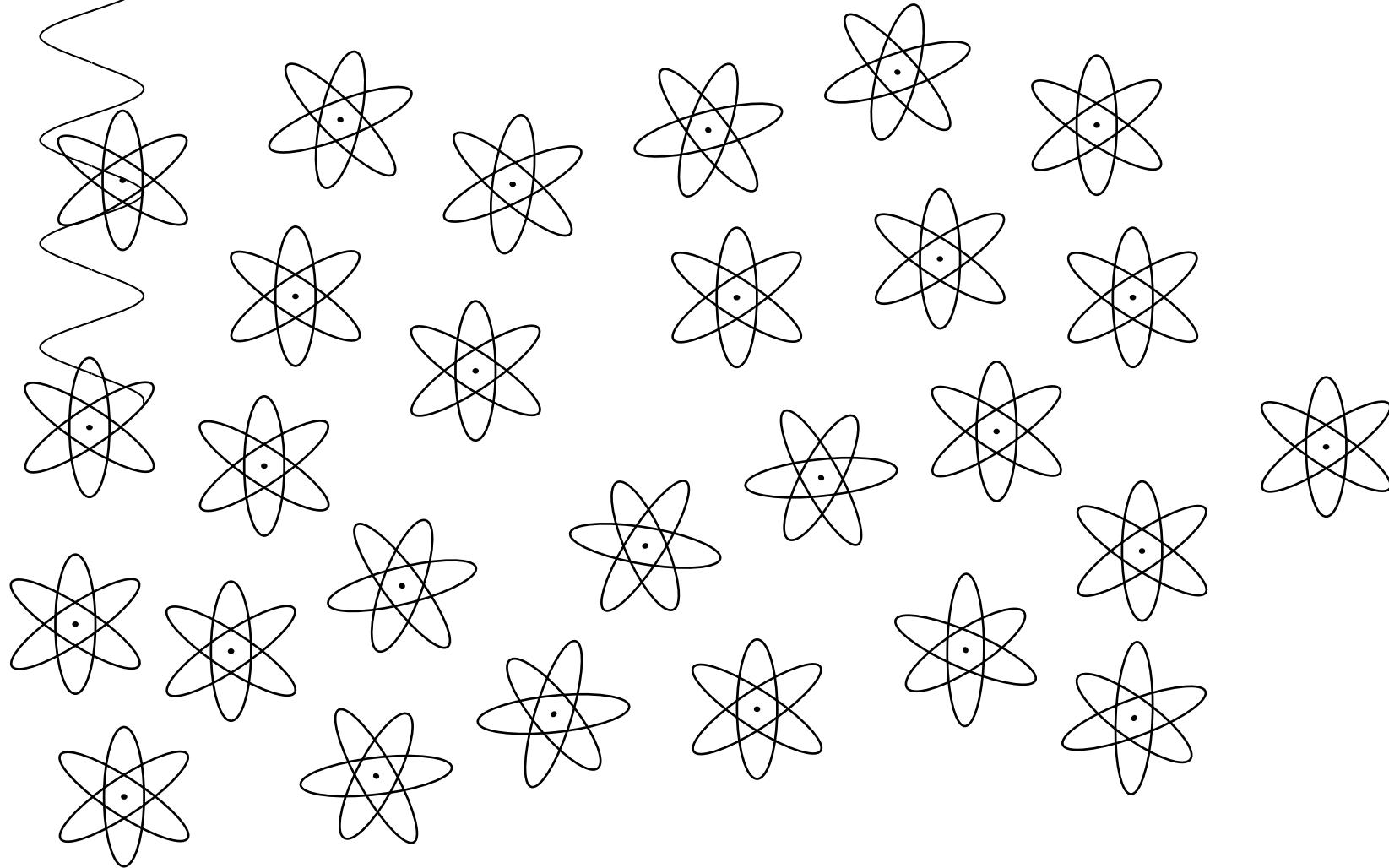


Excitation transfer

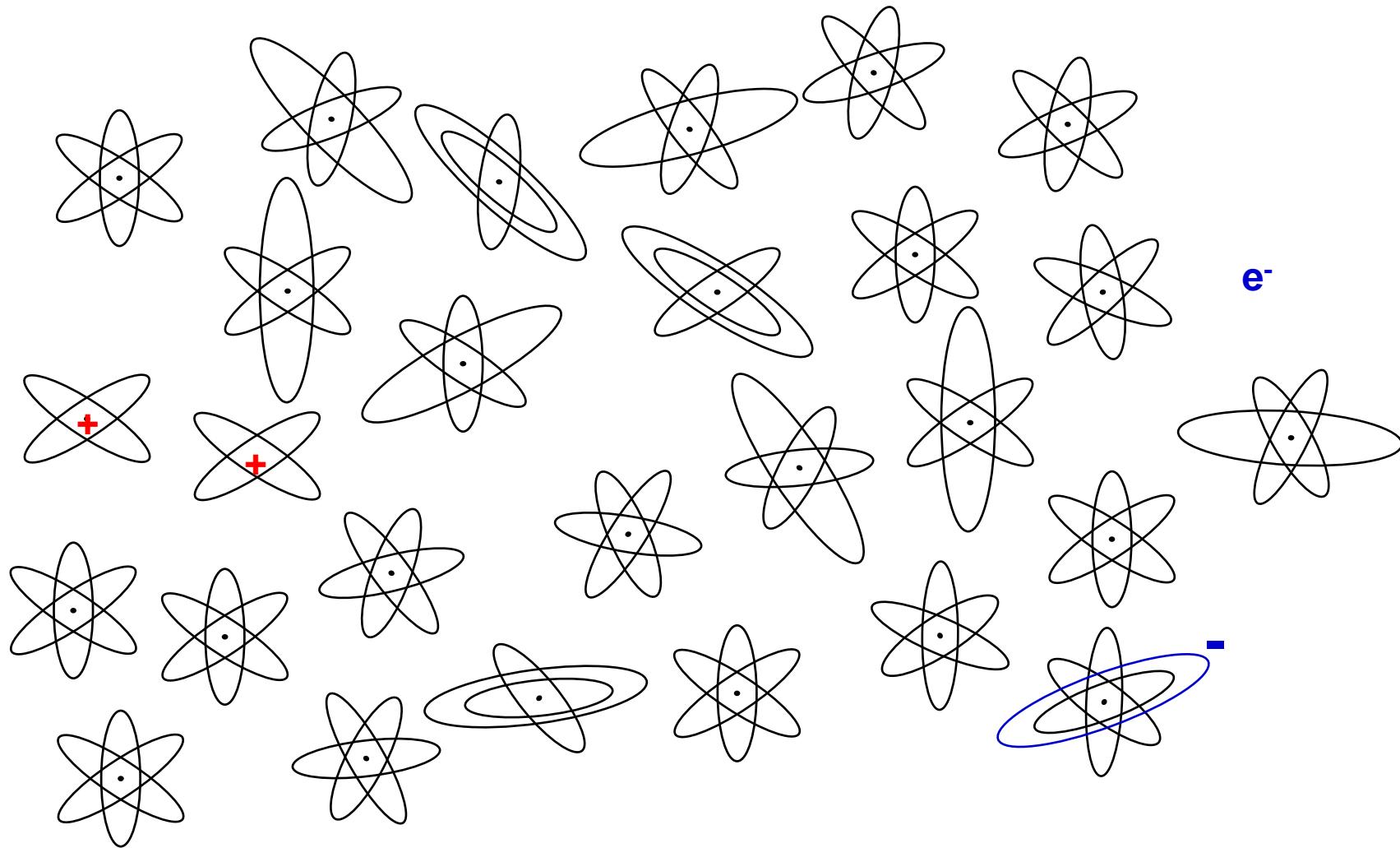




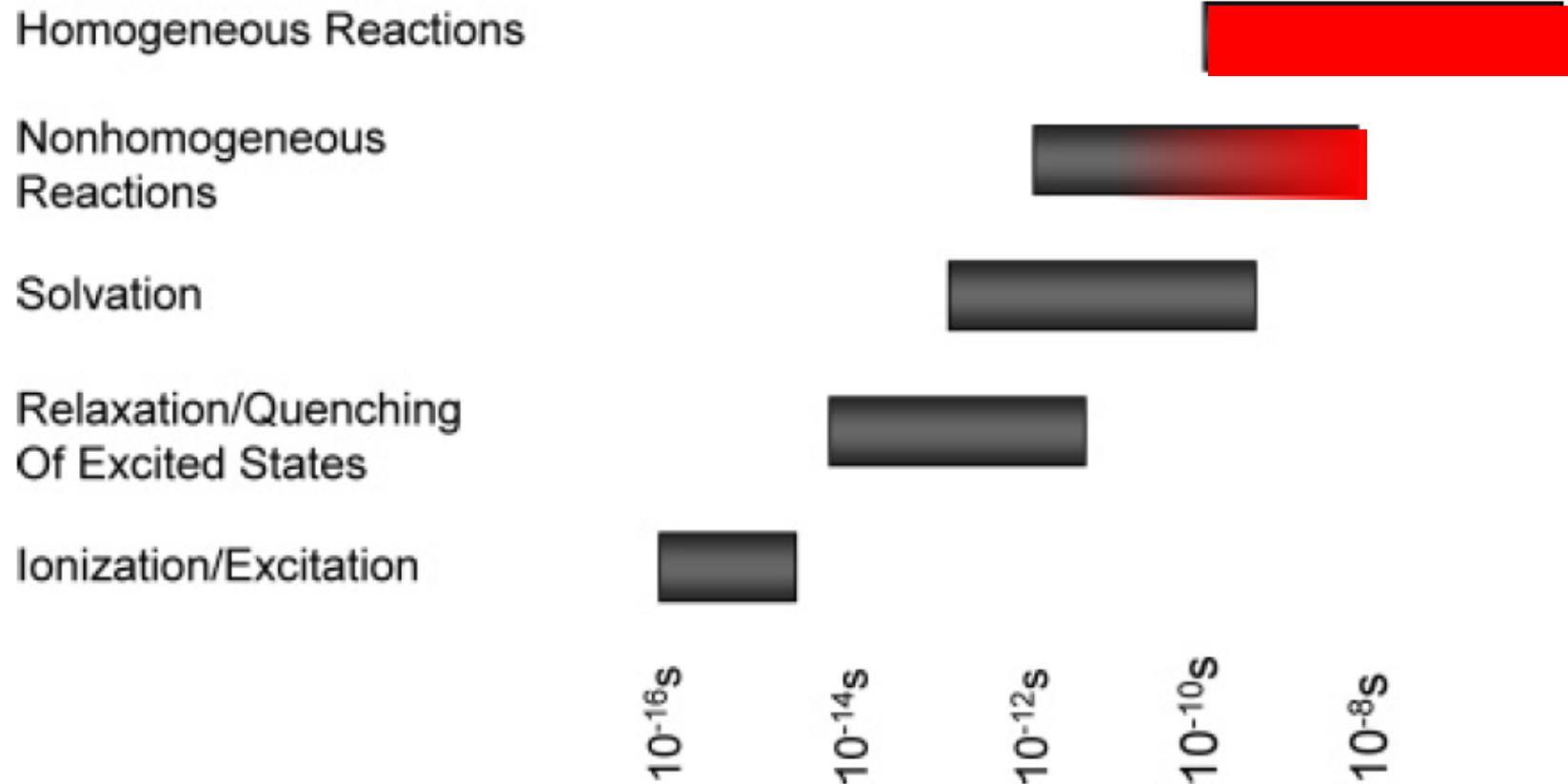
$t = 0 \text{ s}$



$$t \approx 10^{-14} \text{ s}$$



Timescales of radiation damage



Garret et. al. (2005) *Chem. Rev.* **105**, 355-389

Timescales of radiation damage

Homogeneous Reactions



Nonhomogeneous
Reactions



Solvation



LCLS



Relaxation/Quenching
Of Excited States



Ionization/Excitation



ALS
bunch

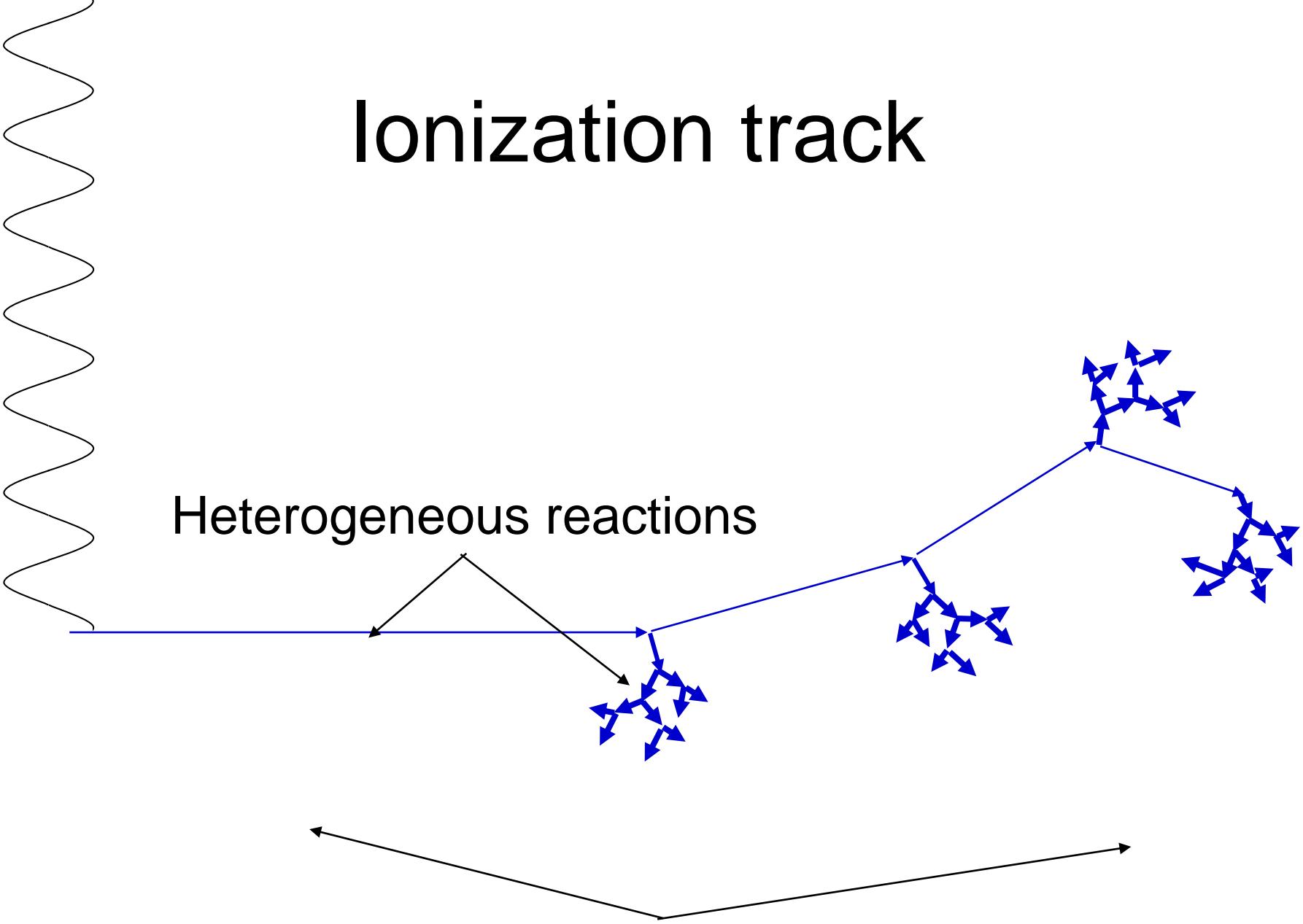
$10^{-16}s$ $10^{-14}s$ $10^{-12}s$ $10^{-10}s$ $10^{-8}s$

Garret et. al. (2005) *Chem. Rev.* **105**, 355-389

Ionization track

Heterogeneous reactions

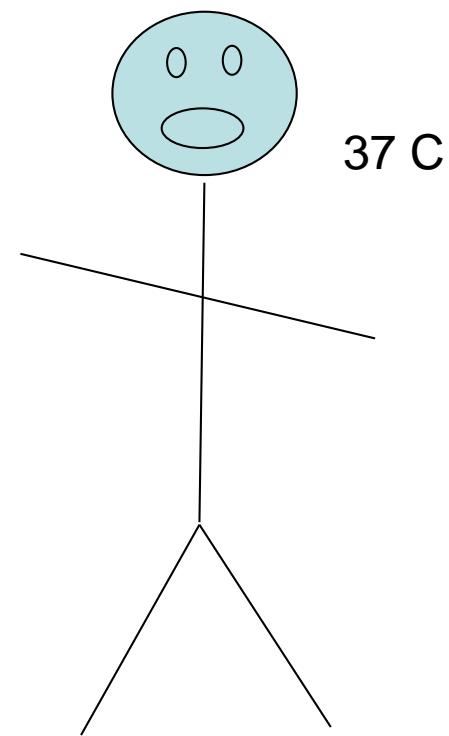
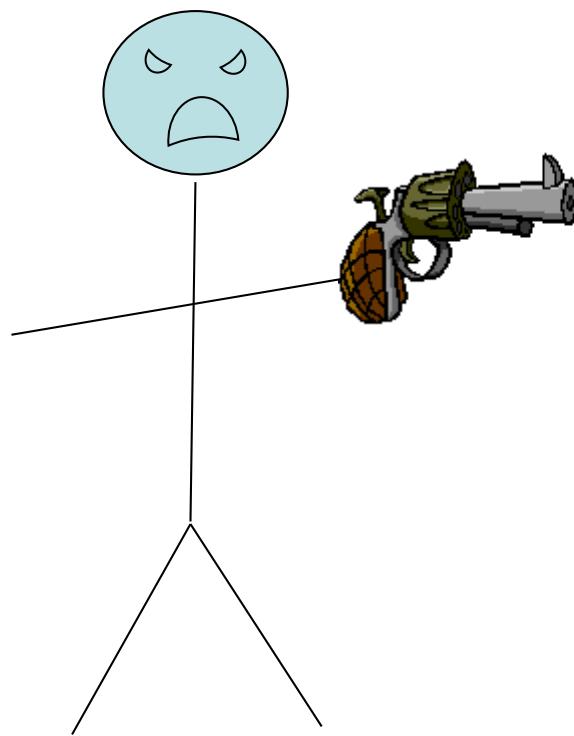
Homogeneous reactions



Rough values of energy quanta

1 MeV	100 GJ/mol	Medical radiation therapy
100 keV	10 GJ/mol	Medical imaging
10 keV	1 GJ/mol	X-ray crystallography
1 keV	100 MJ/mol	S and P K-edges
100 eV	10 MJ/mol	“water window”
10 eV	1 MJ/mol	C≡C bond
1 eV	100 kJ/mol	C-C bond, visible light
100 meV	10 kJ/mol	hydrogen bond
10 meV	1 kJ/mol	heat (~300 K)

Energy Transfer Analogy



Damage is done
by dose (MGy)
proportional to photons/area

not time
not heat

- Sliz P, Harrison SC & Rosenbaum G (2003). *Structure* **11**, 13-19.
Garman EF & McSweeney SM (2006). *J. Sync. Rad.* **14**, 1-3.
Owen RL, Rudino-Pinera E & Garman EF (2006). *PNAS* **103**, 4912-4917.
Leiros et al. (2006). *Acta Cryst. D* **62**, 125-132.
Holton JM (2007). *J. Synch Rad.* **14**, 51-72.

what the is a MGy?

$$1 \text{ MGy} = 10^6 \text{ J/kg}$$

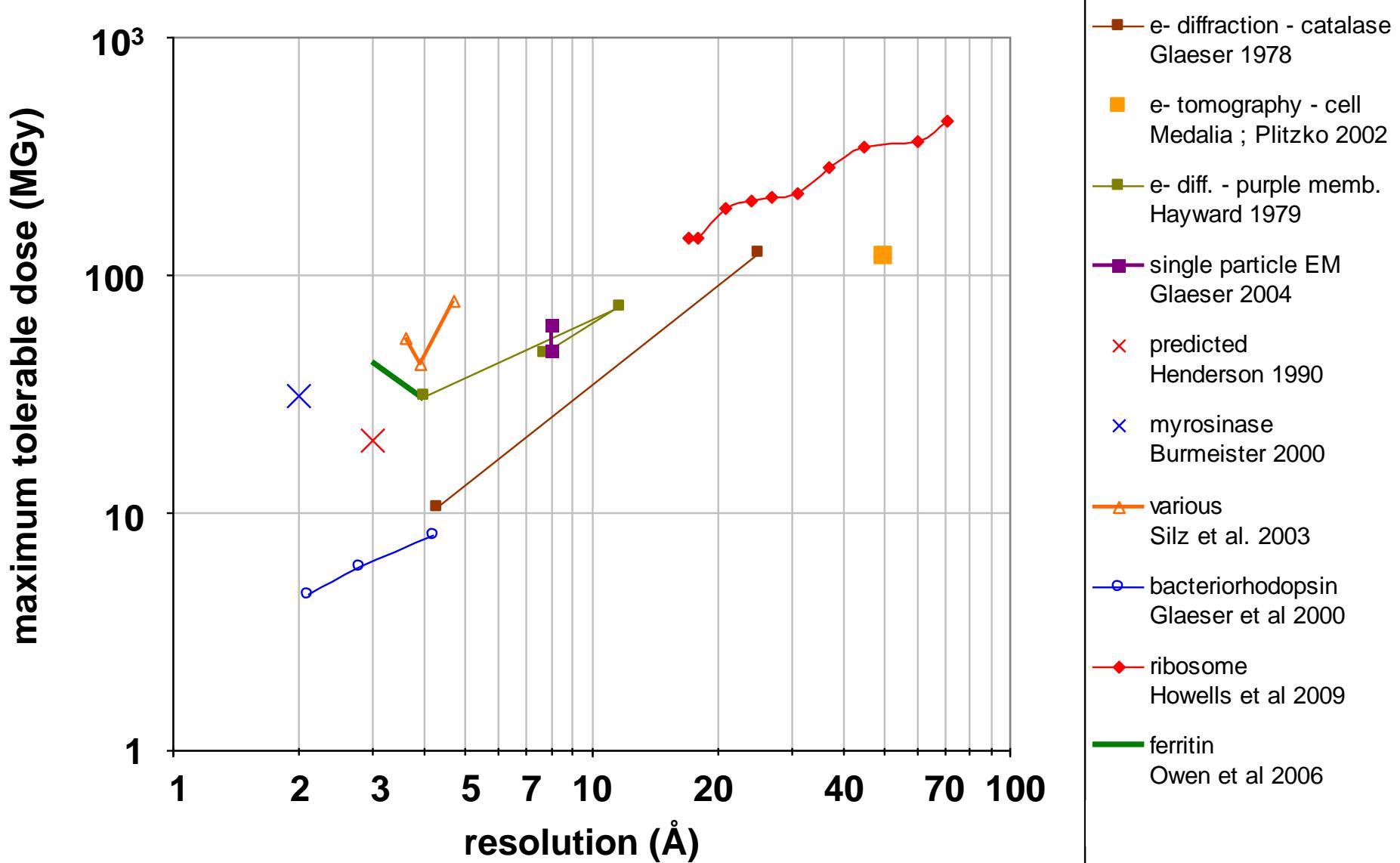
<http://bl831.als.lbl.gov/>

damage_rates.pdf

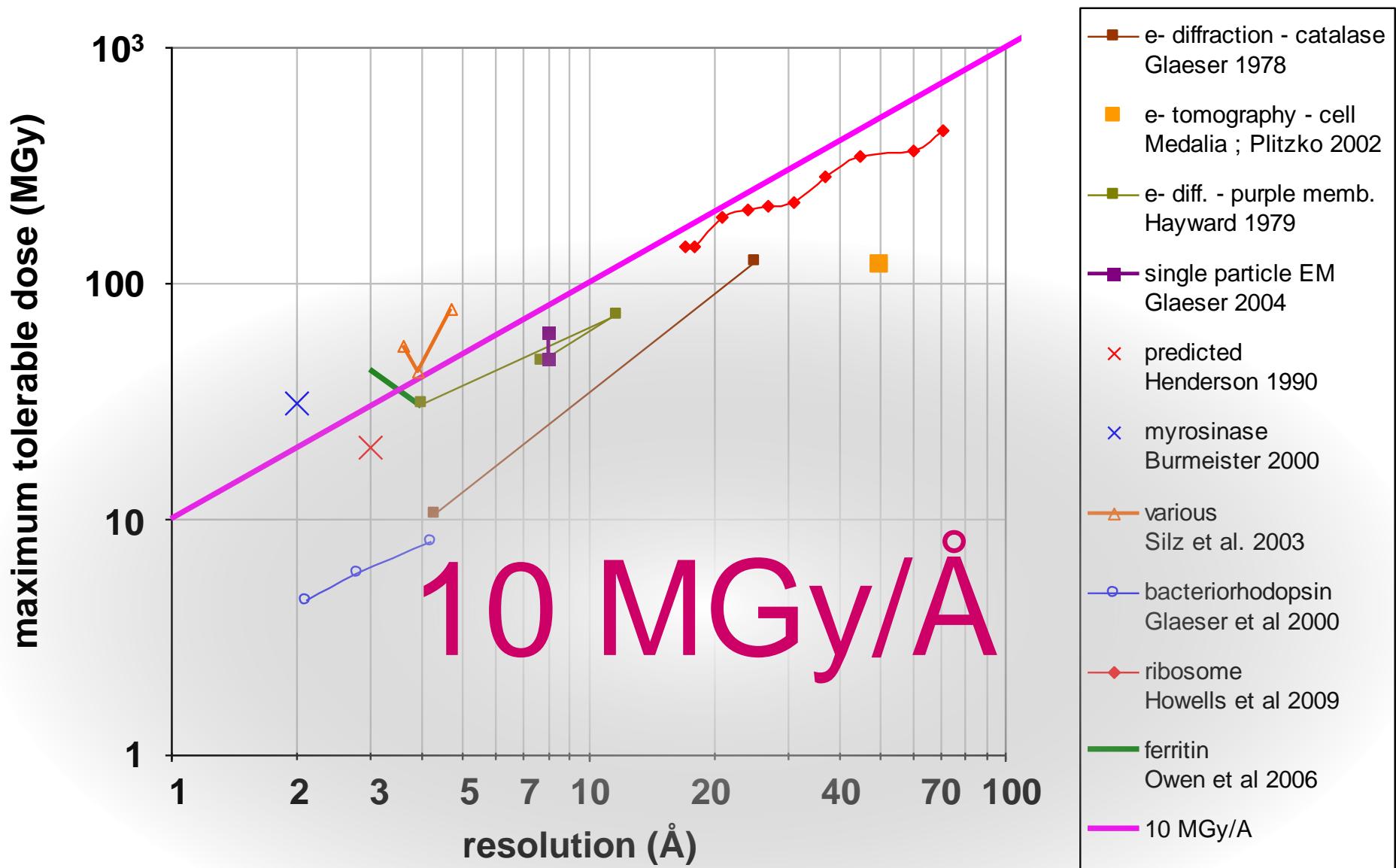
How long will my crystal last?

synch	line	type	flux ph/s	beamsize μm	flux density ph/μm ² /s	dose rate	max xtal lifetime	min site lifetime
ALS	4.2.2	MAD	2.2e11	75x80	3.7e+07	27.3 kGy/s	18 m	73 s
ALS	5.0.1	mono	1.6e11	100	2.0e+07	10.4 kGy/s	48 m	3.2 m
ALS	5.0.2	MAD	8e11	100	1.0e+08	51.8 kGy/s	9.6 m	39 s
ALS	5.0.3	mono	1.7e11	100	2.2e+07	11 kGy/s	45 m	3 m
ALS	8.2.1	MAD	1.8e11	100	2.3e+07	11.7 kGy/s	43 m	2.9 m
ALS	8.2.2	MAD	2.3e11	100	2.9e+07	14.9 kGy/s	34 m	2.2 m
ALS	8.3.1	MAD	9e11	70	2.3e+08	119 kGy/s	4.2 m	17 s
ALS	8.3.1	typical	6e11	70	1.6e+08	115 kGy/s	4.3 m	17 s
ALS	12.3.1	MAD	1.8e11	100	2.3e+07	11.7 kGy/s	43 m	2.9 m
ALS	12.3.1	ML	4.0e12	100	5.1e+08	513 kGy/s	58 s	3.9 s
APS	8-BM	MAD	1e11	200	2.5e+06	1.27 kGy/s	6.6 h	26 m
APS	14-BM-C	mono	5.8e10	200	1.4e+06	738 Gy/s	11 h	45 m
APS	14-BM-D	MAD	3.3e9	200	8.2e+04	42 Gy/s	8.3 d	13 h
APS	14-ID-B	MAD	6.0e10	200	1.5e+06	763 Gy/s	11 h	44 m
APS	17-BM	MAD	1.1e11	200	2.8e+06	1.4 kGy/s	6 h	24 m
APS	17-ID	MAD	2.3e11	200	5.8e+06	2.93 kGy/s	2.8 h	11 m
APS	19-BM	MAD	2.0e11	70x60	4.8e+07	24.2 kGy/s	21 m	83 s
APS	19-ID	MAD	1.3e13	80x40	4.1e+09	2.07 MGy/s	15 s	0.97 s
APS	19-ID	typical	5.5e11	100x100	5.5e+07	28 kGy/s	18 m	71 s
APS	22-BM	MAD	7e12	80x40	2.2e+09	1.23 MGy/s	24 s	1.6 s
APS	22-ID	MAD	7e12	80x40	2.2e+09	1.23 MGy/s	24 s	1.6 s
APS	22-ID	typical	1.5e12	80	2.3e+08	119 kGy/s	4.2 m	17 s
APS	23-ID-B	MAD	1e13	75x25	5.3e+09	3.01 MGy/s	10 s	0.66 s
APS	23-ID	typical	1.5e12	80	2.3e+08	119 kGy/s	4.2 m	17 s
APS	24-ID-C	MAD	1.3e13	20x60	1.1e+10	5.23 MGy/s	5.7 s	0.38 s
APS	24-ID-E	MAD	0.5e13	20x100	2.5e+09	1.19 MGy/s	25 s	1.7 s
APS	31-ID	MAD	2e12	70	4.1e+08	194 kGy/s	2.6 m	10 s

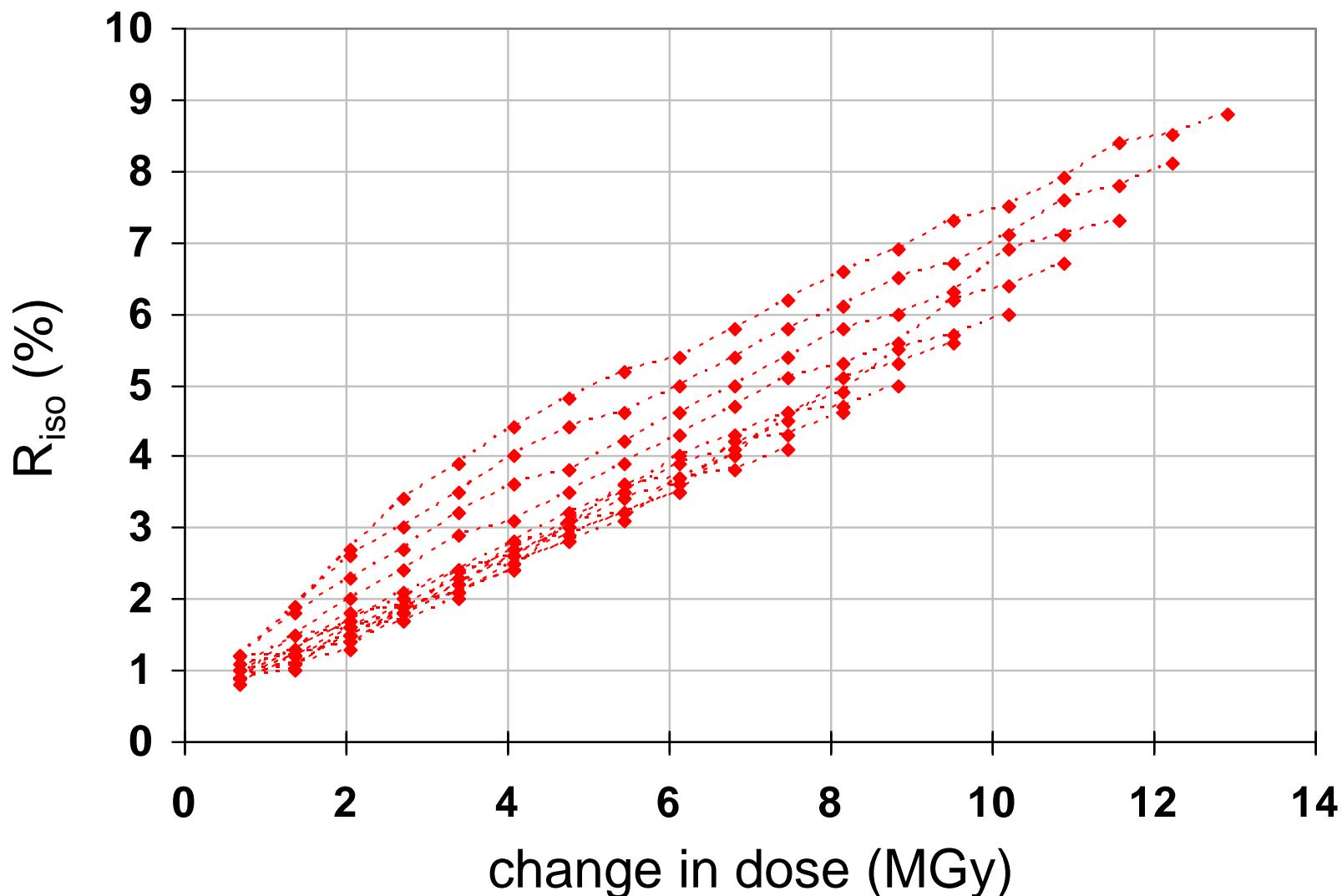
resolution dependence of global damage



resolution dependence of global damage

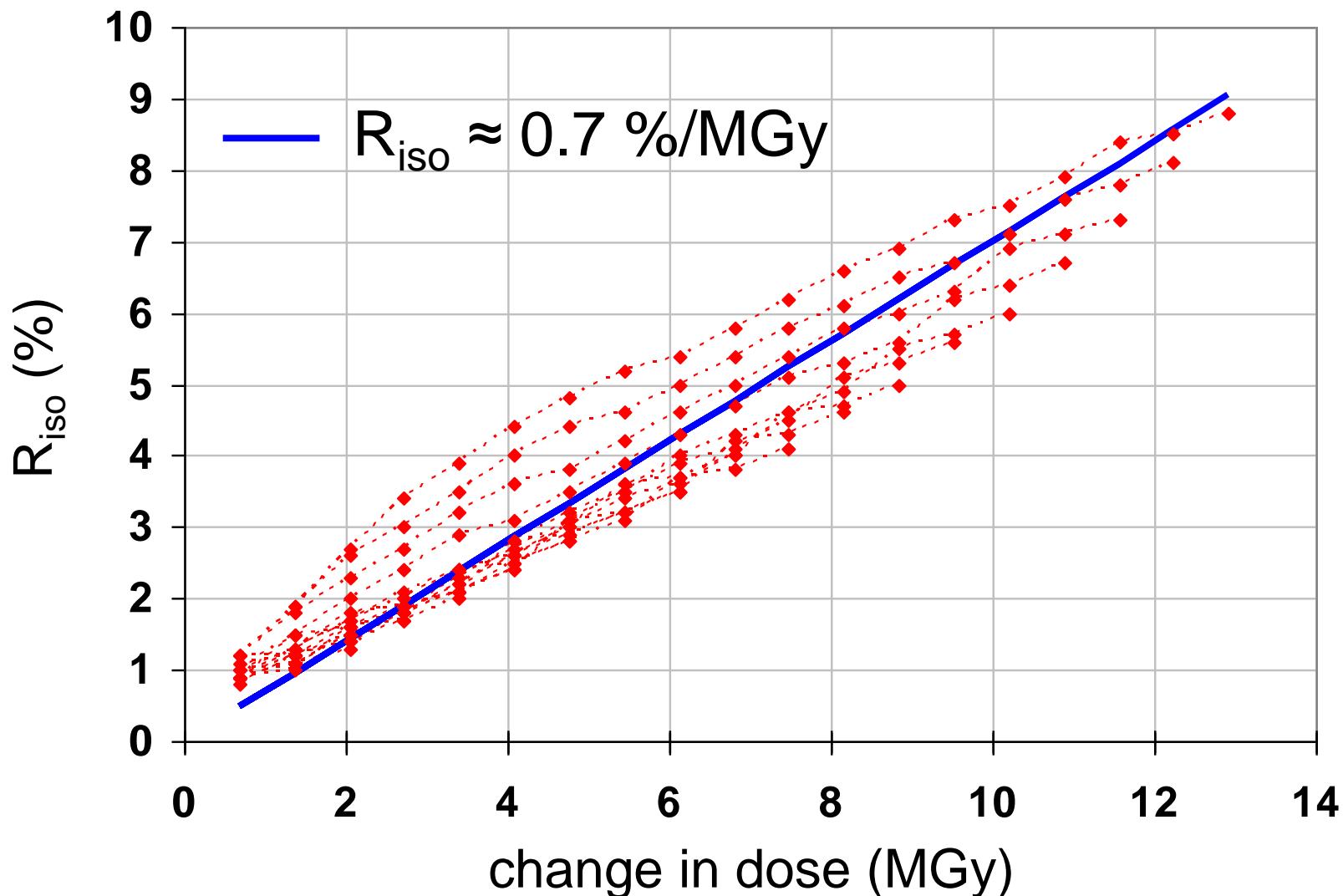


R_{iso} vs dose



data taken from Banumathi, et al. (2004) *Acta Cryst. D* **60**, 1085-1093.

R_{ISO} vs dose



Dose-doubling concentration

molar, at the Se edge based on RADDOSE

Na	44	As	0.44
Mg	27	Se	0.42
P	9	Br	2.9
S	7	I	0.47
Cl	5	Gd	0.21
K	3.3	Ta	0.13
Ca	2.7	Pt	0.17
Fe	1.0	Au	0.16
Cu	0.67	Hg	0.16
Zn	0.58	U	0.23

Dose-doubling concentration

molar, **at the Se edge** based on RADDOSE

Na	44	As	0.44
Mg	27	Se	0.42
P	9	Br	2.9
S	7	I	0.47
Cl	5	Gd	0.21
K	3.3	Ta	0.13
Ca	2.7	Pt	0.17
Fe	1.0	Au	0.16
Cu	0.67	Hg	0.16
Zn	0.58	U	0.23

Specific Damage World Records

MGy	reaction	reference
~45	global damage	Owen <i>et al.</i> (2006)
5	Se-Met	Holton (2007)
4	Hg-S	Ramagopal <i>et al.</i> (2004)
3	S-S	Murray <i>et al.</i> (2002)
1	Br-RNA	Olieric <i>et al.</i> (2007)
?	Cl-C	???
0.5	Mn in PS II	Yano <i>et al.</i> (2005)
0.02	Fe in myoglobin	Denisov <i>et al.</i> (2007)

expected crystal lifetime calculator

source =

full flux =

7.0e+12 photons/s

attenuation =

transmittance =

beam size_{horiz} =

40.0 microns

beam size_{vert} =

80.0 microns

wavelength =

1 Ang

k_{dose} =2000 photons/micron²/Gy

dose rate =

1.1e+6 Gy/s

experiment goal =

resolution =

3 Ang

dose limit =

30 MGy

exposure time =

1 seconds/image

xtal size_{horiz} =

50 microns

xtal size_{vert} =

50 microns

translation during dataset =

0 microns

rotisserie factor

1

 disable warnings

max images =

28 at damage limit

inverse beam =

number of wavelengths =

images/wedge =

required number of crystals calculator - Mozilla Firefox

File Edit View History Bookmarks Tools Help

<http://bl831.als.lbl.gov/xtalsize.html>

wikipedia (en)

Required crystal number or size calculator

$$n_{\text{xtals}} = \langle I_{\text{DL}} \rangle / 20 * f_{\text{NH}} * \text{MW} * V_M^2 / \exp(-0.5 * B/\text{reso}^2) / \text{xtalsize}^3 / (\text{reso}^3 - 1.53)$$

Enter values:

experiment goal =	subtle differences (MAD/SAD)		
number of sites =	1	in asymmetric unit	
fpp =	4	electrons	Bijvoet ratio = 1.75 %
molecular weight =	30	kDa in asymmetric unit	
resolution =	3.4	Ang	signal to noise = 81 at this resolution
reso on snapshot =	2.4	Ang	→ Wilson B = 35 Ang ²
background level =	100	ADU/pixel	multiplicity = 7.3
spot size =	5	pixels	
detector type =	ADSC Q210/315r (hwbin)		
solvent content =	50	%	
xtal size _{beam} =	20	microns	beam size _{vert} = 100 microns
xtal size _{vert} =	20	microns	beam size _{spindle} = 100 microns
xtal size _{spindle} =	20	microns	
<input type="button" value="Calculate n_xtals"/> ↓	<input type="button" value="Calculate size"/> ↑		
n _{xtals} =	1.4	xtals you will need to merge	← <I _{DL} > 11000 photons/hkl

Done

The number of photons you get
before a given crystal dies

is independent
of data collection time

$$1 \text{ um}^3 = 10^6 \text{ photons}$$

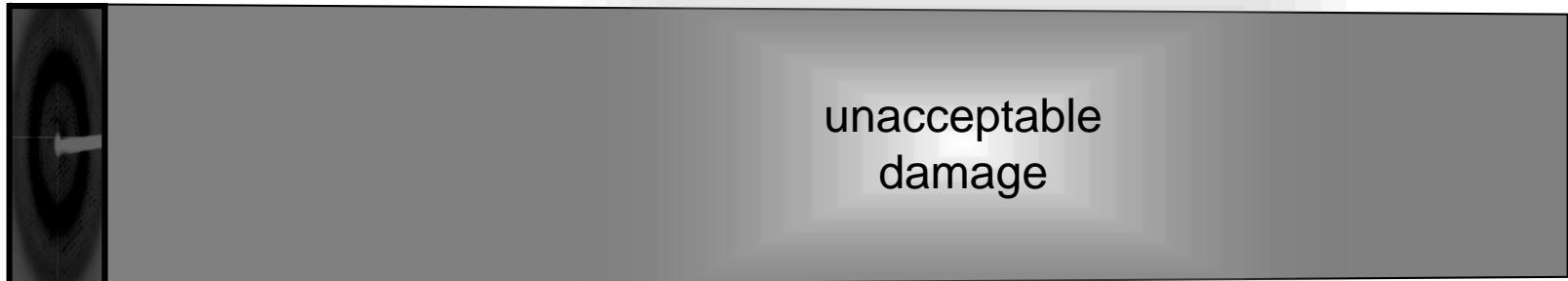
Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

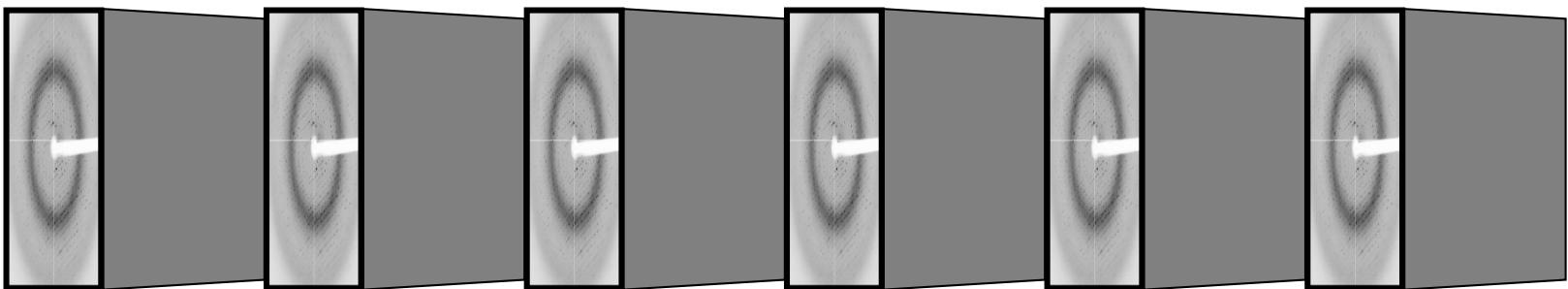
Dose slicing

N
photons



unacceptable
damage

N
photons



unacceptable
read noise

N
photons



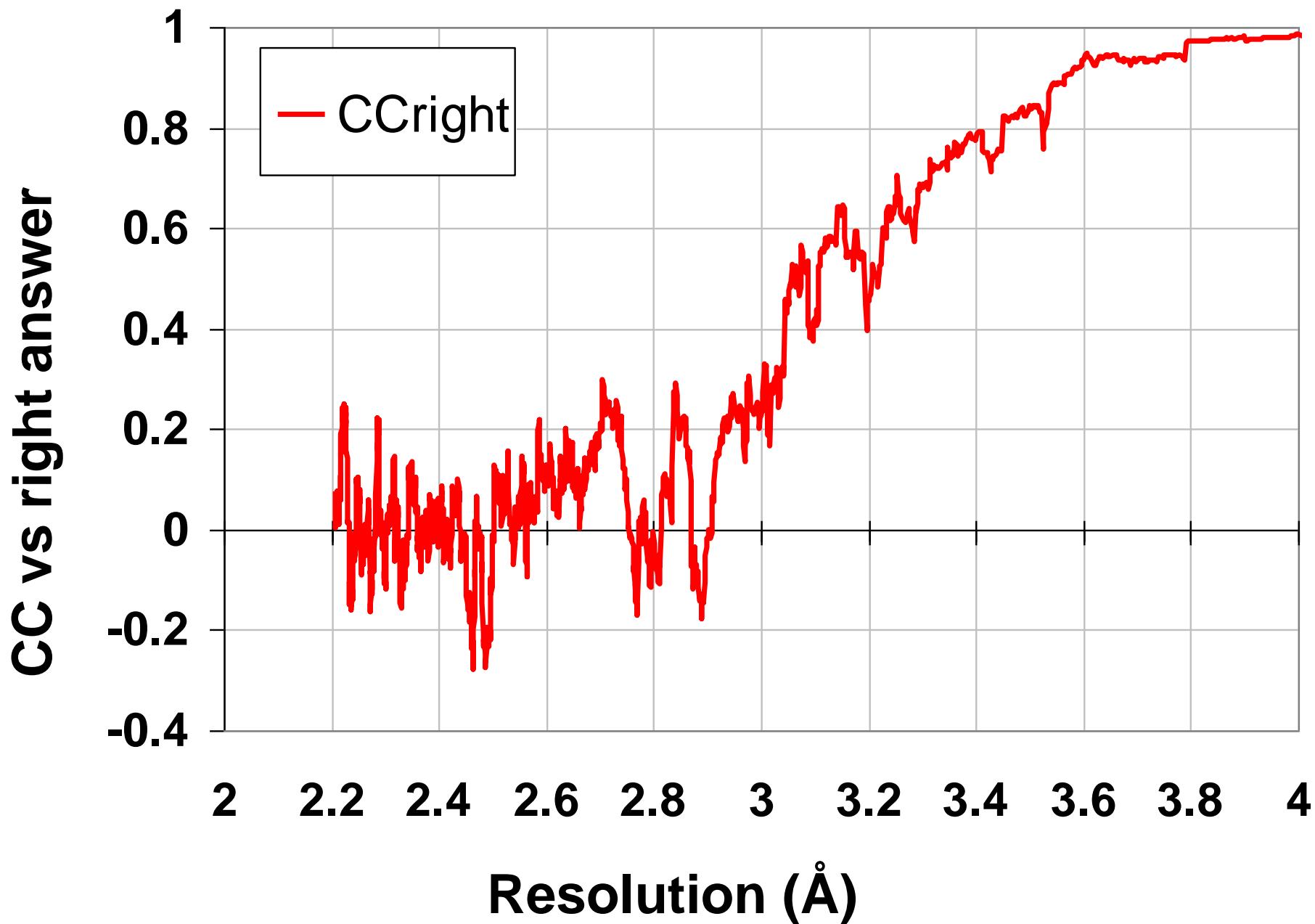
What if...

- You only have a few small crystals?

Should you:

- a) Collect 360° from each?
- b) Collect 10° from each at 36x exposure?
- c) Glue 36 xtals together, then collect 360° ?
- d) Glue together and do 12960° ?

