



Methods for High Resolution Refinement in Single Particle Processing

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Single Particle Reconstruction with EMAN

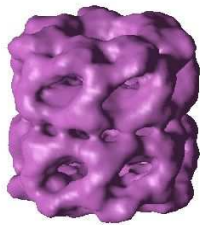
GroEL

Donghua Chen
Joanita Jakana
Wah Chiu

Jiu-Li Song (UT-SW Med)
David Chuang (UT-SW Med)

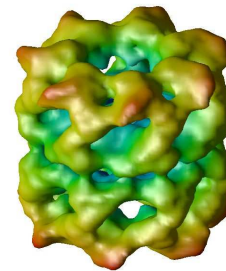
EMAN: <http://ncmi.bcm.tmc.edu/eman>

GroEL 2000 (15 Å)



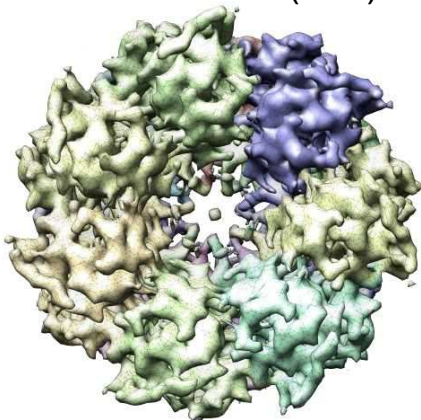
5000 particles, JEOL 4000

GroEL 2001 (11.5 Å)



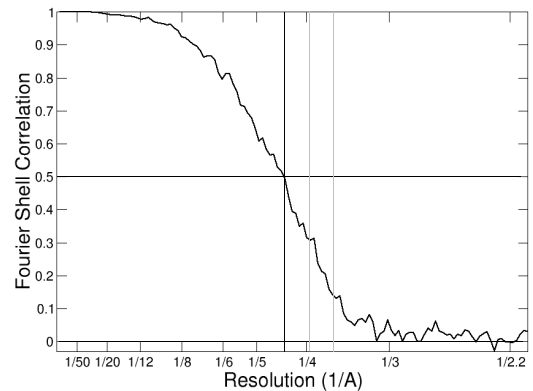
5000 particles, JEOL 4000

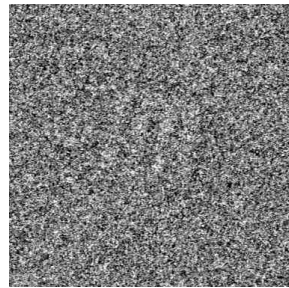
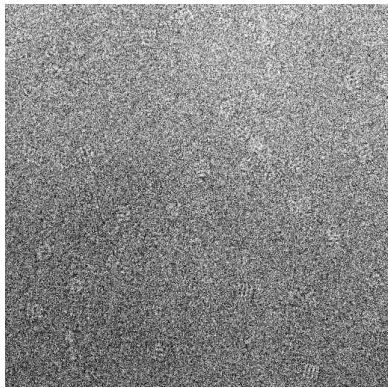
GroEL 2003 (6 Å)



30,000 particles, JEOL 2010F

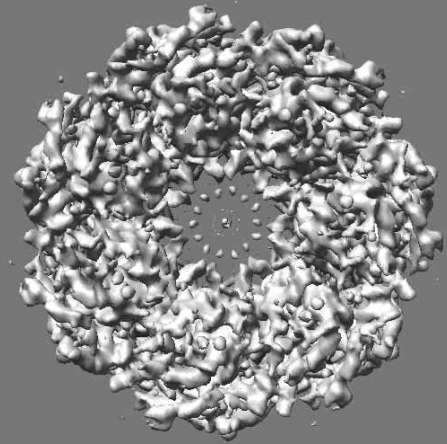
2005



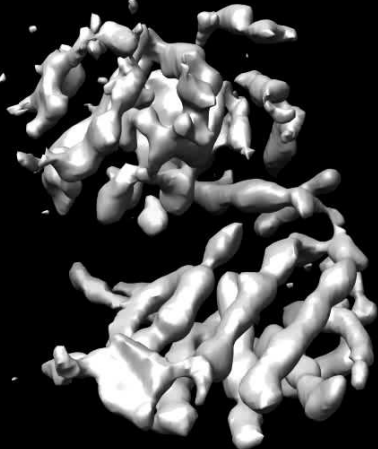


Jeol 3000
7 Days of imaging, 910 micrographs
1.06 Å/pix, Nikon 9000 scanner
135 used, 34,868 particles

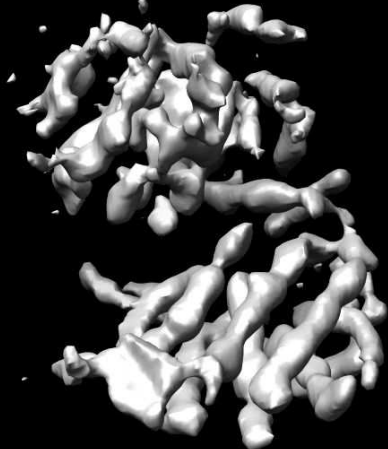
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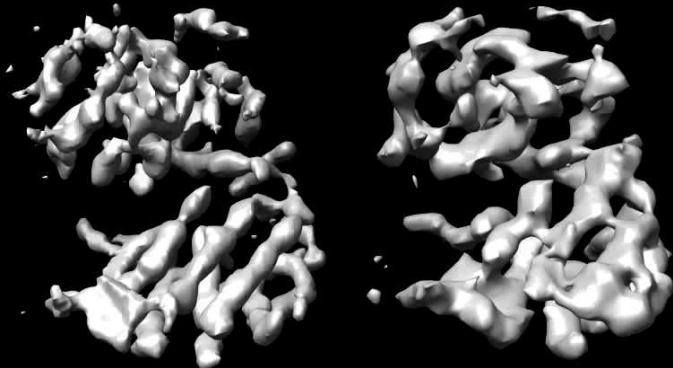
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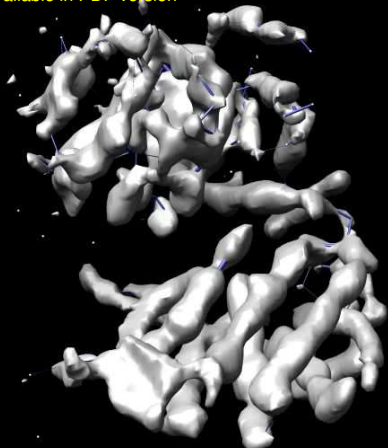
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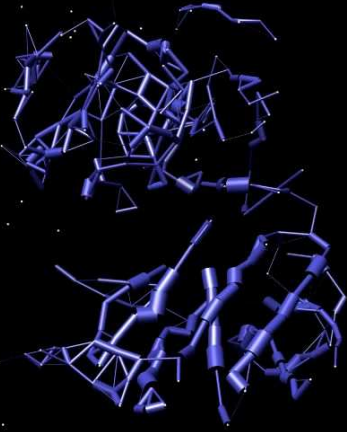
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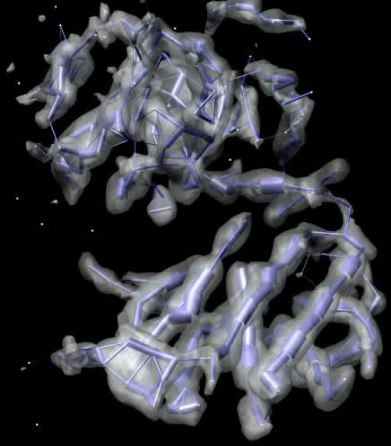