“true” resolution limit



“true” resolution vs strategy

processing procedure	Dose slice collection scenario			
	fine	coarse	glue	gluefine
xds/xscale	3.03874	3.02970	3.0270	3.02948
xds/aimless	3.13351	3.03531	3.02638	3.05292
xds/noscale	3.01449	3.03048	3.02576	3.31013

301,640,334 photons

Self-calibrated damage limit

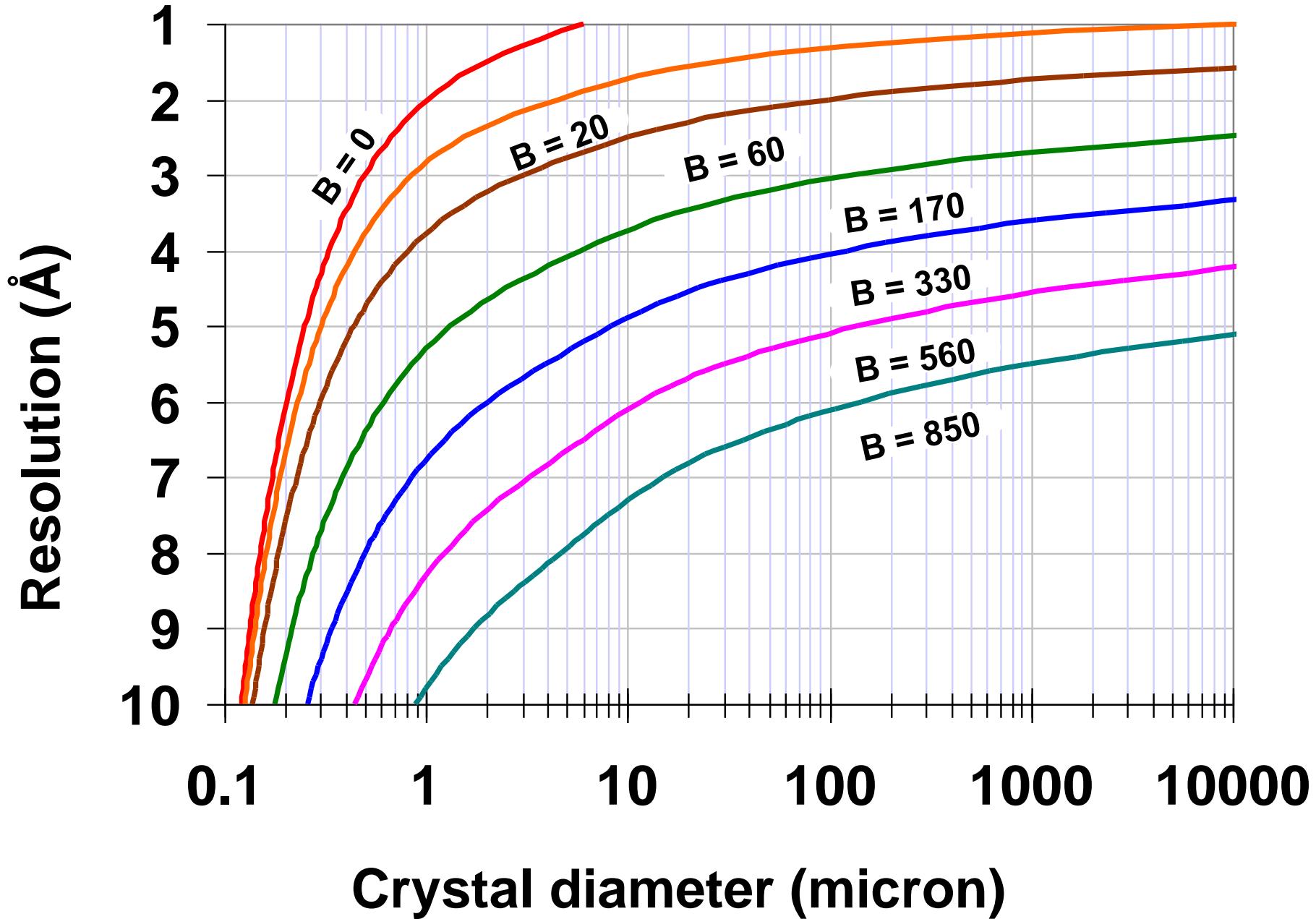
$$\langle I \rangle_{DL} = \frac{2\pi}{9} \frac{10^5 r_e^2}{hc} \frac{f_{decayed} \rho R^4 \lambda^4}{f_{NH} n_{ASU} M_r V_M^2} \frac{0.5 \lambda H}{\ln(2) \sin \theta} \frac{T_{sphere}(2\theta, \mu, R)}{(1 - T_{sphere}(0, \mu_{en}, R))} \frac{(3 + \cos 4\theta) \langle f_a^2 \rangle}{\sin \theta} \frac{\langle M_a \rangle}{\exp \left(-2B \left(\frac{\sin \theta}{\lambda} \right)^2 \right)}$$

Where:

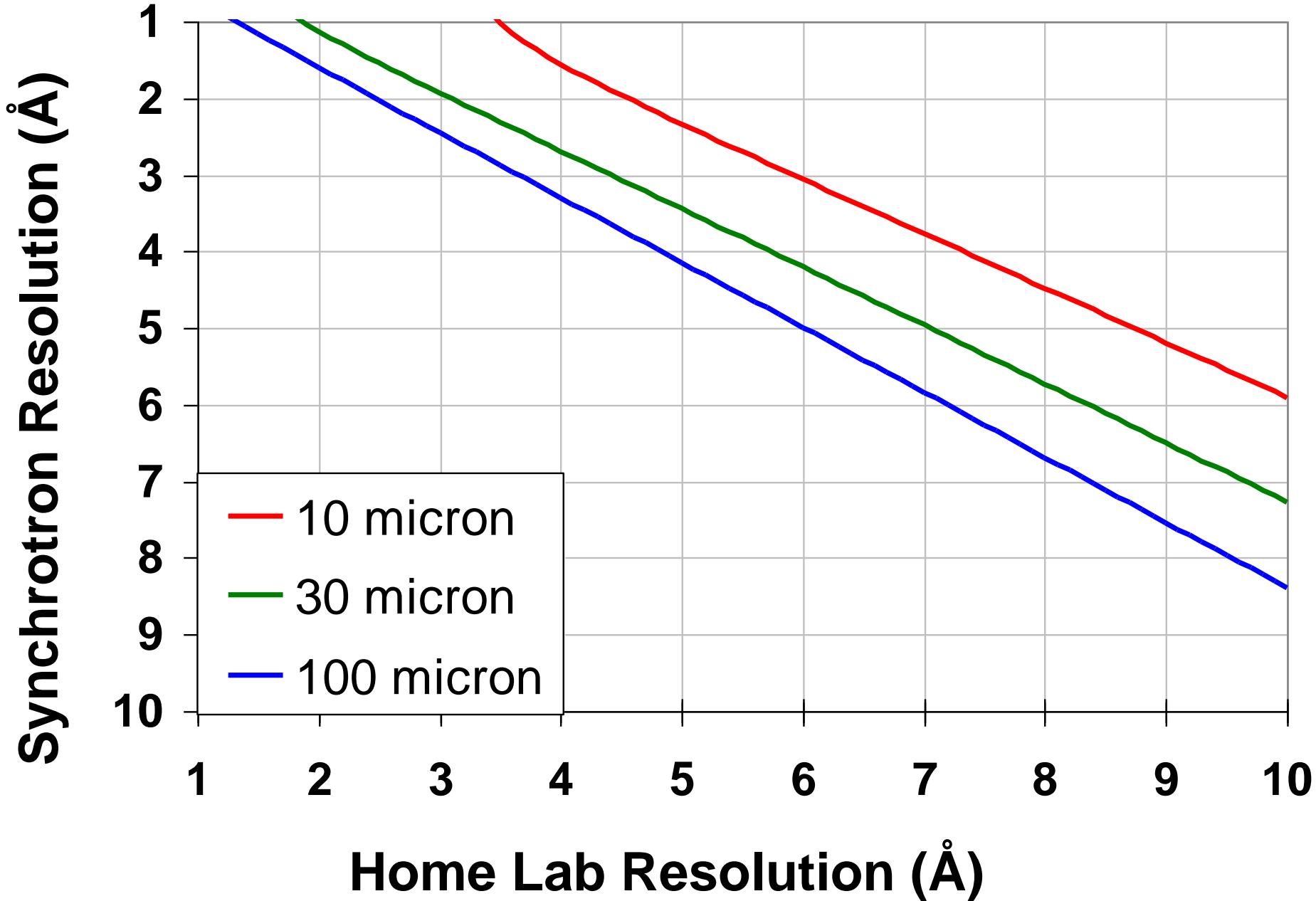
$\langle I \rangle_{DL}$	- average damage-limited intensity (photons/hkl) at a given resolution
10^5	- converting R from μm to m , r_e from m to \AA , ρ from g/cm^3 to kg/m^3 and MGy to Gy
r_e	- classical electron radius ($2.818 \times 10^{-15} \text{ m/electron}$)
h	- Planck's constant ($6.626 \times 10^{-34} \text{ J}\cdot\text{s}$)
c	- speed of light (299792458 m/s)
$f_{decayed}$	- fractional progress toward completely faded spots at end of data set
ρ	- density of crystal ($\sim 1.2 \text{ g/cm}^3$)
R	- radius of the spherical crystal (μm)
λ	- X-ray wavelength (\AA)
f_{NH}	- the Nave & Hill (2005) dose capture fraction (1 for large crystals)
n_{ASU}	- number of proteins in the asymmetric unit
M_r	- molecular weight of the protein (Daltons or g/mol)
V_M	- Matthews's coefficient ($\sim 2.4 \text{ \AA}^3/\text{Dalton}$)
H	- Howells's criterion ($10 \text{ MGy}/\text{\AA}$)
θ	- Bragg angle
$\langle f_a^2 \rangle$	- number-averaged squared structure factor per protein atom (electron ²)
$\langle M_a \rangle$	- number-averaged atomic weight of a protein atom (~ 7.1 Daltons)
B	- average (Wilson) temperature factor (\AA^2)
μ	- attenuation coefficient of sphere material (m^{-1})
μ_{en}	- mass energy-absorption coefficient of sphere material (m^{-1})

No flux
No symmetry

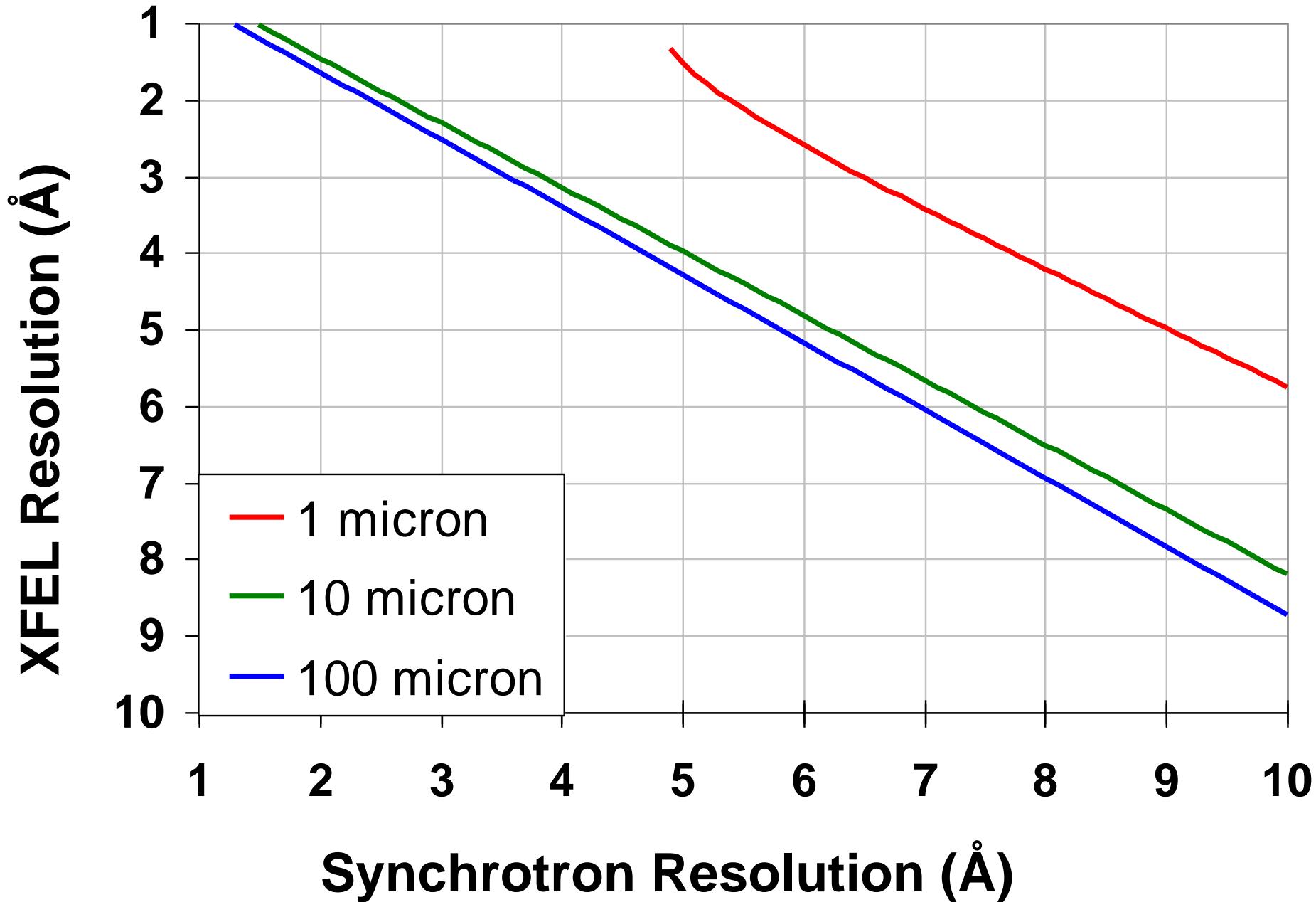
Bigger is better, but not by much



Predicting resolution limits

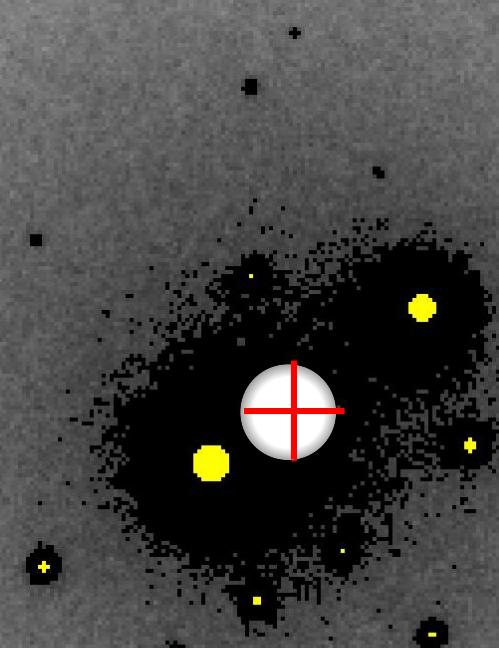


Predicting resolution limits



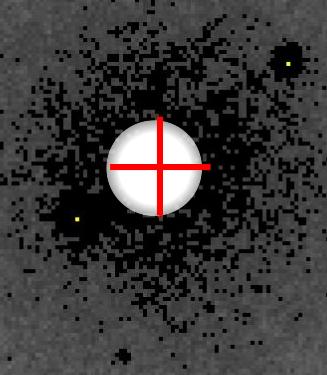
B factor from image analysis

$B = 500$



B factor from image analysis

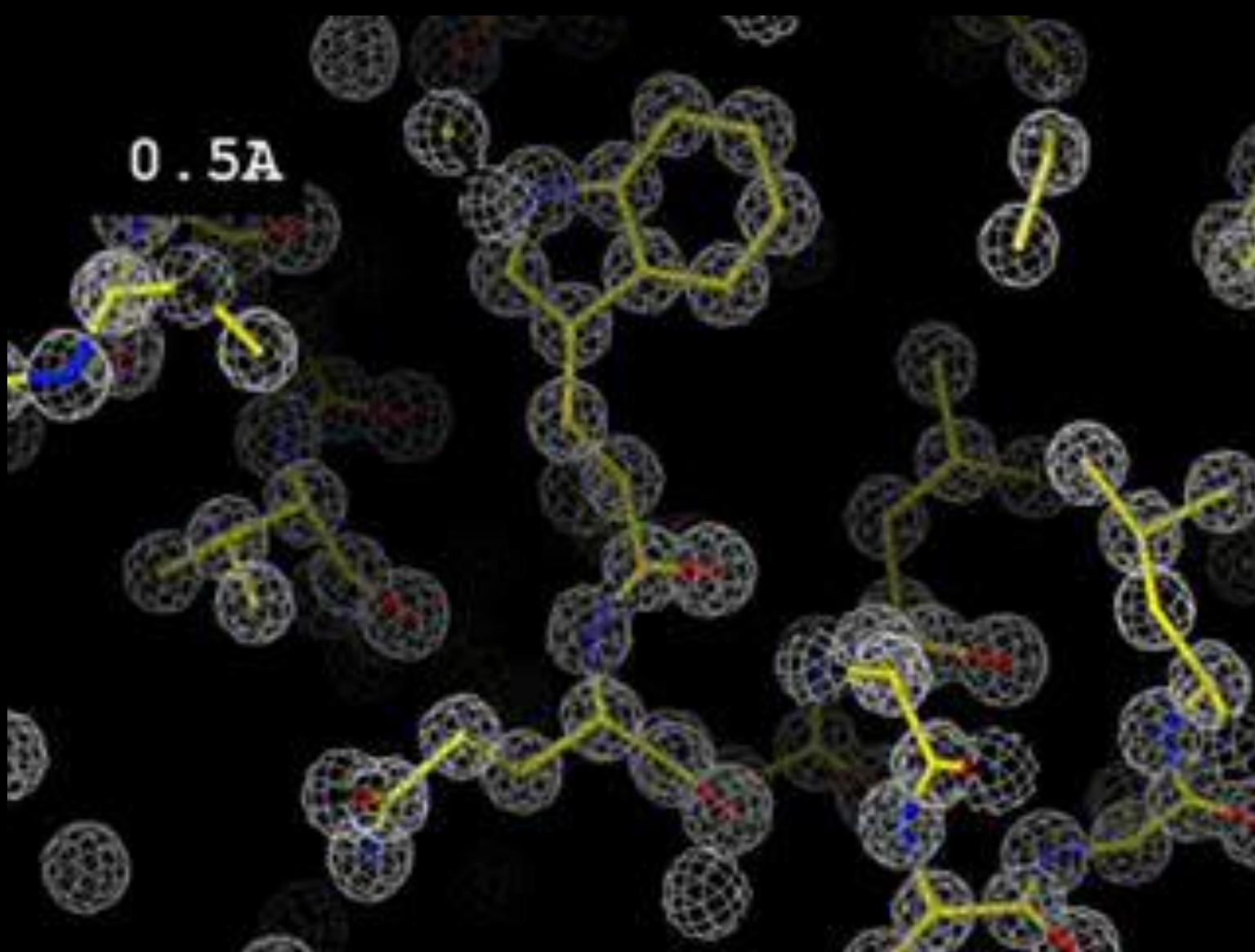
B = 20



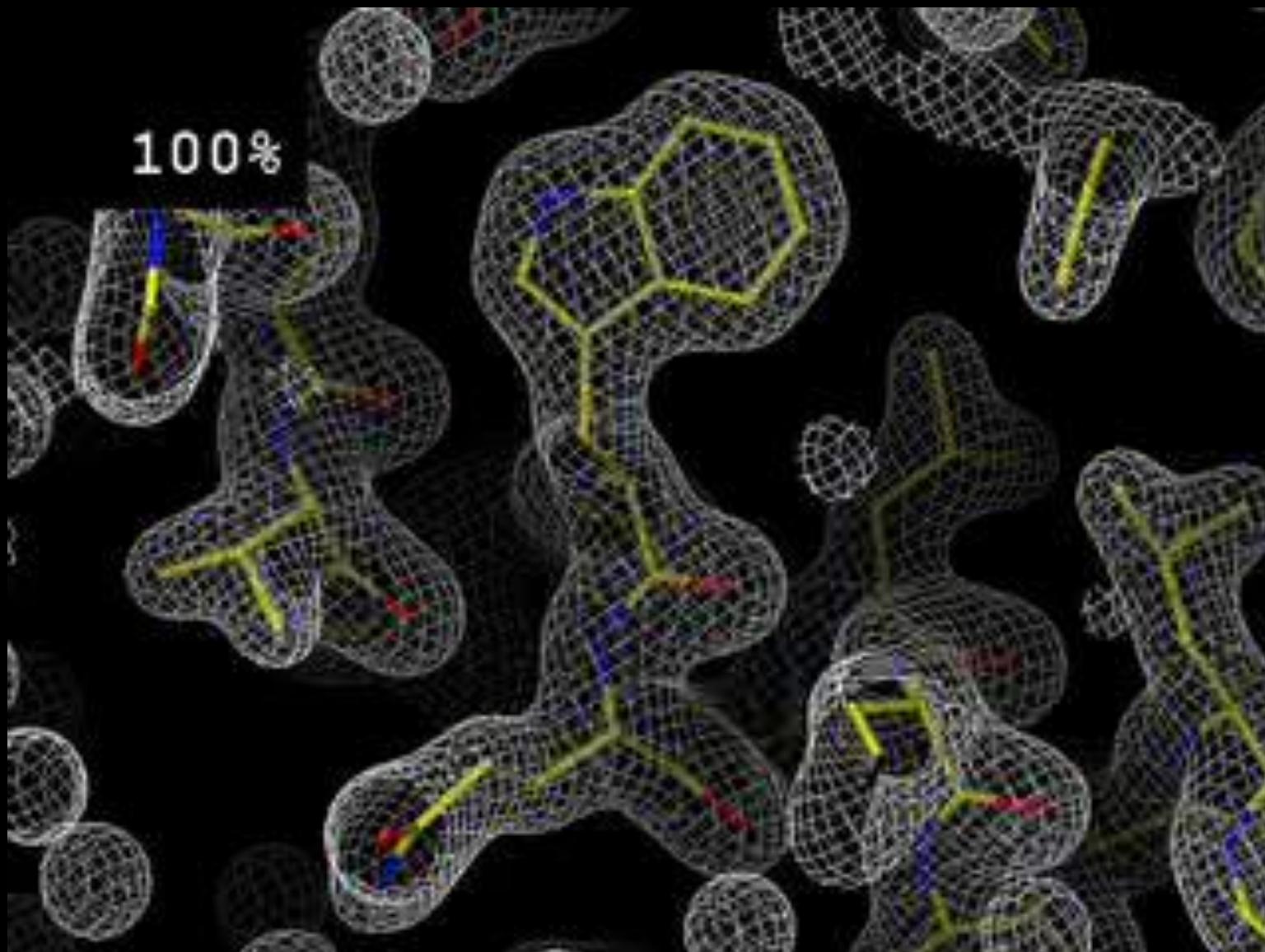
At the beamline...

- Resolution
 - problem: background
 - solution: use as few pixels as possible
- Phases
 - problem: fractional errors
 - solution: use as many pixels as possible

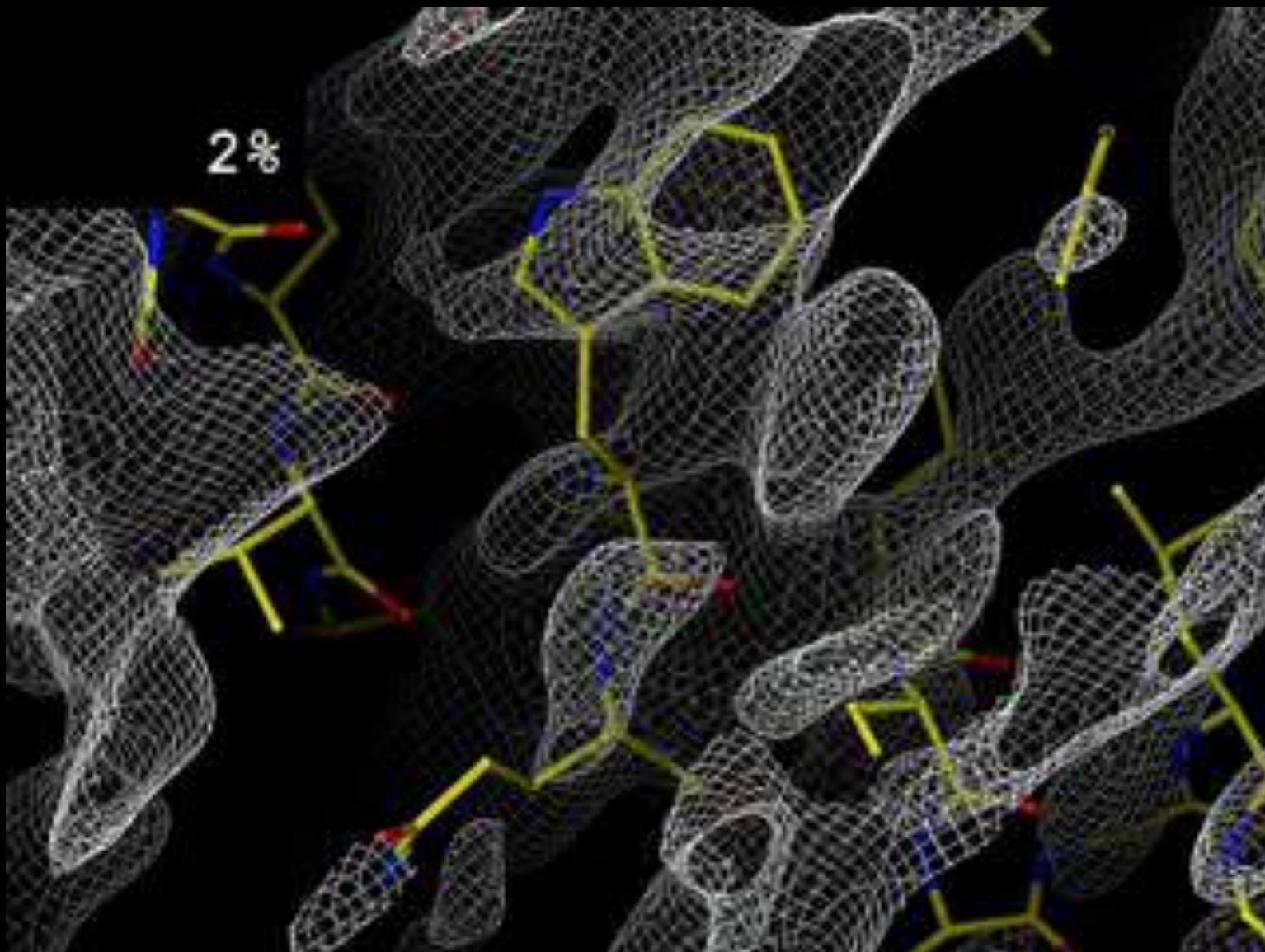
Resolution



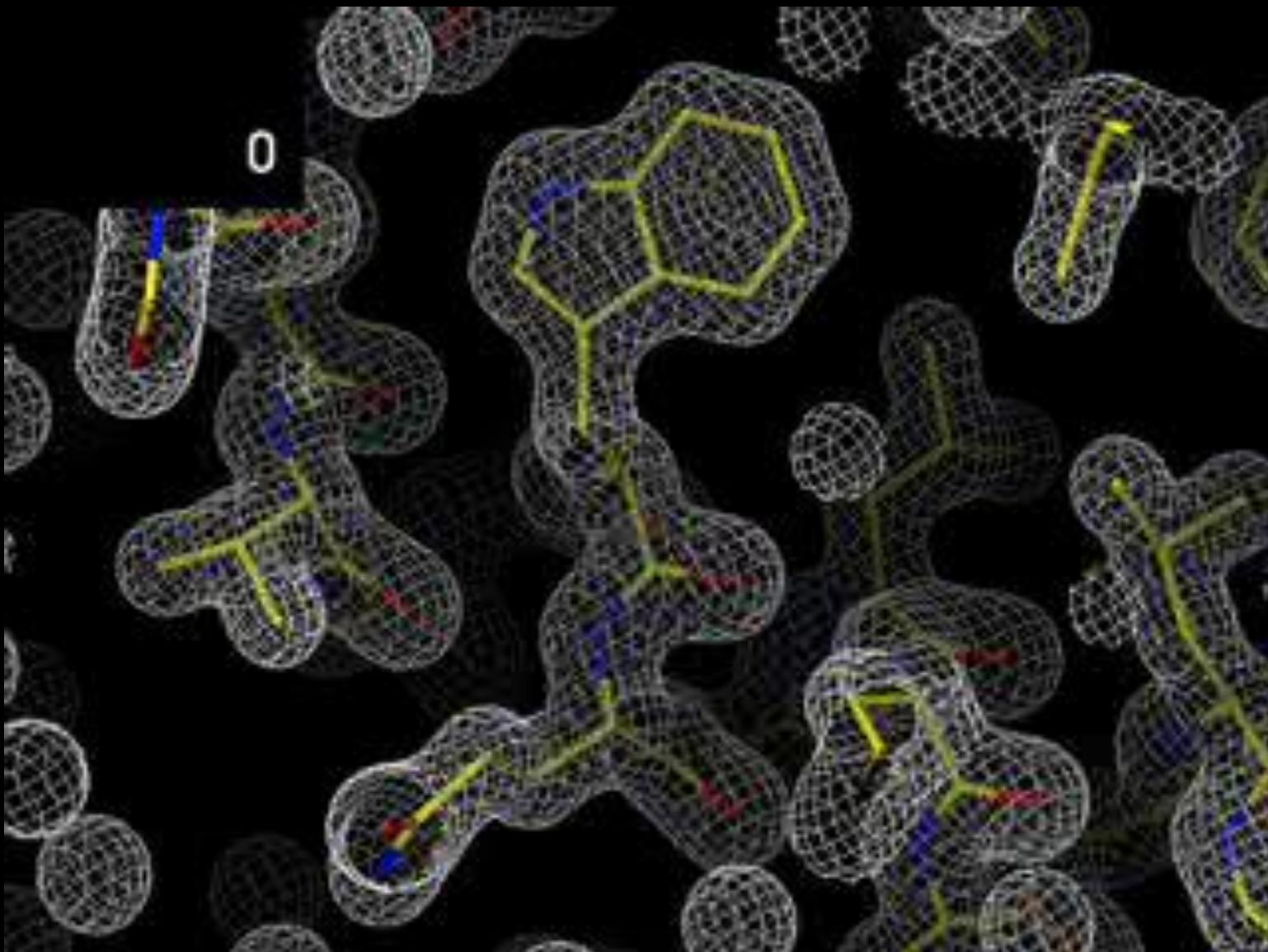
Completeness: random deletion



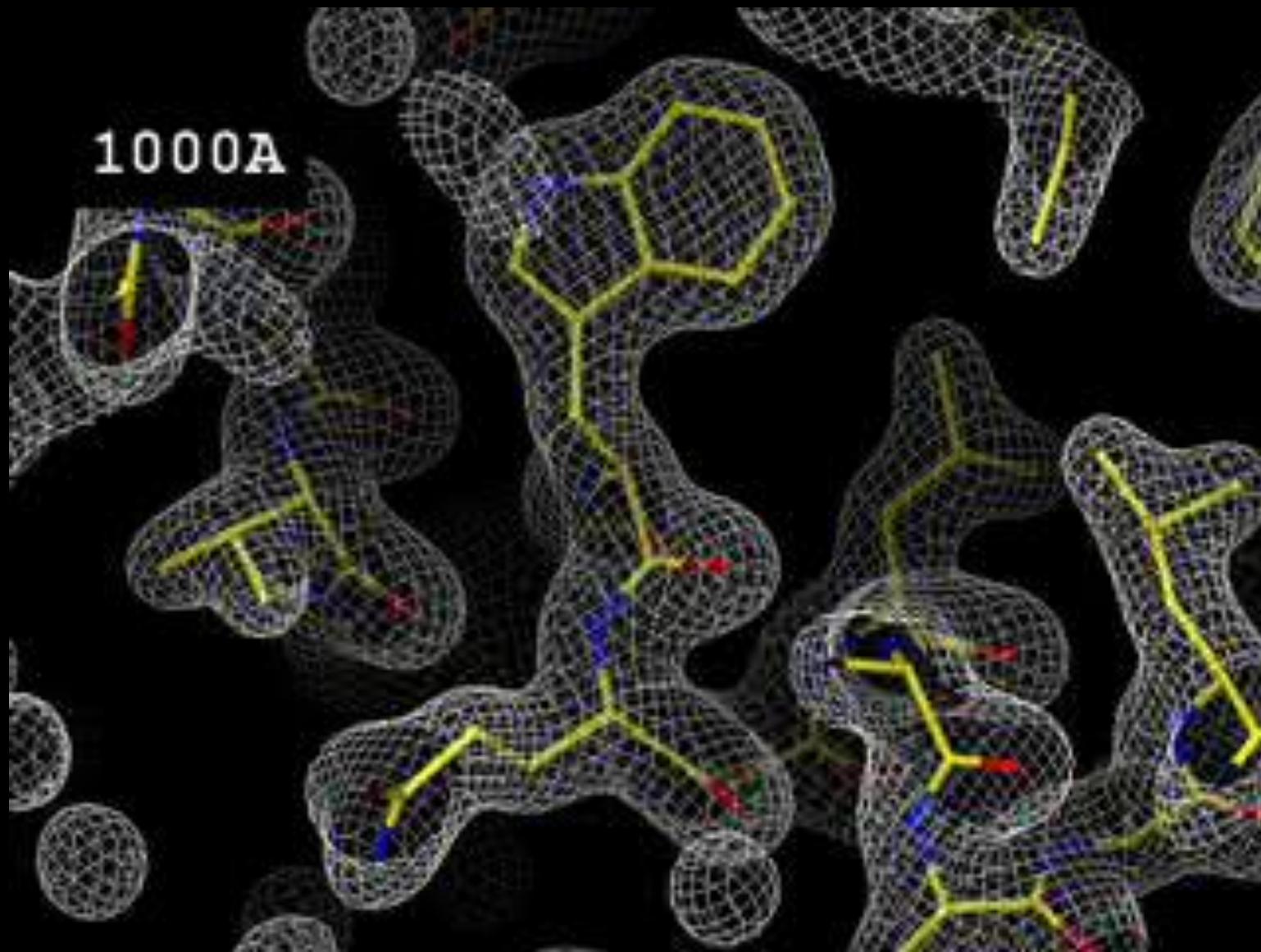
Completeness: missing wedge



Overloads



Resolution: low-angle cutoff



The truth about x-ray beams

Term	units	significance
Flux	photons/s	duration of experiment
Beam Size	μm	match to crystal
Divergence	mrad	spot size vs distance
Wavelength	\AA	resolution and absorption
Dispersion	$\Delta\lambda/\lambda$	spot size
Flux density	ph/s/area	scattering/damage rate
Fluence	ph/area	scattering/damage

The truth about x-ray beams

Term	units	significance
Flux	photons/s	duration of experiment
Beam Size	μm	match to crystal
Divergence	mrad	spot size vs distance
Wavelength	\AA	resolution and absorption
Dispersion	$\Delta\lambda/\lambda$	spot size
Flux density	ph/s/area	scattering/damage rate
Fluence	ph/area	scattering/damage

The truth about x-ray beams

quantity	units	home source	MX2
flux	Photons/second	2.5×10^9	2×10^{12}
exposure	seconds	400	1

Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, **flux**, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

The truth about x-ray beams

quantity	units	home source	MX2
flux	Photons/second	2.5×10^9	2×10^{12}
exposure	seconds	400	1
Dispersion	wavelength range / wavelength	0.2% ($K\alpha_1 - K\alpha_2$)	0.014% (Si111)
Divergence	milliRadian	4.8	1.8 (h) 0.18 (v)
Beam size	microns	100	24 (h) 12 (v)
Spectral brightness	Photons/s/mm ² /mR ² /0.1%BW	5.4×10^9	1.5×10^{17}

The truth about x-ray beams

quantity	units	home source	MX2
flux	Photons/second	72×10^6	2×10^{12}
exposure	time	4 hours	1 second
Dispersion	wavelength range / wavelength	0.2% ($K\alpha_1 - K\alpha_2$)	0.014% (Si111)
Divergence	milliRadian	4.8	1.8 (h) 0.18 (v)
Beam size	microns	17	24 (h) 12 (v)
Spectral brightness	Photons/s/mm ² /mR ² /0.1%BW	5.4×10^9	1.5×10^{17}

The truth about x-ray beams

quantity	units	home source	MX2
flux	Photons/second	1×10^6	2×10^{12}
exposure	time	10 days	1 second
Dispersion	wavelength range / wavelength	0.2% ($K\alpha_1 - K\alpha_2$)	0.014% (Si111)
Divergence	milliRadian	0.6	1.8 (h) 0.18 (v)
Beam size	microns	17	24 (h) 12 (v)
Spectral brightness	Photons/s/mm ² /mR ² /0.1%BW	5.4×10^9	1.5×10^{17}

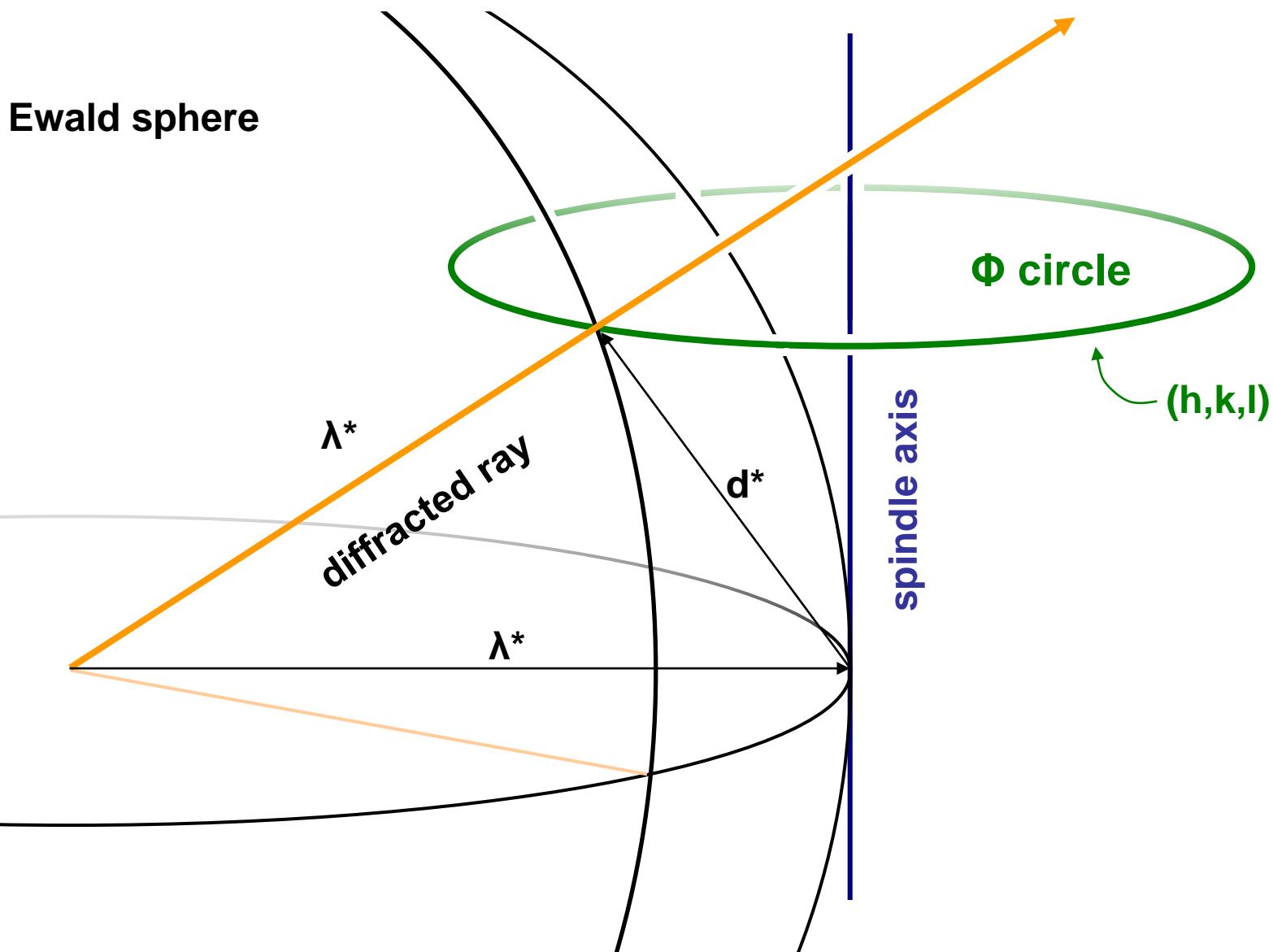
The truth about x-ray beams

quantity	units	home source	MX2
flux	Photons/second	8×10^4	2×10^{12}
exposure	time	5 months	1 second
Dispersion	wavelength range / wavelength	0.014% (Si111)	0.014% (Si111)
Divergence	milliRadian	0.6	1.8 (h) 0.18 (v)
Beam size	microns	17	24 (h) 12 (v)
Spectral brightness	Photons/s/mm ² /mR ² /0.1%BW	5.4×10^9	1.5×10^{17}

The truth about x-ray beams

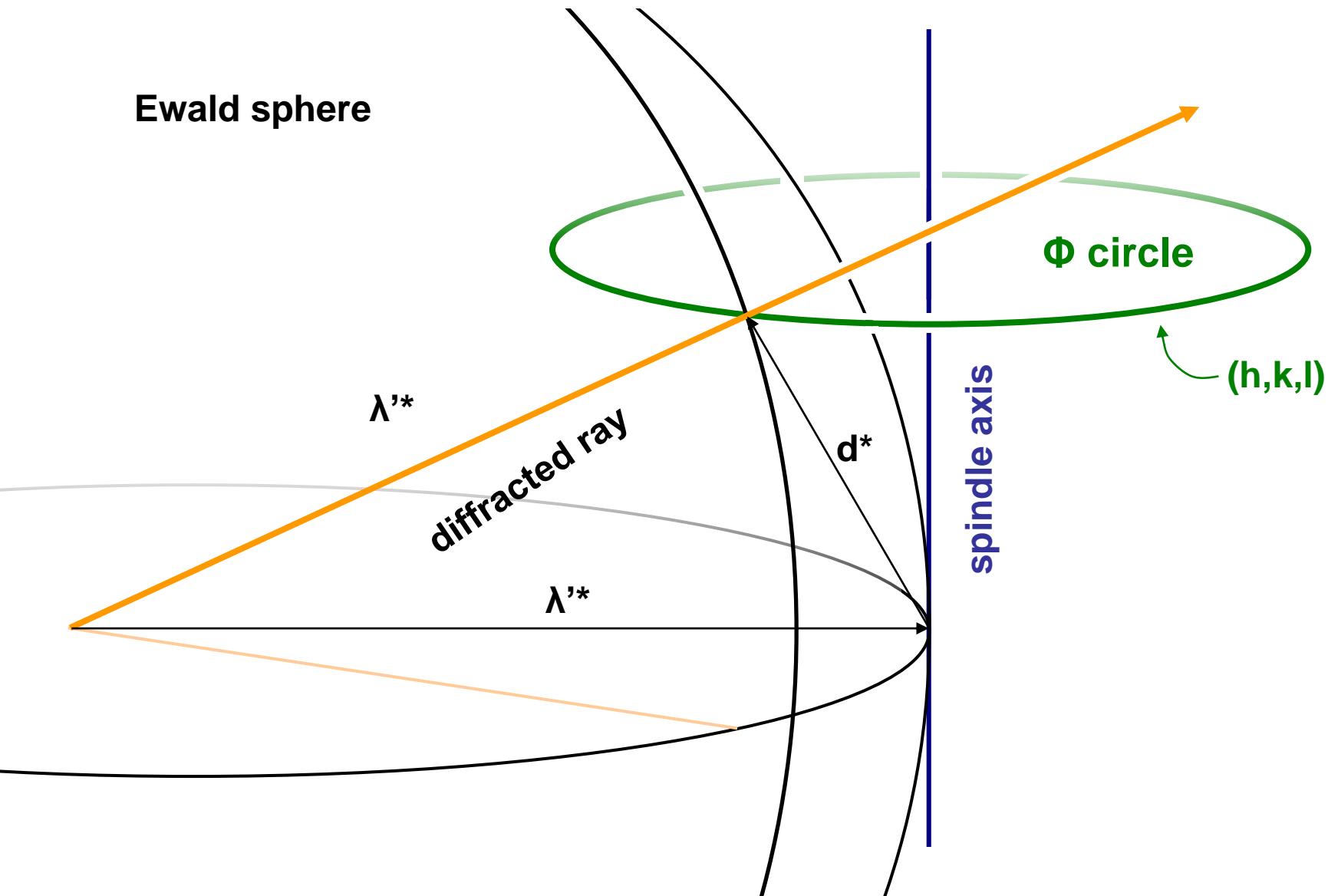
Term	units	significance
Flux	photons/s	duration of experiment
Beam Size	μm	match to crystal
Divergence	mrad	spot size vs distance
Wavelength	\AA	resolution and absorption
Dispersion	$\Delta\lambda/\lambda$	spot size
Flux density	ph/s/area	scattering/damage rate
Fluence	ph/area	scattering/damage

spectral dispersion



spectral dispersion

Ewald sphere



Si(111) vs multilayers

0.014%
Si(111)

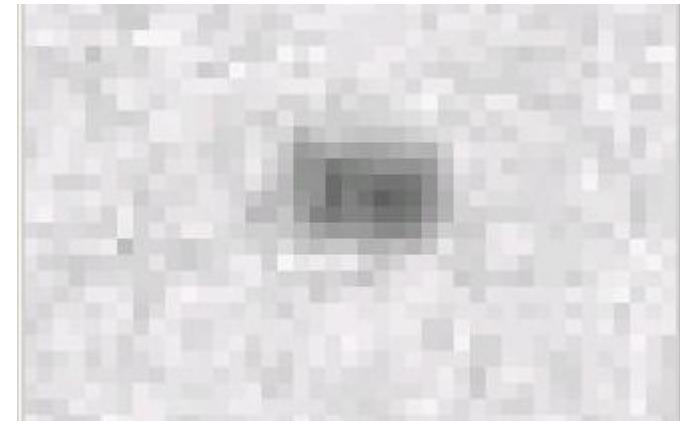
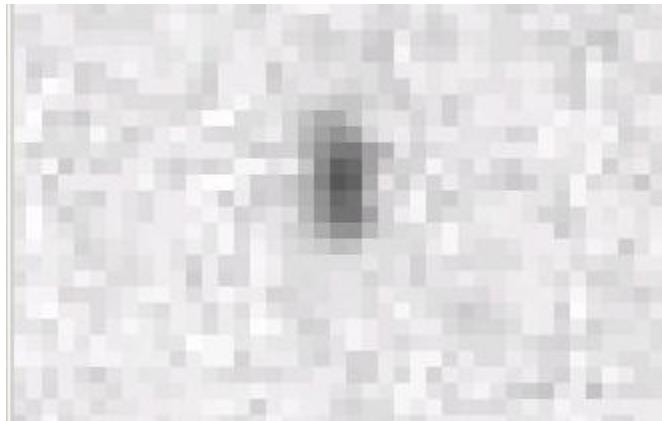
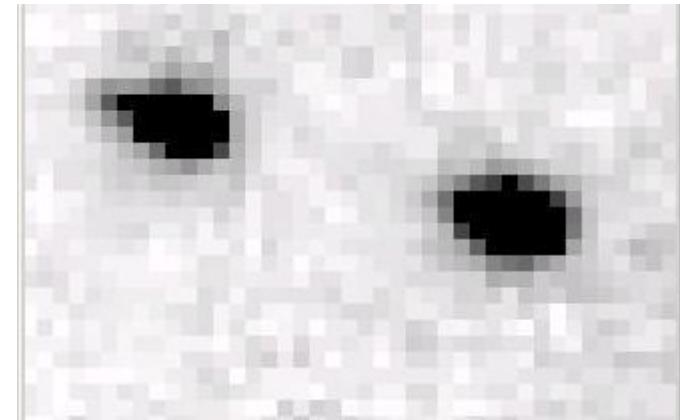
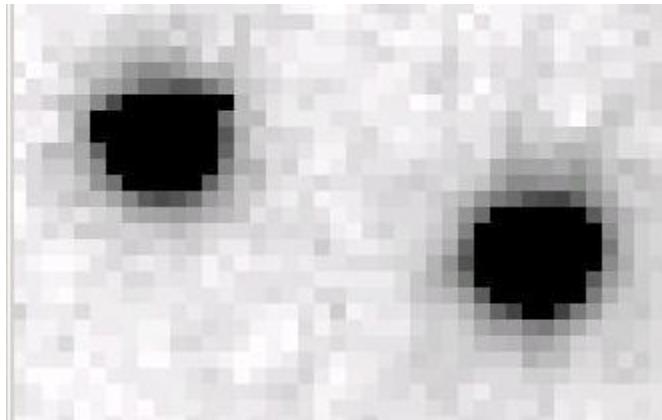
spectral
dispersion