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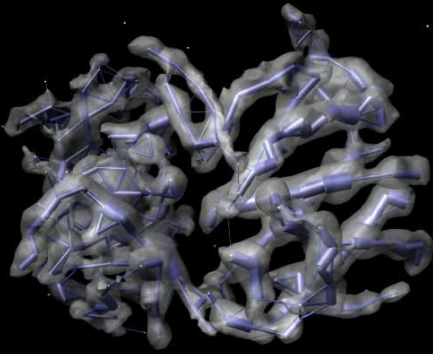
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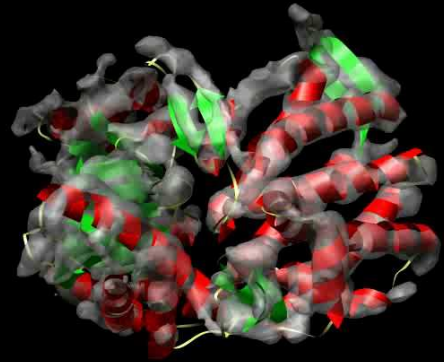
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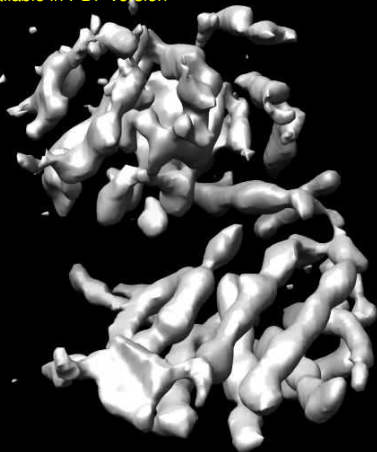
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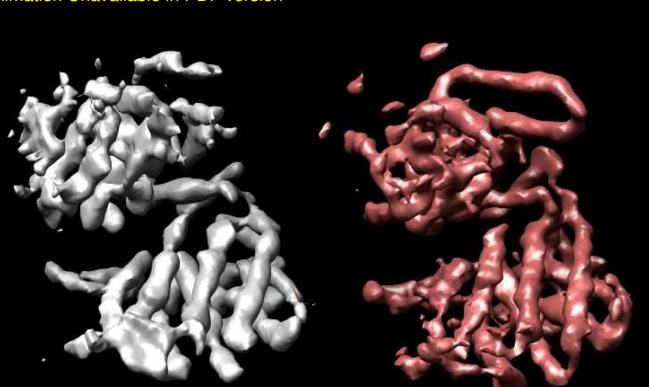
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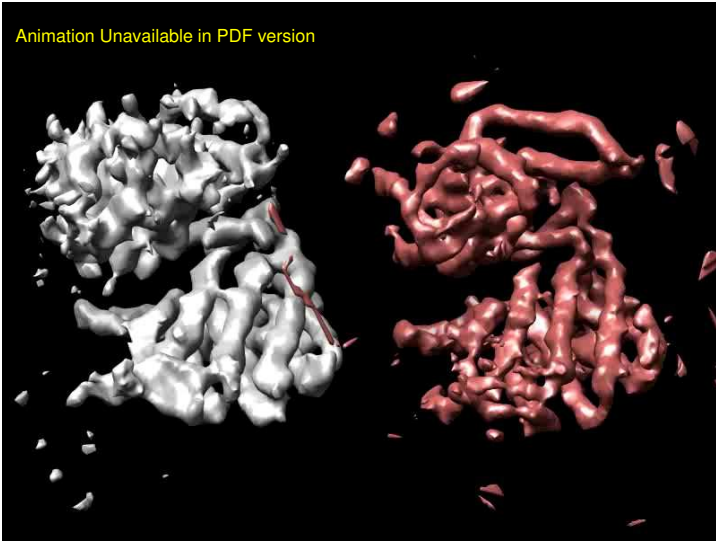
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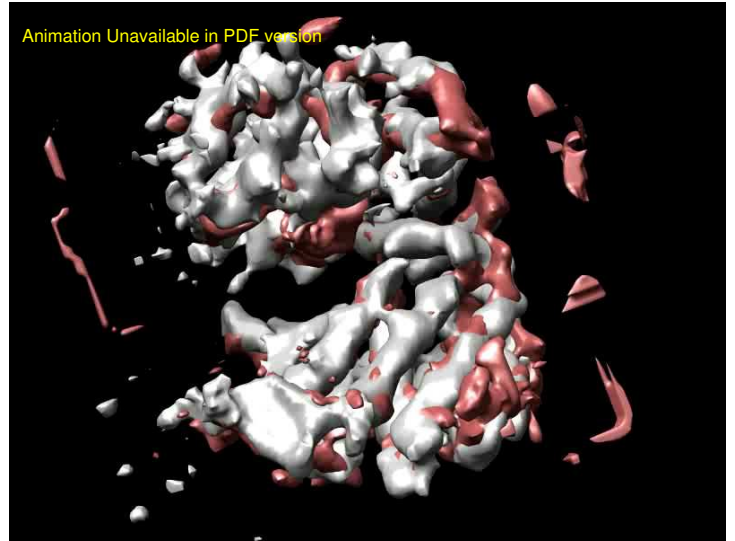
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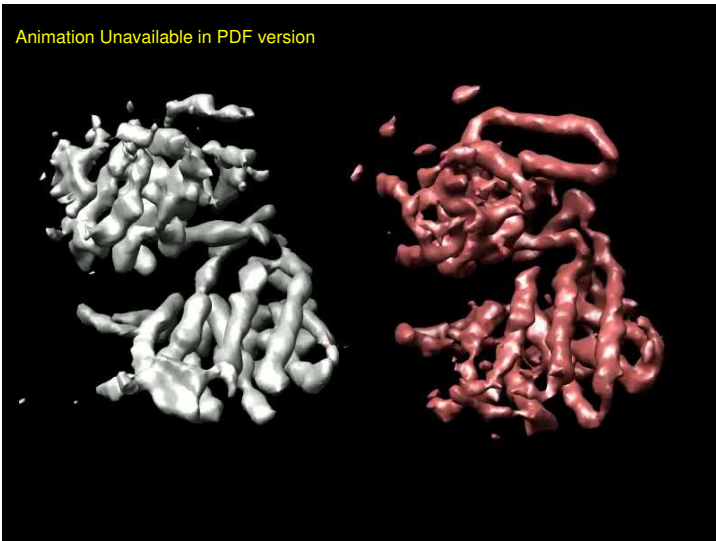
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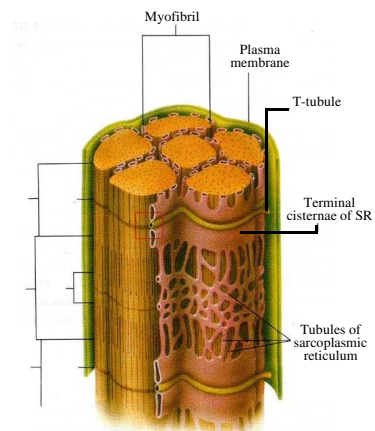