1%
multilayer

4 Å

resolution

1.9 Å



Decisions, Decisions, Decisions

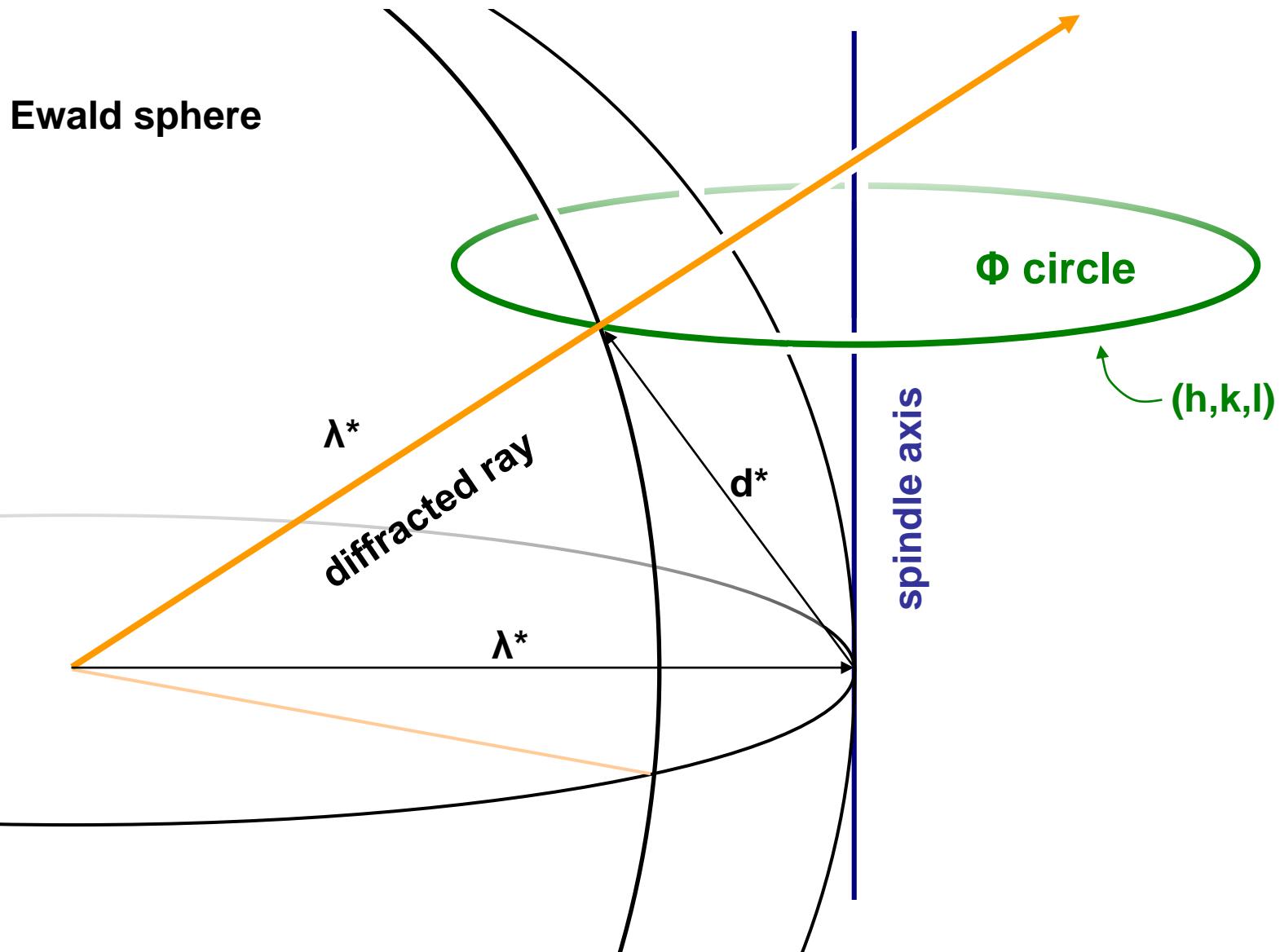
- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, **bandpass**
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

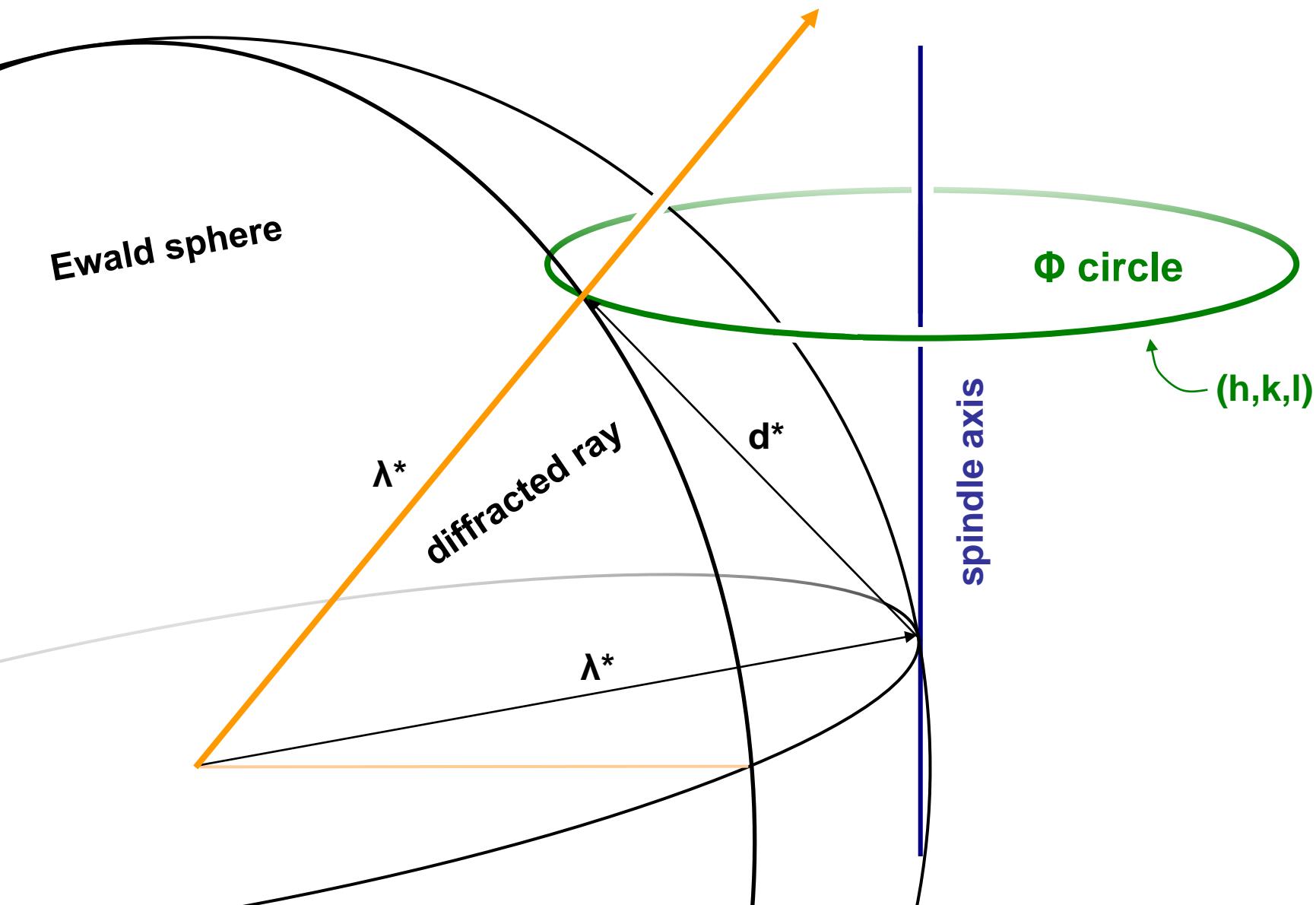
The truth about x-ray beams

Term	units	significance
Flux	photons/s	duration of experiment
Beam Size	μm	match to crystal
Divergence	mrad	spot size vs distance
Wavelength	\AA	resolution and absorption
Dispersion	$\Delta\lambda/\lambda$	spot size
Flux density	ph/s/area	scattering/damage rate
Fluence	ph/area	scattering/damage

beam divergence



beam divergence



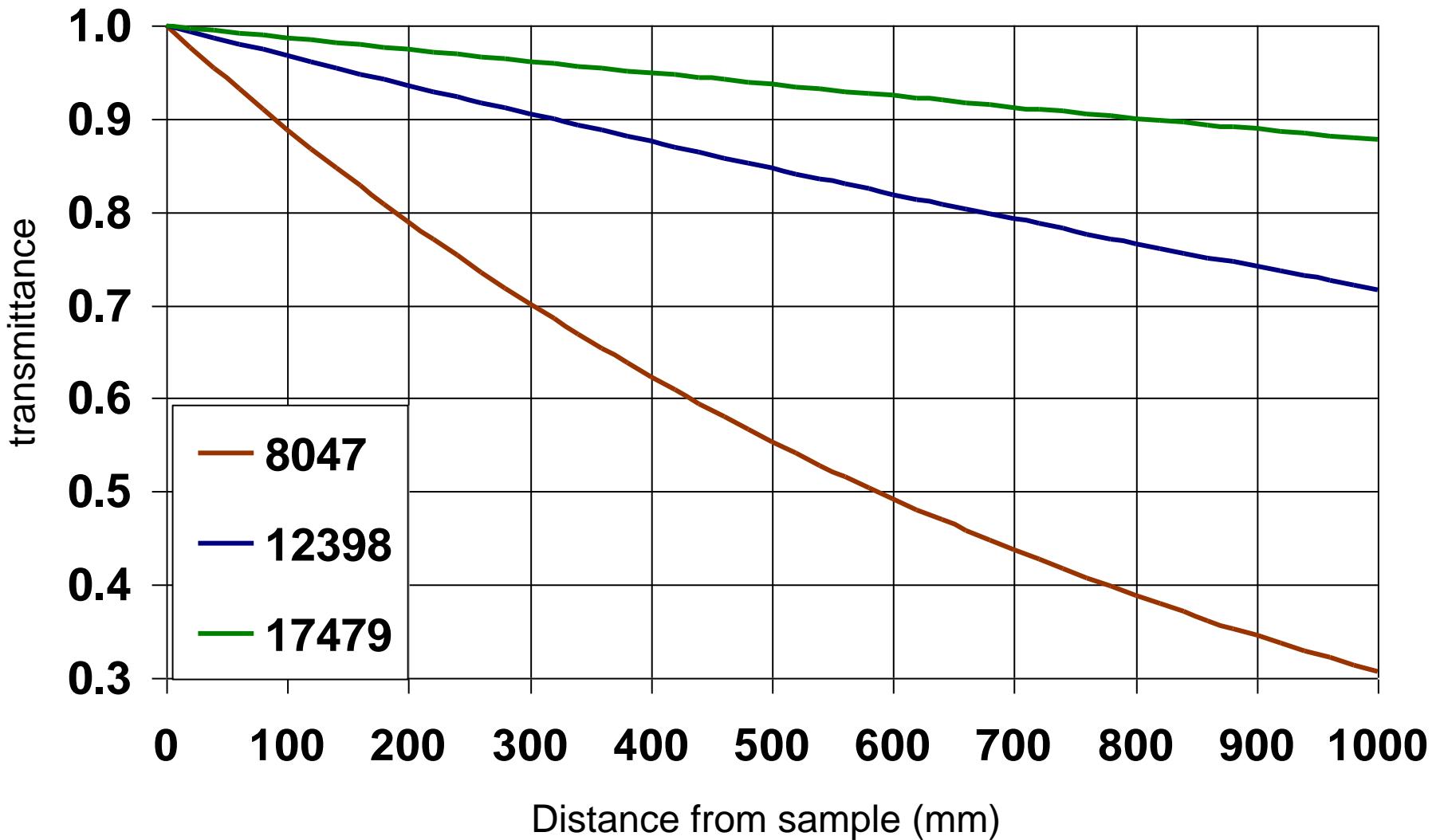
divergence = 0 °

divergence = 0.3 °

The truth about x-ray beams

Term	units	significance
Flux	photons/s	duration of experiment
Beam Size	μm	match to crystal
Divergence	mrad	spot size vs distance
Wavelength	\AA	resolution and absorption
Dispersion	$\Delta\lambda/\lambda$	spot size
Flux density	ph/s/area	scattering/damage rate
Fluence	ph/area	scattering/damage

Air absorption



The truth about x-ray beams

Term	units	significance
Flux	photons/s	duration of experiment
Beam Size	μm	match to crystal
Divergence	mrad	spot size vs distance
Wavelength	\AA	resolution and absorption
Dispersion	$\Delta\lambda/\lambda$	spot size
Flux density	ph/s/area	scattering/damage rate
Fluence	ph/area	scattering/damage

Decisions, Decisions, Decisions

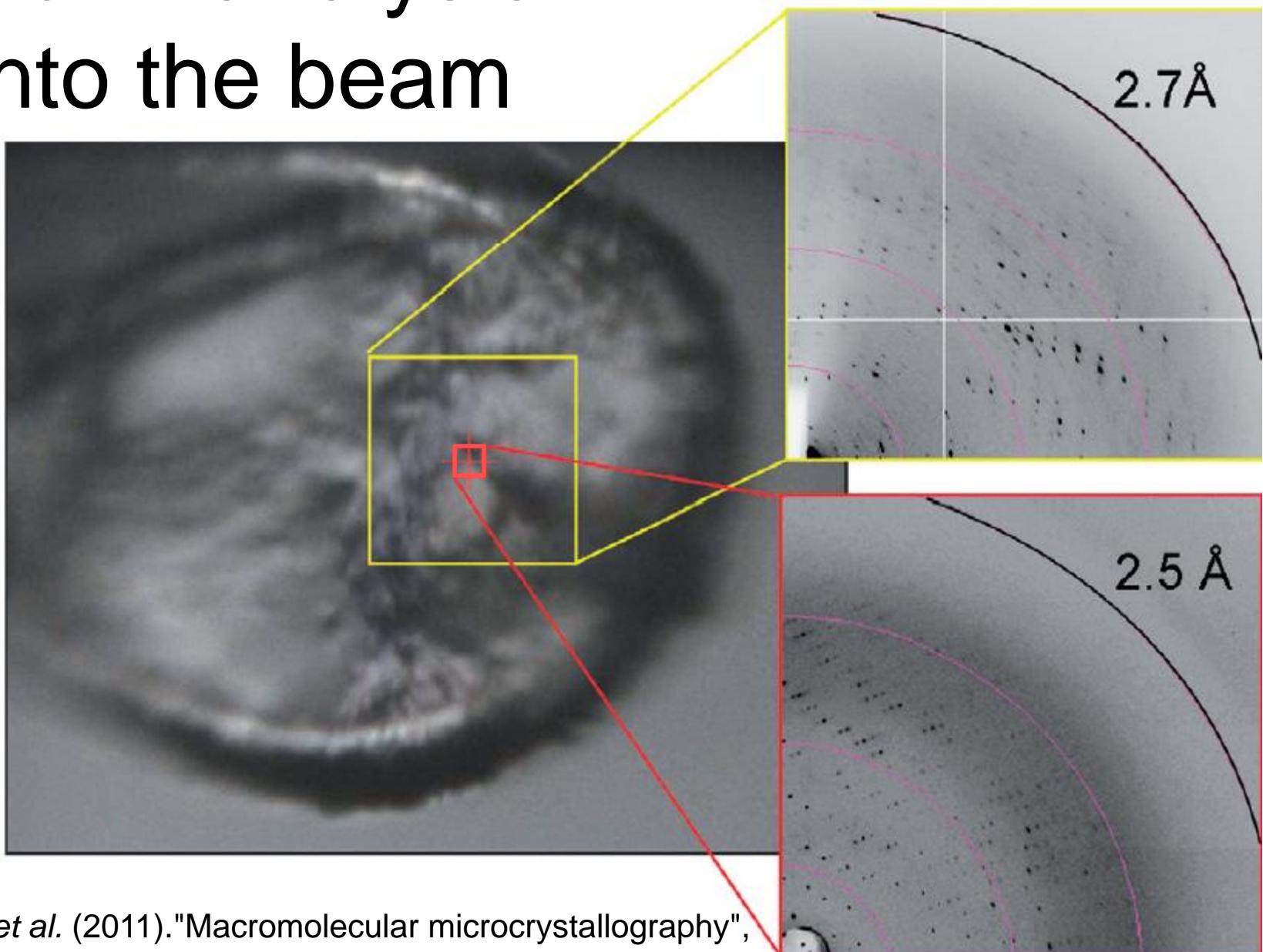
- Exposure time
- Number of images
- Wavelengths
- Beam: **size**, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

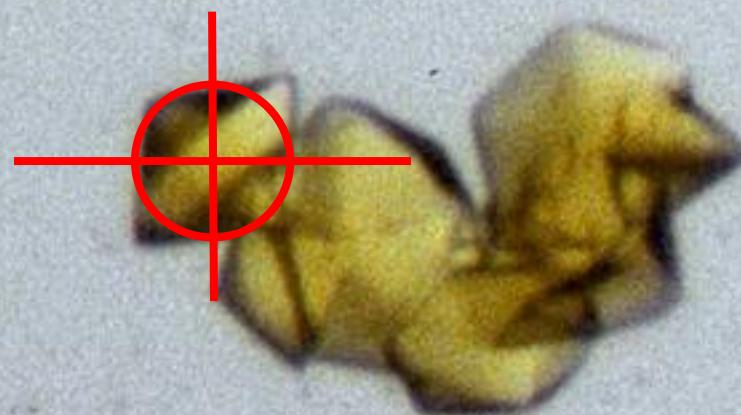
beam size vs xtal size

1. Put your crystal into the beam

Put the “crystal” into the beam



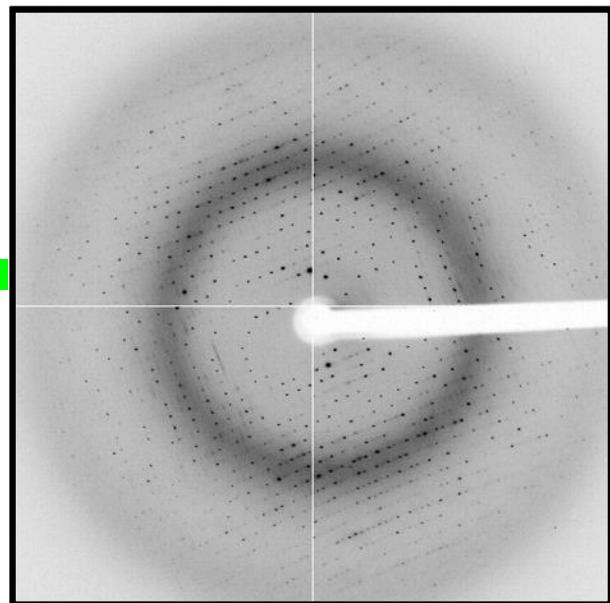
“crystal” = thing you want to shoot



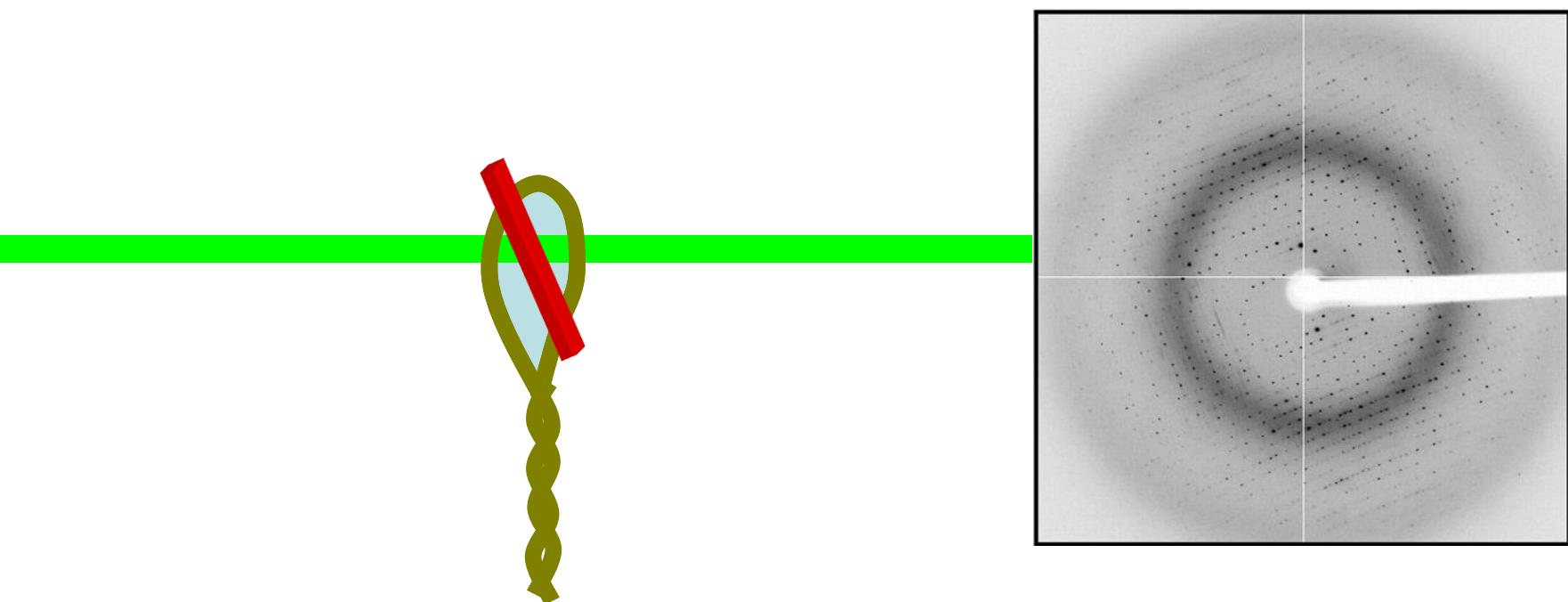
beam size vs xtal size

1. Put your crystal into the beam
2. Shoot the whole crystal

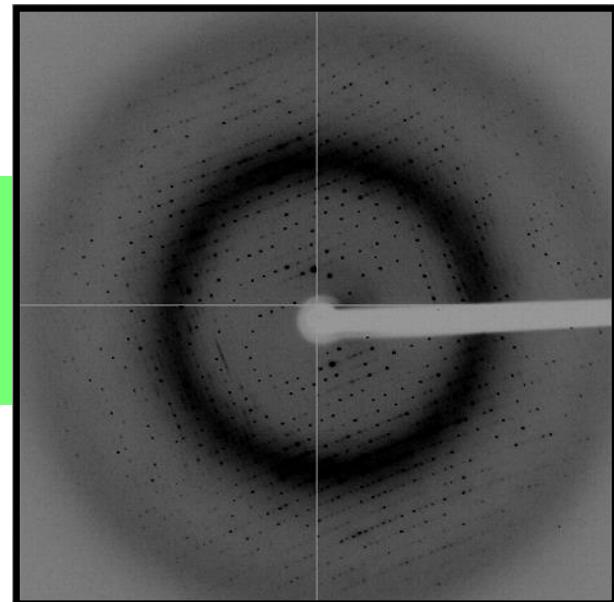
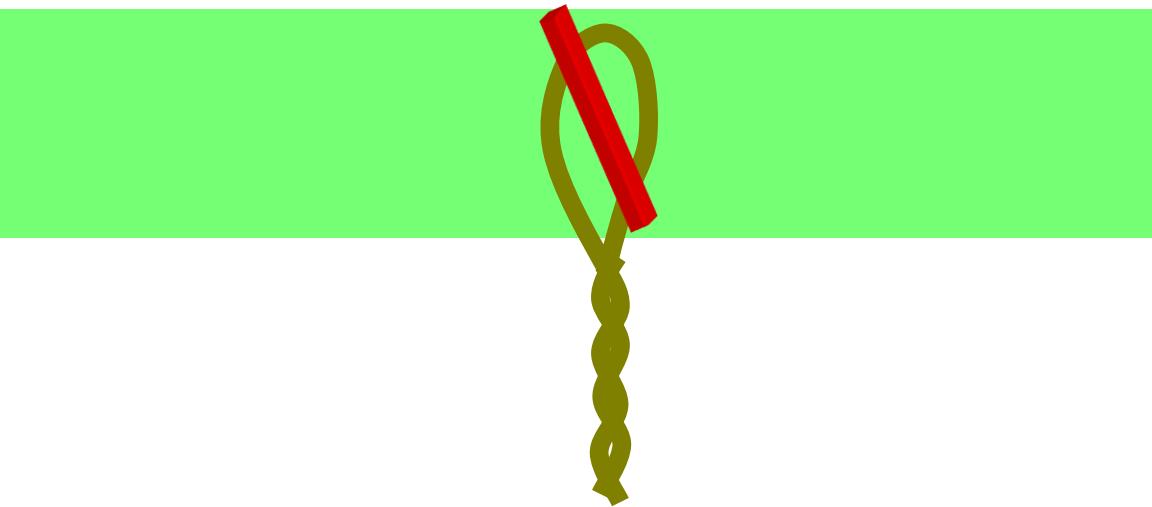
shoot the whole crystal



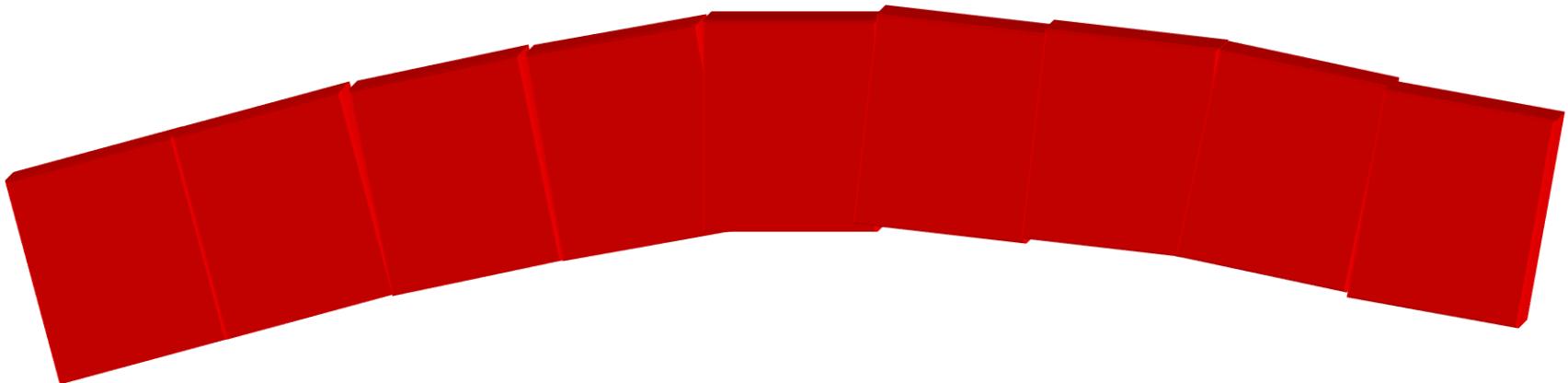
shoot the whole crystal



shoot the whole crystal

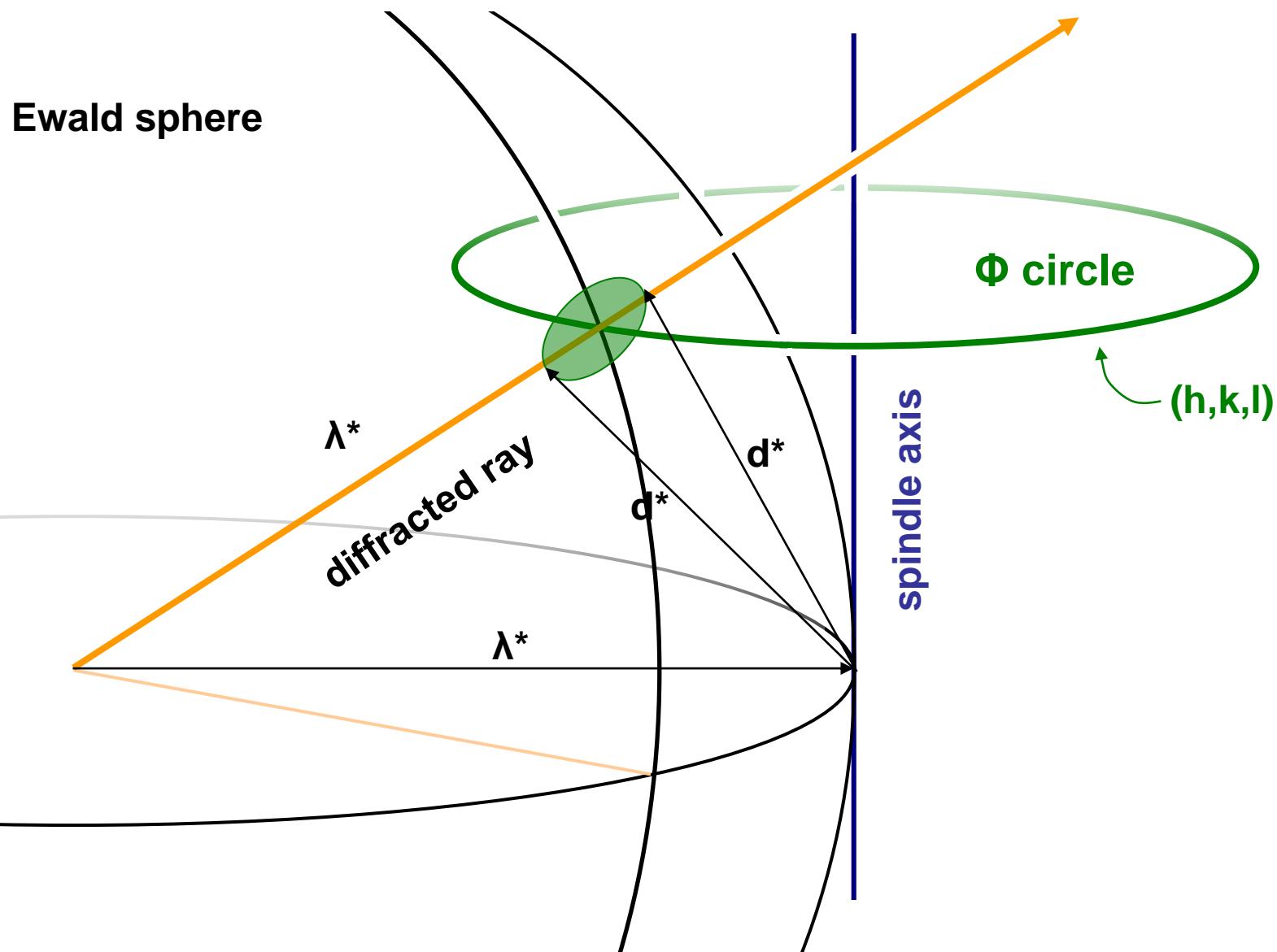


How many crystals do you see?



Shoot the “**crystal**” (singular)

mosaic spread



mosaic spread

Ewald sphere

diffracted rays

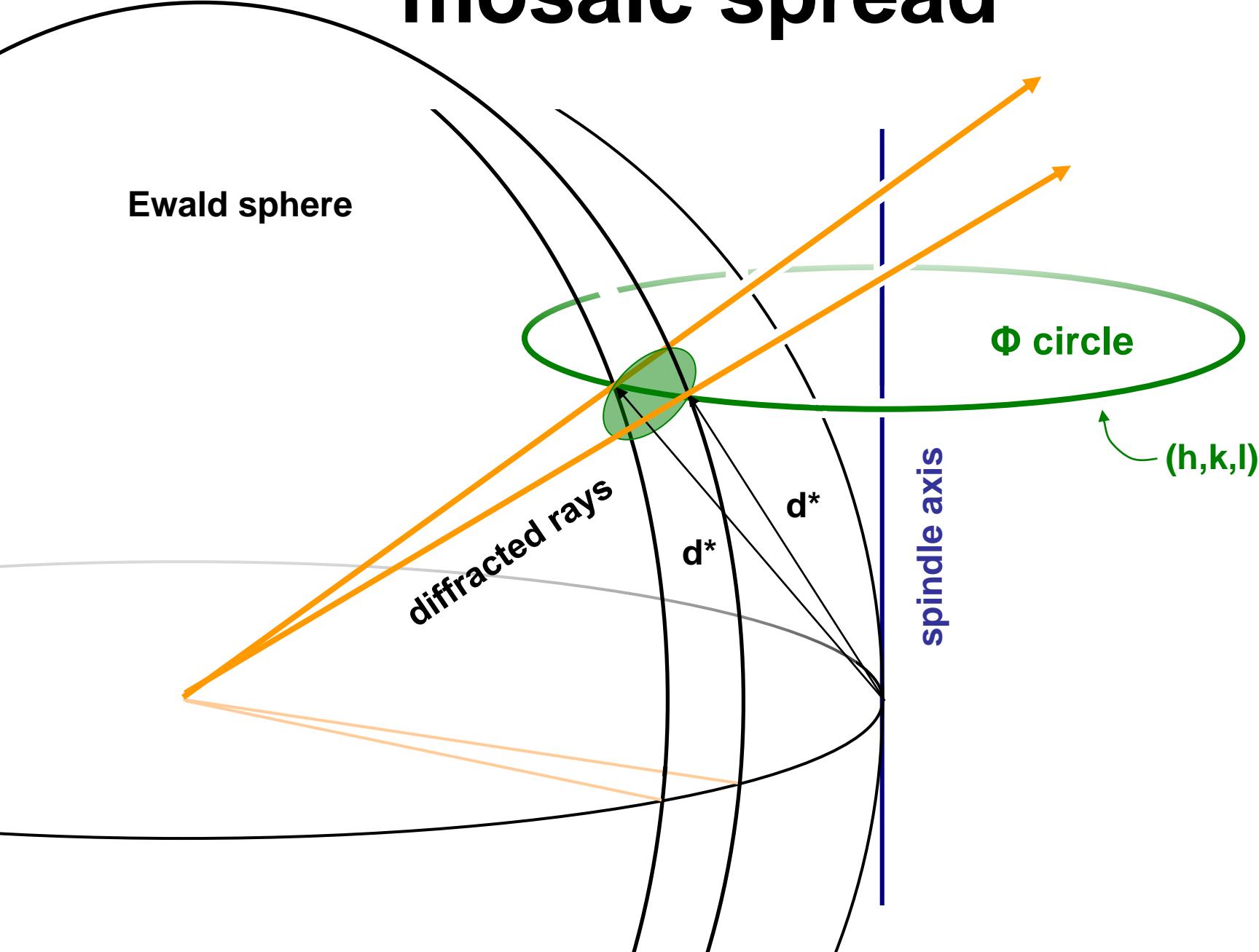
d^*

d^*

spindle axis

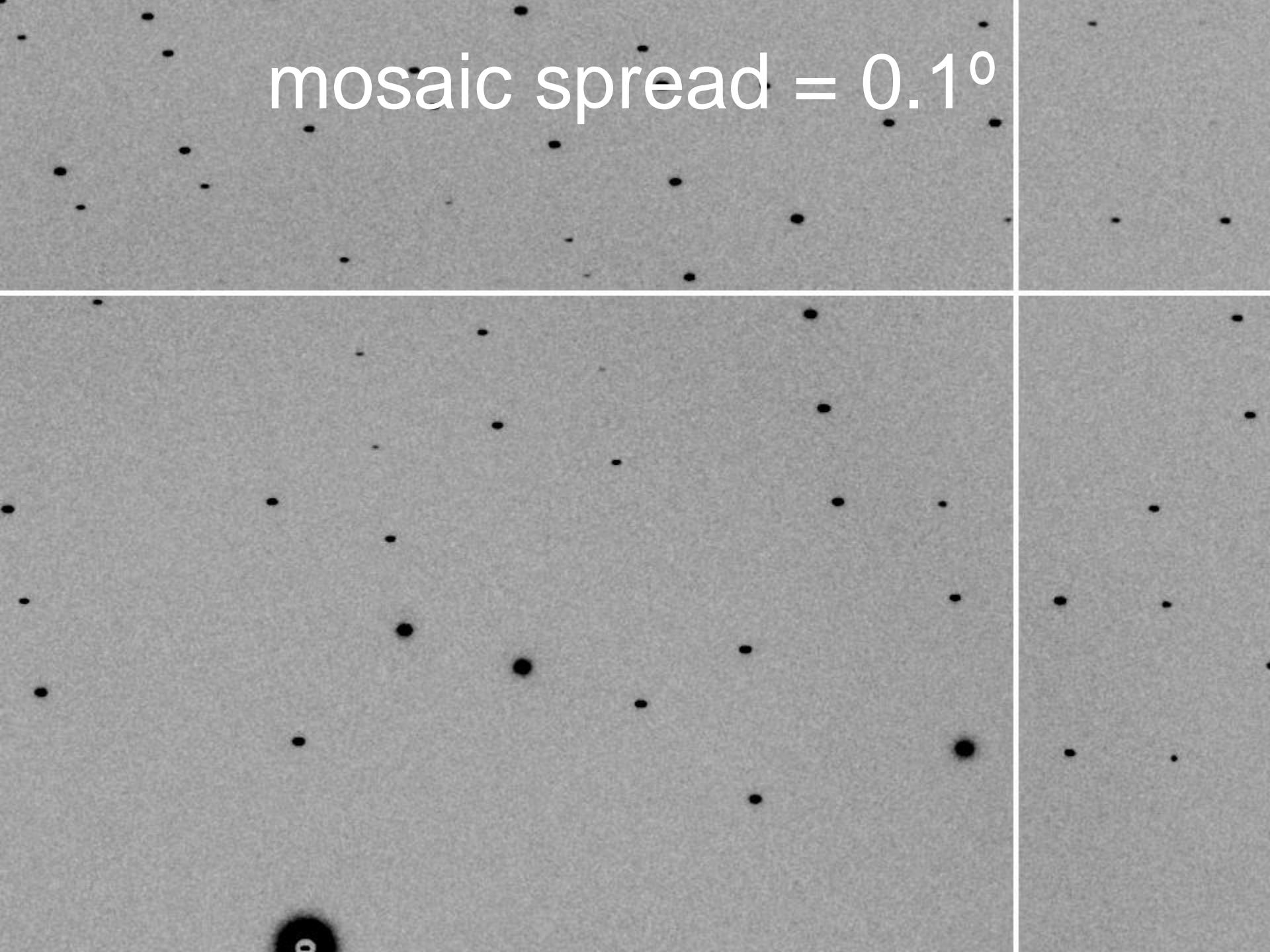
Φ circle

(h, k, l)

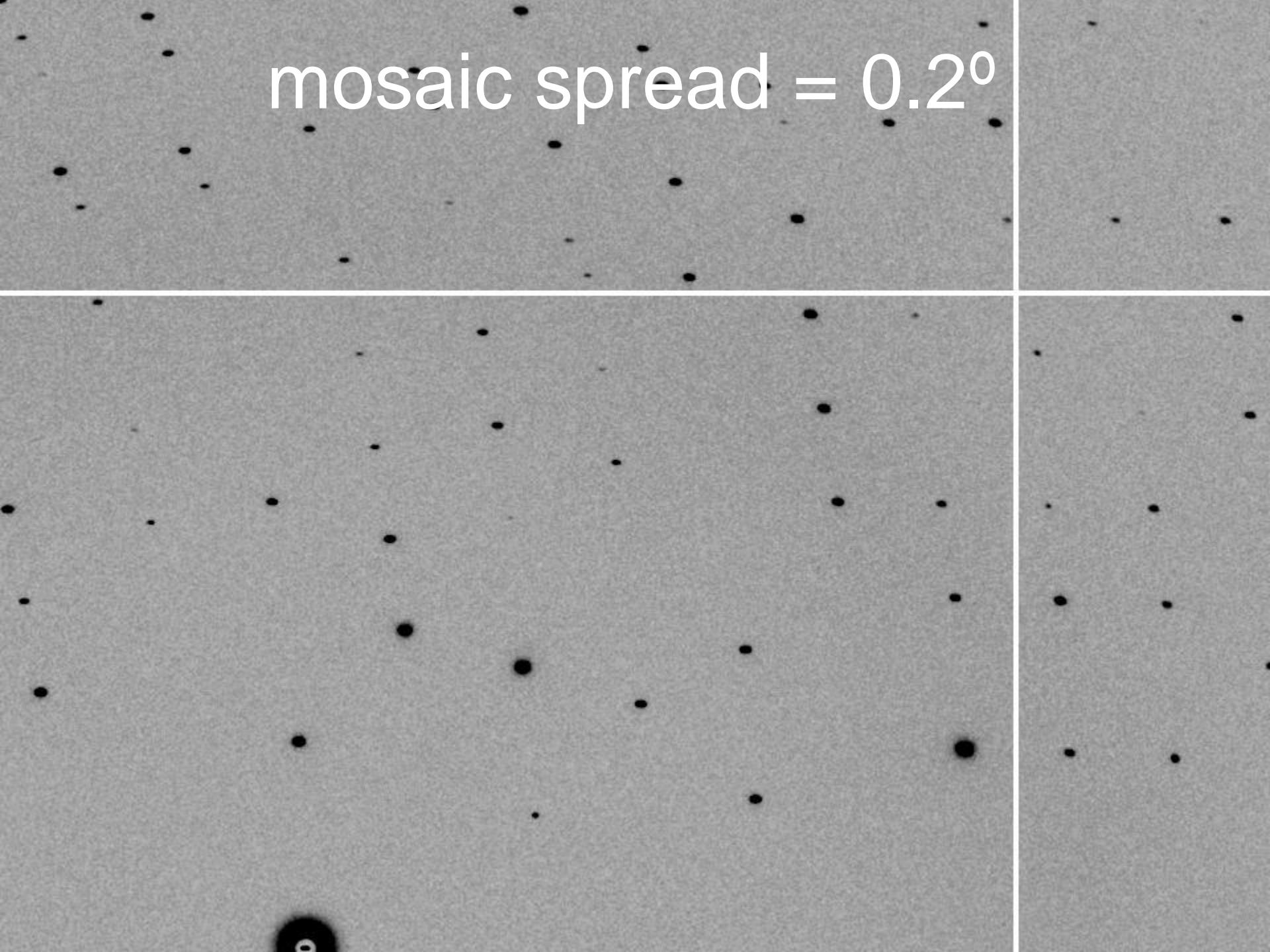


mosaic spread = 0 °

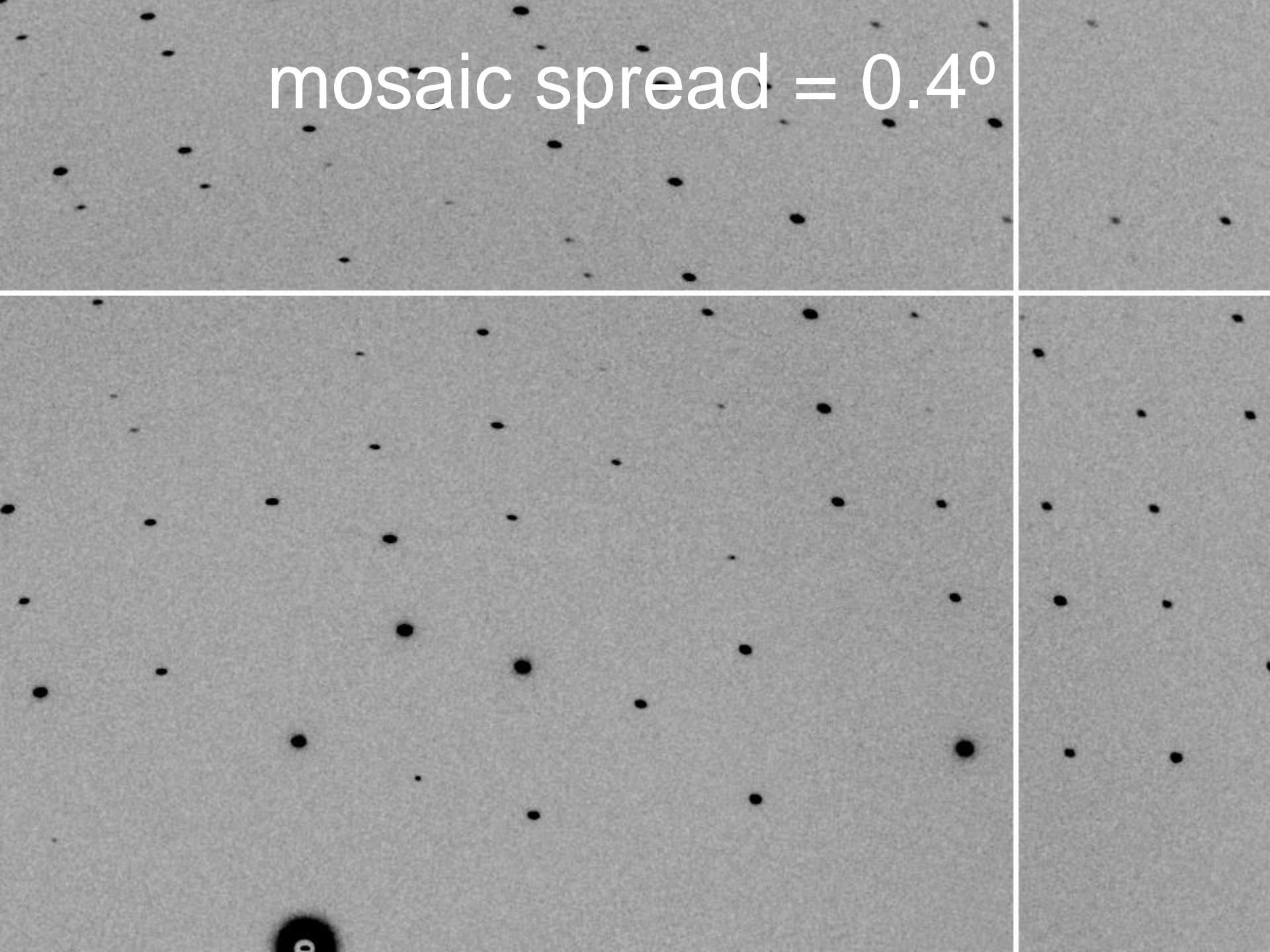
mosaic spread = 0.1°

A grayscale image showing a grid of small black dots on a gray background. The dots are arranged in a regular pattern, creating a sense of a mosaic or a grid. The overall effect is similar to a grayscale version of the 'Brennan's Grid' test pattern.