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Ca²⁺ Release Channel

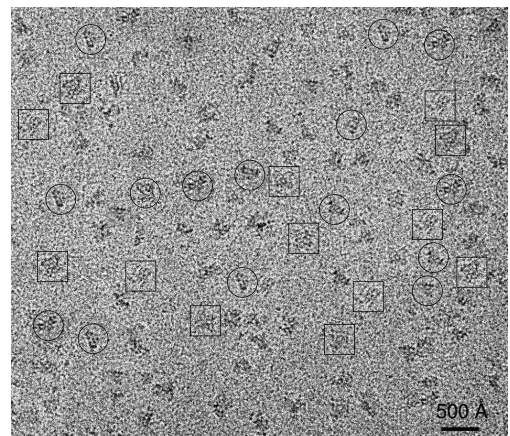
Irina Serysheva
Wah Chiu
Susan Hamilton



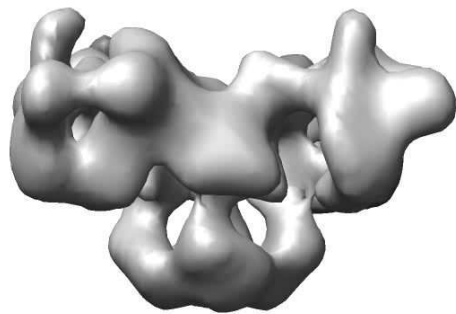
Ca²⁺ Release Channel

- SR membrane, triggered by DHPR in T-tubule
- Homotetramer
- ~2200 kDa
- Releases Ca⁺⁺ which initiates cross-bridge cycle

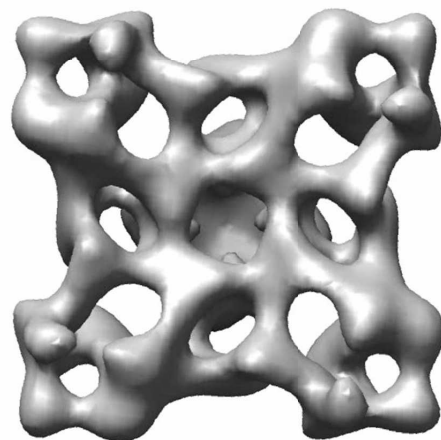
200 kV image of ice-embedded RyR1 (no continuous CF)



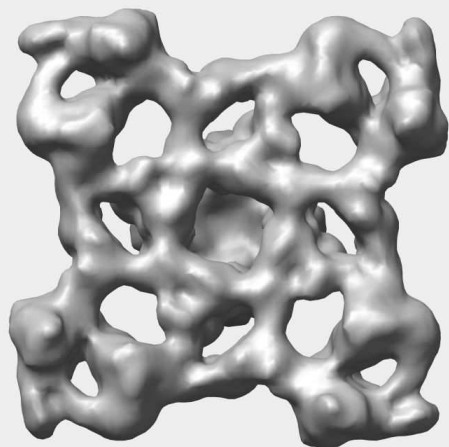
~30 Å Resolution



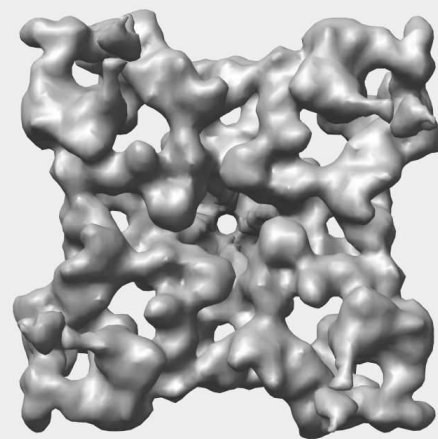
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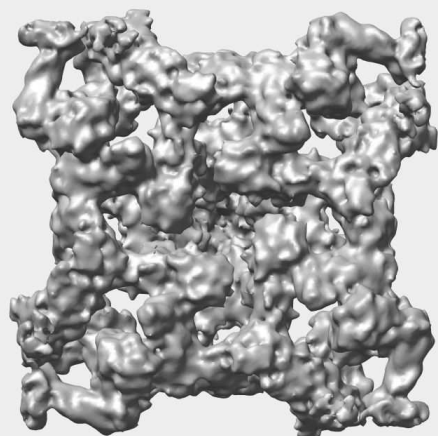
20 Å Resolution



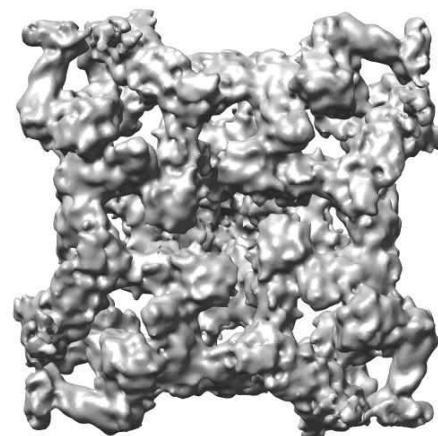
14 Å Resolution



9.6 Å Resolution

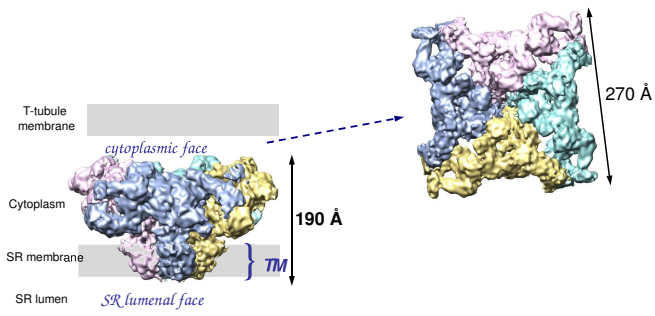
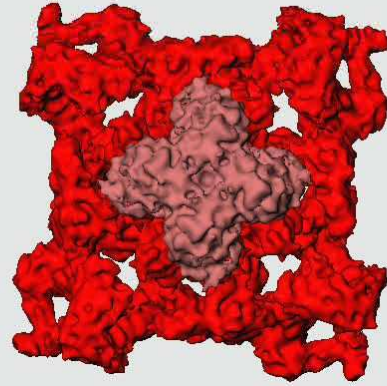


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Calcium Release Channel @9.6Å



Sequence assignment of observed helices

RyR1: 4864 - NKSEDEDEPMKCCDDMMTCYLFHMVYGVYRAGGGIGDEIEDPAGDEYELYRVVDFITFFVIVILLARIGLIDAFGELRDQOEVKEDMETK- 4957

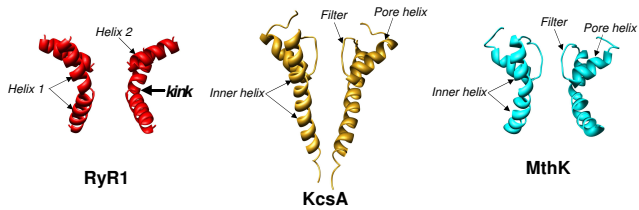
Filter
Hinge
M9 Helix 2
M10 Helix 1

KcsA: 36 - QLITYPRALWWSVETATTYVGGDLVPTLVWGRCAVAVVMVAGITSGFLVTAALATWVFGREG- 119

Filter
Pore helix
Inner helix

MthK: 45 - SWTVSLYWTFVITATVYGGYSPSTPLGMVFTYTLVIGLIGTFAVAVERLLEFLINREQ- 103

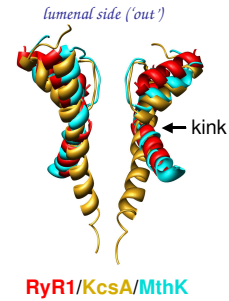
Filter
Pore helix
Inner helix



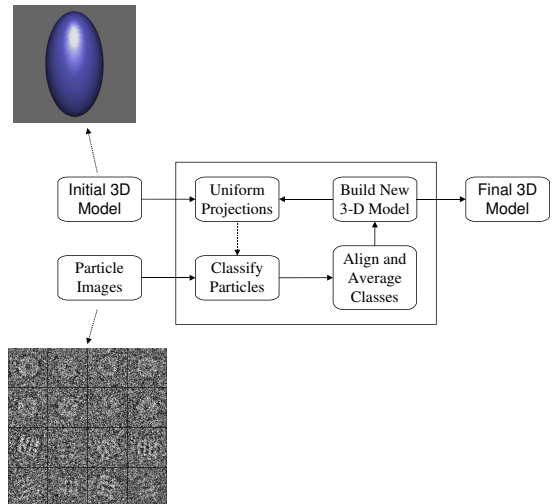
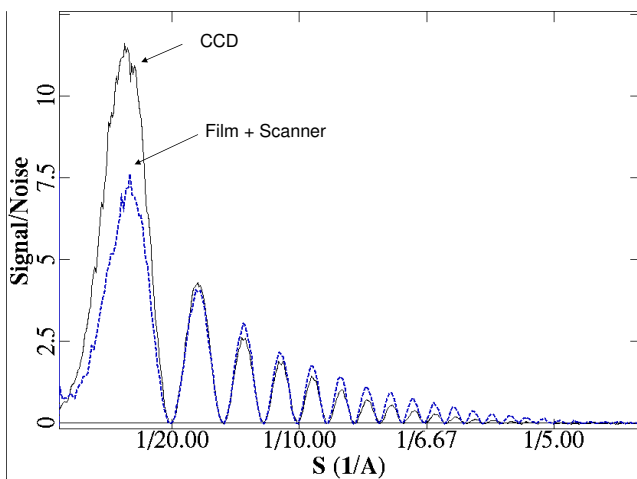
Sequence assignment of observed helices