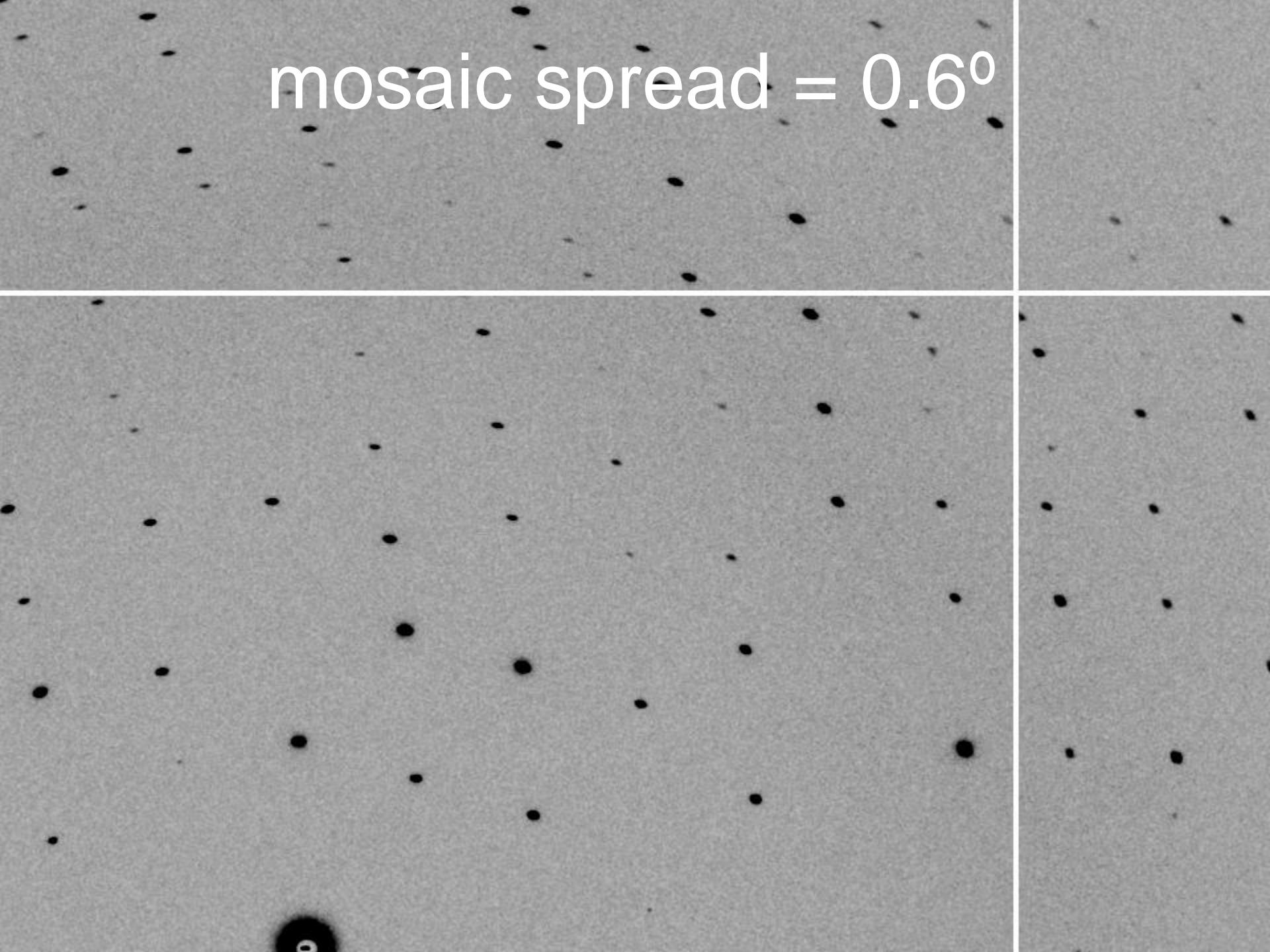
mosaic spread = 0.2°



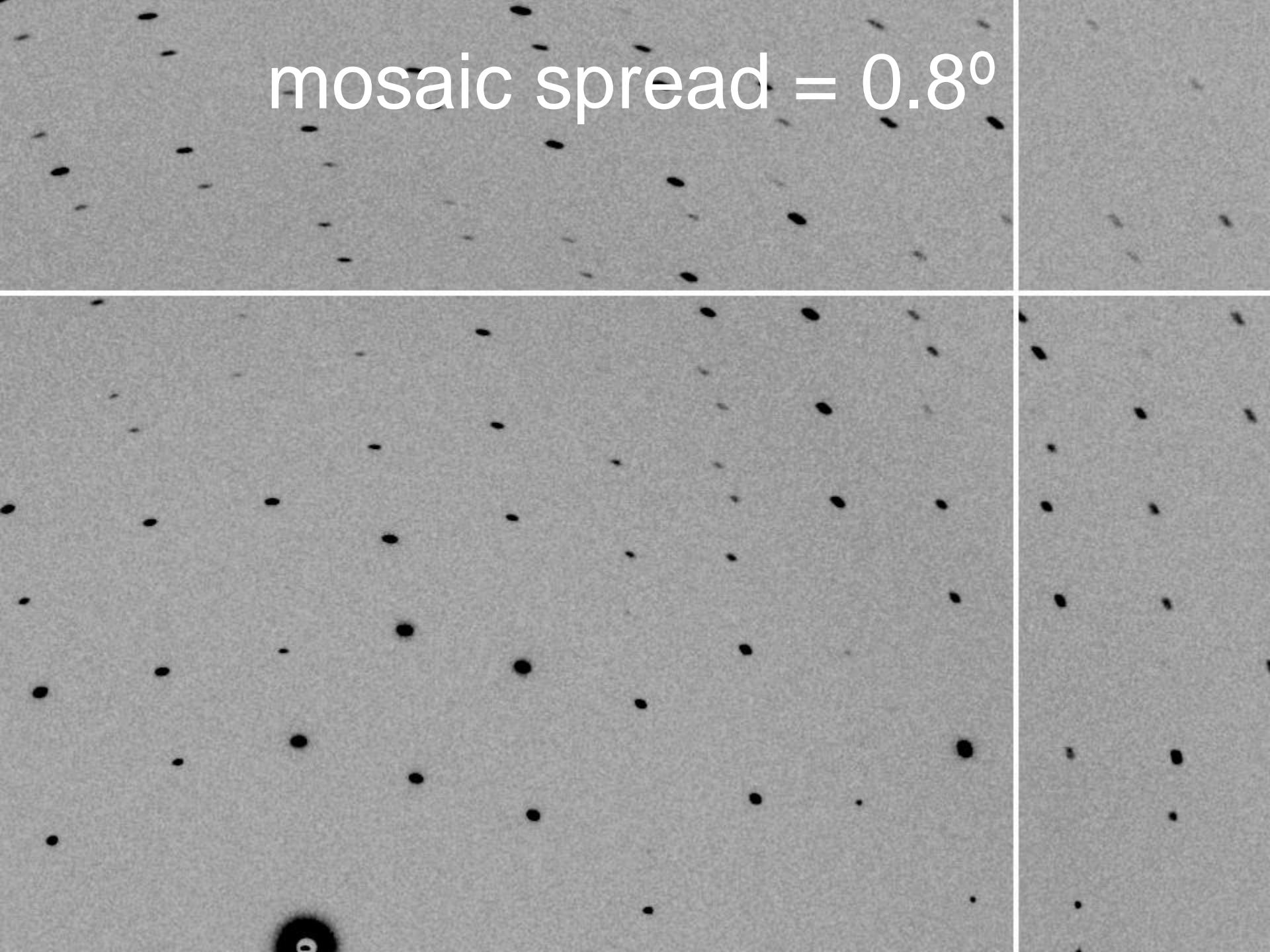
mosaic spread = 0.4°



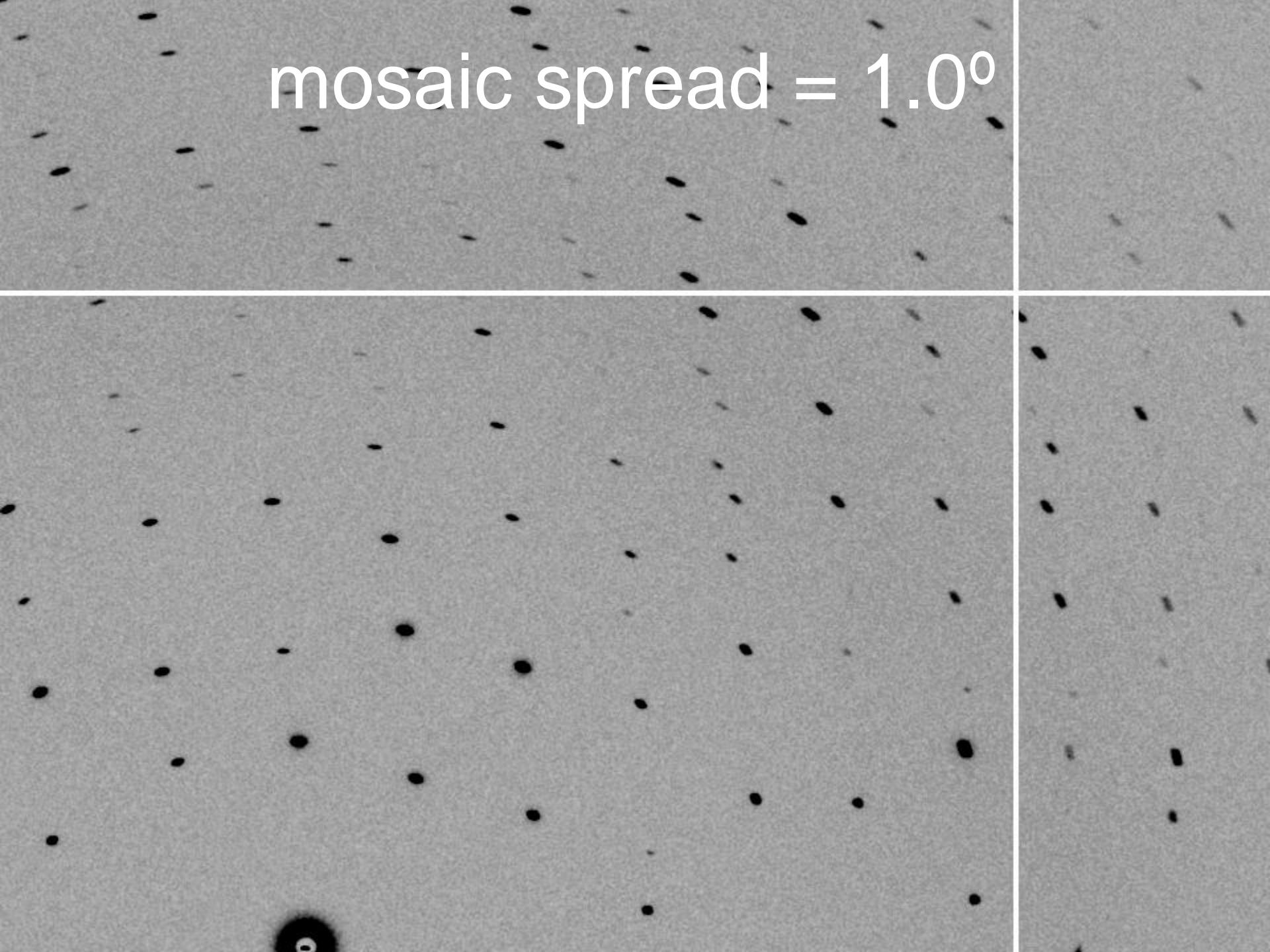
mosaic spread = 0.6°



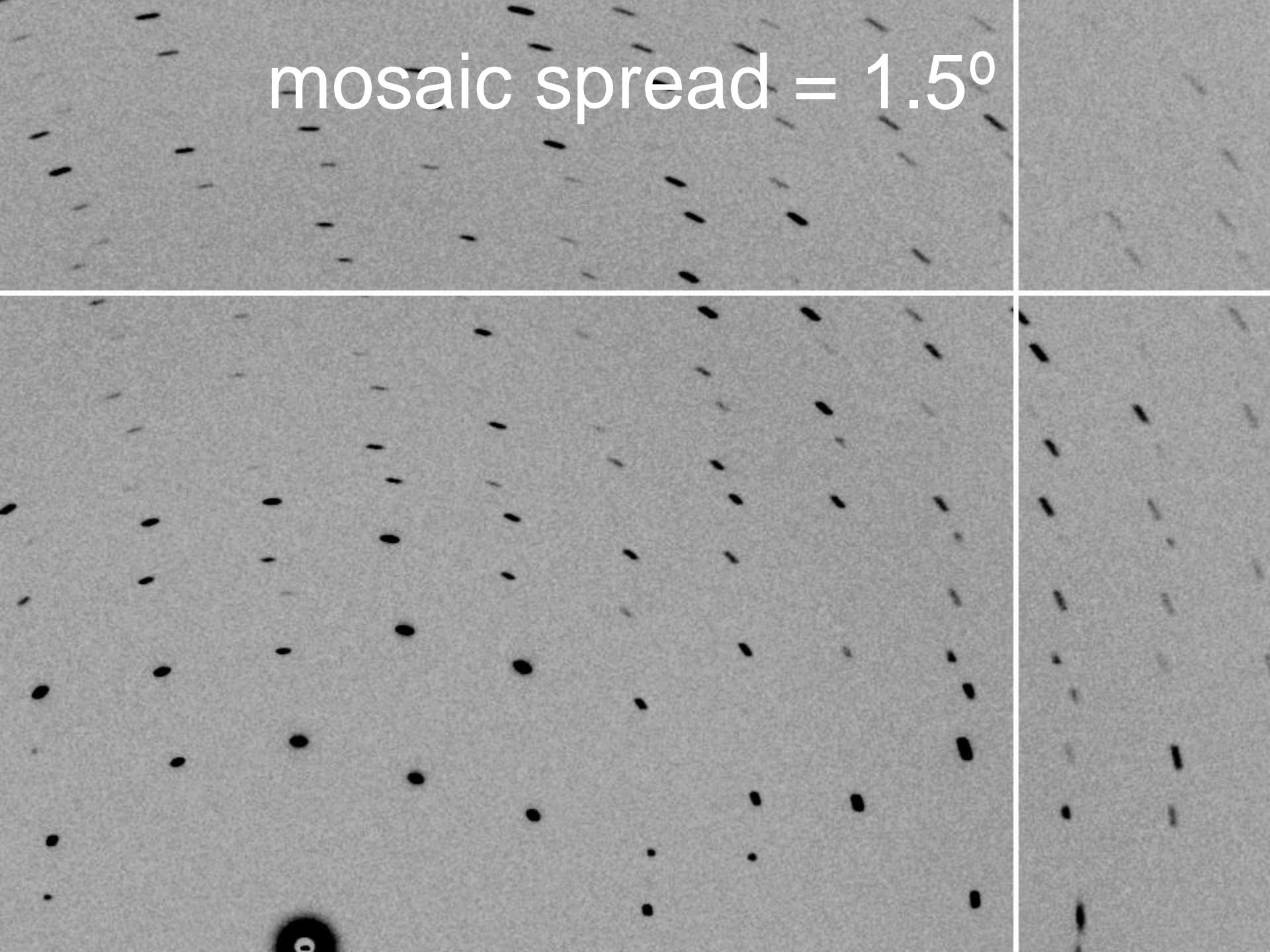
mosaic spread = 0.8°



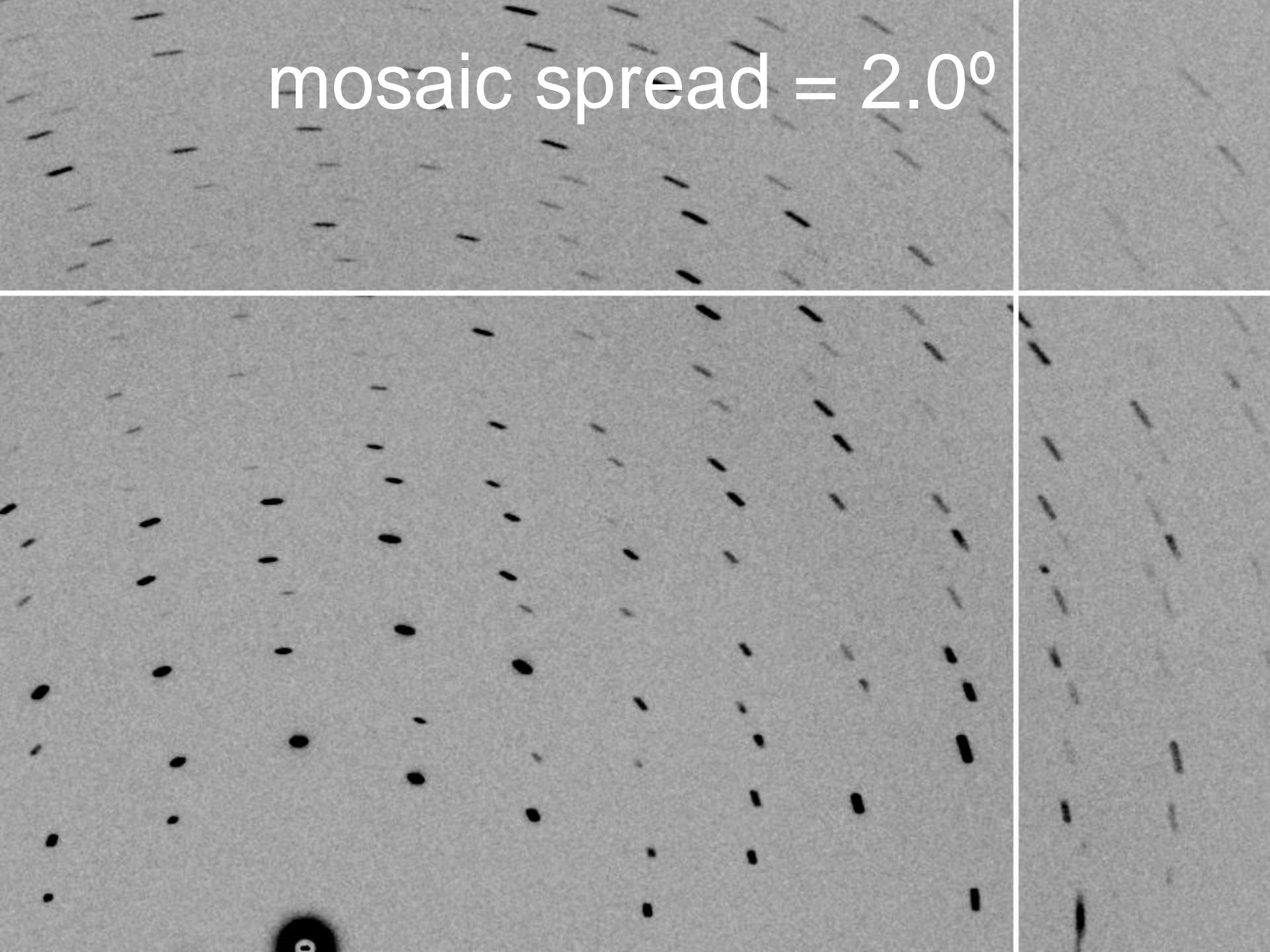
mosaic spread = 1.0°

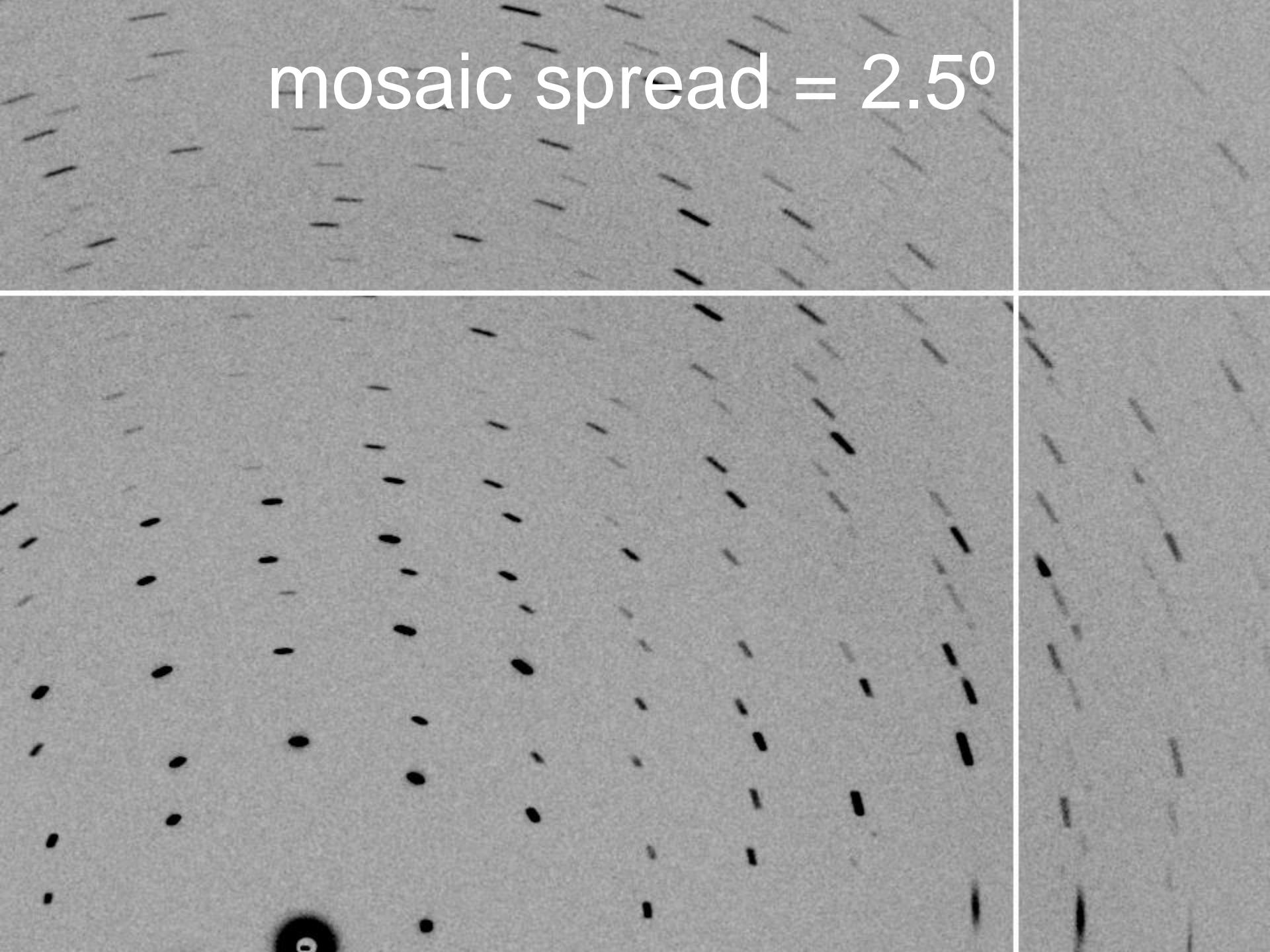


mosaic spread = 1.5°

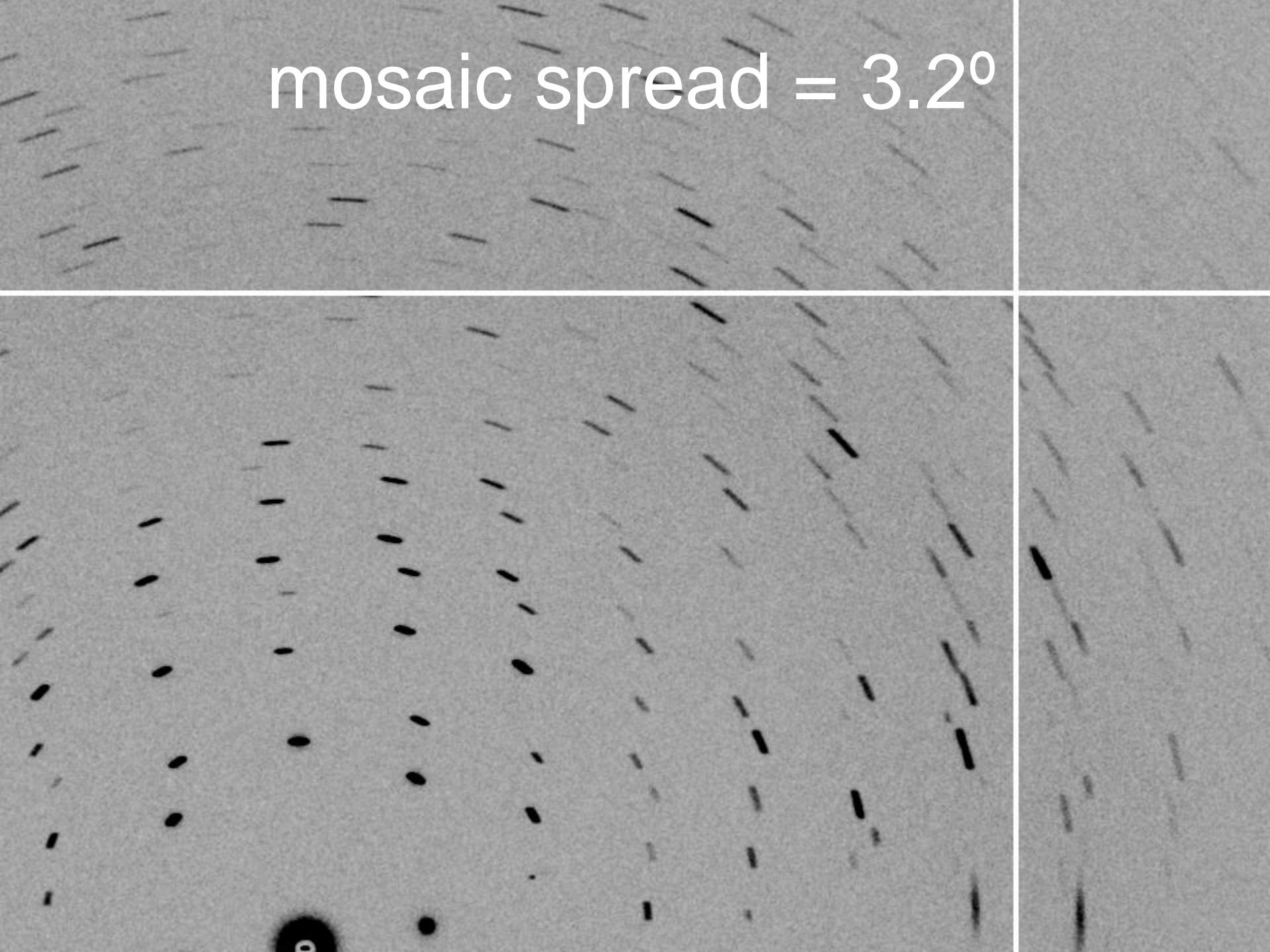


mosaic spread = 2.0°



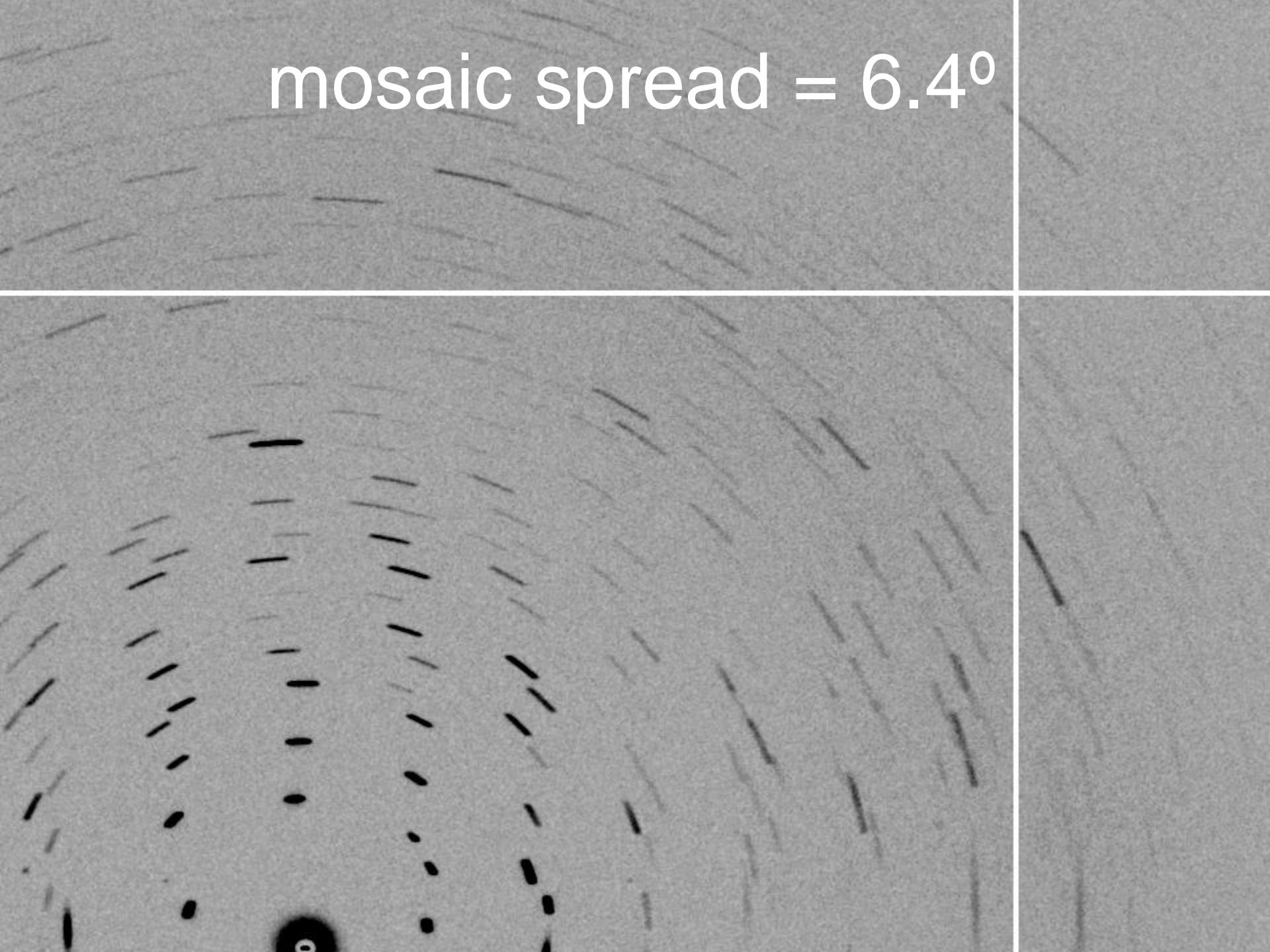


mosaic spread = 2.5°

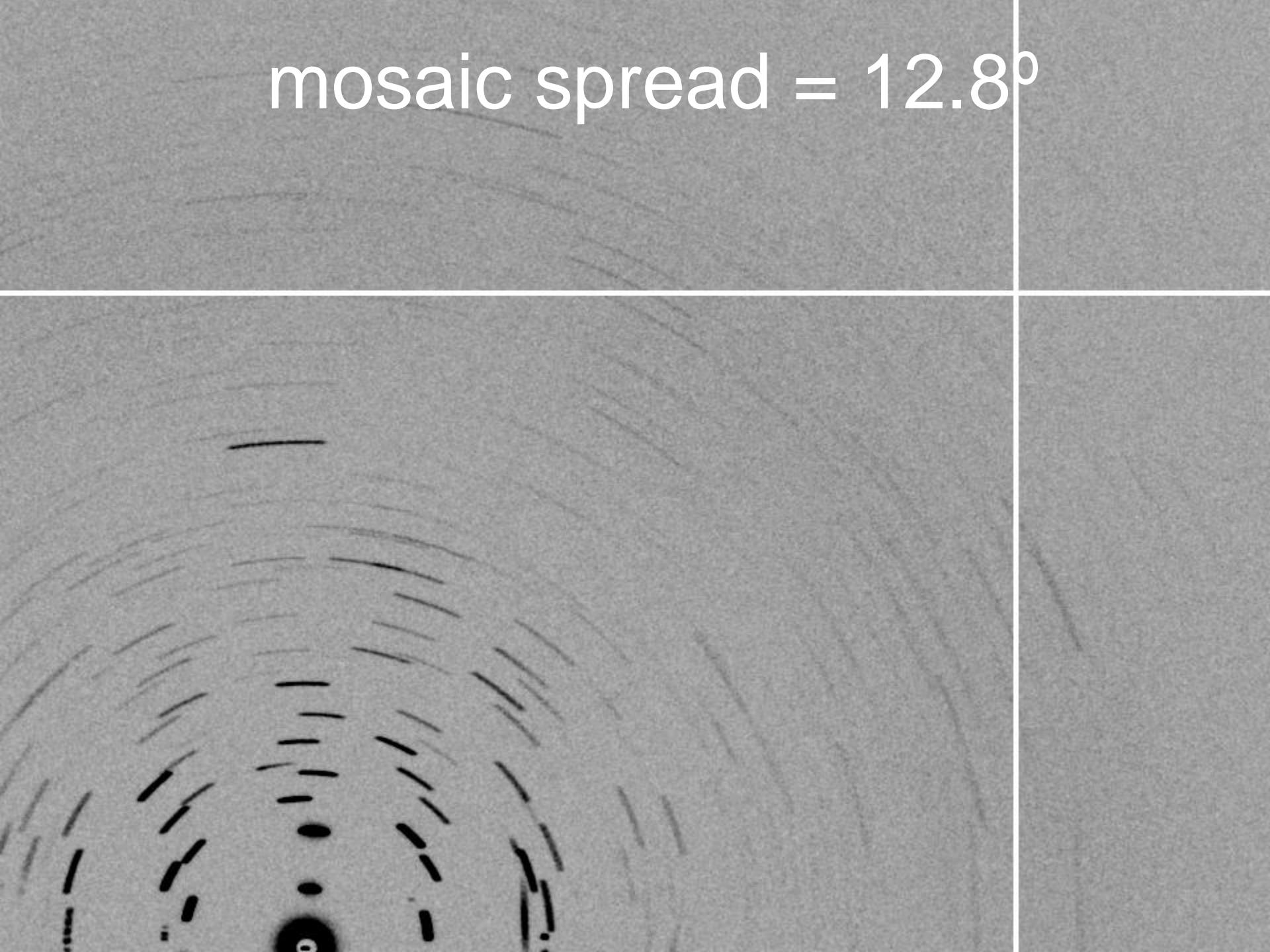


mosaic spread = 3.2°

mosaic spread = 6.4°

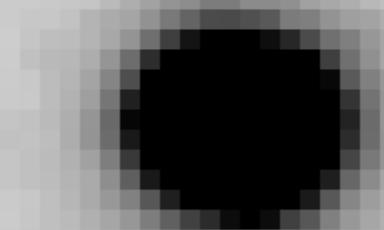


mosaic spread = 12.8°



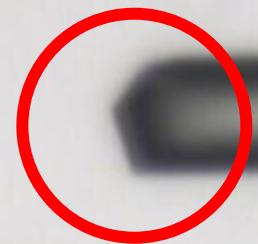


~0 MGy

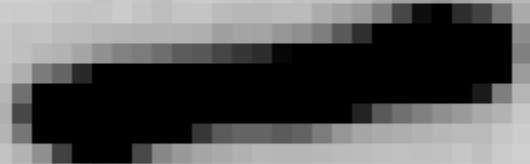
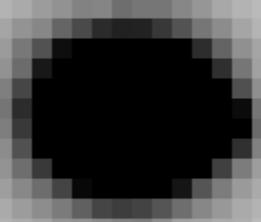




5 MGy

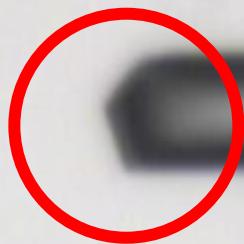


11 MGy

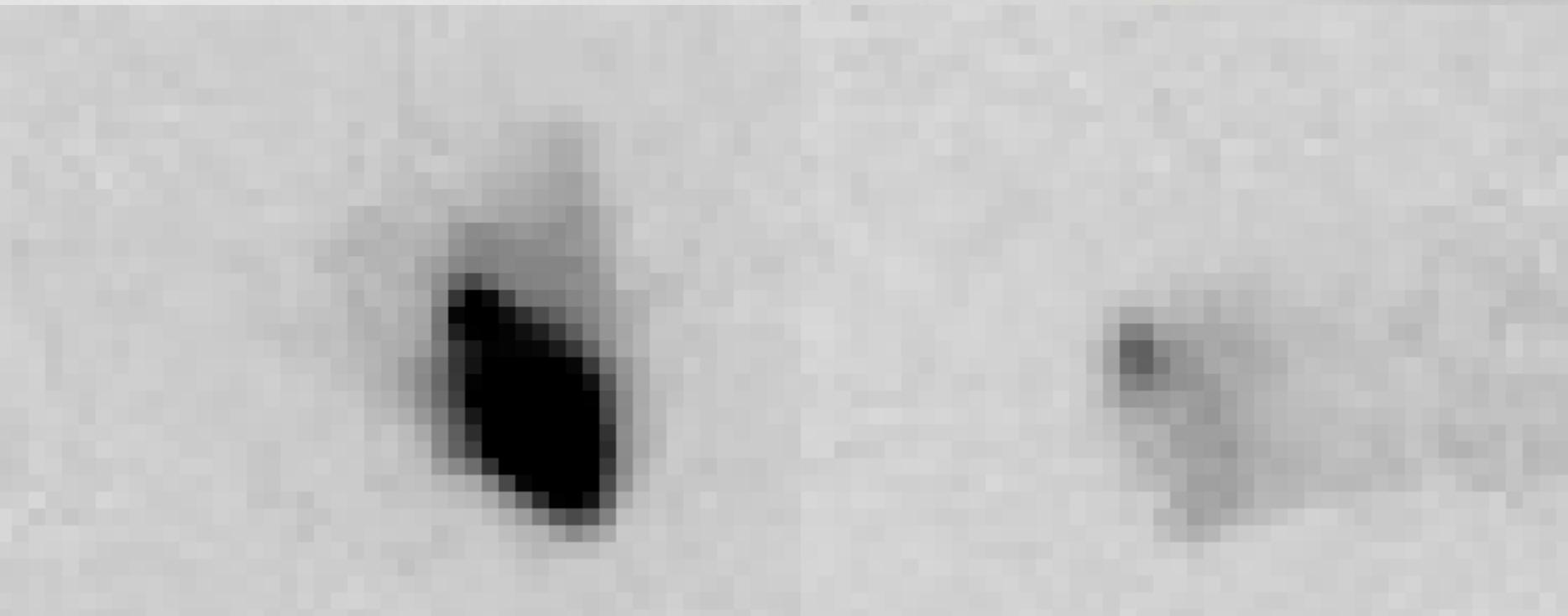




16 MGy



22 MGy

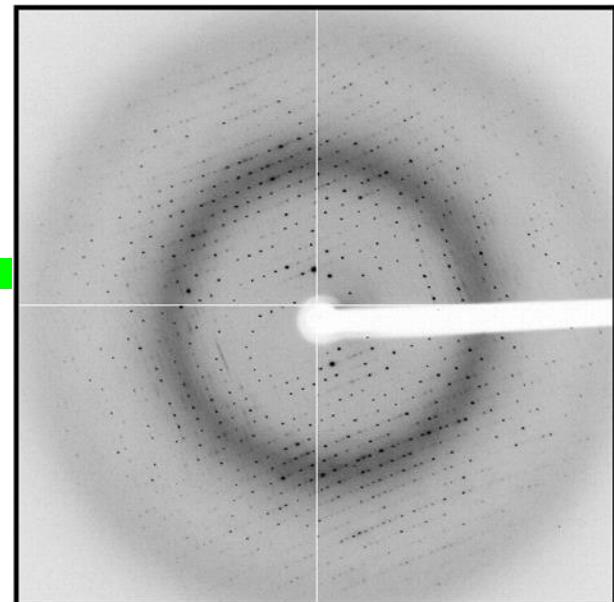
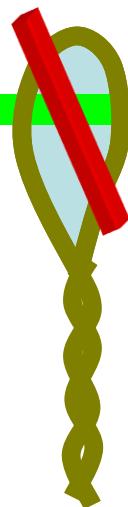




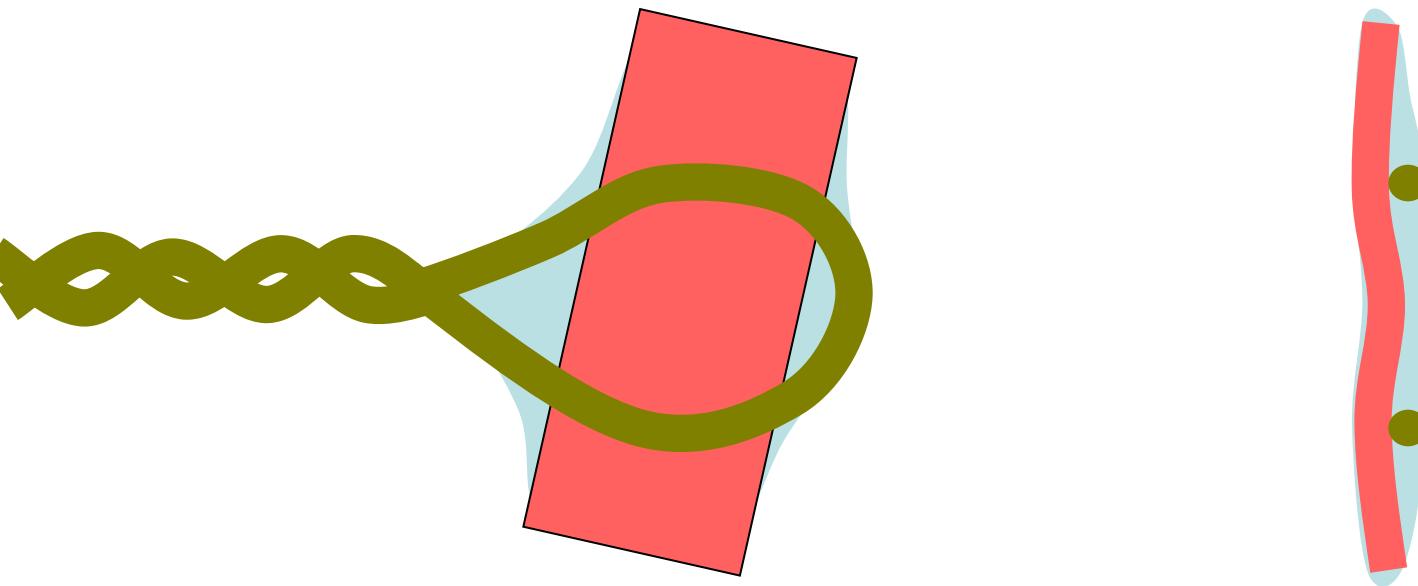
26 MGy

shoot the whole crystal

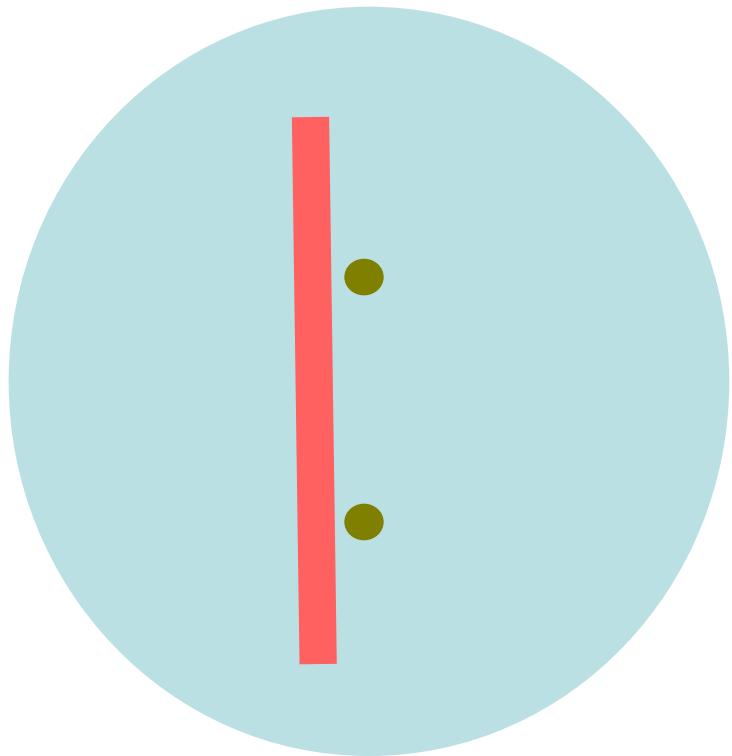
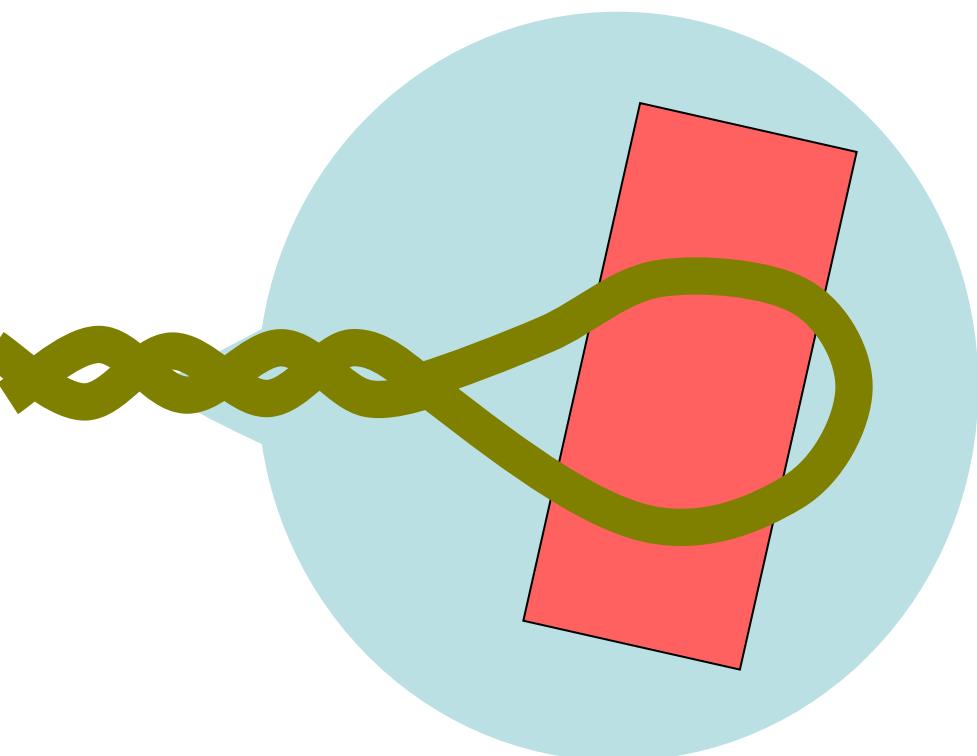
Do not bend!