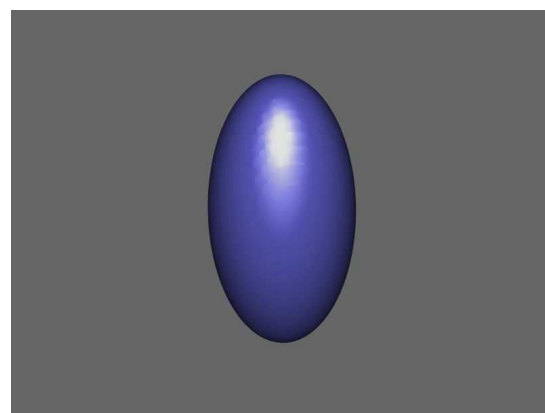
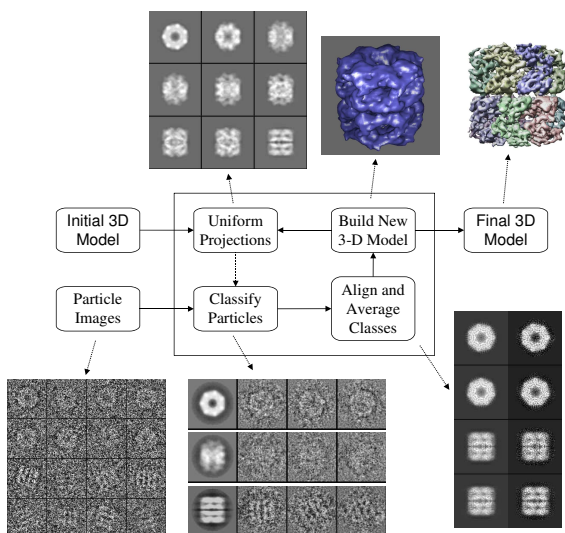
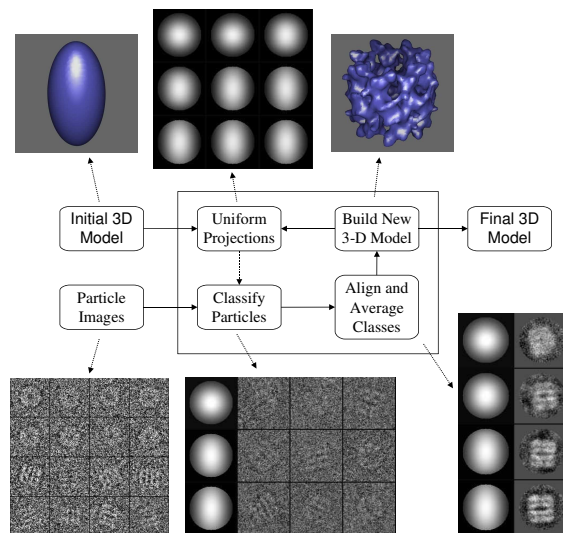
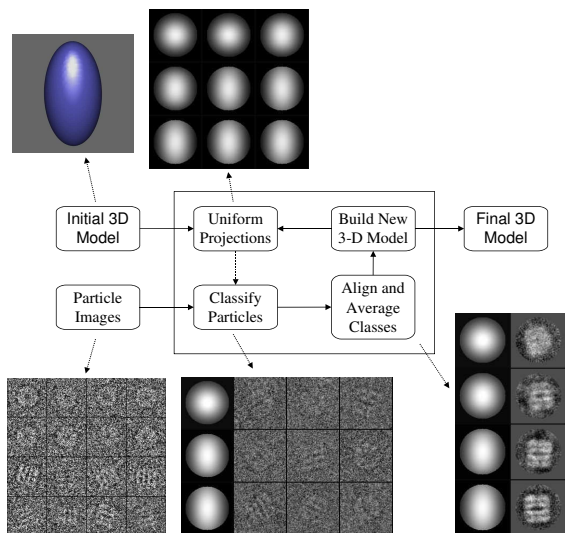
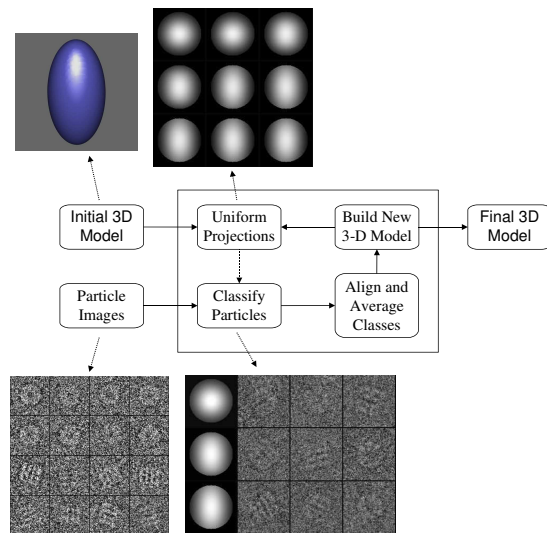