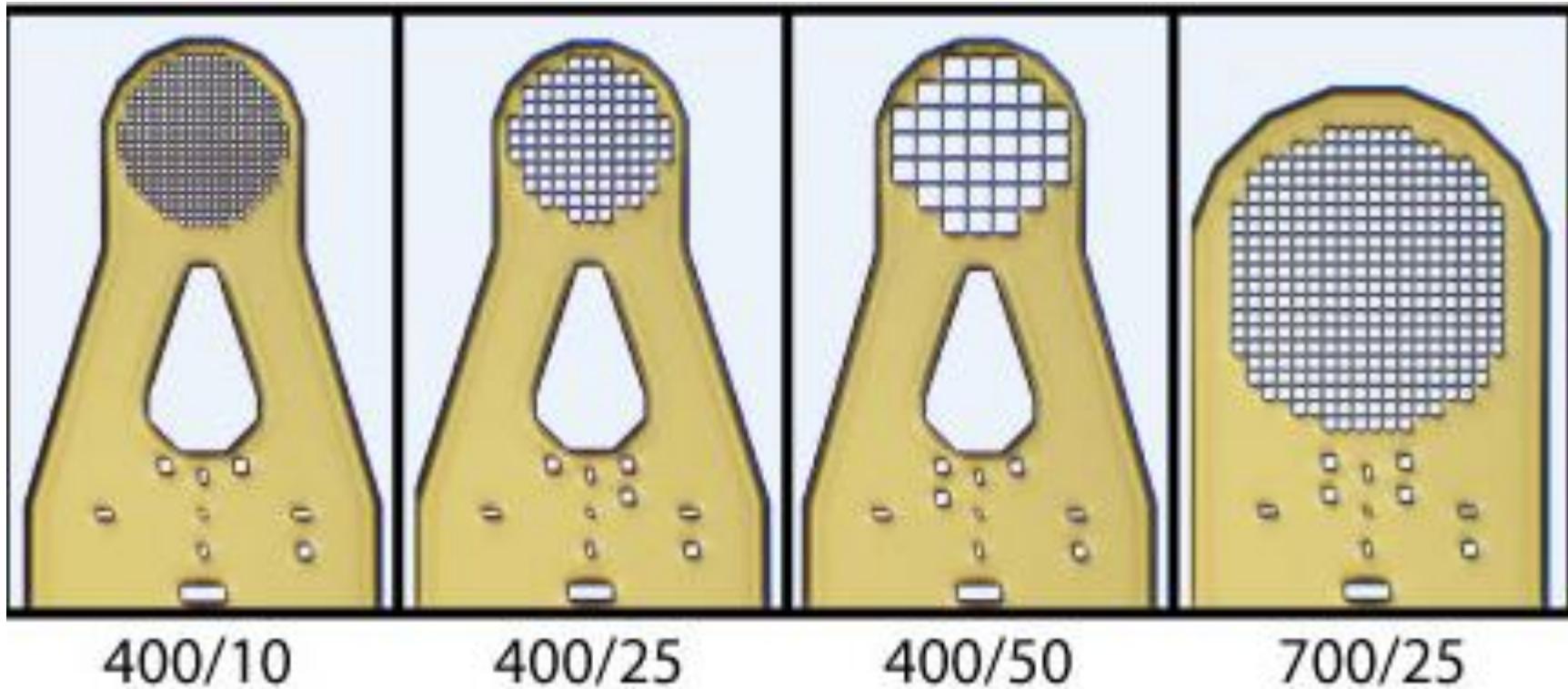
platy crystals



platy crystals

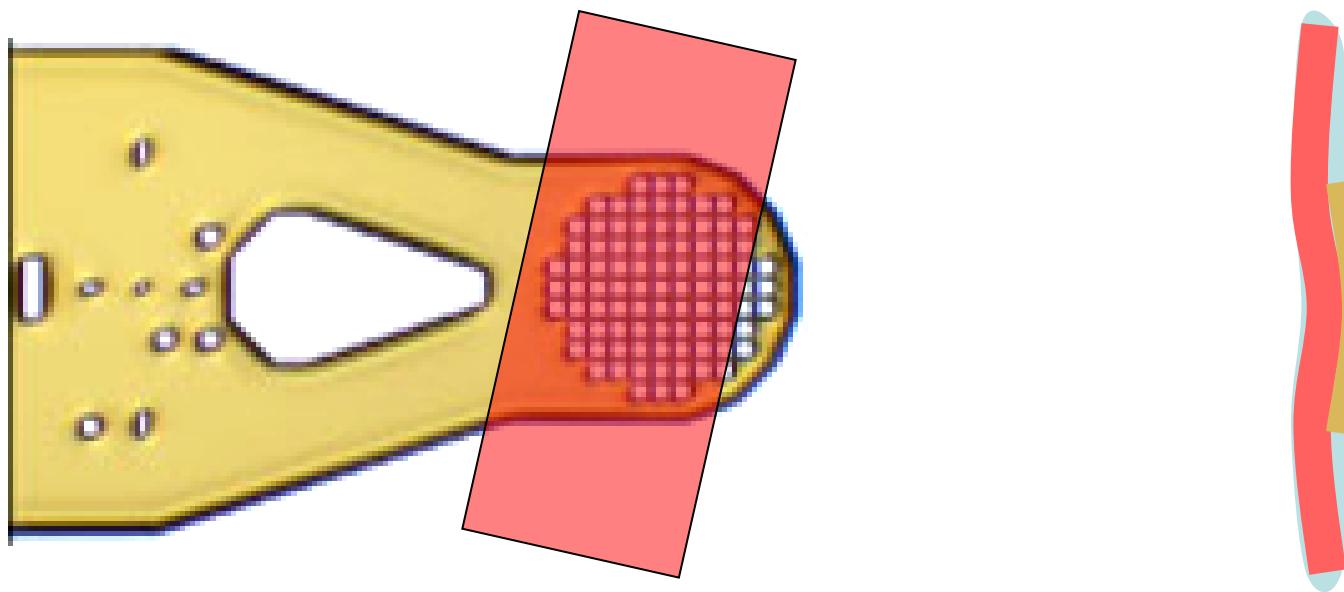


MiTeGen MicroMesh grids

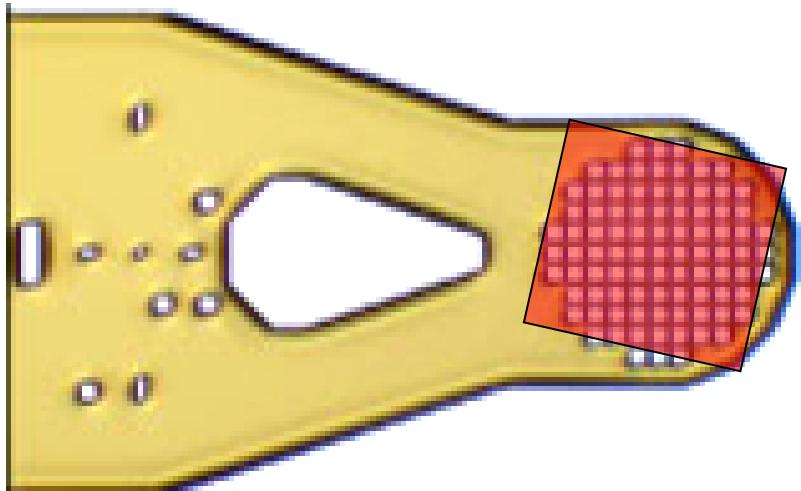


<http://www.mitegen.com/products/micromeshes/micromeshes.shtml>

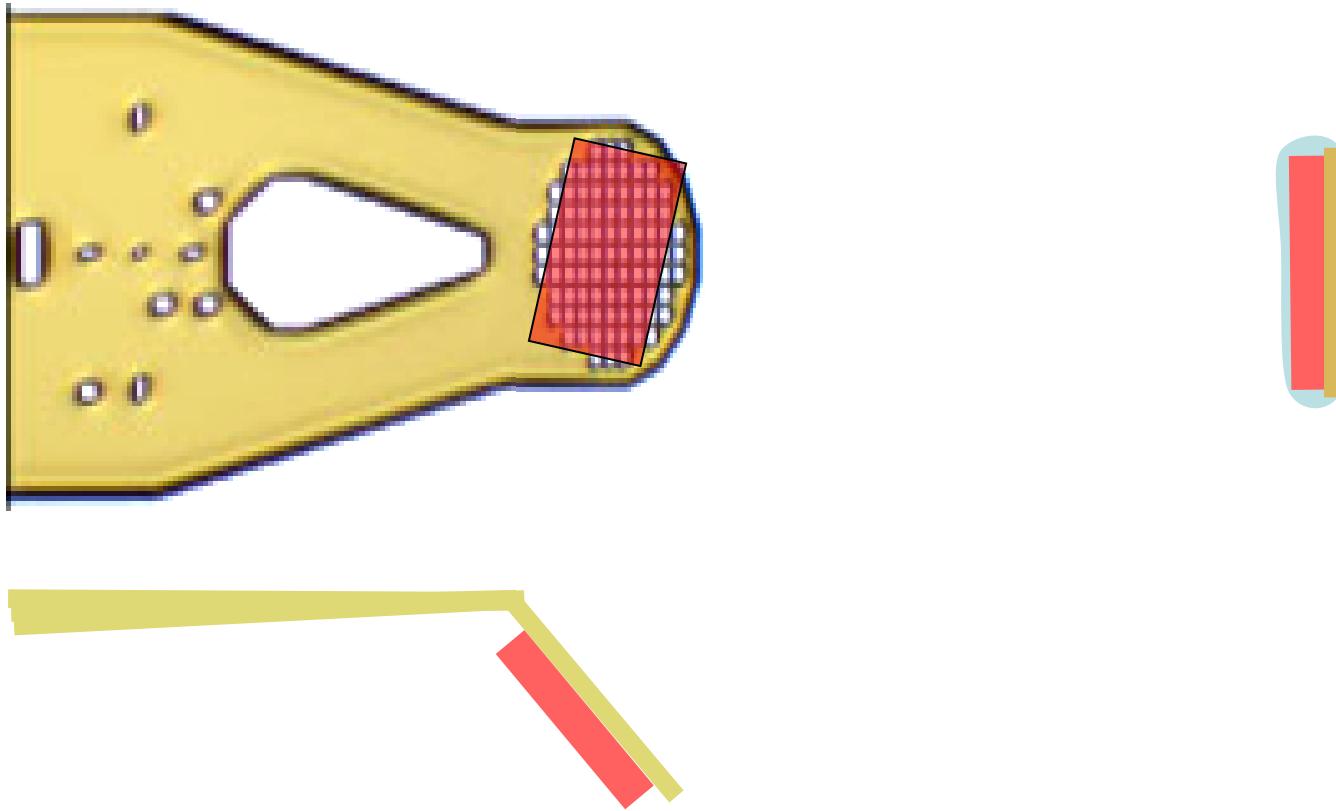
platy crystals



platy crystals



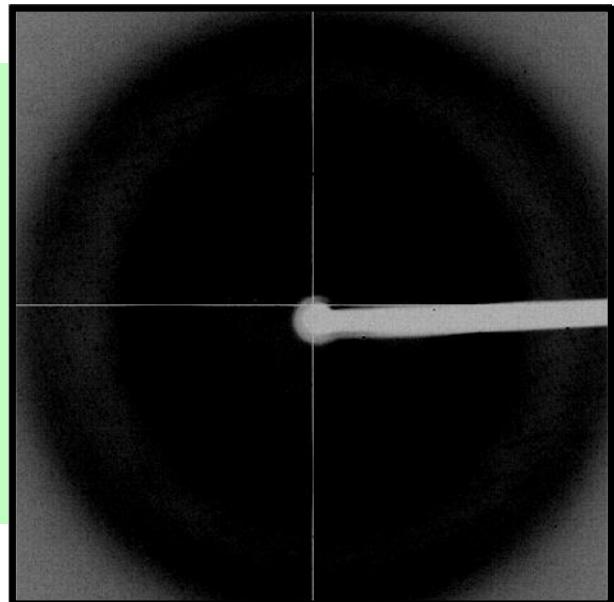
platy crystals



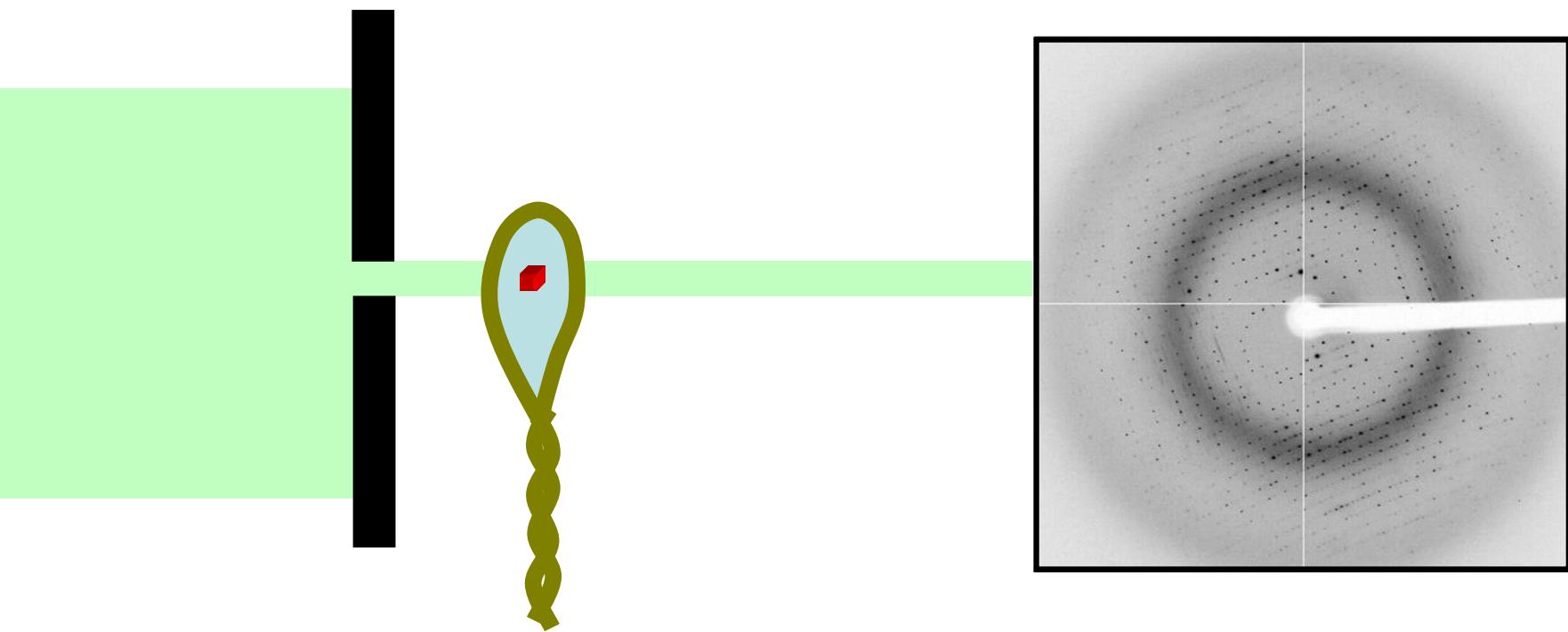
beam size vs xtal size

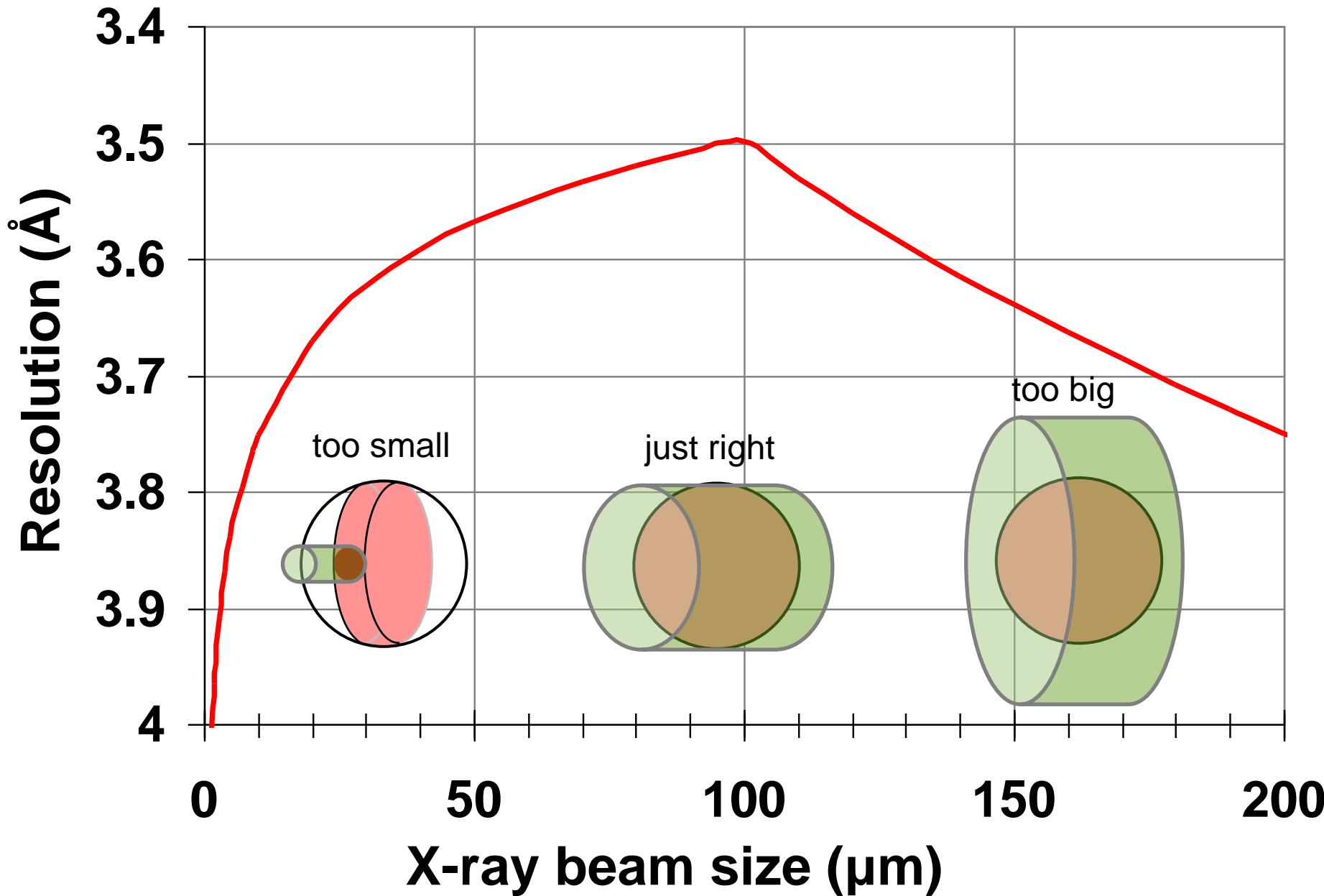
1. Put your crystal into the beam
2. Shoot the whole crystal
3. Shoot nothing but the crystal

shoot nothing but the crystal

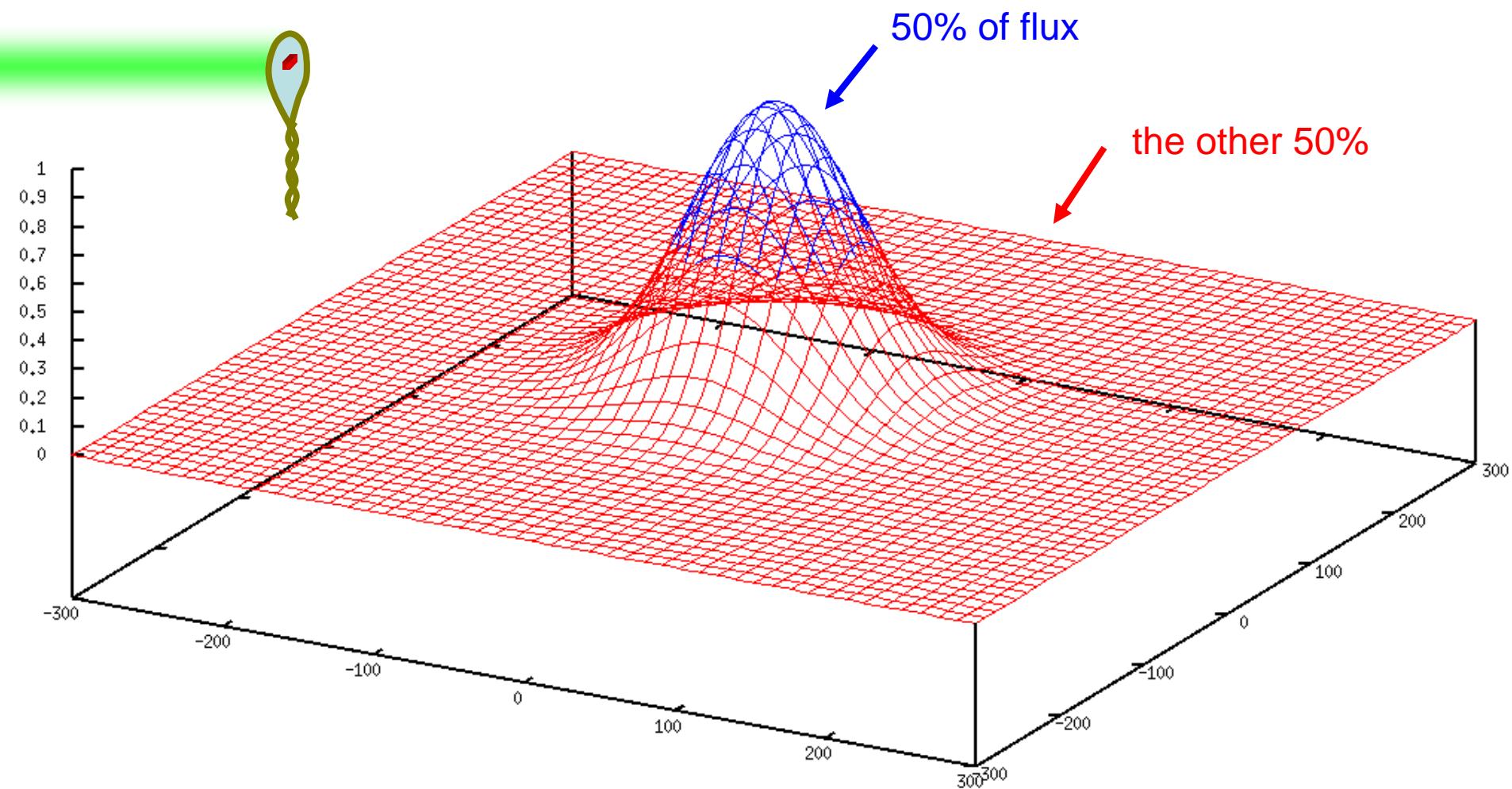


shoot nothing but the crystal

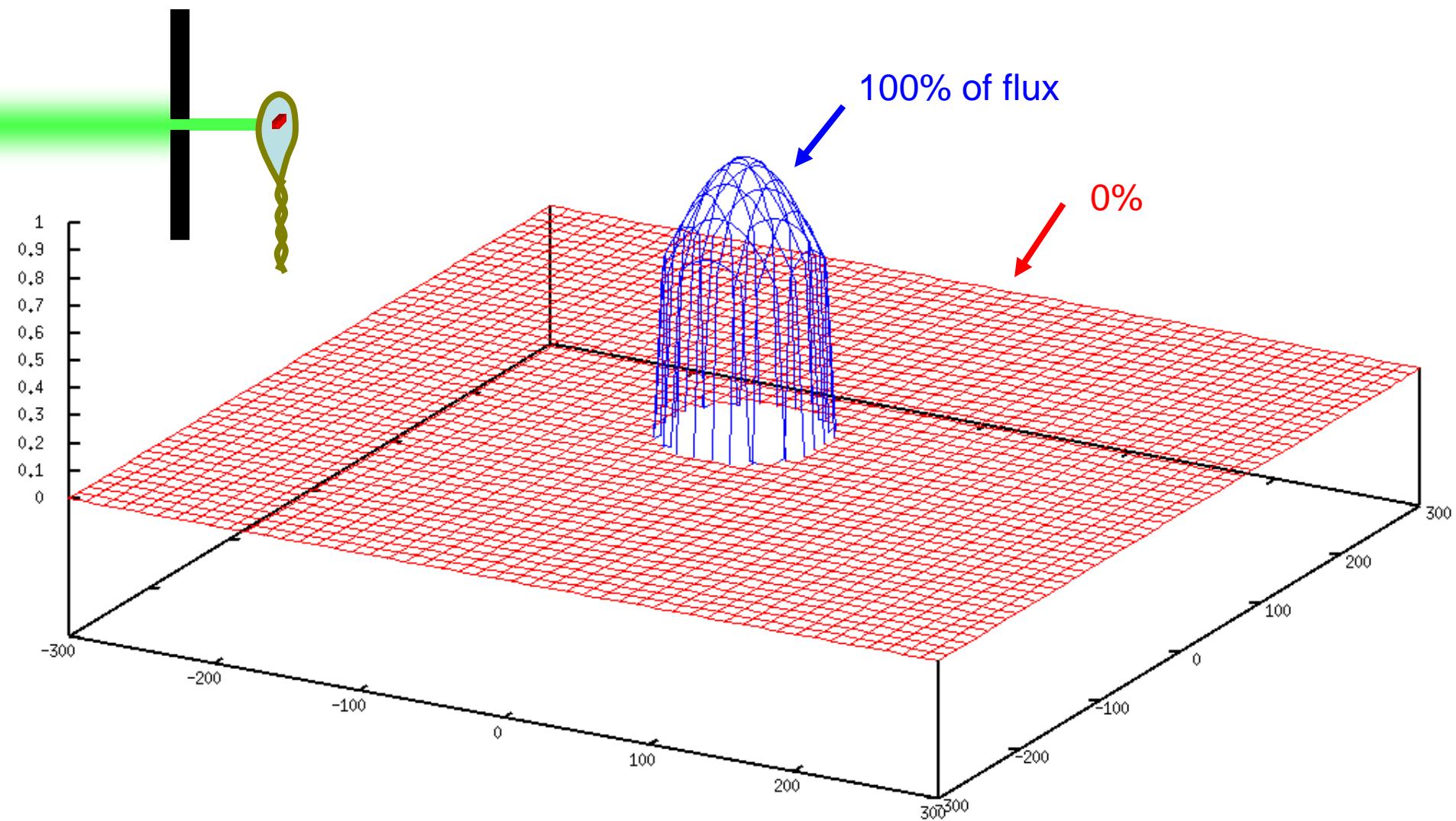




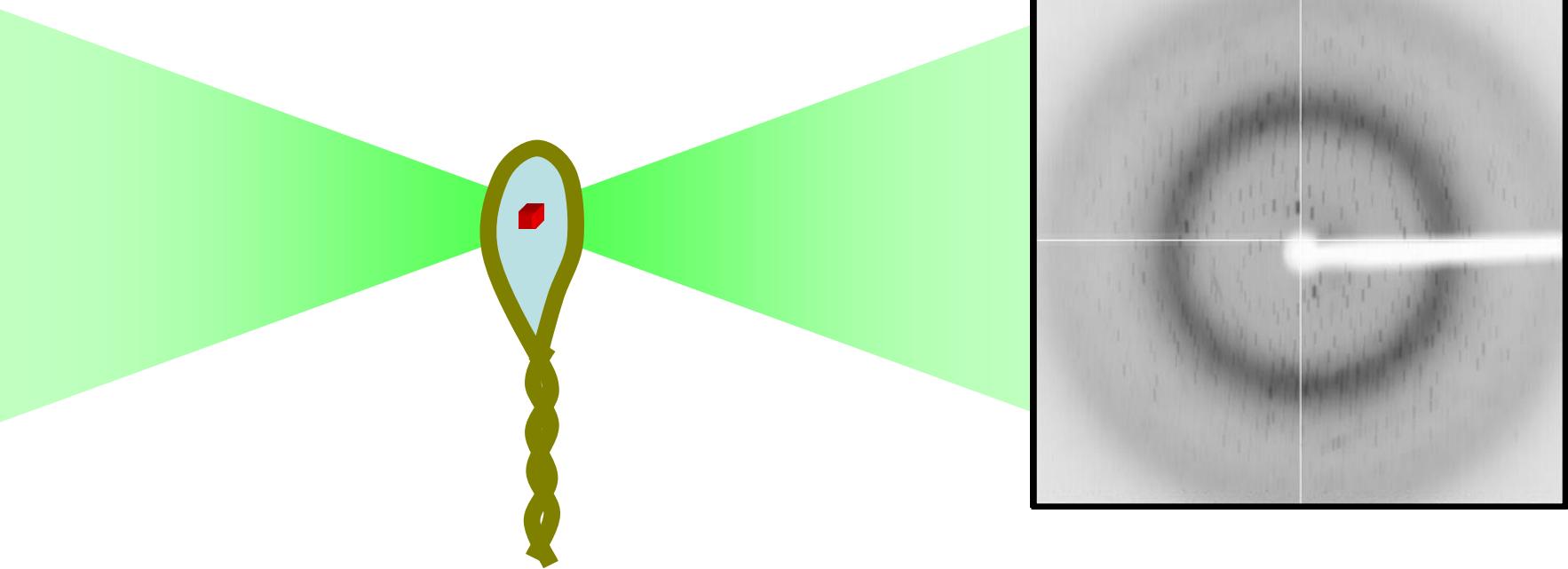
Gaussian beam



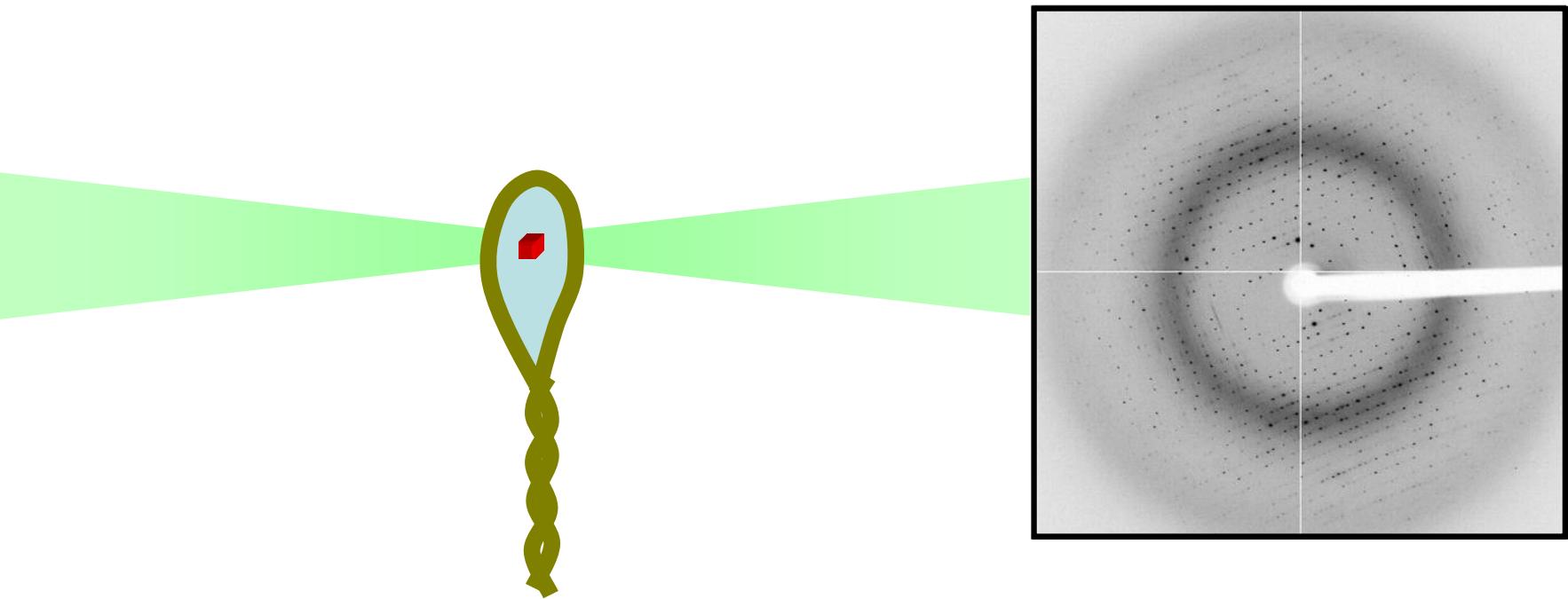
Collimated beam



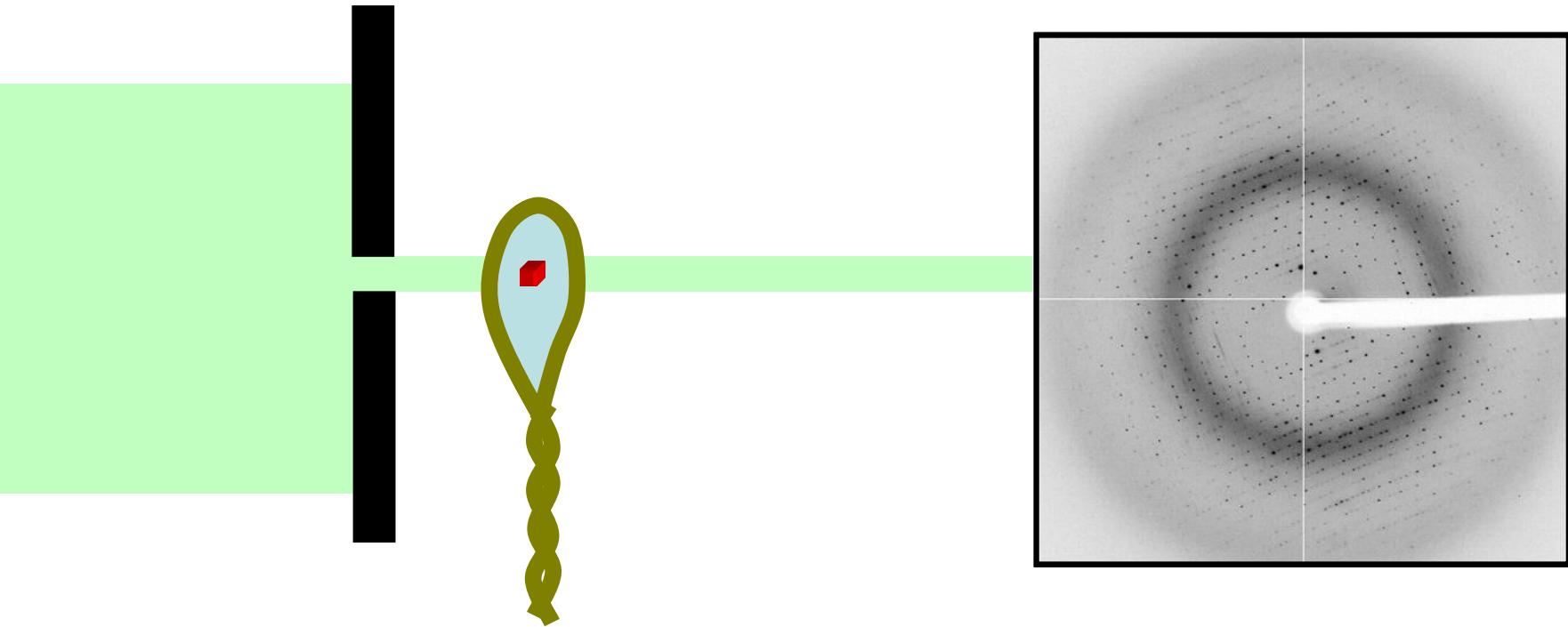
shoot nothing but the crystal



shoot nothing but the crystal



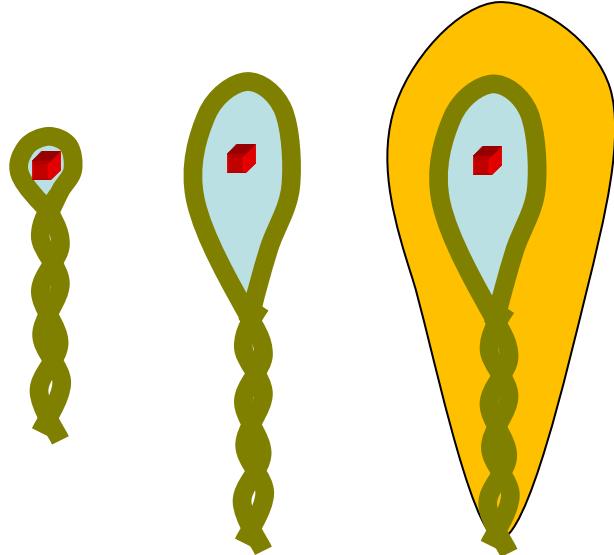
shoot nothing but the crystal



X-ray scattering “rules”:

Optimum: support thickness = crystal thickness

1 μm crystal



\approx 1 μm water

\approx 1 μm plastic

\approx 0.1 μm glass

\approx 1000 μm air

\sim 50 mm He

\approx 370 mm sat water vapor @ 4C

beam size vs xtal size

1. Put your crystal into the beam
2. Shoot the whole crystal
3. Shoot nothing but the crystal
4. Back off!

Mosquito tips cost \$0.15 each

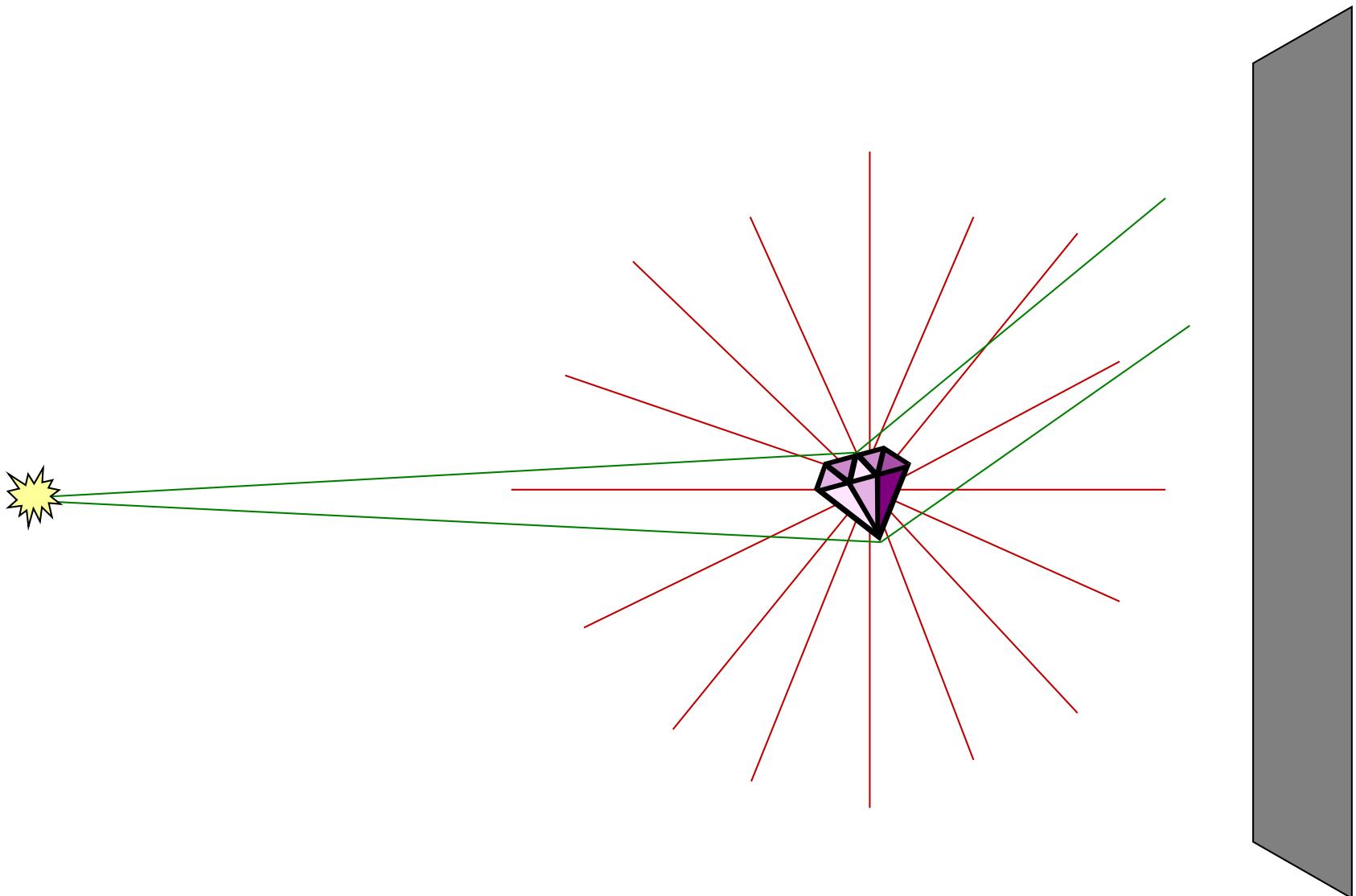
So do Pilatus pixels!

We bought 6,224,001

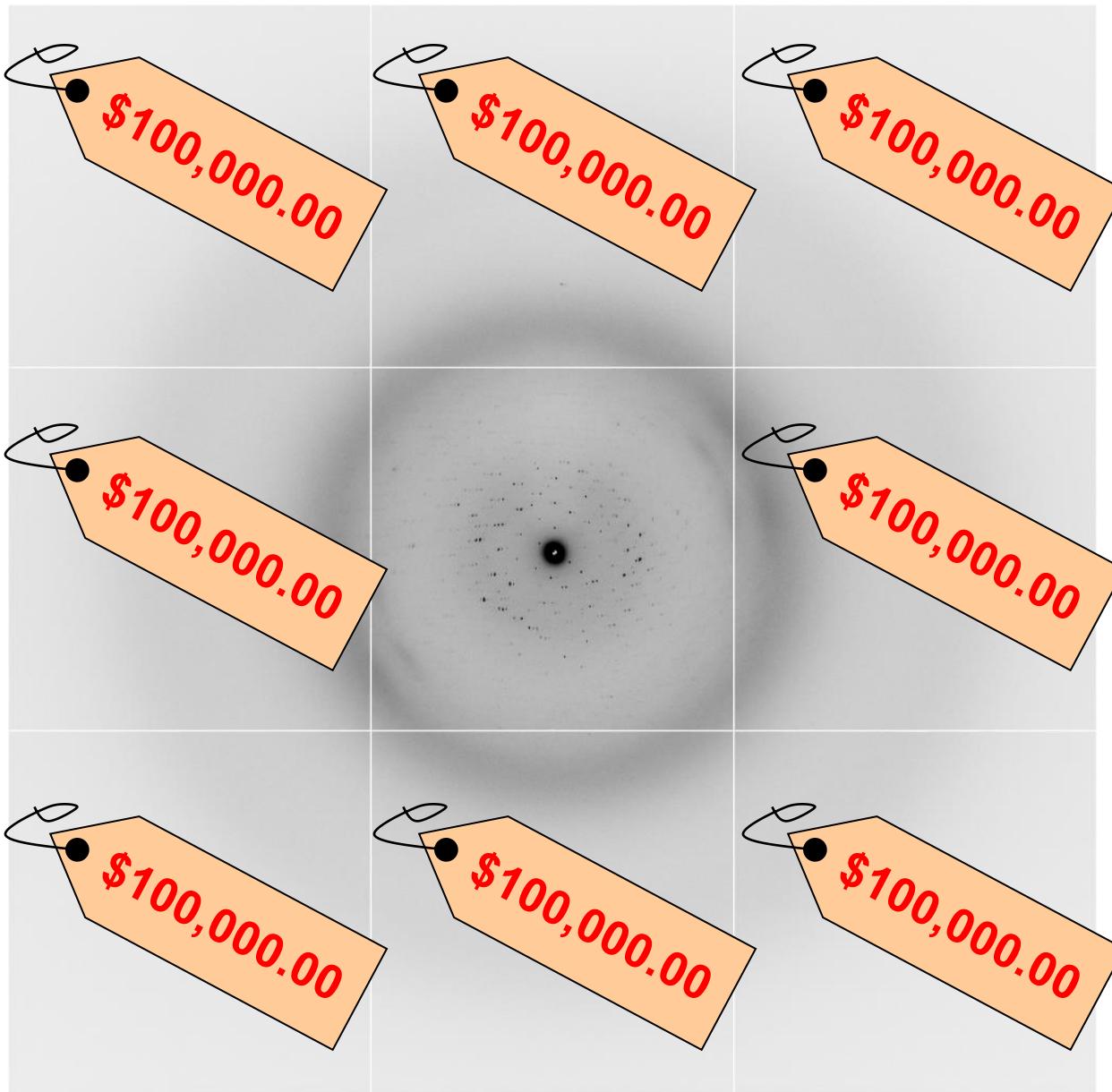
Use them!

(they are reusable)

Background scattering



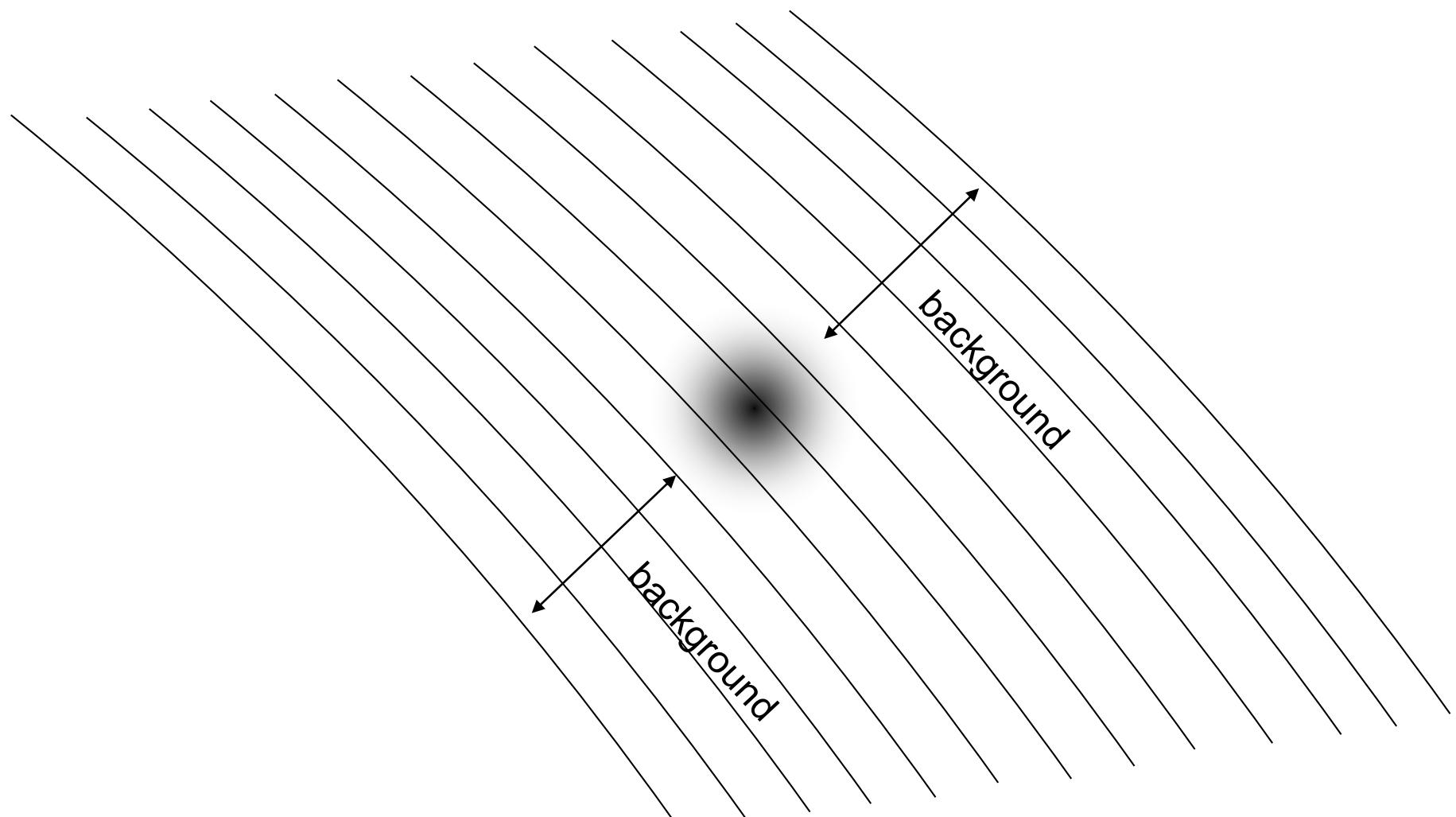
Background scattering



real estate is
expensive

use it!

Fine Slicing



Pflugrath, J. W. (1999). "The finer things in X-ray diffraction data collection", *Acta Cryst. D* 55, 1718-1725.

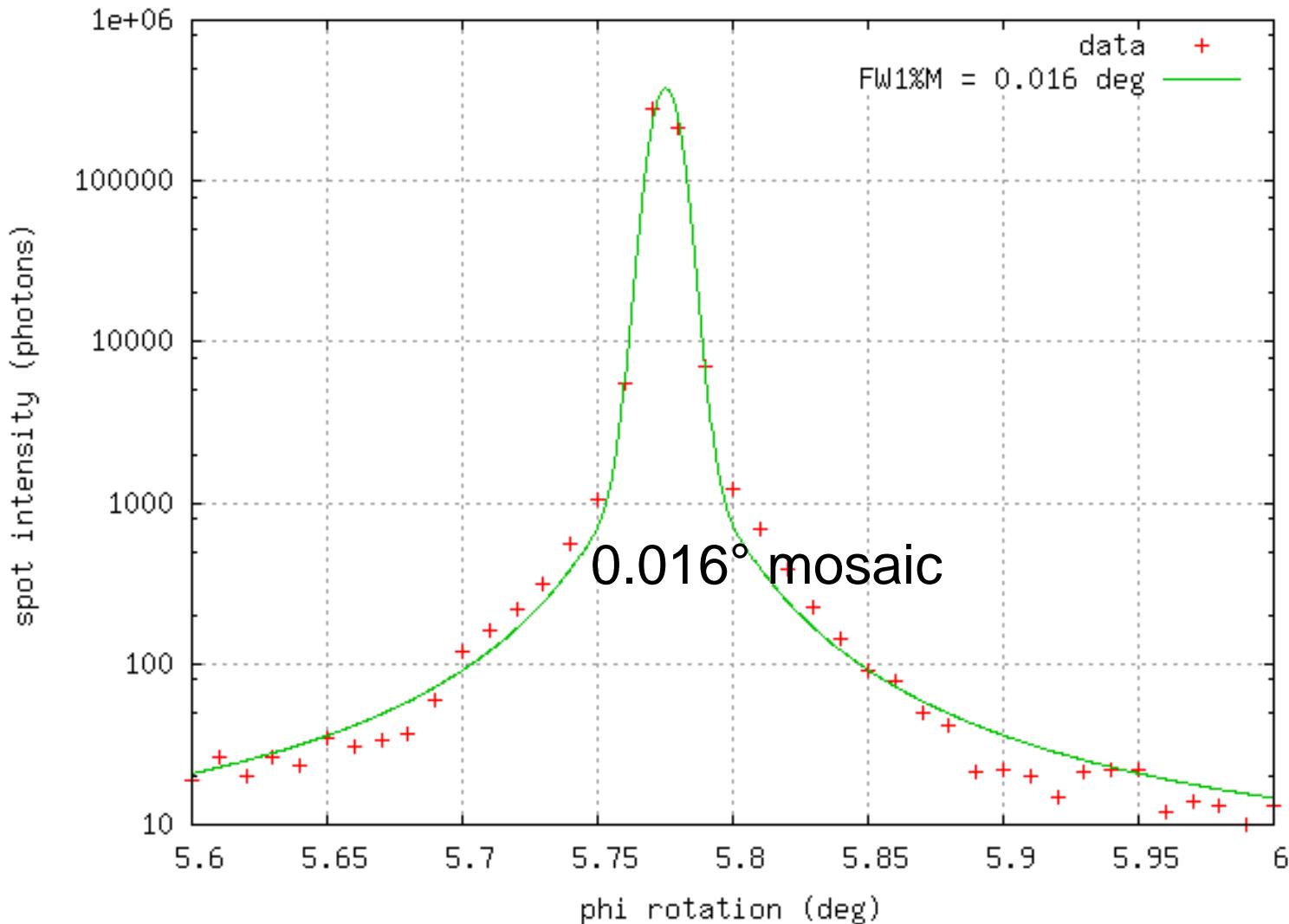
Pilatus pile-up for RT MX? same photons, different speeds

1	4	5	1	2	6	5	2	14	17	13	9	9	7	1	2	2	1	1	3	2			
1	1	2	2	4	3	4	3	29	30	25	22	20	12	5	7	2	0	0	3	3			
1	3	1	3	3	4	5	5	58	79	71	85	95	78	19	4	3	2	3	2	2			
4	6	3	2	3	5	6	6	44	419	685	747	890	1007	777	171	8	2	0	1	4	1		
1	3	8	4	11	7	38	724	12483	18639	20506	18841	16832	9766	1662	18	11	2	3	3	1			
4	4	2	6	13	16	74	3063	31927	26499	17788	10175	5929	2021	264	14	8	2	3	1	2			
3	1	1	1	3	9	12	130	1042	734	445	260	158	78	20	1	2	0	4	0	3			
1	1	2	1	1	4	8	17	102	93	65	65	55	28	10	2	3	2	1	2	3			
1	1	3	0	3	5	3	20	74	73	53	36	35	9	8	3	4	4	1	3	0			
0	0	2	2	4	4	8	7	23	20	10	10	9	2	4	4	0	3	1	1	3			
Sum of 193 shots with 193-fold attenuation										4	16	11	9	11	5	8	1	3	0	2	1	2	1

Pilatus pile-up for RT MX? same photons, different speeds

1	0	0	1	2	2	3	5	22	23	14	16	13	9	4	2	3	4	2	2	2	
1	2	2	2	4	1	2	10	37	38	37	31	31	18	4	5	2	4	1	1	2	
1	3	0	2	2	2	3	15	88	105	101	83	106	95	19	4	2	3	2	4	1	
4	3	1	1	2	9	14	60	585	930	980	1014	1115	893	226	5	2	0	1	2	3	
1	2	4	6	4	14	66	738	1680	1557	1398	1278	1306	1174	1052	31	9	7	7	5	2	
5	1	2	6	7	17	84	1198	1452	1340	1281	1305	1442	1094	211	13	6	2	4	4	5	
5	2	4	5	4	3	14	132	1112	831	480	255	147	71	14	5	4	6	3	3	0	
2	0	2	2	4	1	5	21	137	147	94	109	53	31	5	4	1	1	4	3	1	
0	3	2	2	5	6	6	17	108	104	64	48	39	12	5	3	4	2	4	0	0	
4	3	2	2	2	1	2	4	17	19	10	16	9	6	2	3	3	2	0	6	1	
1 shot with no attenuation											1	0	3	15	12	5	7	4	4	1	2

Pilatus pile-up for RT MX? same photons, different speeds



Classes of error in MX

Dependence on signal

none

sqrt

proportional

none

CCD
Read-out

Photon
counting

Detector calibration
attenuation
partiality
Non-isomorphism
Radiation damage

$1/\sqrt{t}$

Beam flicker

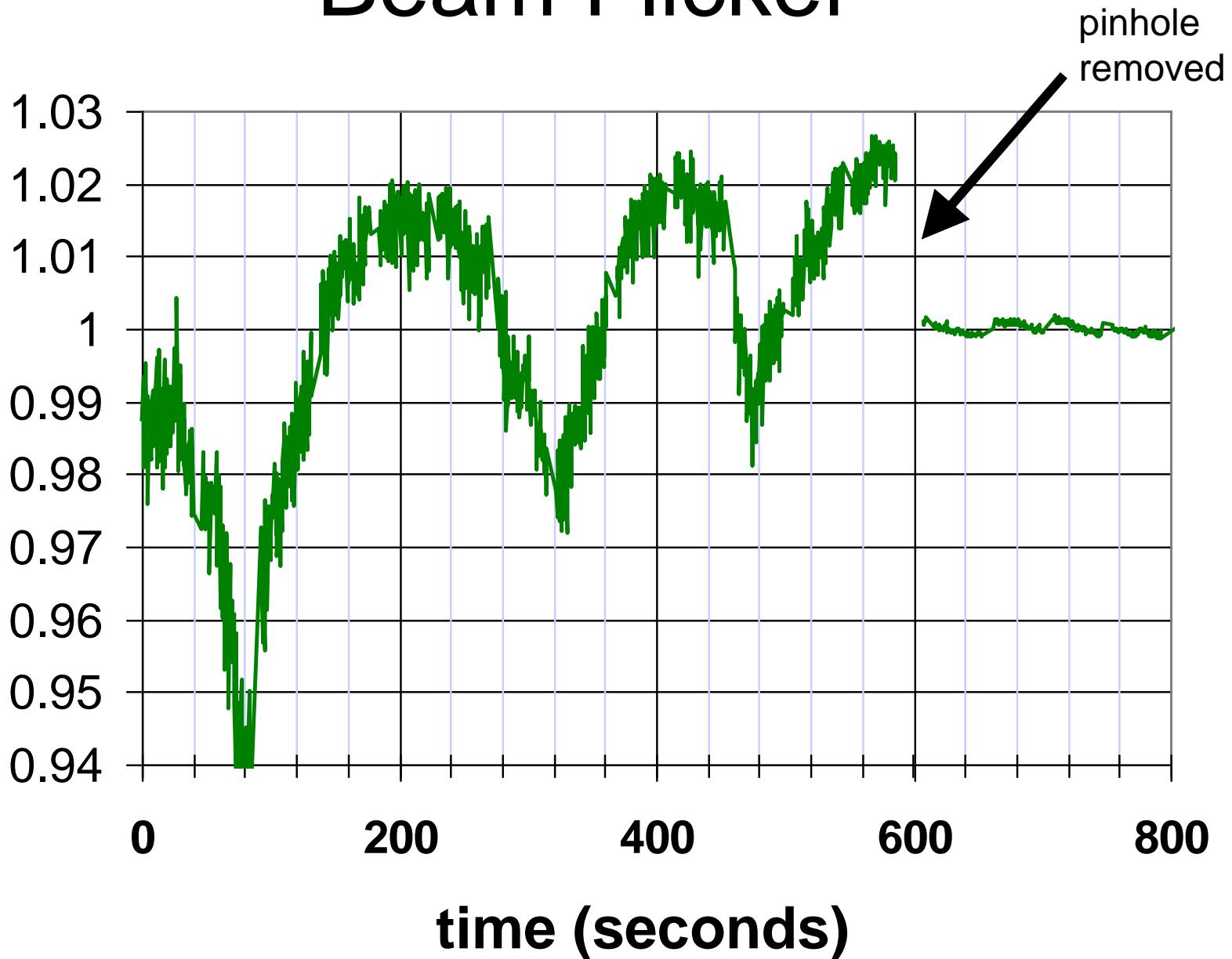
$1/t$

Shutter jitter
Sample vibration
Pile-up

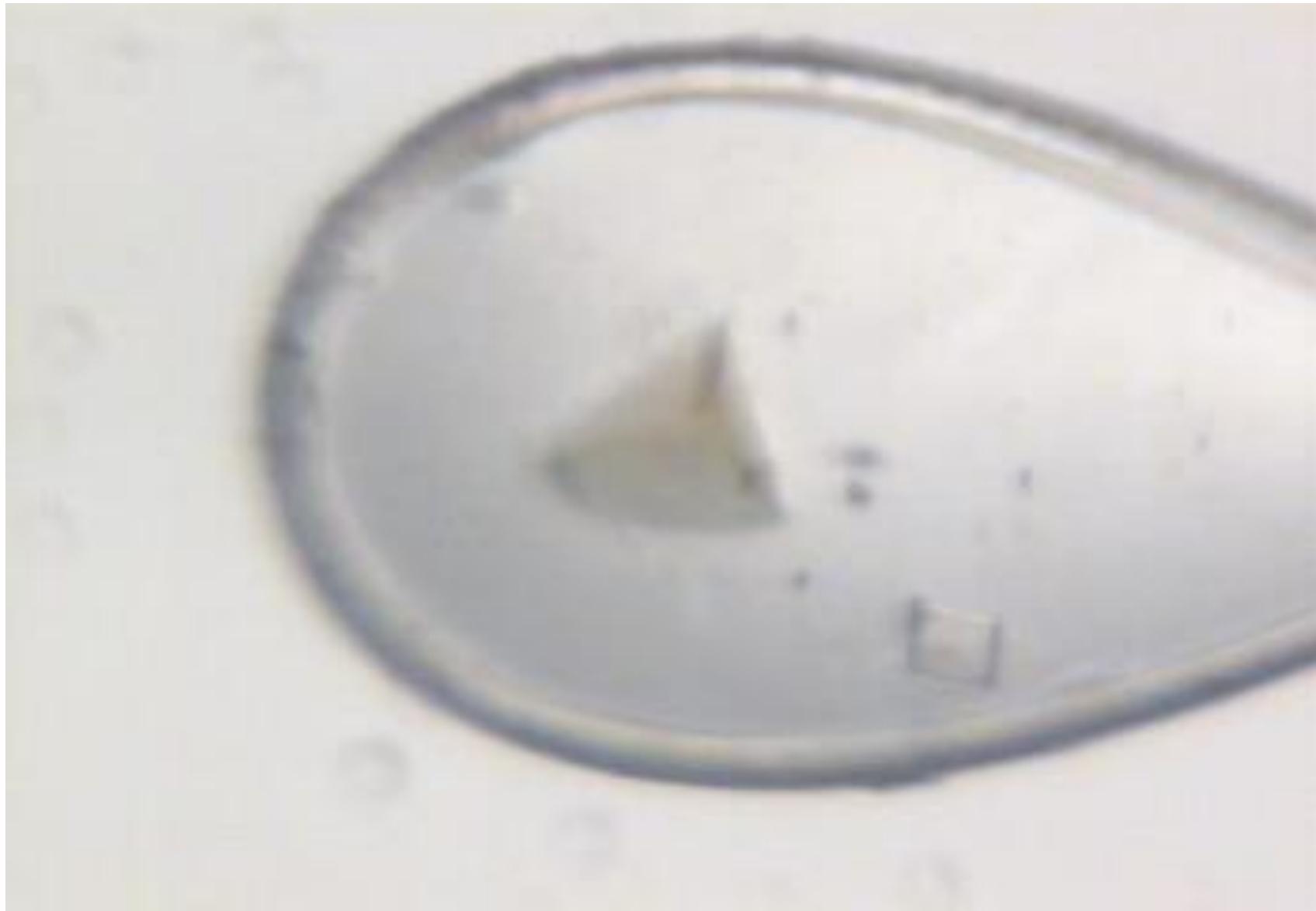
Time

Beam Flicker

normalized flux through pinhole



This is an Aperture!



Classes of error in MX

		Dependence on signal		
		none	sqrt	proportional
Time	none	CCD Read-out	Photon counting	Detector calibration attenuation partiality Non-isomorphism Radiation damage
	1/sqrt			Beam flicker
	1/prop.			Shutter jitter Sample vibration Pile-up

Optimal exposure time

(faint spots on CCD)

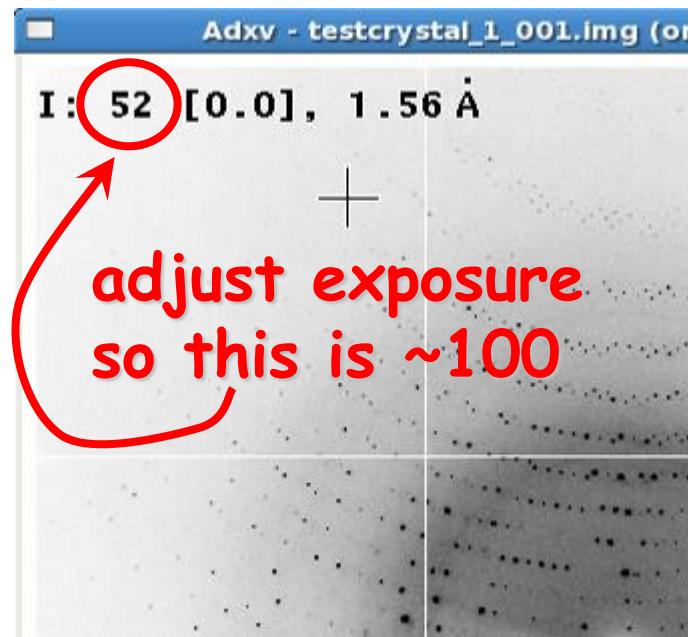
$$t_{hr} = t_{ref} \frac{10 \cdot m \cdot \sigma_0^2}{gain \cdot (bg_{ref} - bg_0)}$$

t_{hr}
 t_{ref}
 bg_{ref}

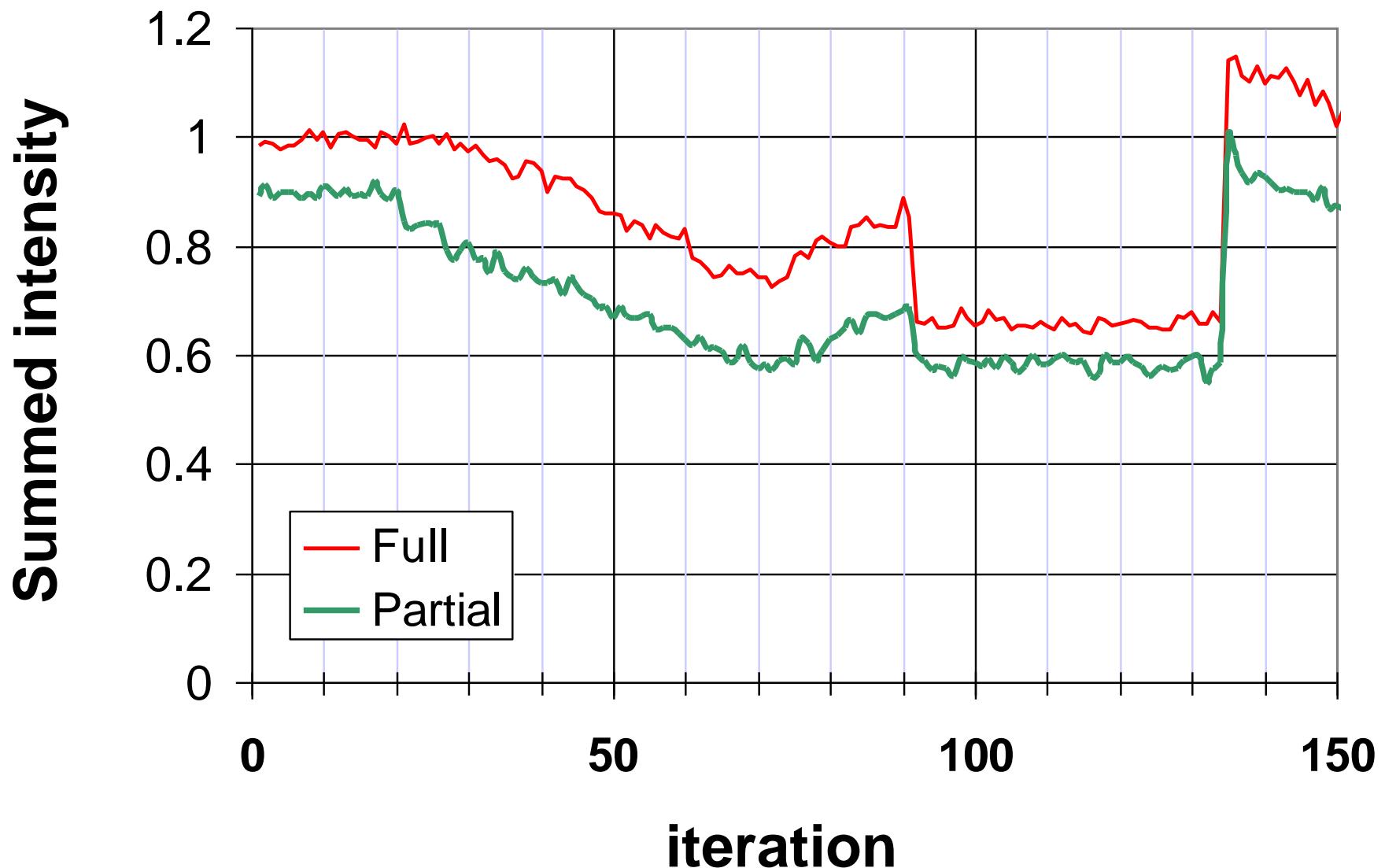
Optimal exposure time for data set (s)
exposure time of reference image (s)
background level near weak spots on
reference image (ADU)

bg_0
 bg_{hr}
 σ_0
 $gain$
 m

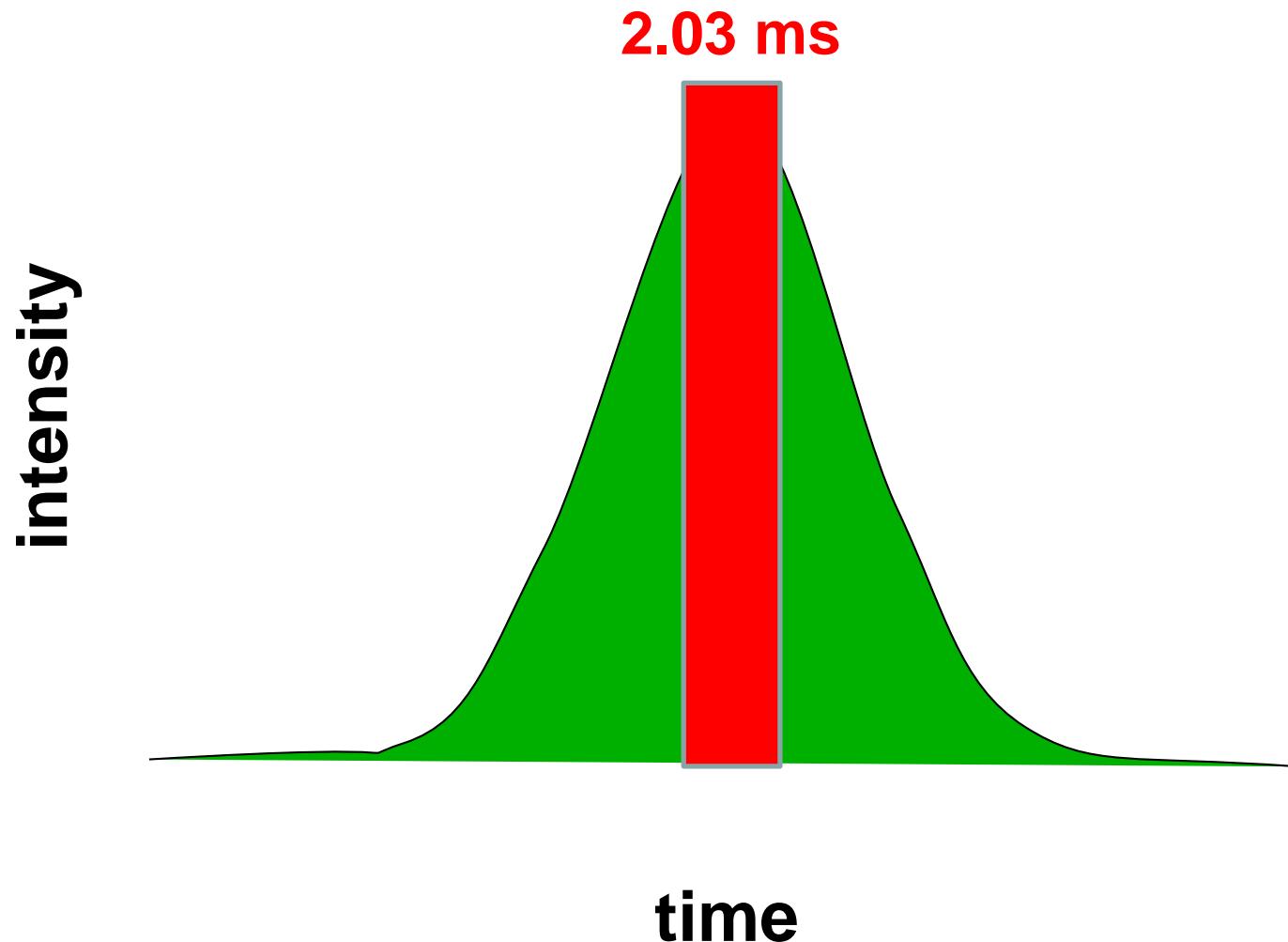
ADC offset of detector (ADU)
optimal background level (via t_{hr})
rms read-out noise (ADU)
ADU/photon
multiplicity of data set (including partials)



PAD full vs partial



PAD full vs partial



Optimal exposure time

(faint spots on PAD)

$$t_{hr} > t_{ro} / 5\%$$

- t_{hr} Optimal exposure time (s)
read-out time (s)
2.03 ms on PILATUS3 S
3 μ s on EIGER

Optimal exposure time

(anomalous on PAD)

$$t_{ano} > t_{ro} \frac{\langle F \rangle}{\langle \Delta F \rangle}$$

t_{ano}	Optimal exposure time (s)	0.2 s
t_{ro}	read-out time (s)	
	2.03 ms on PILATUS3 S	
	3 μ s on EIGER	

Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

How to avoid
time-dependent error?

Attenu-wait!

Apply 10x attenuation

Expose 10x longer

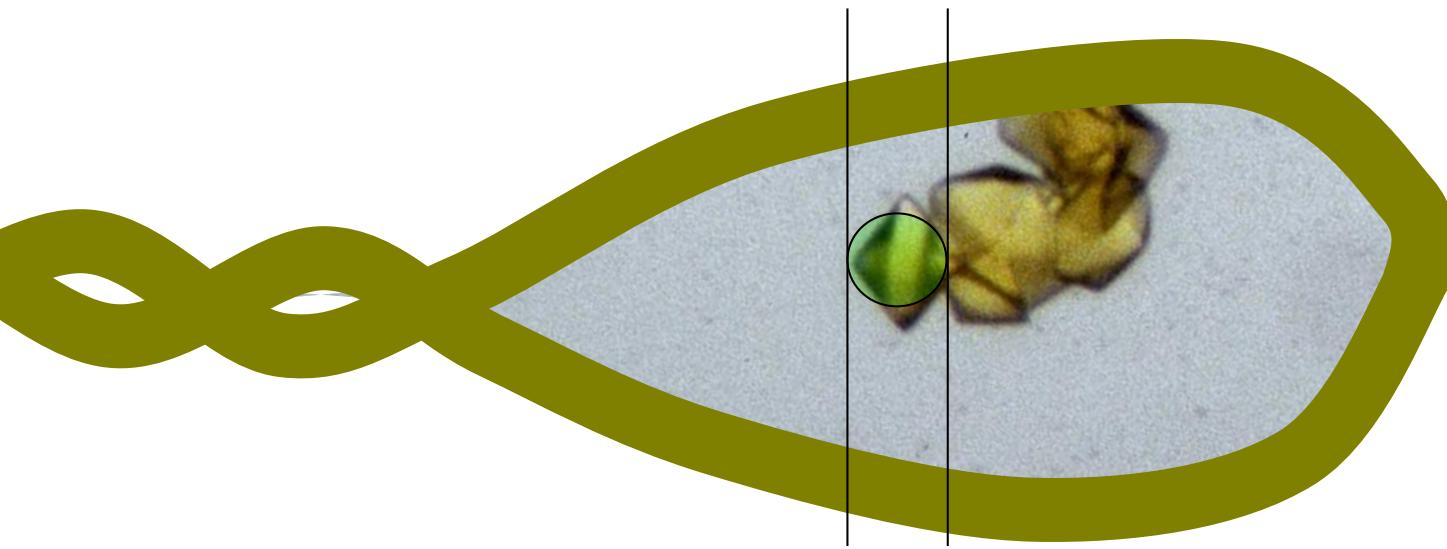
Number of photons should be the same!

Damage will also be the same!

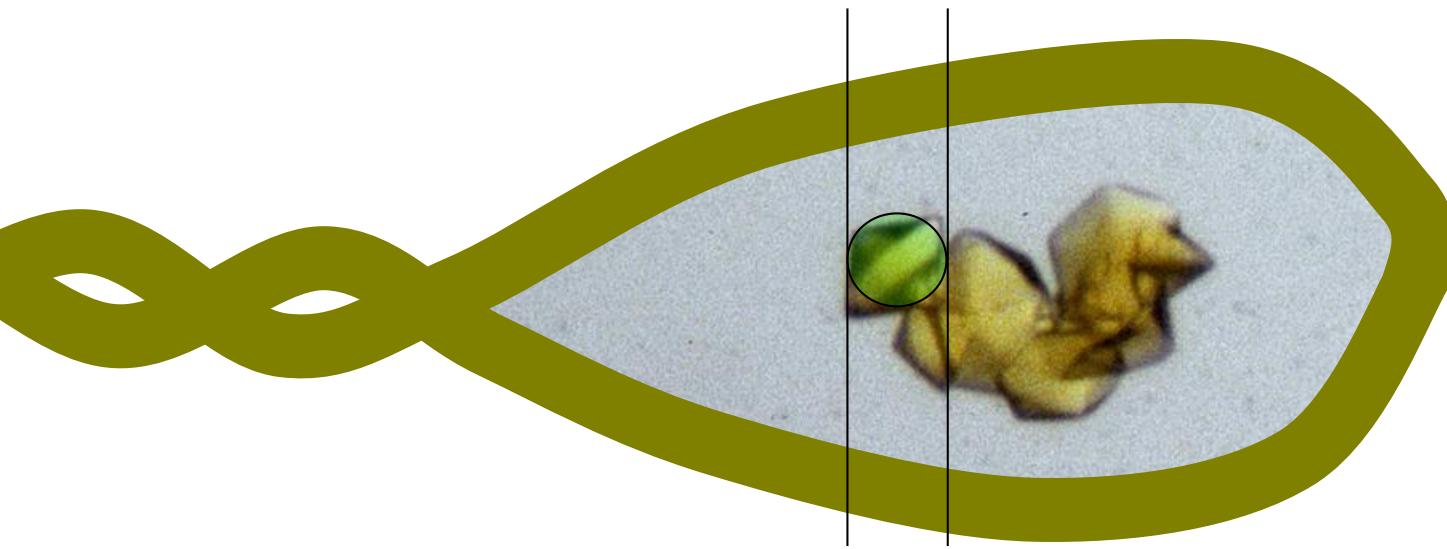
beam size vs xtal size

1. Put your crystal into the beam
2. Shoot the whole crystal
3. Shoot nothing but the crystal
4. Back off!
5. The crystal must rotate

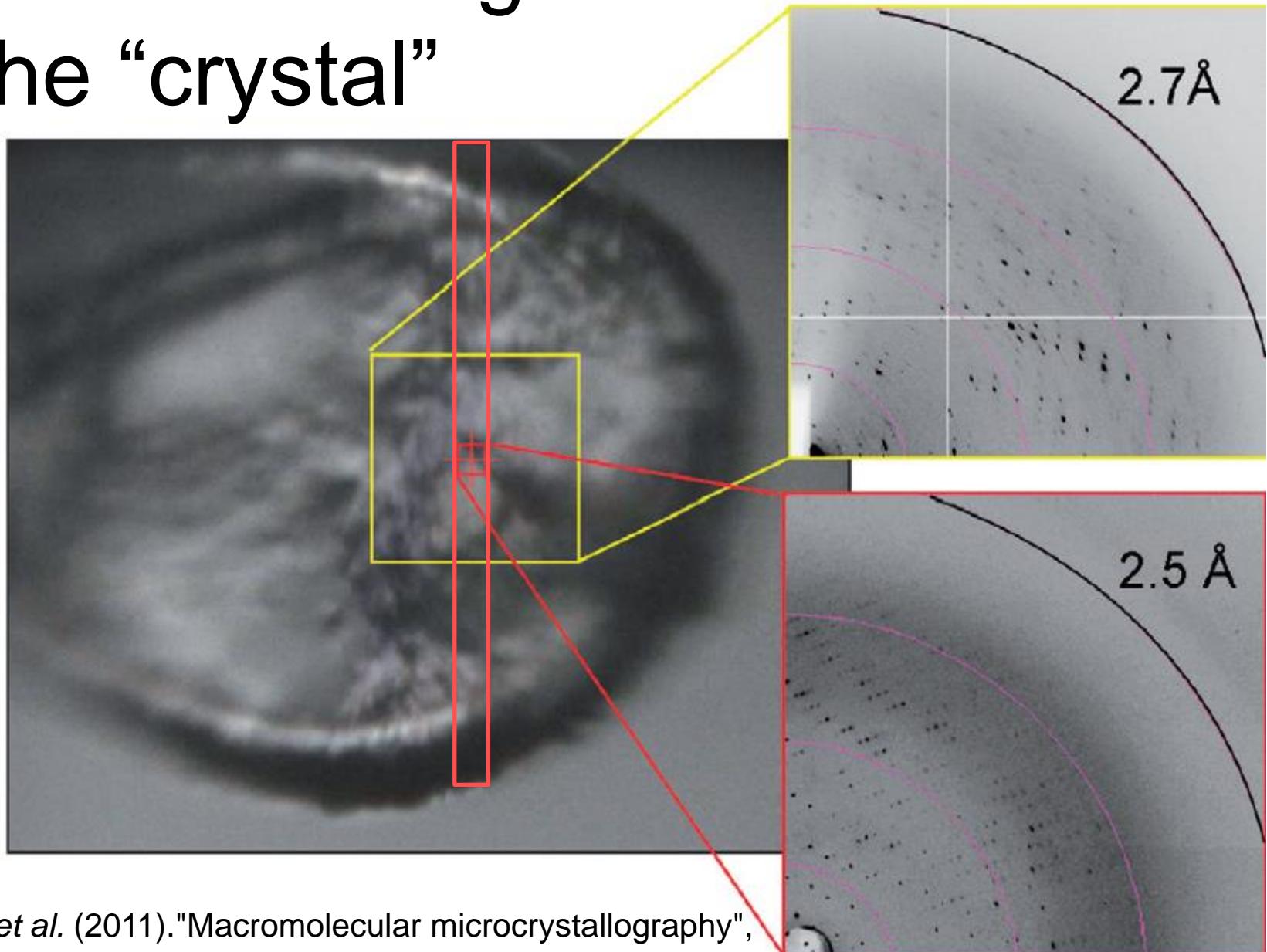
The crystal rotates!



The crystal rotates!



Shoot nothing but the “crystal”



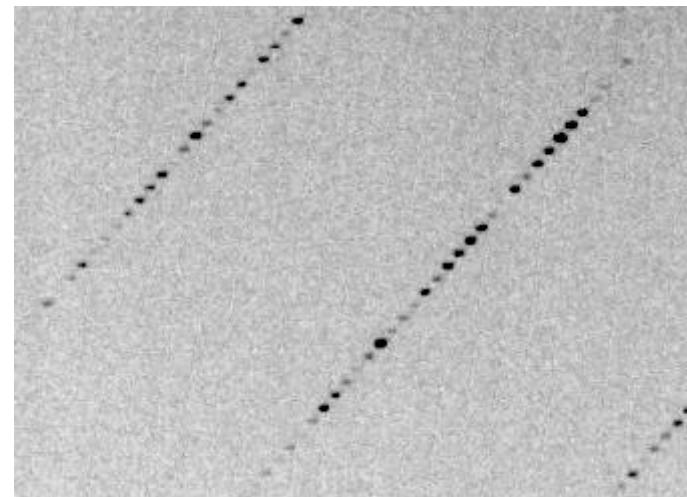
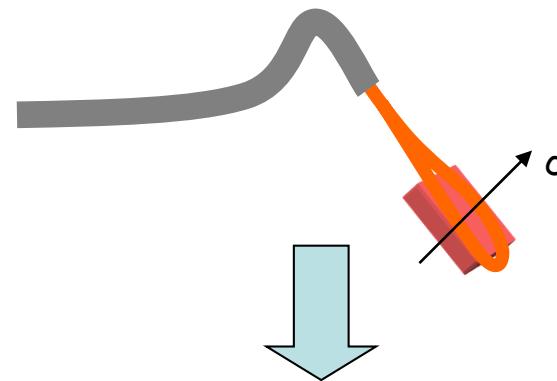
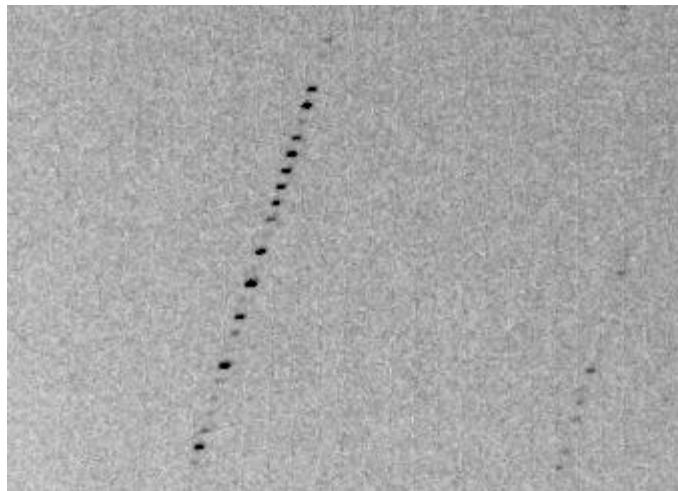
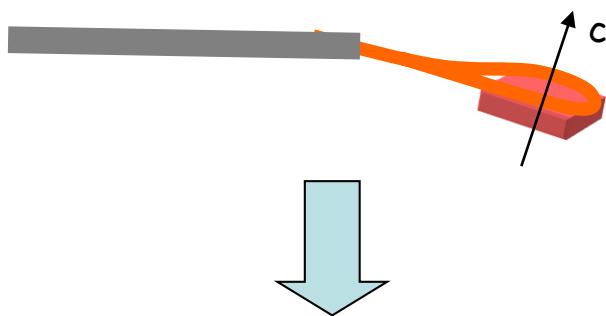
Evans et al. (2011). "Macromolecular microcrystallography",
Crystallography Reviews 17, 105-142.

Decisions, Decisions, Decisions

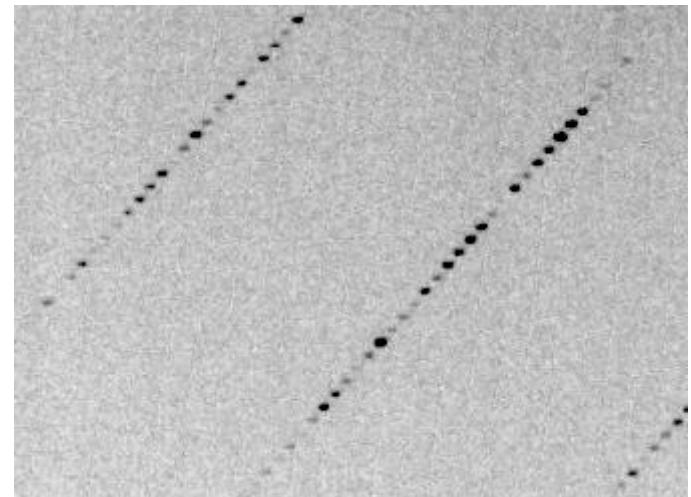
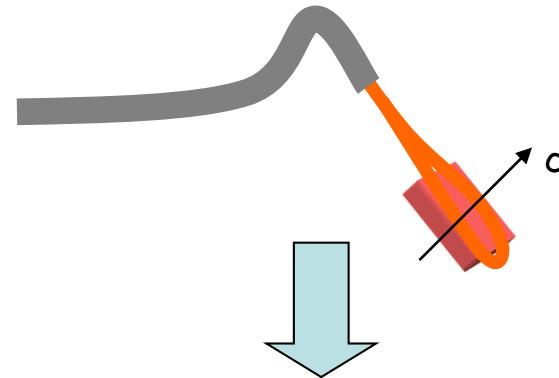
- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

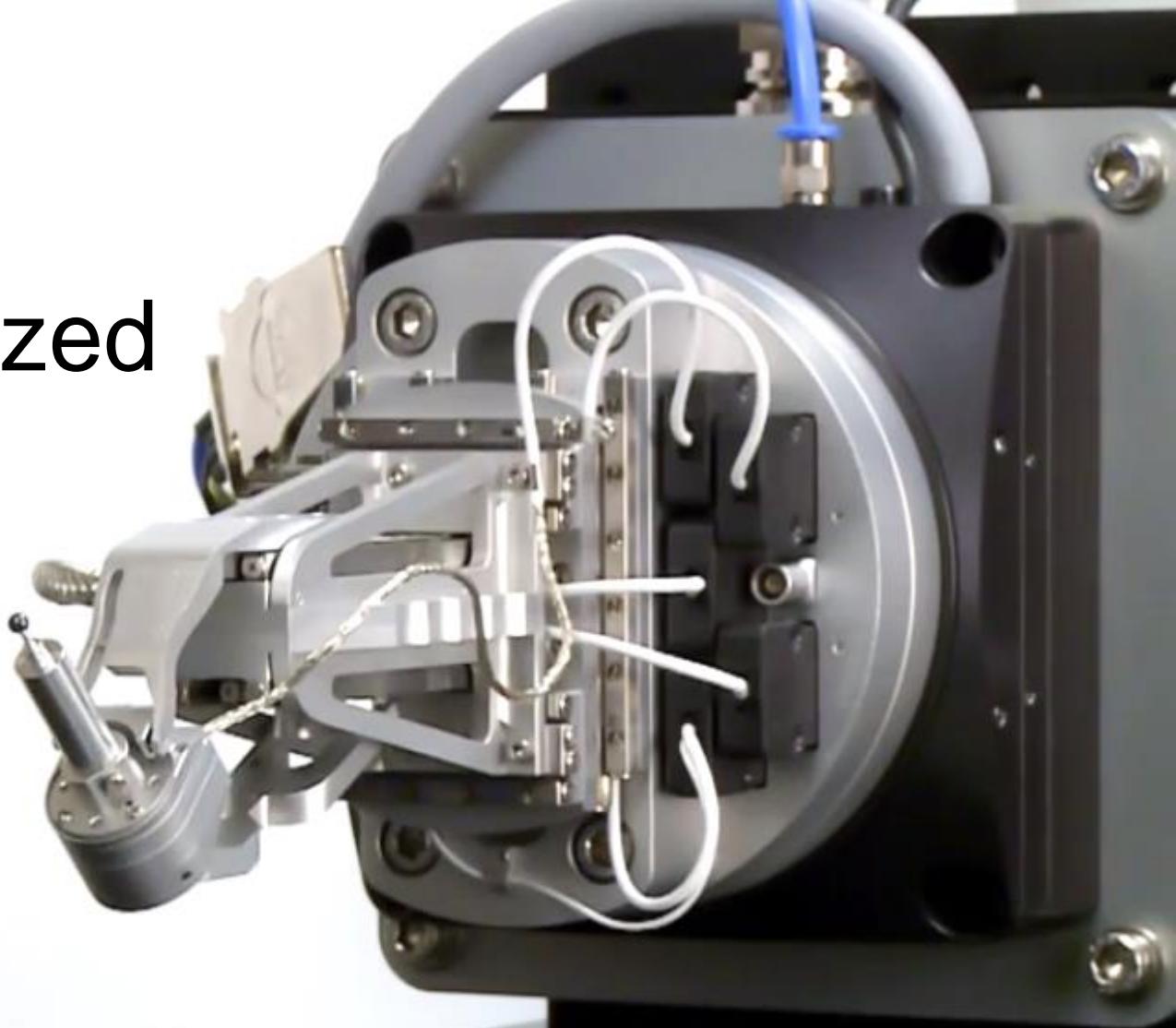
avoiding overlaps



avoiding overlaps



SmarAct's “SmarGon” Commercialized PRIGo



At the beamline...

- Resolution
 - problem: background
 - solution: use as few pixels as possible
- Phases
 - problem: fractional errors
 - solution: use as many pixels as possible

R-factor

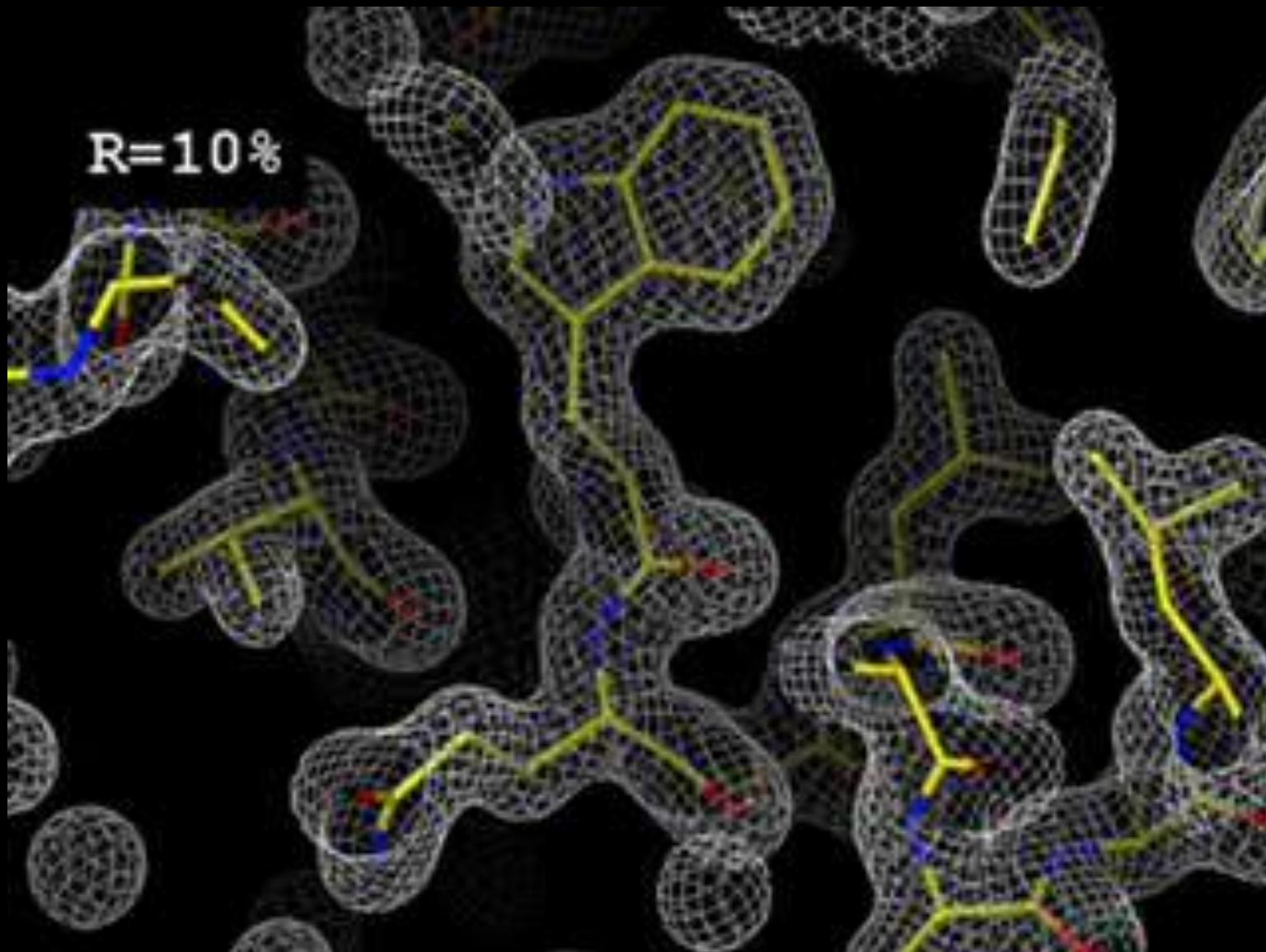
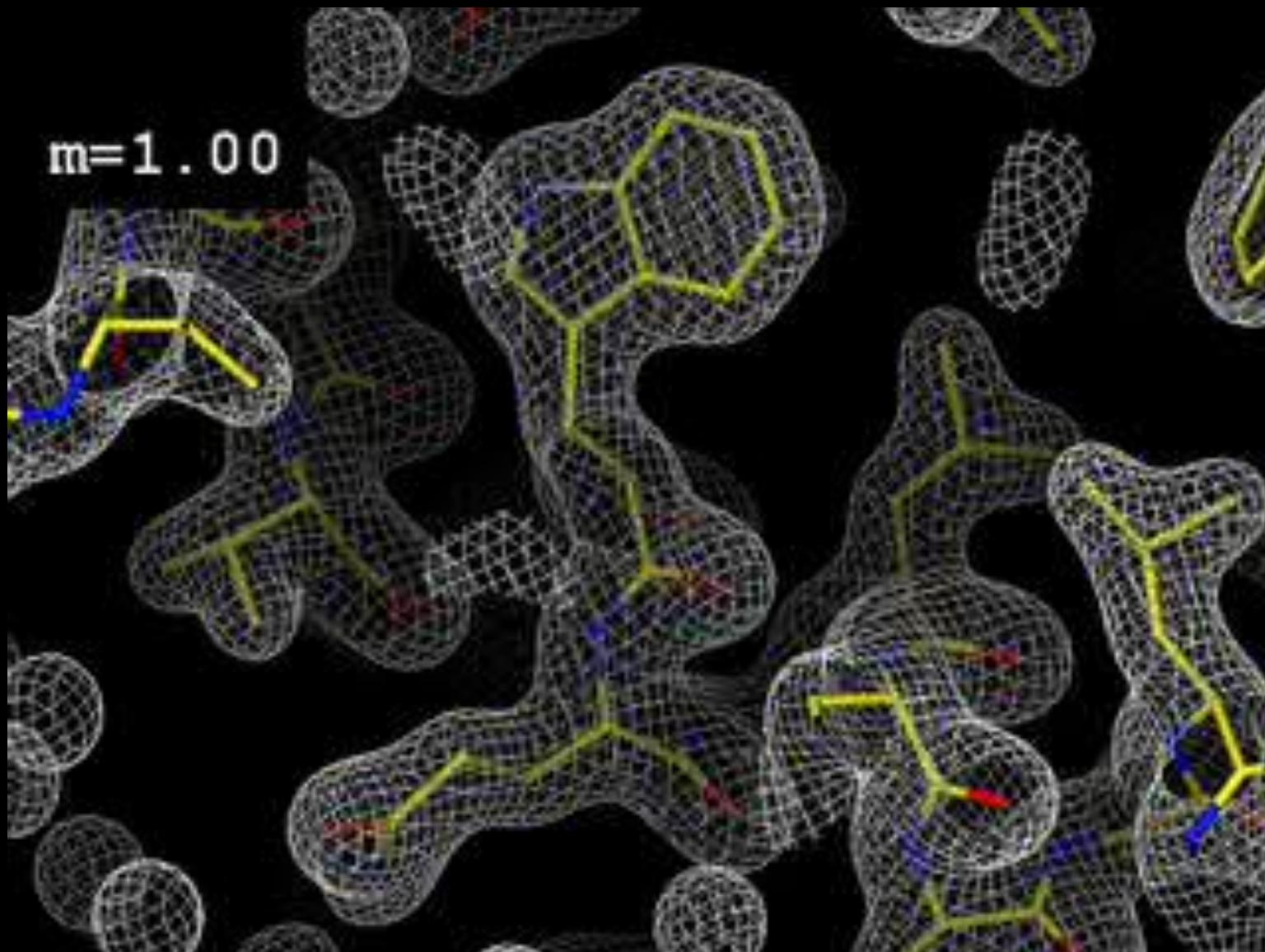


Figure of Merit



anomalous signal

$$\frac{\Delta F}{F} \approx 1.2 f'' \sqrt{\frac{\# \text{ sites}}{\text{MW (Da)}}}$$

World record!
 $\Delta F/F = 0.5\%$

Wang, Dauter &
Dauter (2006)
Acta Cryst. D
62, 1475-1483.

Crick, F. H. C. & Magdoff, B. S. (1956) *Acta Crystallogr.* **9**, 901-908.
Hendrickson, W. A. & Teeter, M. M. (1981) *Nature* **290**, 107-113.

Fractional error

$$\text{mult} > \left(\frac{\sim 3\%}{\langle \Delta F/F \rangle} \right)^2$$

required number of crystals calculator - Mozilla Firefox

File Edit View History Bookmarks Tools Help

<http://bl831.als.lbl.gov/xtalsize.html>

wikipedia (en)

required number of crystals calculator

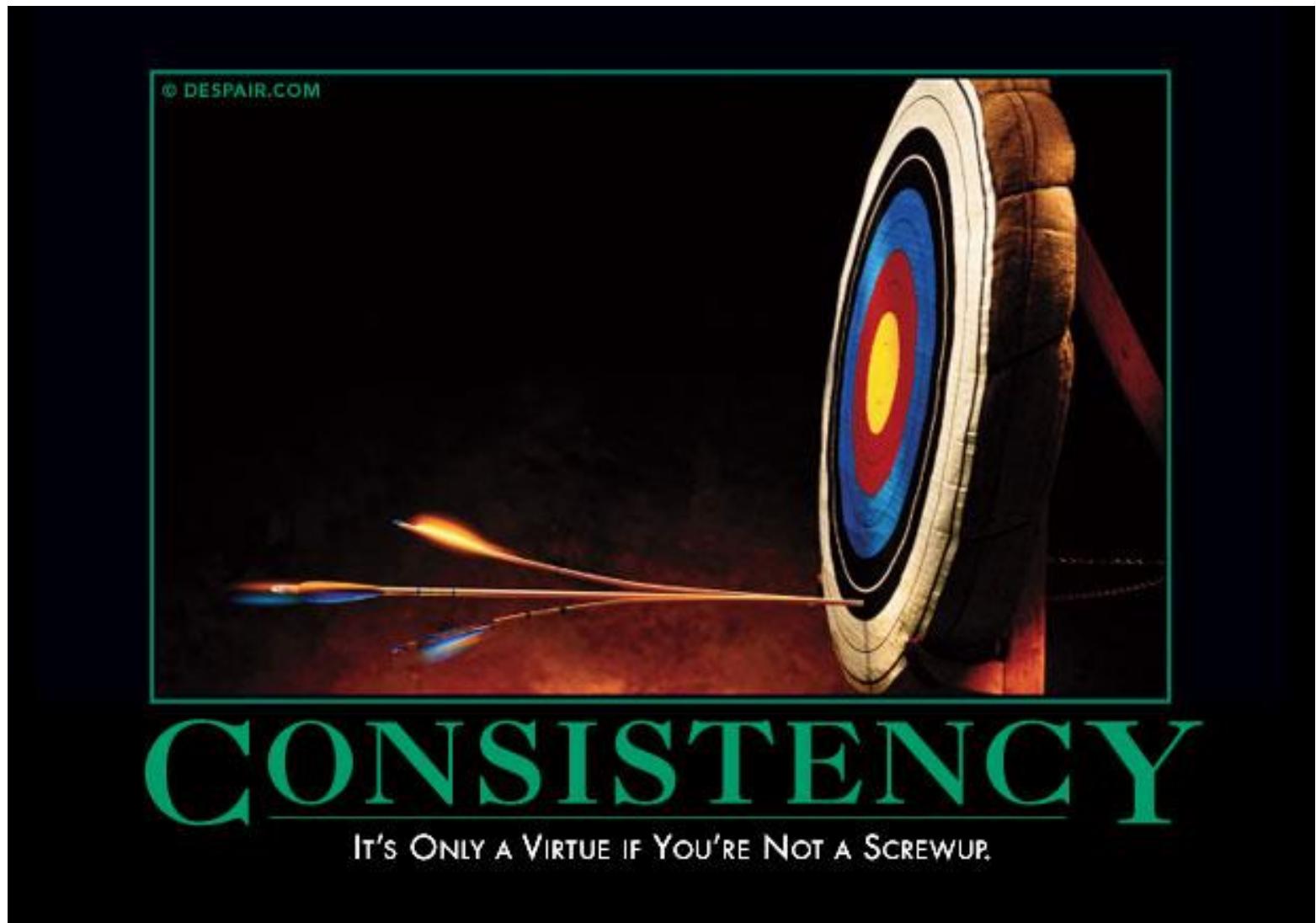
Required crystal number or size calculator

$$n_{\text{xtals}} = \langle I_{\text{DL}} \rangle / 20 * f_{\text{NH}} * \text{MW} * V_M^2 / \exp(-0.5 * B/\text{reso}^2) / \text{xtalsize}^3 / (\text{reso}^3 - 1.53)$$

Enter values:

experiment goal =	subtle differences (MAD/SAD)		
number of sites =	1	in asymmetric unit	
fpp =	4	electrons	Bijvoet ratio = 1.75 %
molecular weight =	30	kDa in asymmetric unit	
resolution =	3.4	Ang	signal to noise = 81 at this resolution
reso on snapshot =	2.4	Ang	→ Wilson B = 35 Ang ²
background level =	100	ADU/pixel	multiplicity = 7.3
spot size =	5	pixels	
detector type =	ADSC Q210/315r (hwbin)		
solvent content =	50	%	
xtal size _{beam} =	20	microns	beam size _{vert} = 100 microns
xtal size _{vert} =	20	microns	beam size _{spindle} = 100 microns
xtal size _{spindle} =	20	microns	
<input type="button" value="Calculate n_xtals"/> ↓	<input type="button" value="Calculate size"/> ↑		
n _{xtals} =	1.4	xtals you will need to merge	← <I _{DL} > 11000 photons/hkl

Systematic error does not “average out”



Can you count to 1,000,000 ?

Theoretically: $\frac{\sqrt{1,000,000}}{1,000,000} = 0.1\%$

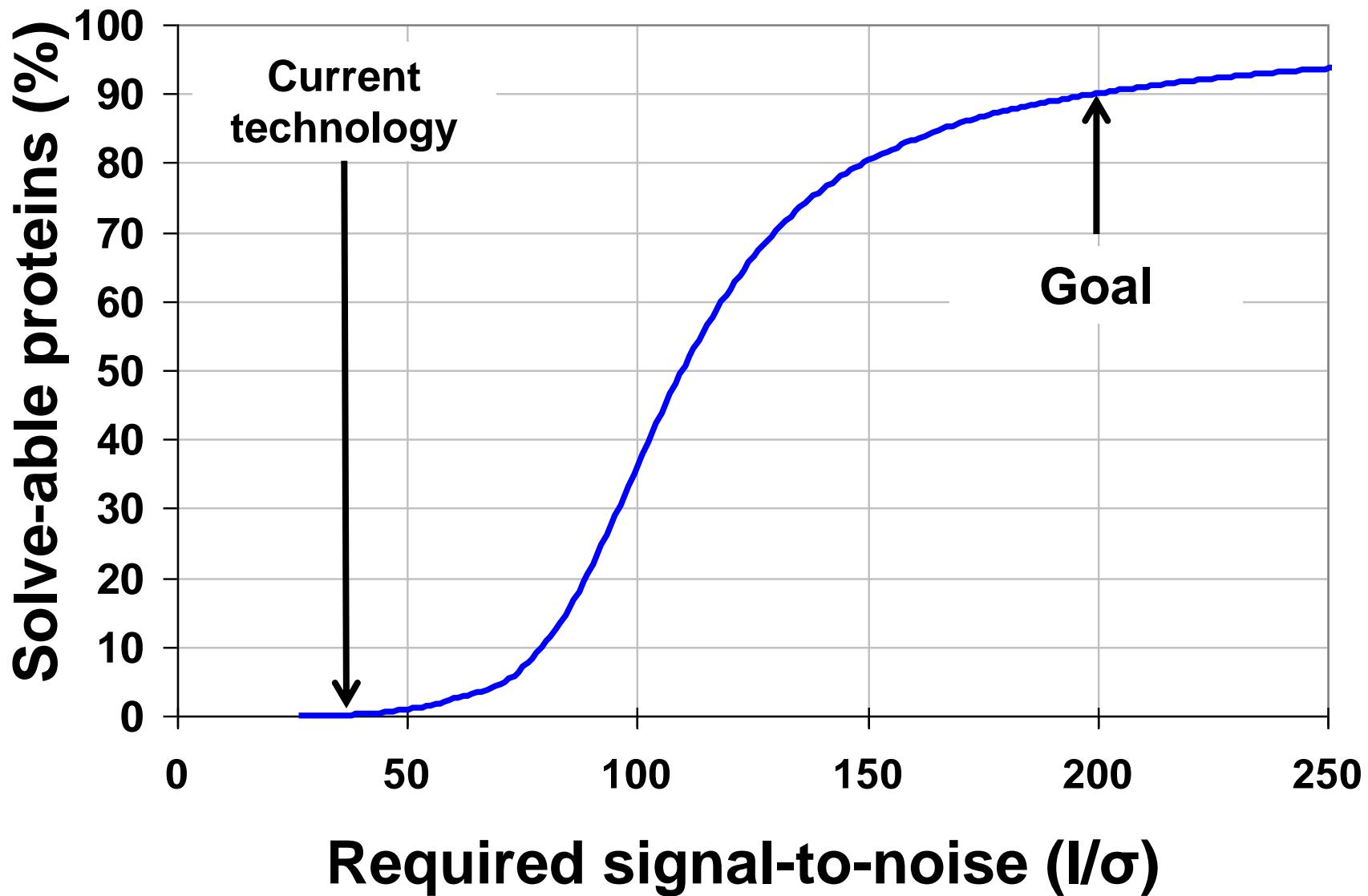
$R_{\text{meas}} \approx 0.1\% ? \longrightarrow ISa = 1000$

In reality: $ISa \sim 33 \longrightarrow R_{\text{meas}} = \approx 3\%$

$$\frac{\sqrt{1,000}}{1,000} = 3\%$$

> 1000 $\frac{\text{photon}}{\text{spot}}$ is a waste!

Threshold of a revolution in phasing

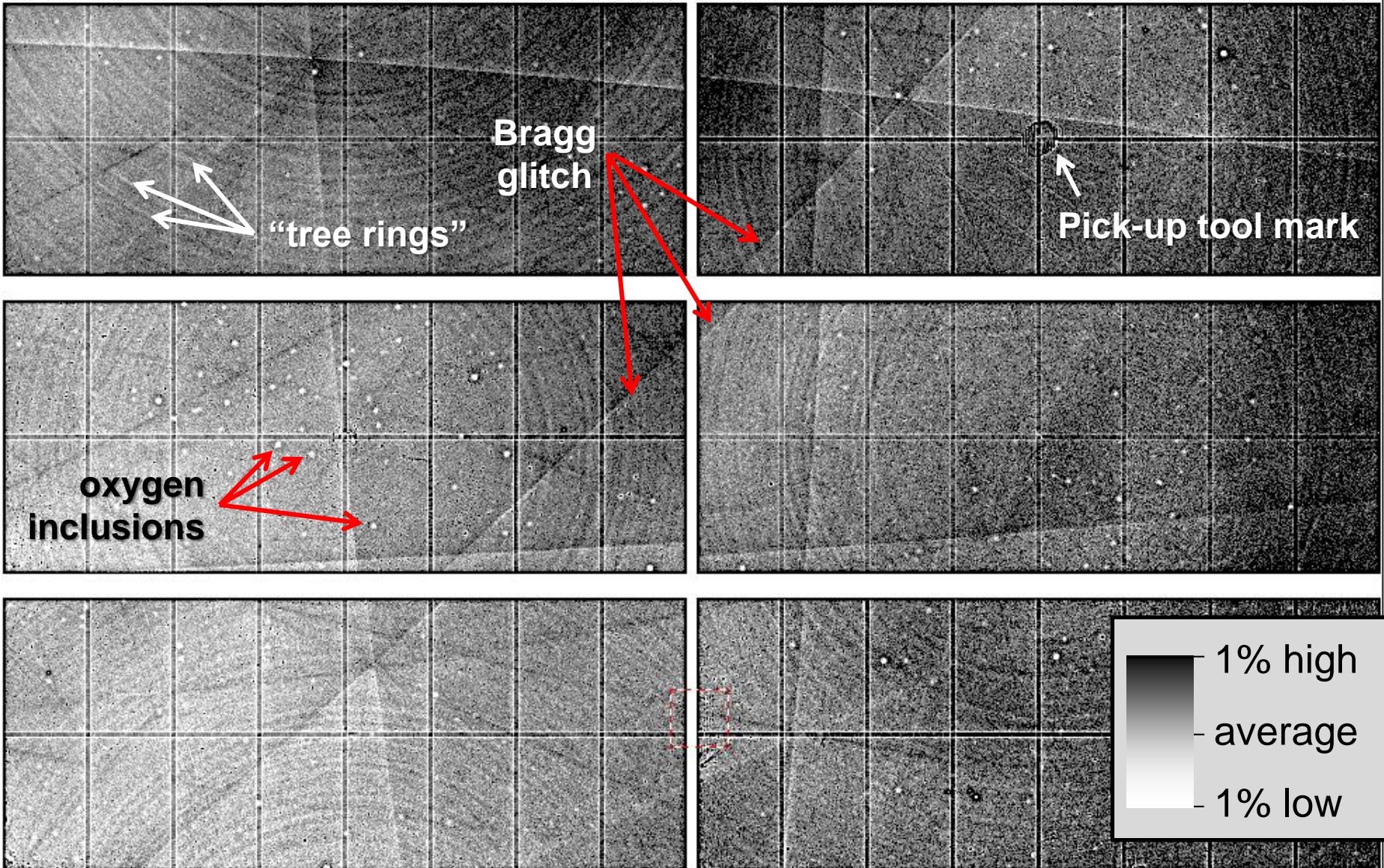


Threshold of a revolution in phasing

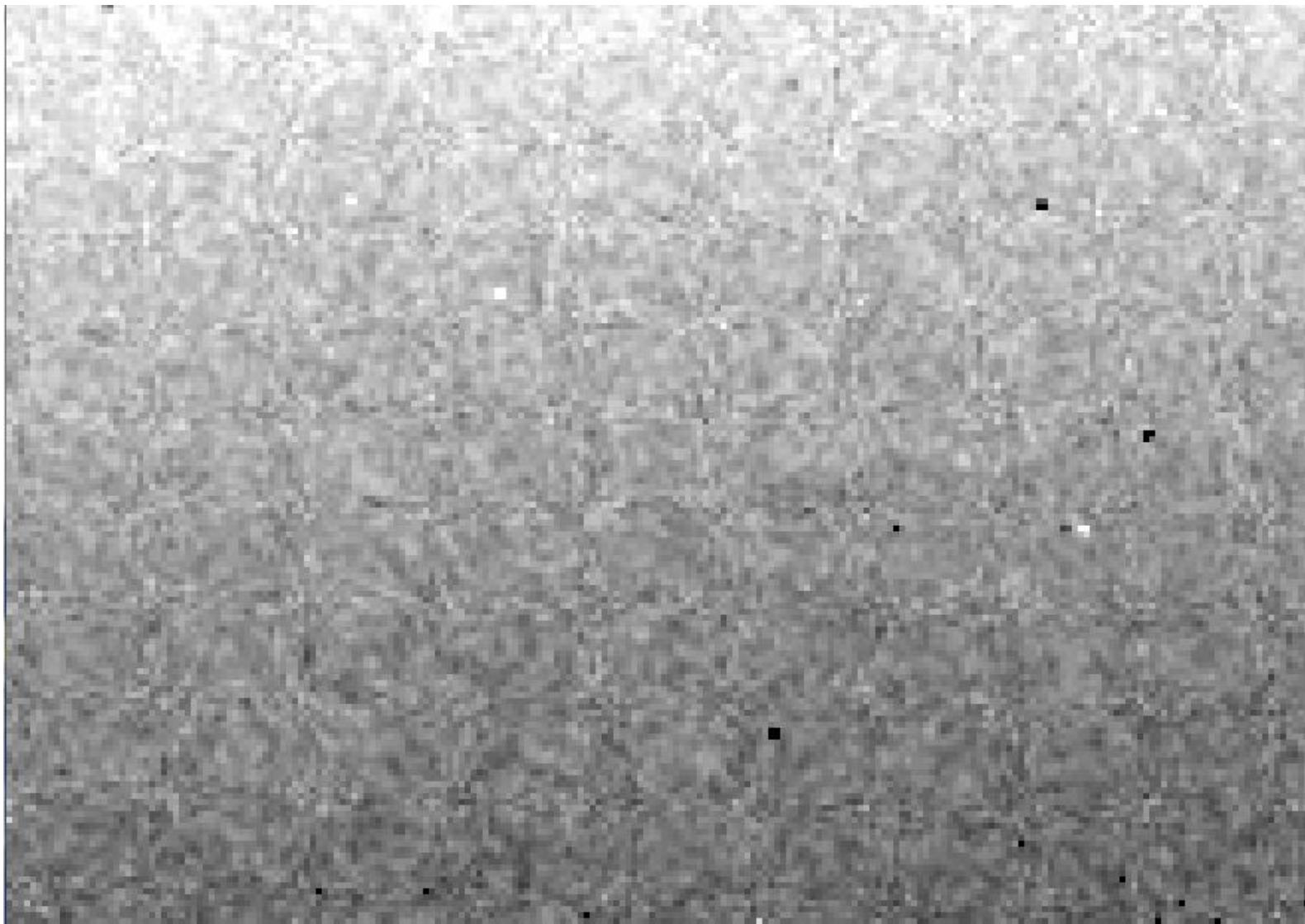
Source of error	realistic simulation
Photon counting	+
Shutter jitter	+
Beam flicker	+
Sample absorption	+
Radiation damage	+
Imperfect spindle	+
vignette	+
Corner correction	+
SHSSS	+
$R_{\text{meas}} (\infty-10 \text{ \AA})$	2.8%
I/σ asymptotic	26.8

Pilatus: subtract smooth baseline

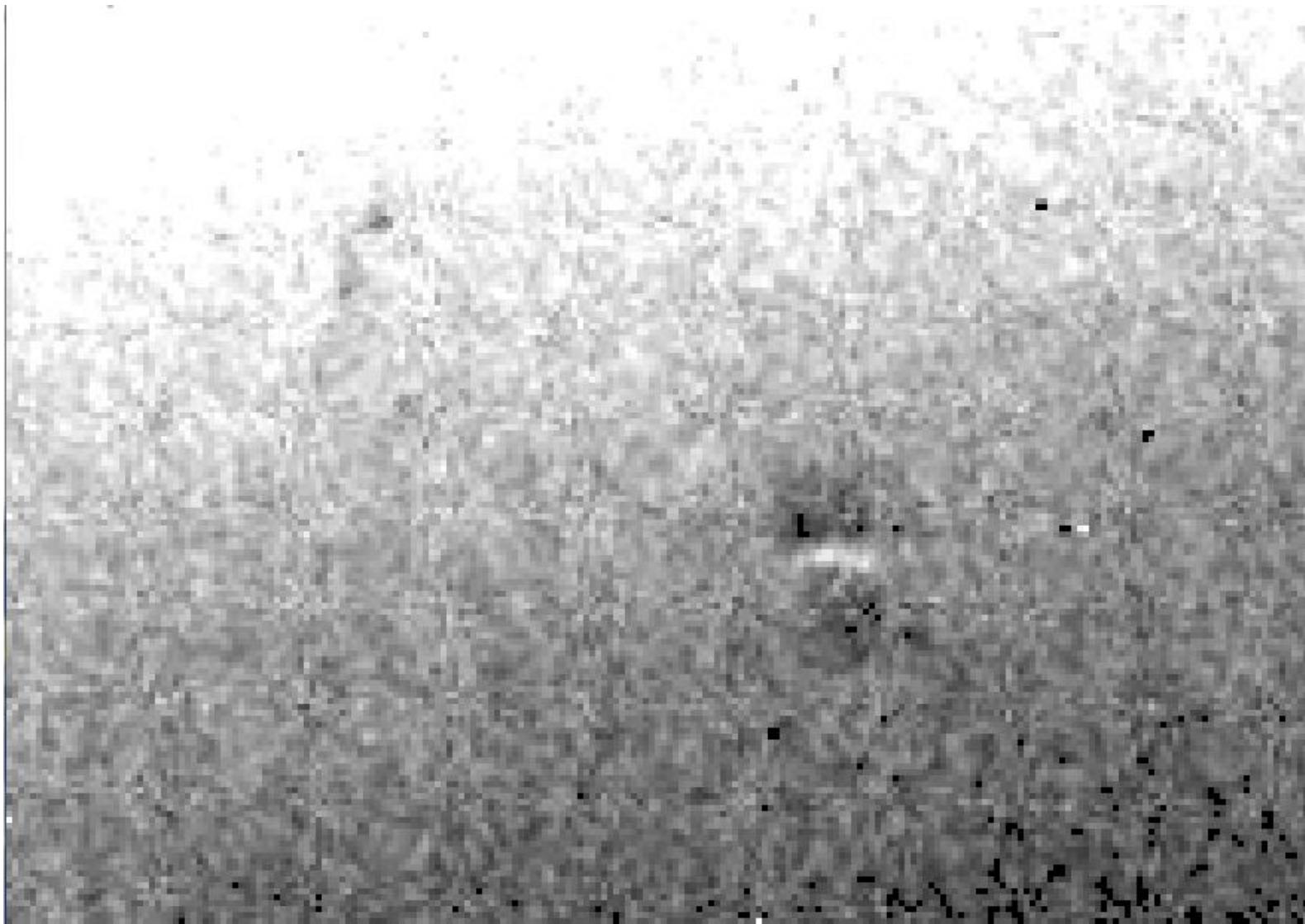
$\sim 3 \times 10^5$ photon/pixel



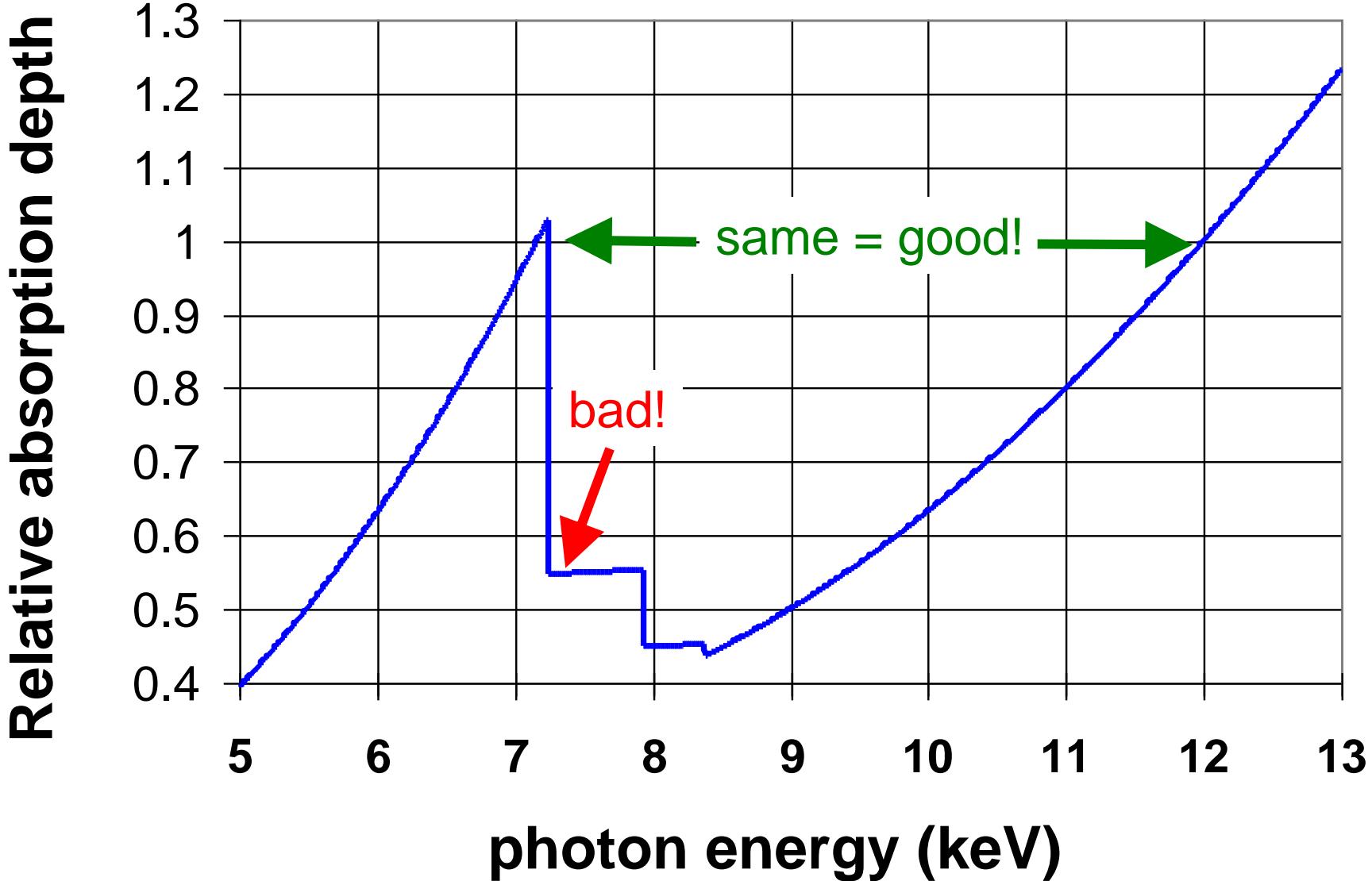
CCD calibration: 7235 eV



CCD calibration: 7247 eV



Gadox calibration vs energy



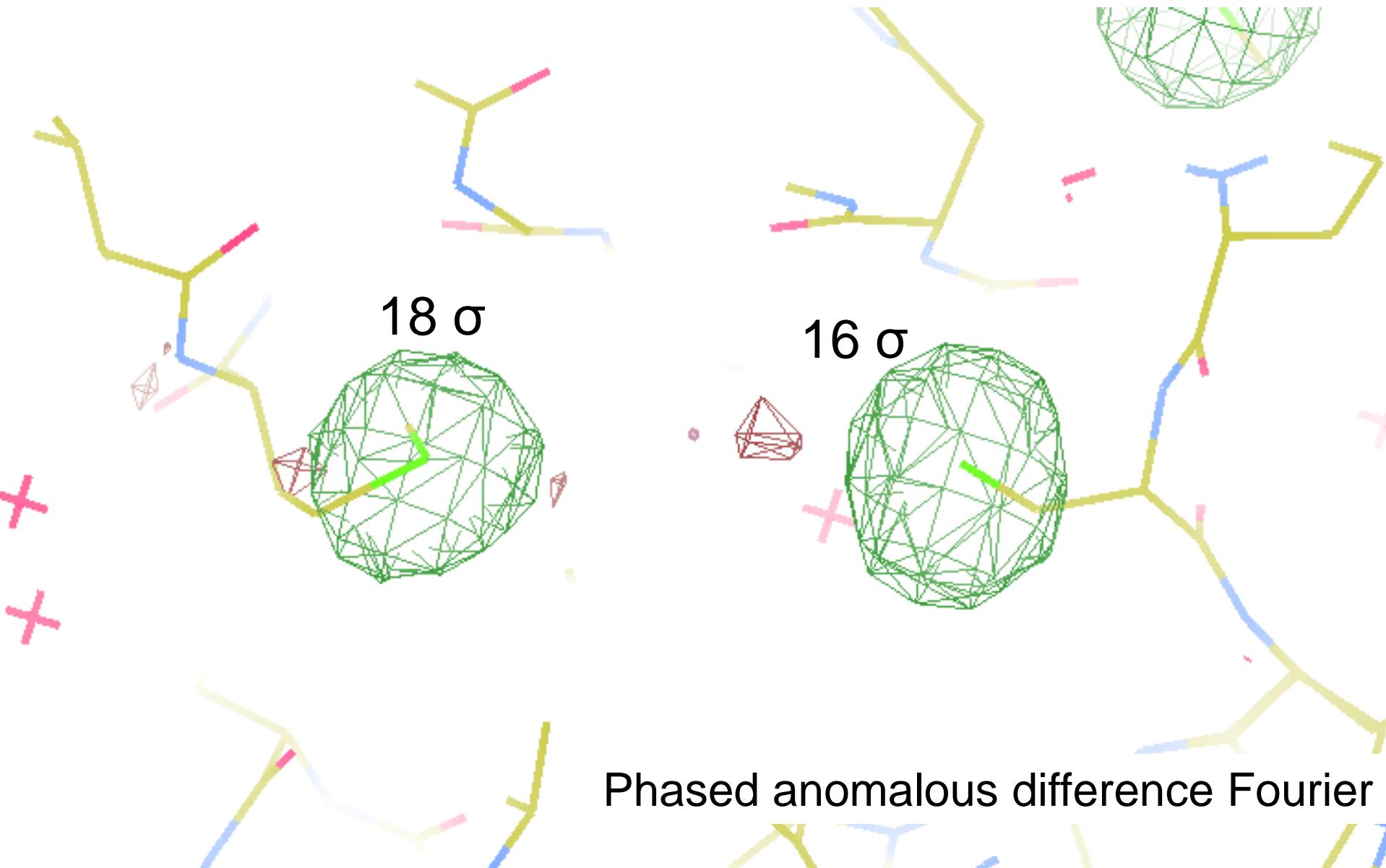
Data collection parameters:

- 16 crystals
- 360° each, inverse beam
- 7235 eV photon energy
- < 1 MGy per xtal
- Australian Synchrotron MX1
 - 35 kGy/s into 100 µm x 100 µm

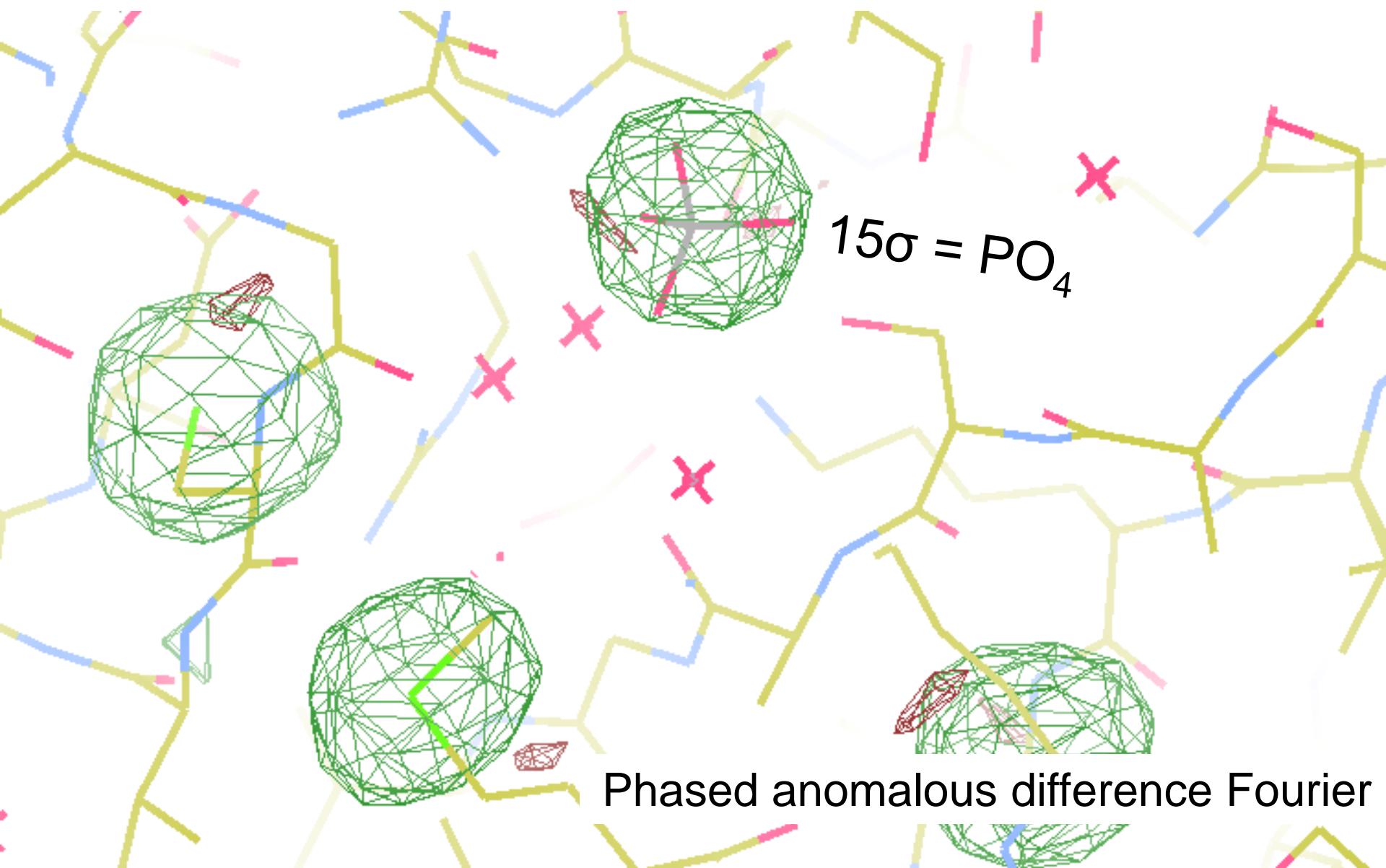
140-fold multiplicity: 16 crystals, 360° each, inverse beam, 7235 eV

RESOLUTION LIMIT	COMPLETENESS OF DATA	R-FACTOR observed	I/SIGMA	R-meas	CC (1/2)	Anomal Corr	SigAno	Nano
9.17	99.1%	3.9%	257.47	3.9%	100.0*	91*	5.024	450
6.49	100.0%	5.2%	214.33	5.2%	100.0*	86*	3.836	882
5.30	100.0%	7.2%	165.13	7.2%	100.0*	76*	3.257	1175
4.59	100.0%	7.2%	175.42	7.3%	100.0*	67*	2.589	1403
4.10	99.9%	7.7%	174.13	7.7%	100.0*	59*	2.264	1594
3.74	99.9%	9.4%	143.09	9.4%	100.0*	49*	1.953	1783
3.47	100.0%	11.2%	120.17	11.2%	100.0*	39*	1.696	1942
3.24	100.0%	14.1%	91.14	14.1%	100.0*	30*	1.333	2103
3.06	99.9%	19.5%	65.79	19.5%	100.0*	23*	1.117	2214
2.90	99.9%	29.0%	44.85	29.1%	99.9*	17*	1.008	2369
2.77	99.9%	40.5%	32.58	40.6%	99.8*	11*	0.901	2493
2.65	99.9%	52.8%	25.16	52.9%	99.8*	10*	0.866	2605
2.54	100.0%	67.4%	19.47	67.6%	99.6*	2	0.804	2705
2.45	100.0%	88.9%	14.58	89.2%	99.2*	4	0.831	2859
2.37	100.0%	109.3%	9.97	109.7%	98.1*	5	0.829	2925
2.29	100.0%	138.2%	6.87	138.9%	96.1*	1	0.760	3037
2.22	100.0%	197.1%	4.03	198.6%	83.5*	-1	0.721	3159
2.16	100.0%	227.3%	2.41	230.8%	46.9*	-1	0.677	3224
2.10	61.2%	154.4%	1.28	163.6%	47.0*	-2	0.660	1999
2.05	47.9%	170.1%	0.68	196.5%	25.7*	3	0.629	1578
total	93.3%	15.7%	54.30	15.8%	100.0*	12*	1.217	42499

140-fold multiplicity



140-fold multiplicity

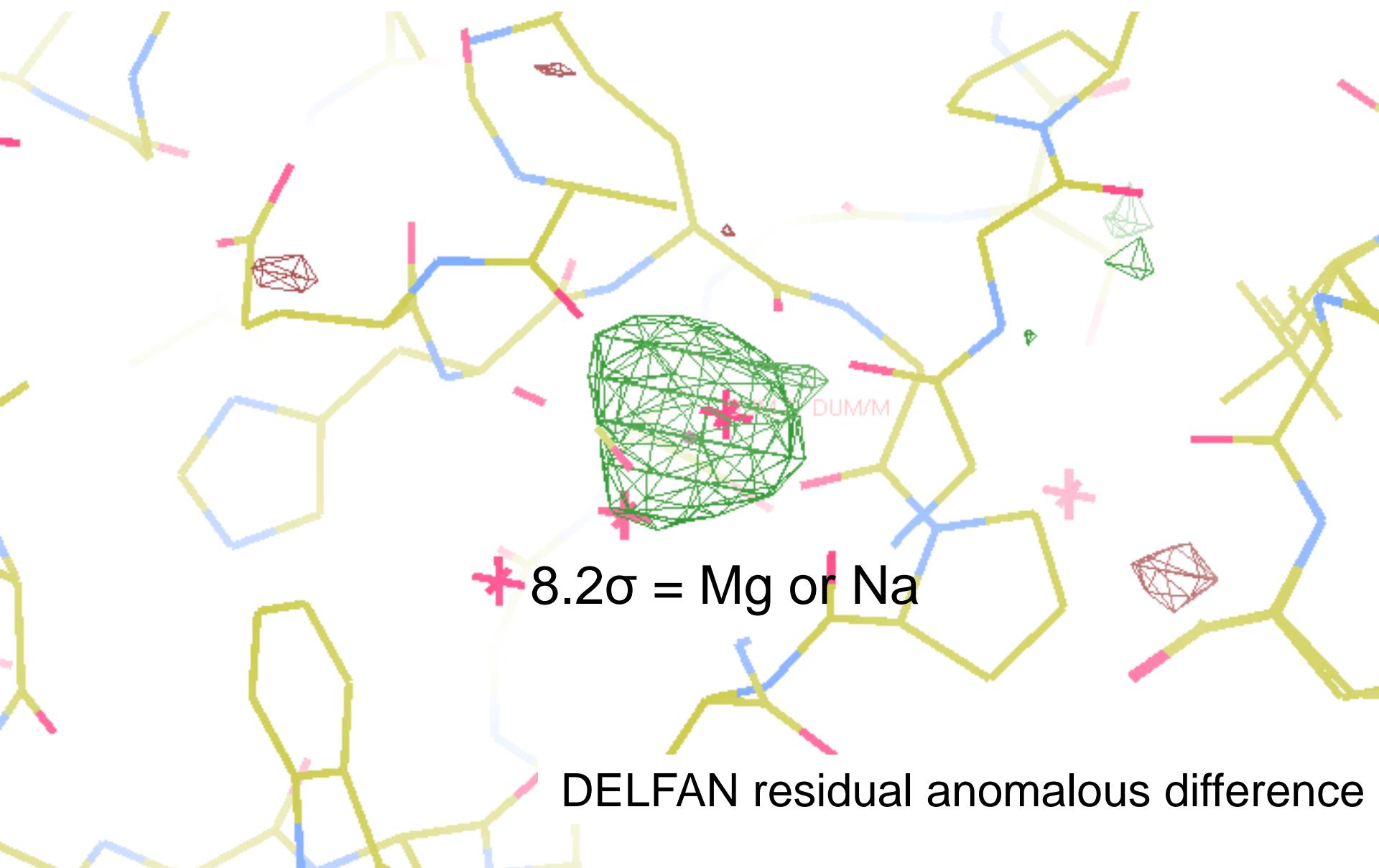


140-fold multiplicity

≈ 2σ = Mg?

Phased anomalous difference Fourier

140-fold multiplicity



Why doesn't everyone do this?

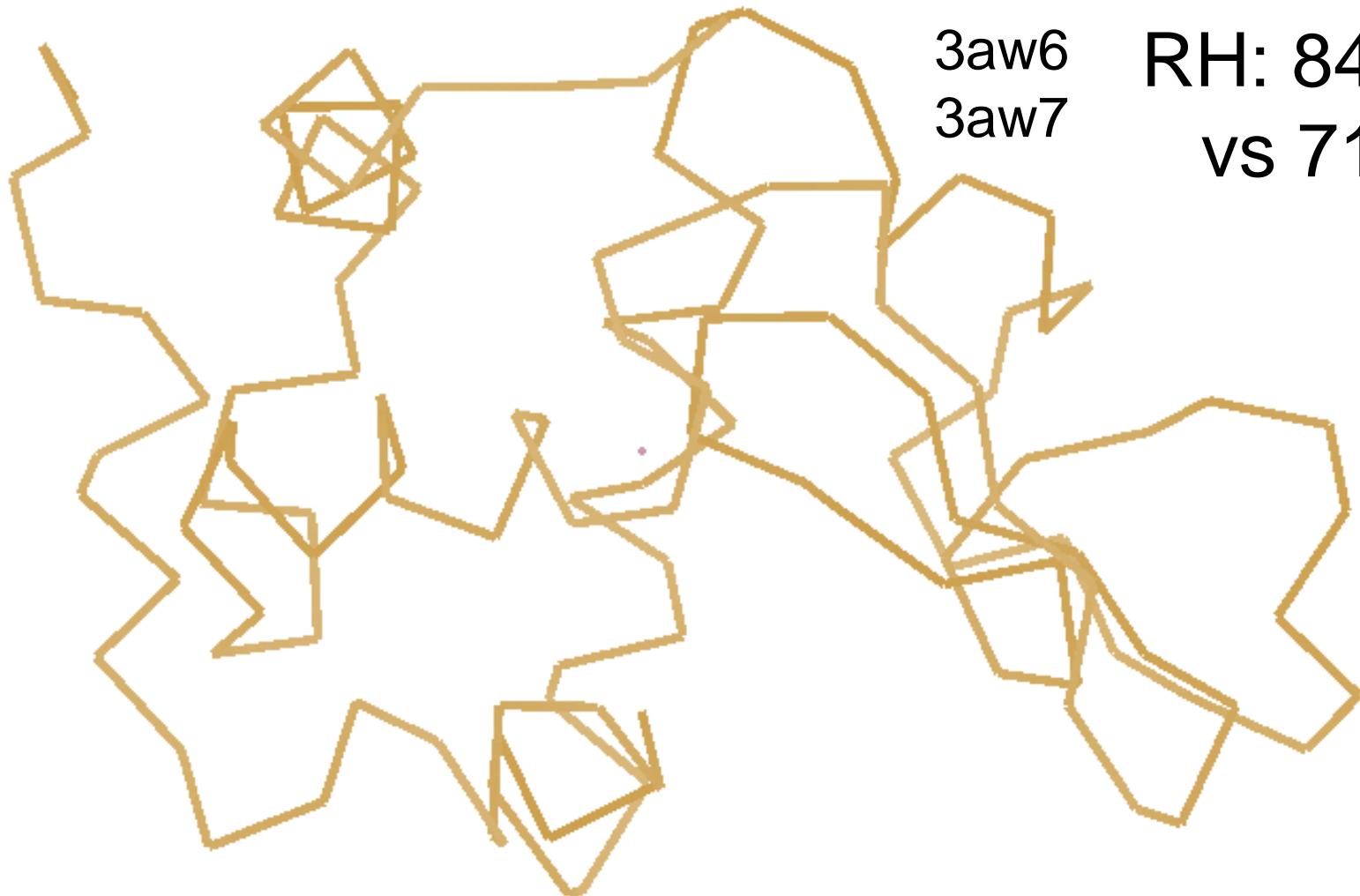
non-isomorphism

Non-isomorphism in lysozyme

RMSD = 0.18 Å

$\Delta\text{cell} = 0.7 \%$

$R_{\text{iso}} = 44.5\%$

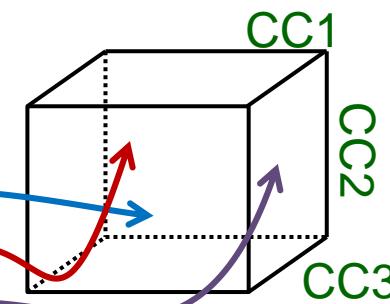
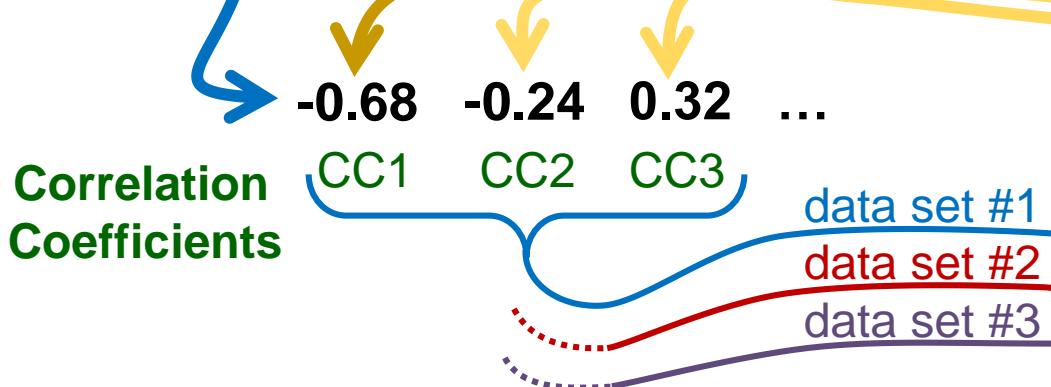


X-ray Data Sets

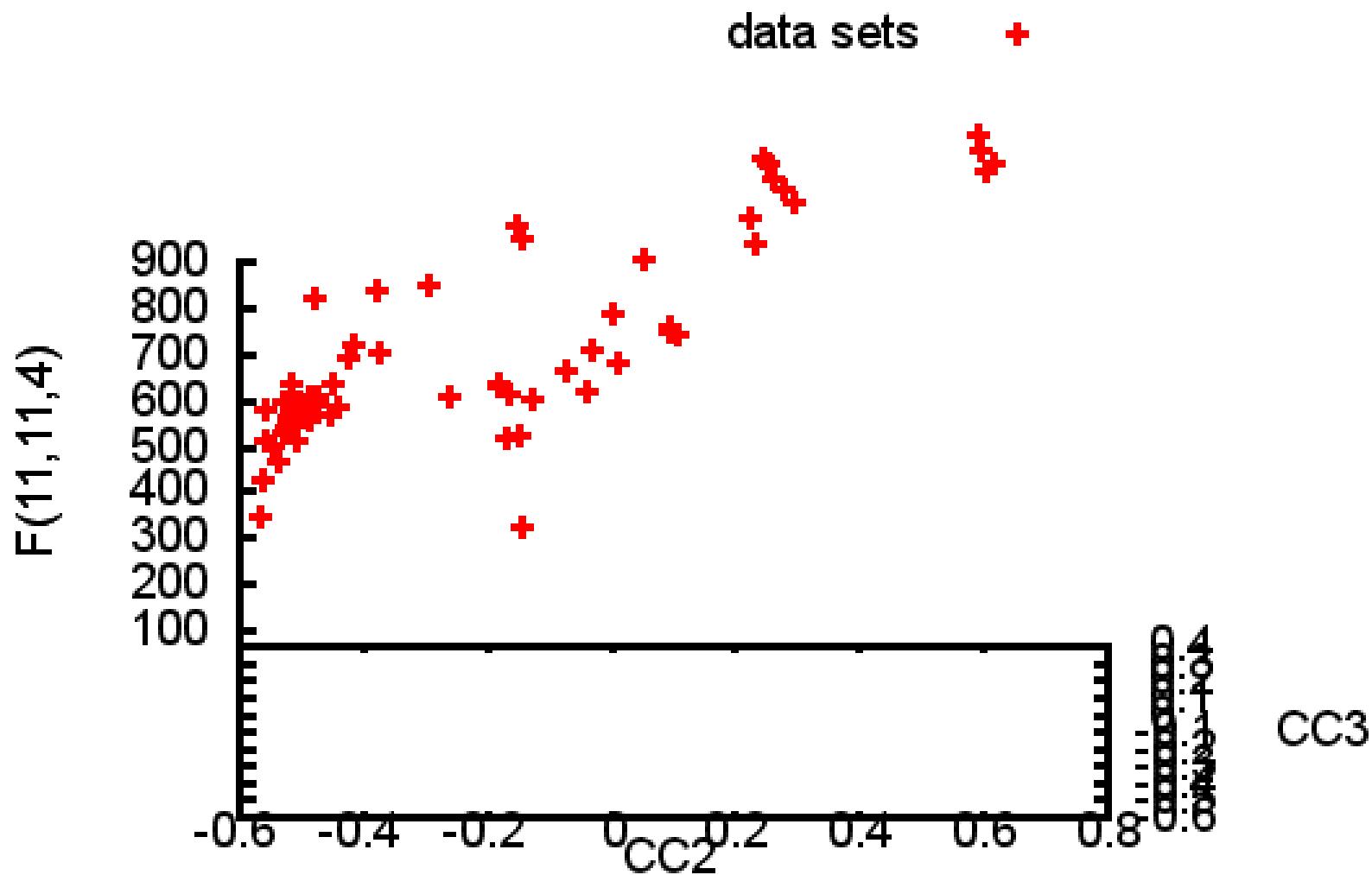
data set h,k,l	structure factors (F)			
	#1	#2	#3	#4
5,3,4	523.7	559.8	579.9	603.2
5,4,4	168.2	166.6	177.2	196.1
5,5,4	34.9	26.4	19.2	17.3
6,1,4	305.7	301.1	298.1	296.3
6,2,4	353.0	353.9	356.2	366.9
6,3,4	300.9	285.3	273.5	259.4
6,3,5	223.8	226.3	234.6	251.4
...

Singular values & vectors

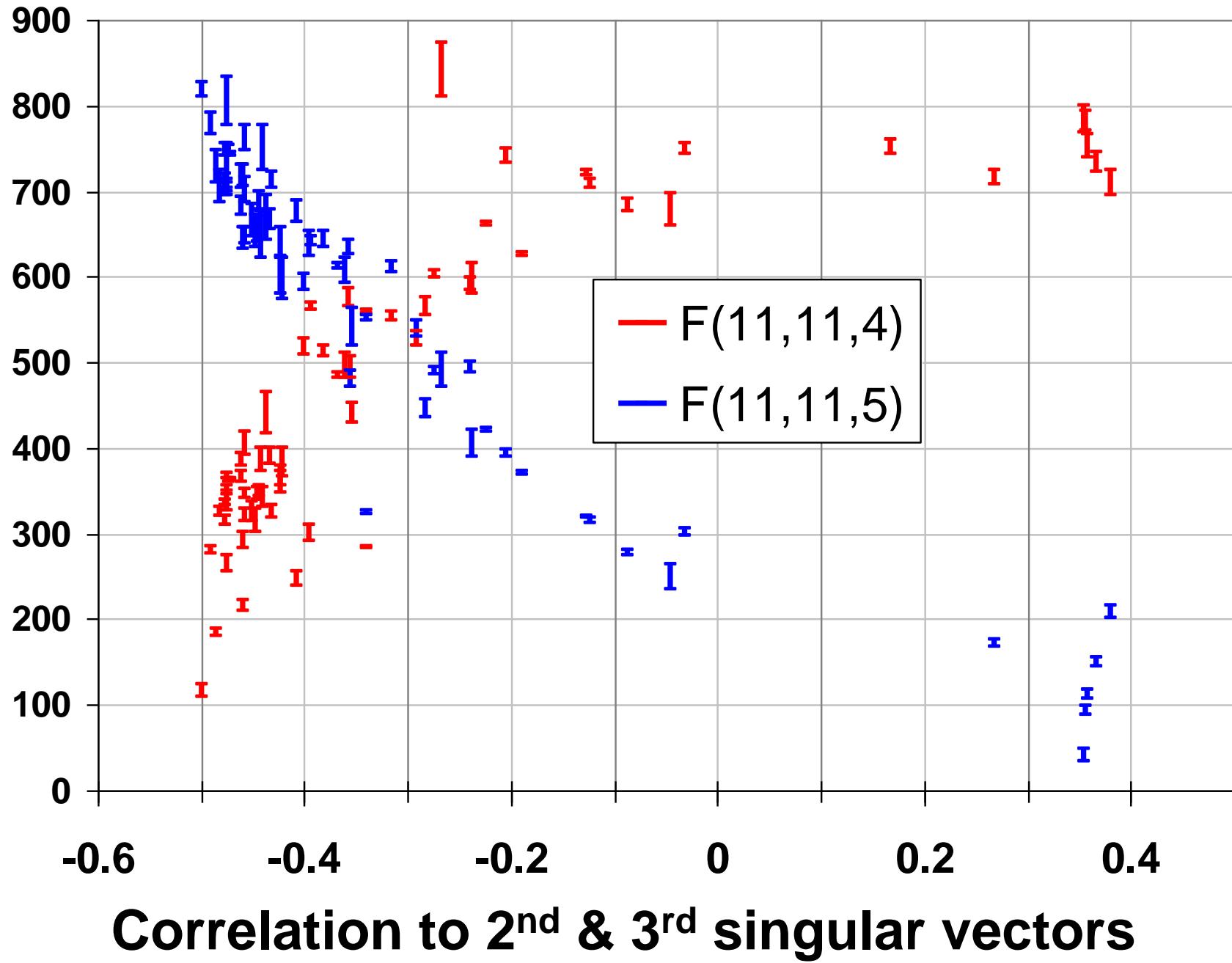
value vector h,k,l	100 %	17 %	13 %	9 %
	1st	2nd	3rd	4th
5,3,4	-1.46	-0.02	1.84	-0.72
5,4,4	-0.88	-0.29	-0.34	0.15
5,5,4	-0.42	-0.65	1.02	1.47
6,1,4	-0.90	-0.85	1.44	-0.40
6,2,4	-1.20	-0.37	0.67	0.01
6,3,4	-0.75	0.31	0.48	0.00
6,3,5	-0.75	-0.85	0.72	-0.82
...



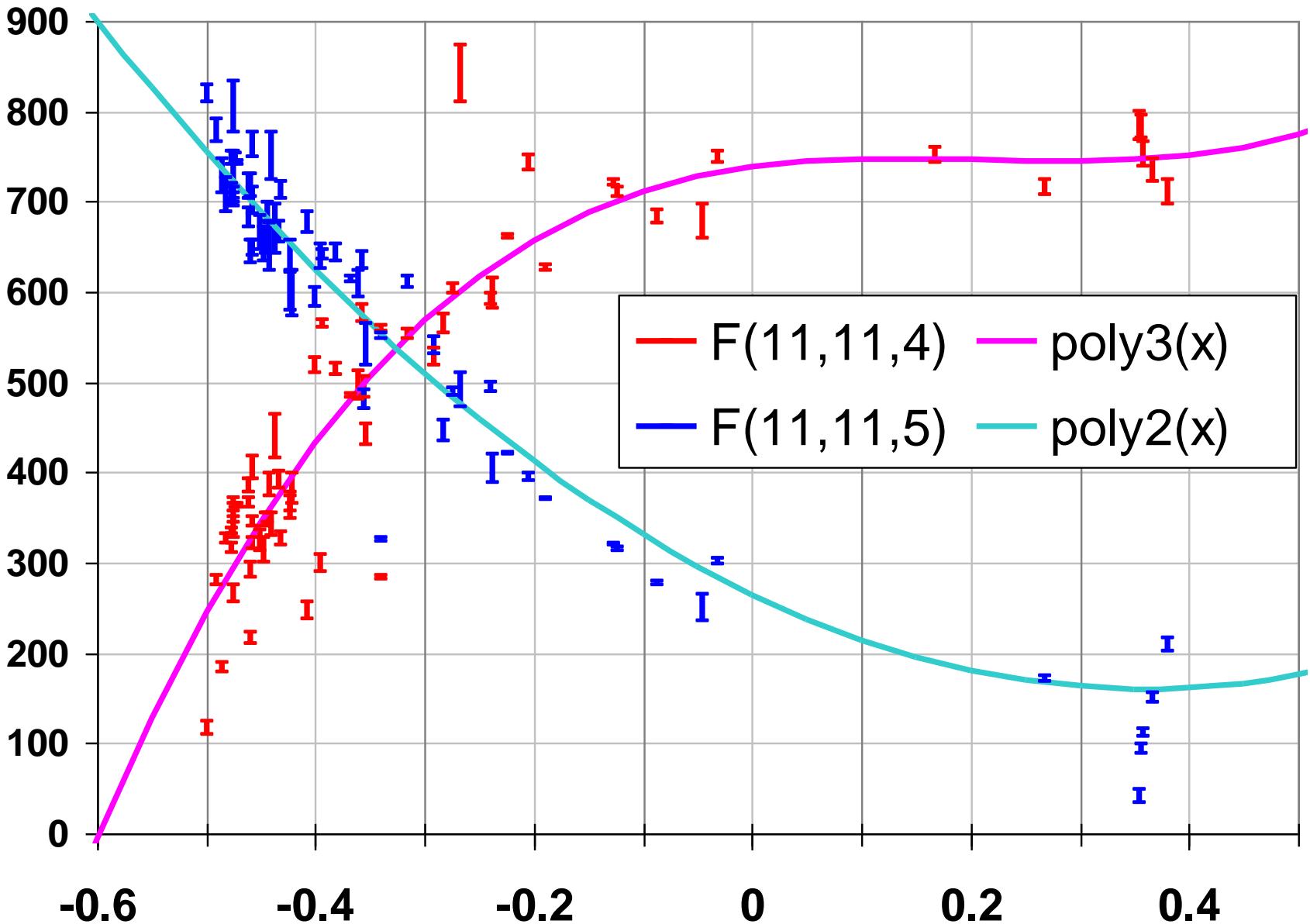
Data sets Positioned in “Correlation Space”



Structure factor (electrons)

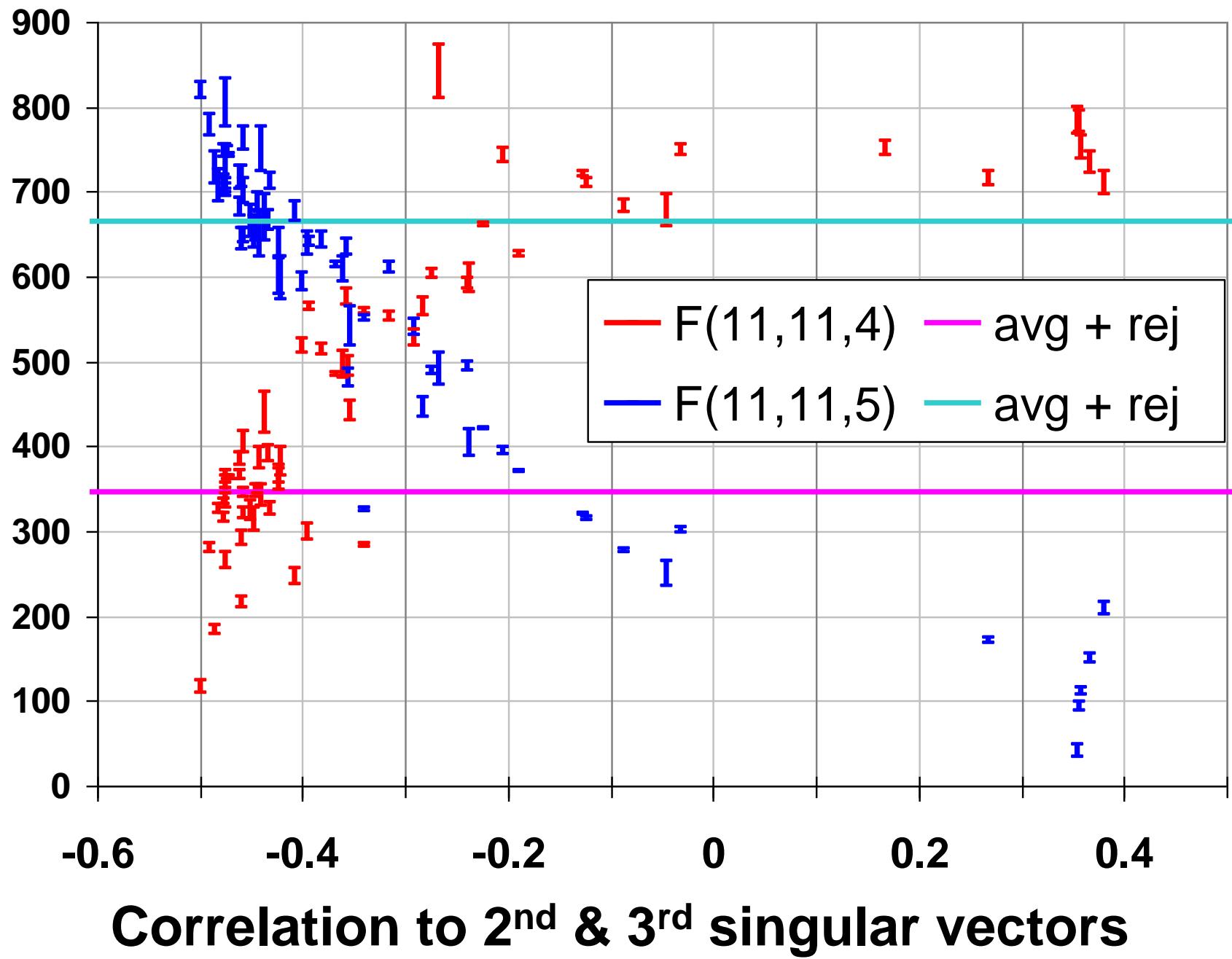


Structure factor (electrons)

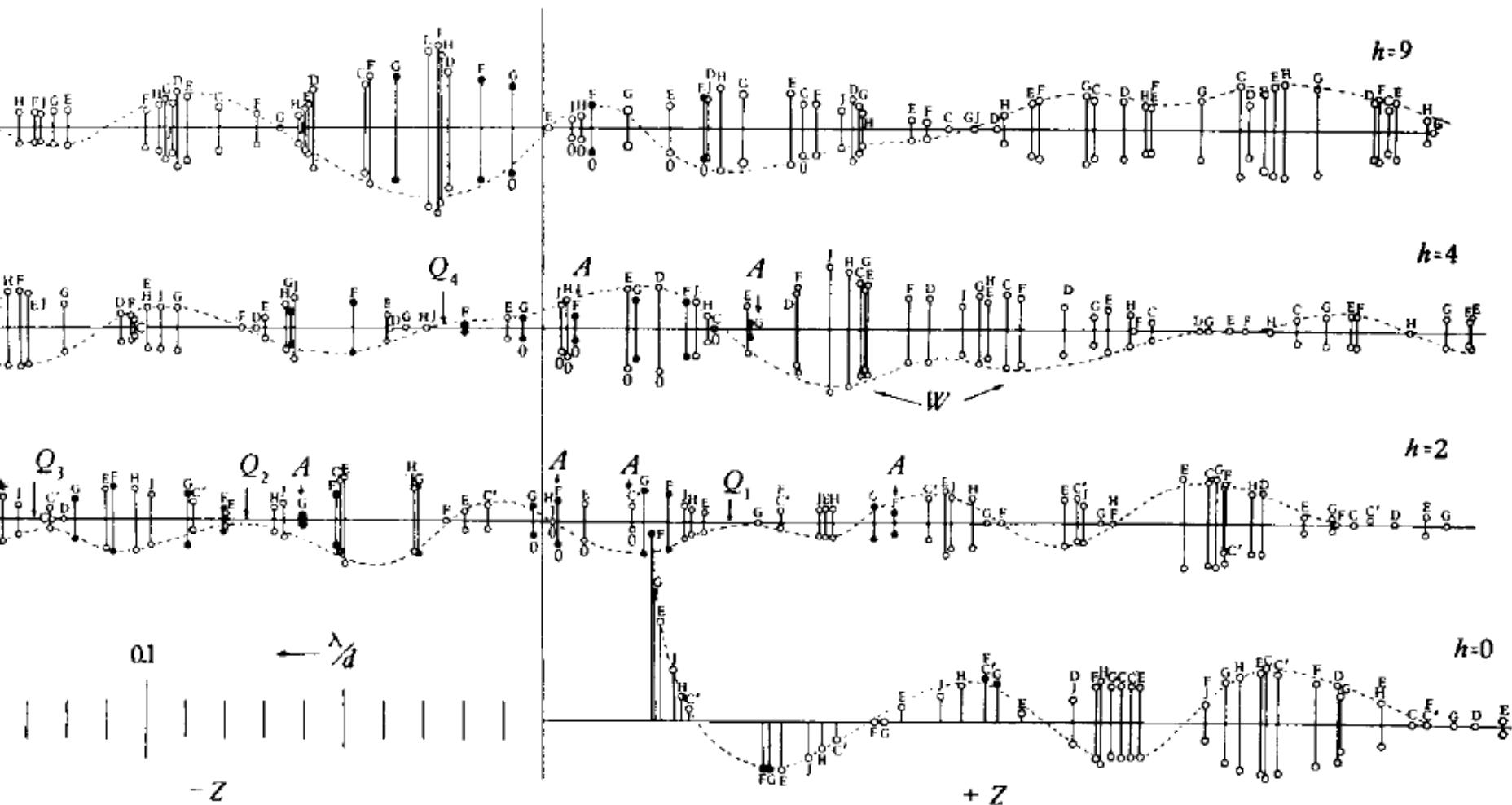


Correlation to 2nd & 3rd singular vectors

Structure factor (electrons)



Bragg's "minimum wavelength principle"

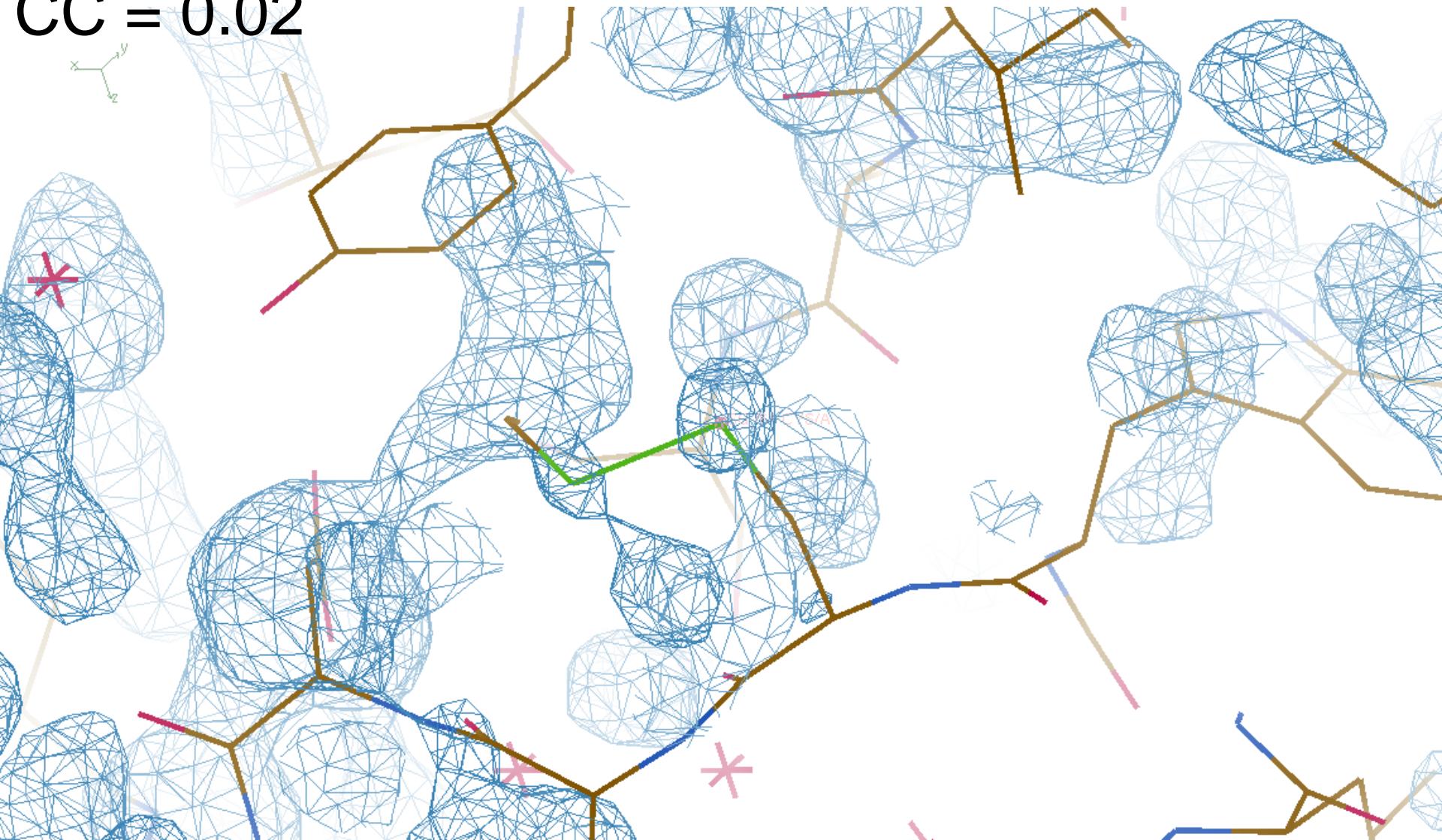


Perutz (1985). "Early Days of Cryst..." *Methods in Enzymology*, Vol. 114, 3-18.
Bragg & Perutz (1952). "external form haemoglobin I", *Acta Cryst.* **5**, 277-283.

DMMULTI – fake data

4 deg rotation: 8 “xtals”: before

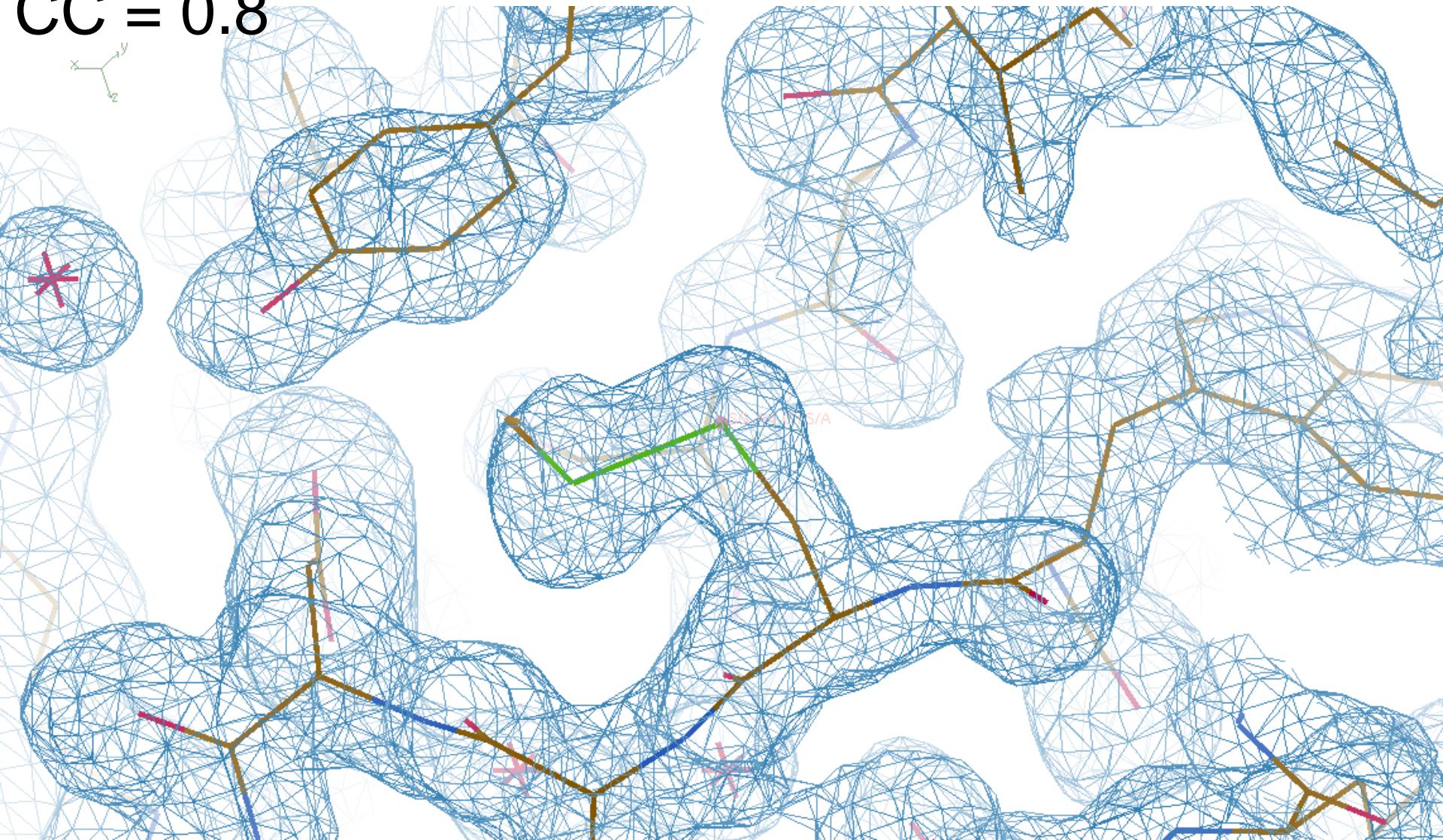
CC = 0.02



DMMULTI – fake data

4 deg rotation: 8 “xtals”: after

CC = 0.8



Suggested anomalous protocol:



1. 360° in < 5 MGy
2. move detector
3. 4X exposure
4. goto 2

2 wavelengths are better than 1
- $(\text{peak} + \text{inf})/2$, and remote
MAD, not M-SAD!

Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

Optimal exposure time (CCD)

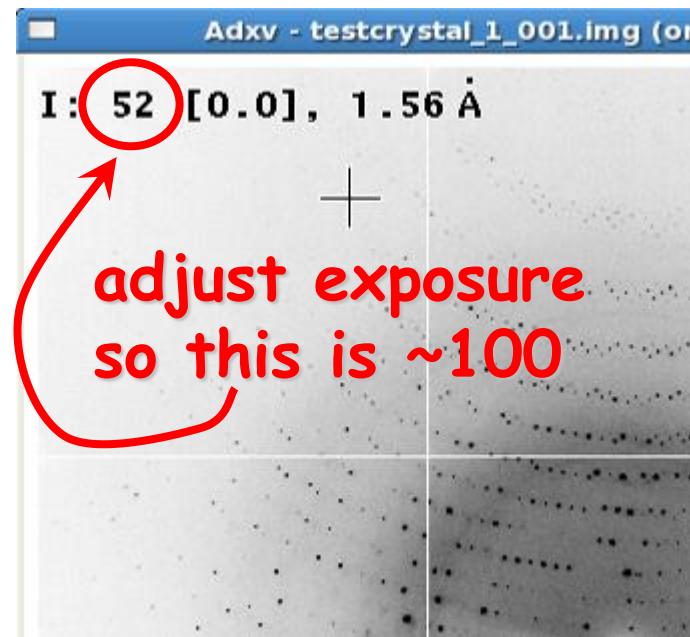
$$t_{hr} = t_{ref} \frac{10 \cdot m \cdot \sigma_0^2}{gain \cdot (bg_{ref} - bg_0)}$$

t_{hr}
 t_{ref}
 bg_{ref}

Optimal exposure time for data set (s)
exposure time of reference image (s)
background level near weak spots on
reference image (ADU)

bg_0
 bg_{hr}
 σ_0
 $gain$
 m

ADC offset of detector (ADU)
optimal background level (via t_{hr})
rms read-out noise (ADU)
ADU/photon
multiplicity of data set (including partials)



Optimal exposure time (PAD)

$$t_{hr} > 100 \times t_{ro}$$

- | | |
|----------|--|
| t_{hr} | Optimal exposure time for data set (s) |
| t_{ro} | read-out time (s) |
| | 2.03 ms on PILATUS3 S |
| | 3 μ s on EIGER |

Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

expected crystal lifetime calculator

source =

full flux =

7.0e+12 photons/s

attenuation =

transmittance =

beam size_{horiz} =

40.0 microns

beam size_{vert} =

80.0 microns

wavelength =

1 Ang

k_{dose} =2000 photons/micron²/Gy

dose rate =

1.1e+6 Gy/s

experiment goal =

resolution =

3 Ang

dose limit =

30 MGy

exposure time =

1 seconds/image

xtal size_{horiz} =

50 microns

xtal size_{vert} =

50 microns

translation during dataset =

0 microns

rotisserie factor

1

 disable warnings

max images =

28 at damage limit

inverse beam =

number of wavelengths =

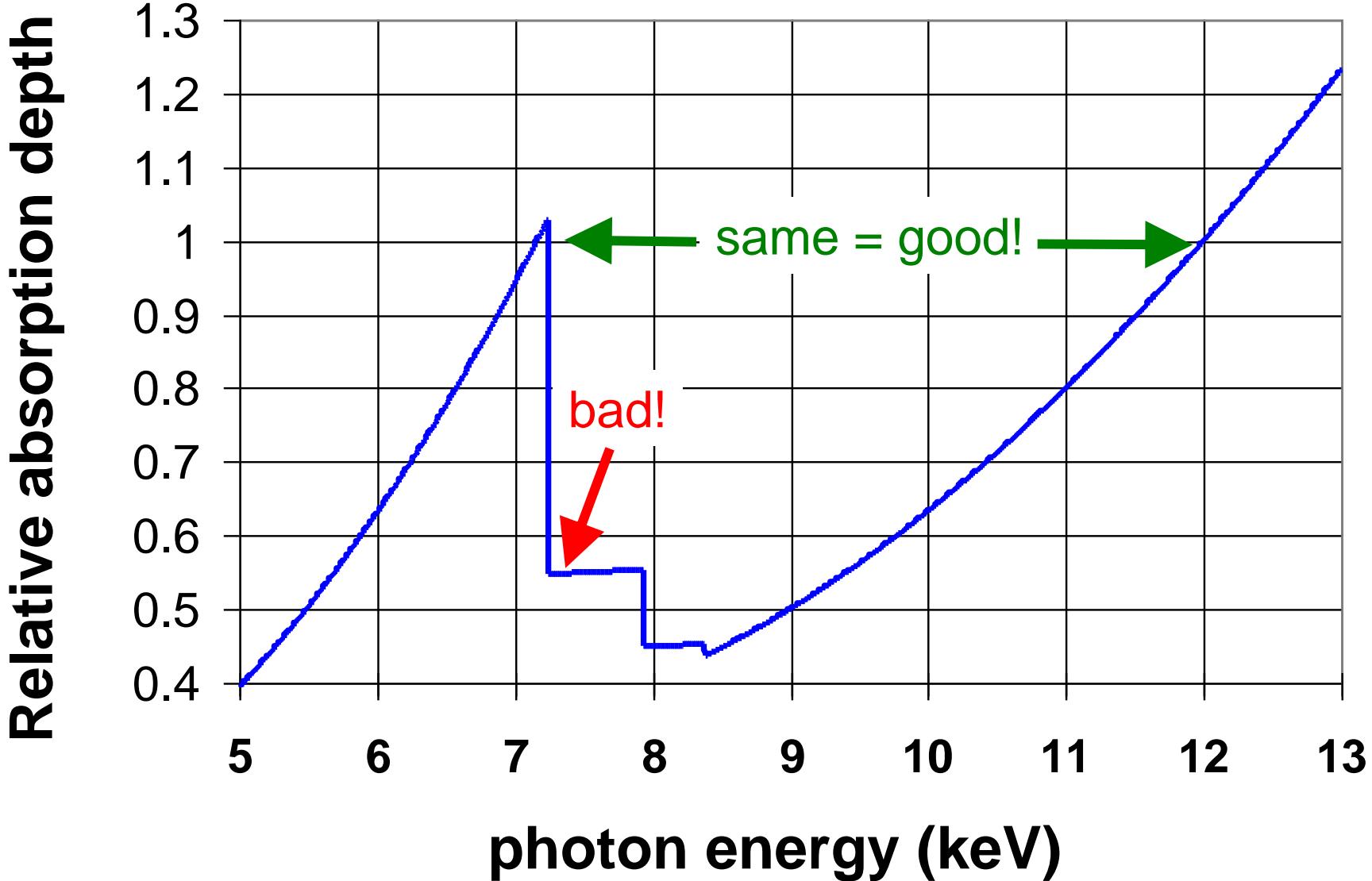
images/wedge =

Decisions, Decisions, Decisions

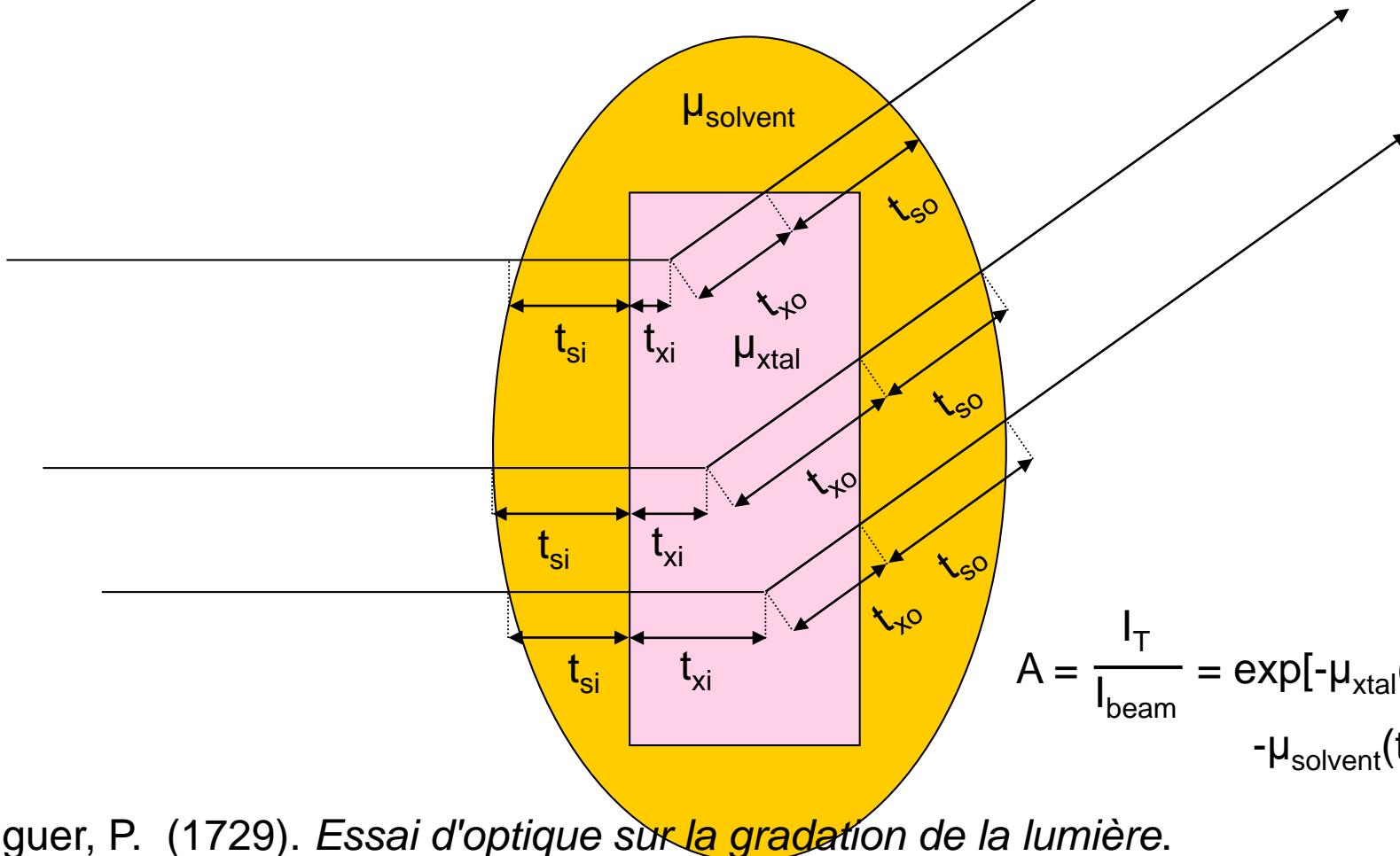
- Exposure time
- Number of images
- **Wavelengths**
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

Gadox calibration vs energy



attenuation factor



$$A = \frac{I_T}{I_{\text{beam}}} = \exp[-\mu_{\text{xtal}}(t_{\text{xi}} + t_{\text{xo}}) - \mu_{\text{solvent}}(t_{\text{si}} + t_{\text{so}})]$$

Bouguer, P. (1729). *Essai d'optique sur la gradation de la lumière*.

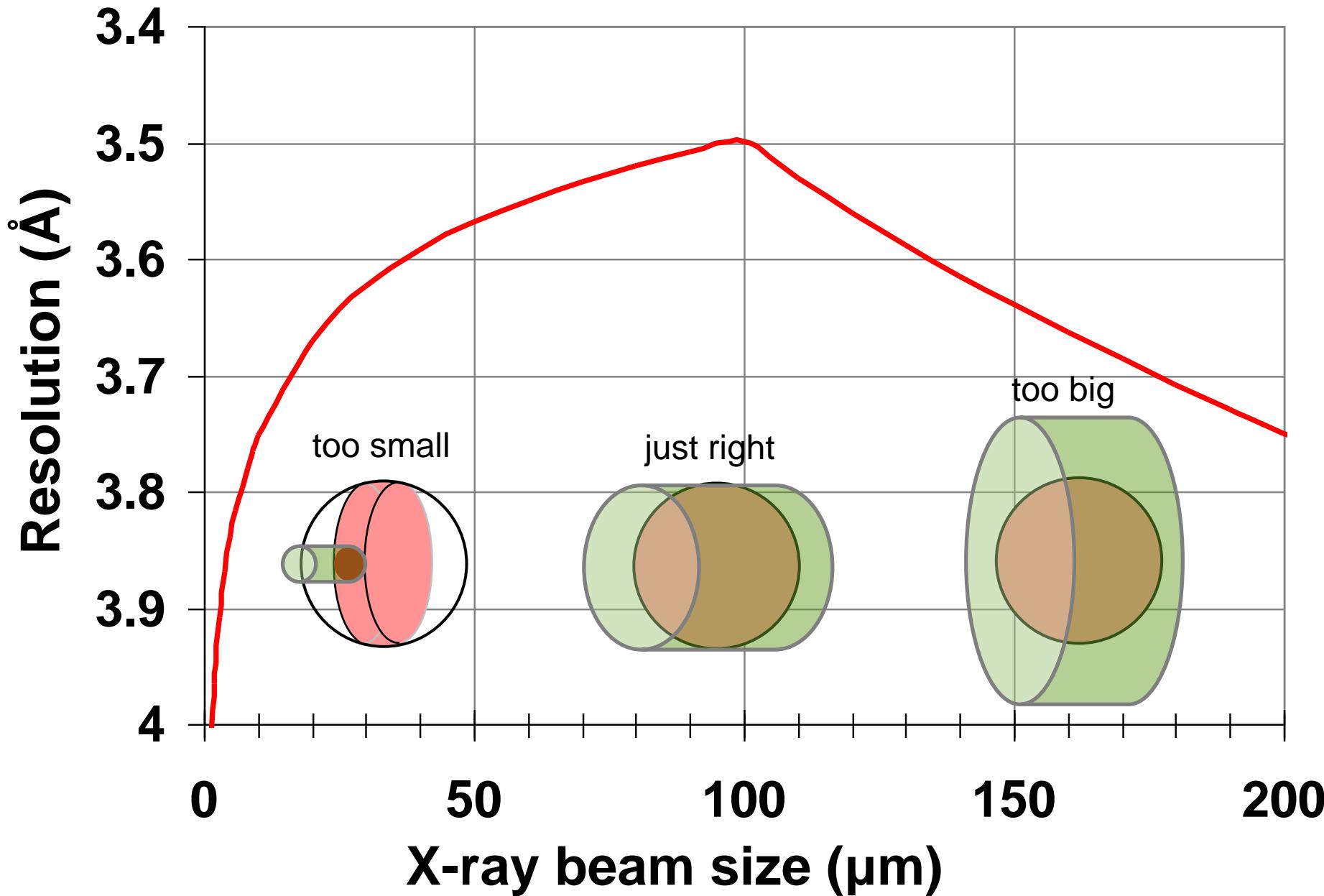
Lambert, J. H. (1760). *Photometria: sive De mensura et gradibus luminis, colorum et umbrae*. E. Klett.

Beer, A. (1852). "Bestimmung der Absorption des rothen Lichts in farbigen Flüssigkeiten", *Ann. Phys. Chem* **86**, 78-90.

Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

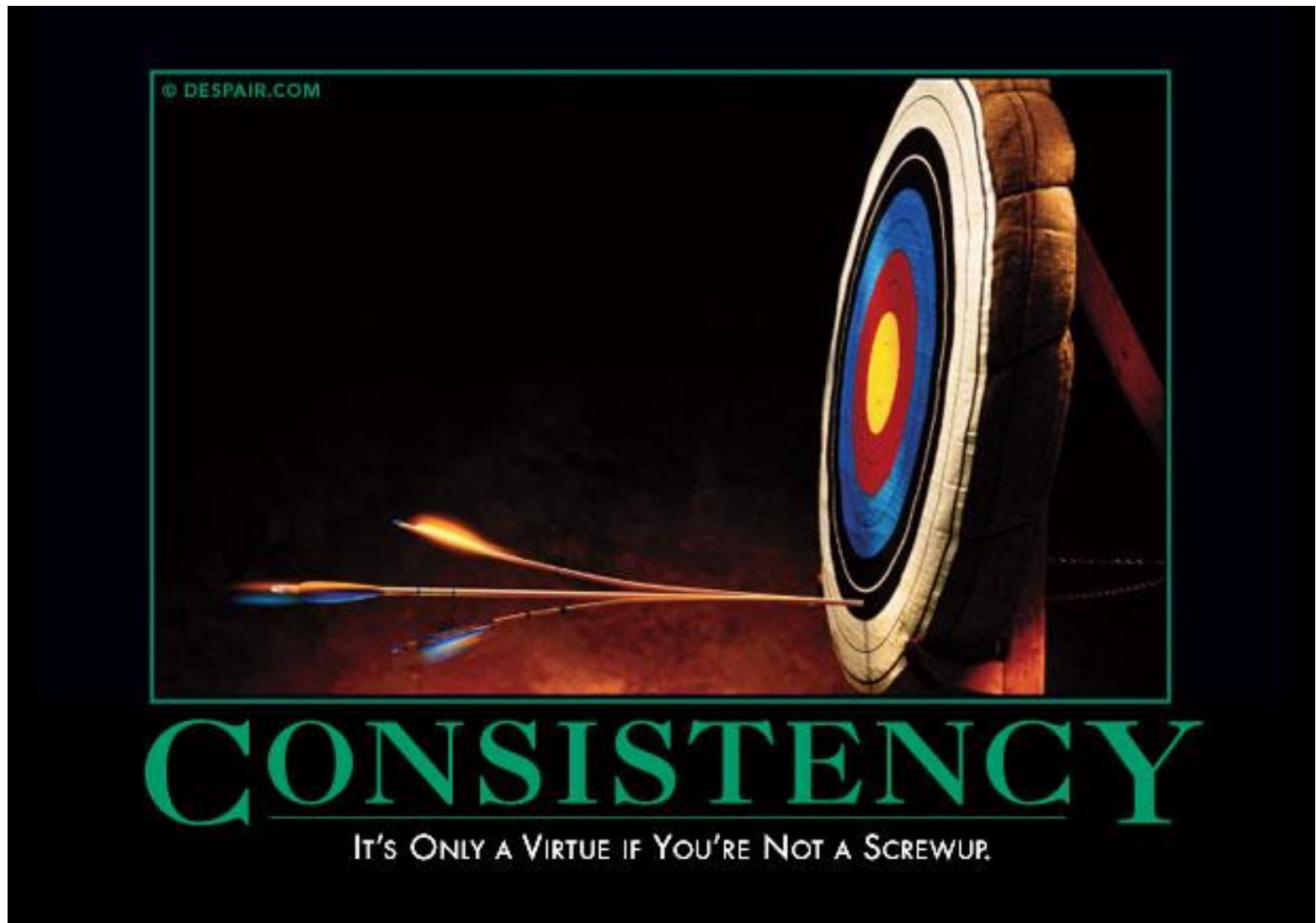


Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

Systematic error does not “average out”

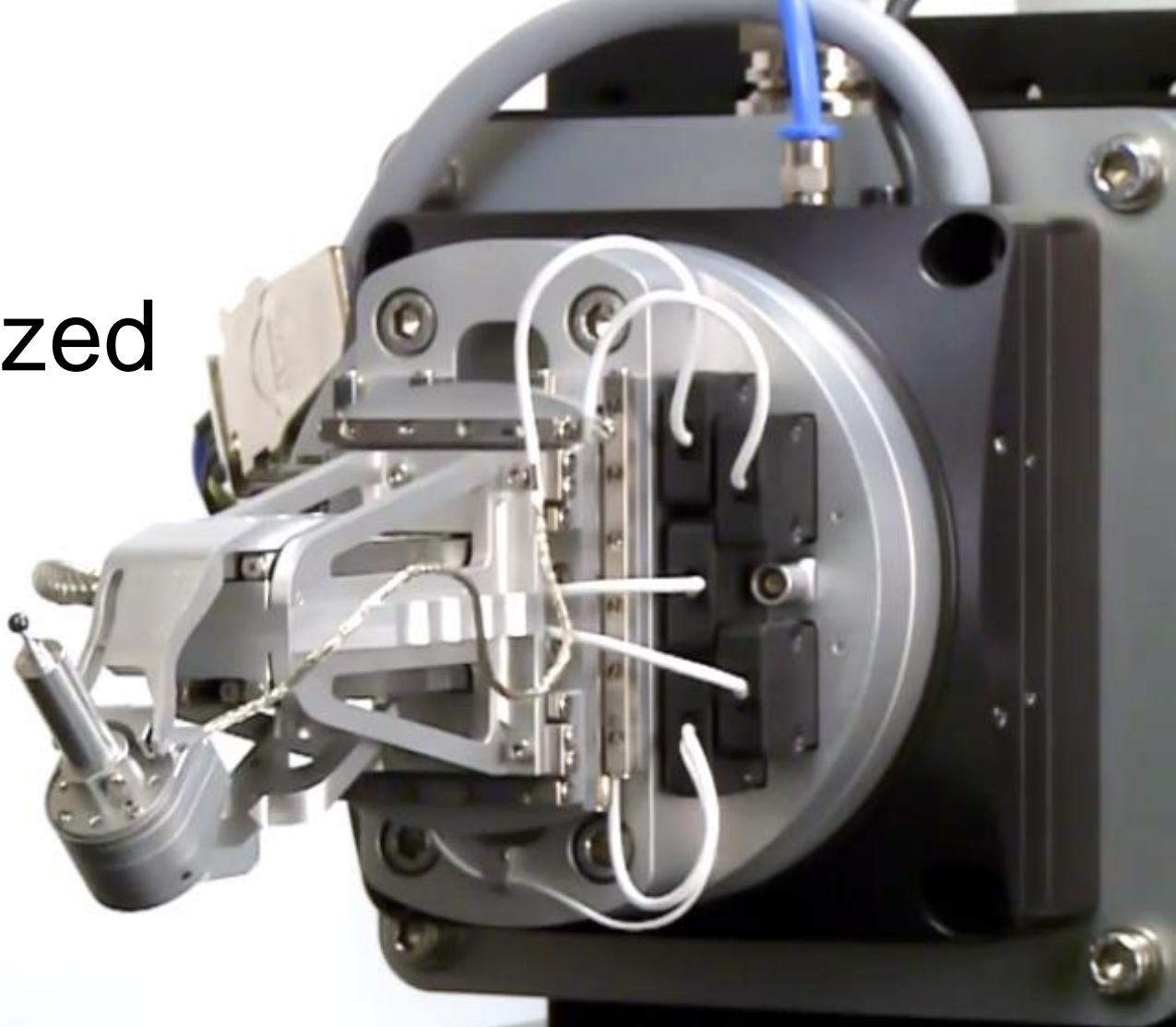


Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

SmarAct's “SmarGon” Commercialized PRIGo

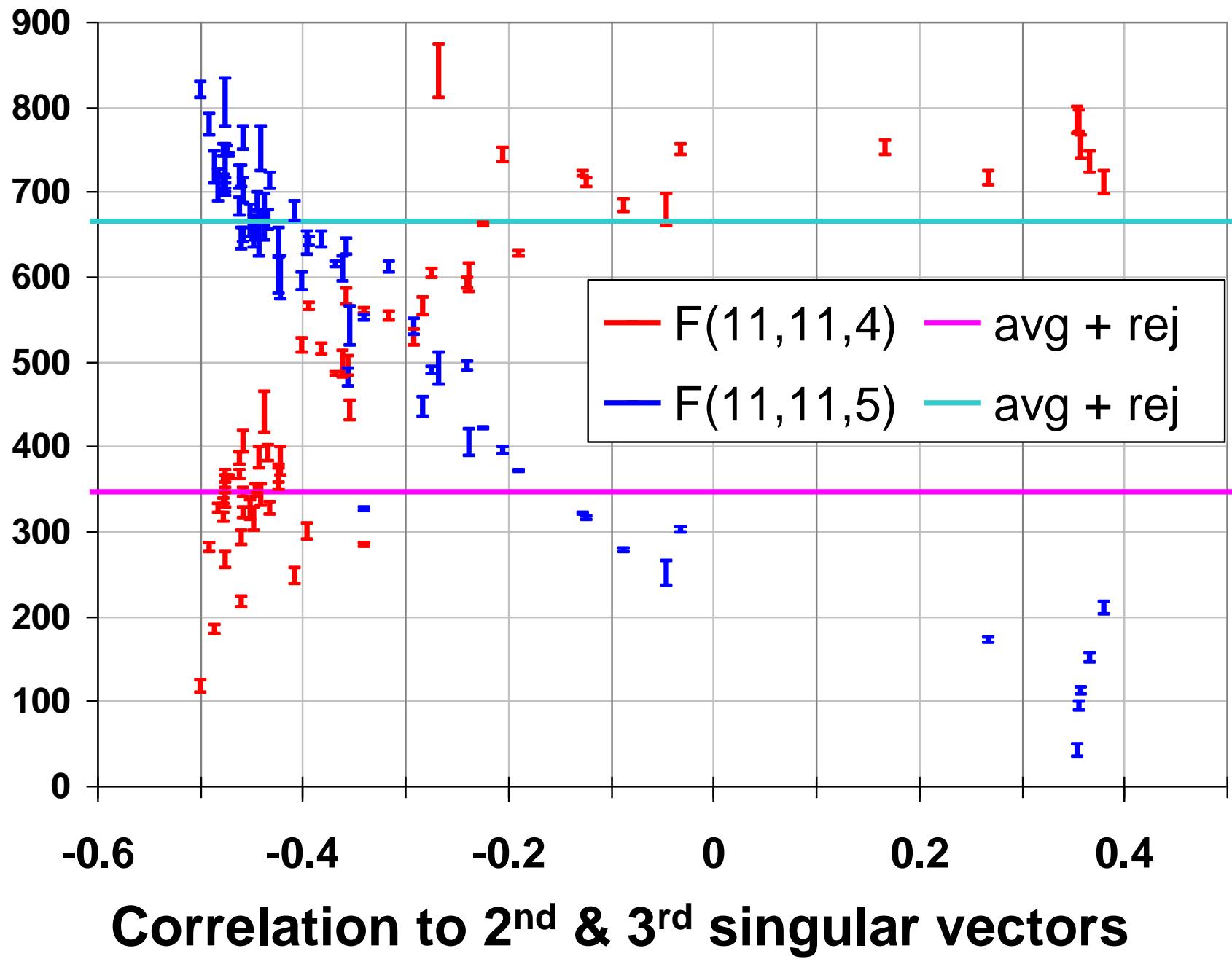


Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

Structure factor (electrons)



Decisions, Decisions, Decisions

- Exposure time
- Number of images
- Wavelengths
- Beam: size, flux, divergence, bandpass
- Inverse beam?
- Kappa? - overlaps
- Multiple crystals? - non-isomorphism

Run Strategy

URL Summary

[http://bl831.als.lbl.gov/~jamesh/powerpoint/
IGBMC_SvN_2016.pptx](http://bl831.als.lbl.gov/~jamesh/powerpoint/IGBMC_SvN_2016.pptx)

<http://bl831.als.lbl.gov/xtalsize.html>

<http://bl831.als.lbl.gov/xtallife.html>

100 ADU/pixel

10 μm for lysozyme

~3% error per spot, 1%/MGy

7235 eV for S-SAD on CCDs

“Attenu-wait” & dose slicing

CCP4: aimless log

\$TABLE: Analysis against resolution, XDSdataset:

\$GRAPHS:I/sigma, Mean Mn(I)/sd(Mn(I)):0|0.216023x0|137.14:2,13,14:

:Rmerge, Rfull, Rmeas, Rpim v Resolution:0|0.216023x0|1.70834:2,4,5,6,7:

:Average I, RMSdeviation and Sd:0|0.216023x0|1650.8:2,10,11,12:

:Fractional bias:0|0.216023x0|0:2,15:

\$\$

N	1/d^2	Dmid	Rmrg	Rfull	Rcum	Rmeas	Rpim	Nmeas	AvI	RMSdev	sd	I/RMS	Mn(I/sd)	Frd
1	0.0064	12.55	0.020	0.020	0.020	0.021	0.006	13115	1651	57	42	29.2	137.1	
2	0.0191	7.24	0.027	0.027	0.024	0.028	0.008	24753	1171	47	42	25.0	105.2	
3	0.0318	5.61	0.038	0.038	0.029	0.040	0.012	32197	857	46	43	18.4	79.6	
4	0.0445	4.74	0.034	0.034	0.031	0.035	0.010	37743	1212	57	53	21.4	91.2	
5	0.0572	4.18	0.036	0.036	0.032	0.038	0.011	42642	1181	59	57	19.9	83.8	
6	0.0699	3.78	0.049	0.049	0.036	0.052	0.015	47224	883	59	57	15.1	65.1	
7	0.0826	3.48	0.065	0.065	0.040	0.068	0.020	51052	685	59	58	11.7	50.9	
8	0.0953	3.24	0.096	0.096	0.045	0.100	0.029	54636	448	56	56	8.0	35.0	
9	0.1080	3.04	0.151	0.151	0.050	0.158	0.046	58072	268	53	53	5.1	22.7	
10	0.1207	2.88	0.229	0.229	0.056	0.240	0.070	60731	171	51	51	3.3	15.4	
11	0.1334	2.74	0.314	0.314	0.063	0.329	0.097	63807	125	51	51	2.4	11.3	
12	0.1461	2.62	0.406	0.406	0.070	0.425	0.125	66241	98	51	52	1.9	8.7	
13	0.1588	2.51	0.537	0.537	0.078	0.562	0.166	68272	76	53	53	1.4	6.5	
14	0.1715	2.41	0.627	0.627	0.081	0.597	0.207	71170	61	52	51	1.1	4.4	

CCP4: aimless log

\$TABLE: Correlations CC(1/2) within dataset, XDSdataset:

\$GRAPHS: Anom & Imean CCs v resolution:0|0.216023x0|1:2,4,7:

: RMS correlation ratio :0|0.216023x0|2.20344:2,6:

\$\$

N	1/d^2	Dmid	CCanon	Nanom	RCRanom	CC1/2	NImean	\$
1	0.0064	12.55	0.659	499	2.203	1.000	669	
2	0.0191	7.24	0.550	975	1.853	1.000	1155	
3	0.0318	5.61	0.527	1295	1.798	1.000	1479	
16	0.1970	2.25	0.037	2123	1.038	0.711	2275	
17	0.2097	2.18	0.043	1682	1.044	0.460	1877	

XDS: CORRECT.LP or XSCALE.LP

ION
COMPARED I/SIGMA R-meas CC (1/2) Anomal SigAno

Corr

3018	33.77	2.3%	99.9*	21*	1.012
4585	22.56	3.6%	99.8*	9	0.914
5327	19.99	4.0%	99.7*	9	0.859
6094	12.27	6.8%	99.3*	-1	0.784
7068	6.01	14.2%	97.8*	-2	0.799
8185	3.10	29.4%	88.8*	-4	0.776
8981	1.90	48.8%	75.9*	2	0.765
5991	1.14	87.3%	53.5*	-2	0.722
2520	0.59	170.4%	21.9*	4	0.693
51769	8.97	9.5%	99.5*	2	0.804

XDS: CORRECT.LP or XSCALE.LP

CORRECTION PARAMETERS FOR THE STANDARD ERROR OF REFLECTION I

The variance $v_0(I)$ of the intensity I obtained from counting statistics is replaced by $v(I)=a*(v_0(I)+b*I^2)$. The model parameters a , b are chosen to minimize the discrepancies between $v(I)$ and the variance estimated from sample statistics of symmetry related reflections. This model is an asymptotic limit $ISa=1/SQRT(a*b)$ for the highest $I/\Sigma(I)$. An experimental setup can produce (Diederichs (2010) Acta Cryst D66: 1026-1033).

a	b
3.806E+00	1.080E-04

ISa
49.32