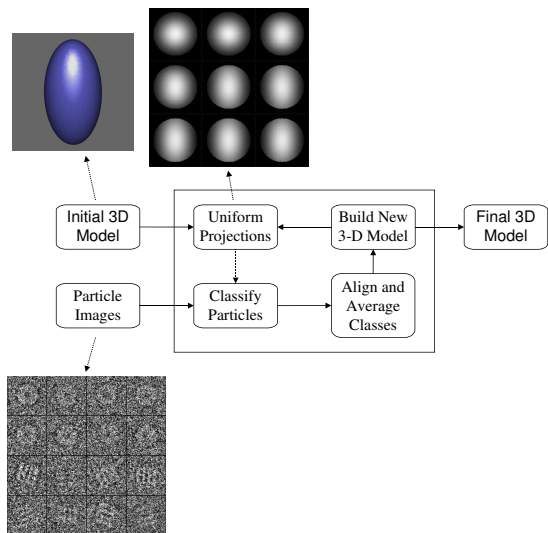
RyR1: 4864 - NKSEDEDEPMKCCDDMMTCYLFHMVYGVYRAGGGIGDEIEDPAGDEYELYRVVDFITFFVIVILLARIGLIDAFGELRDQOEVKEDMETK- 4957

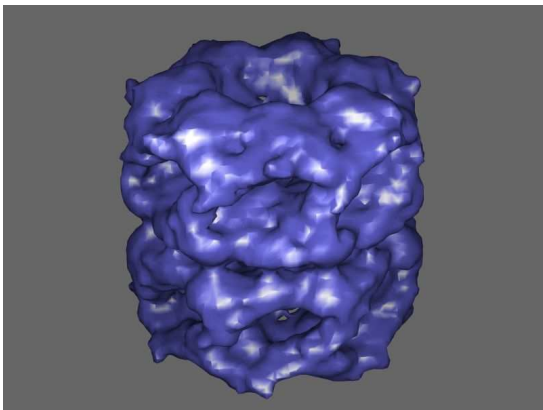
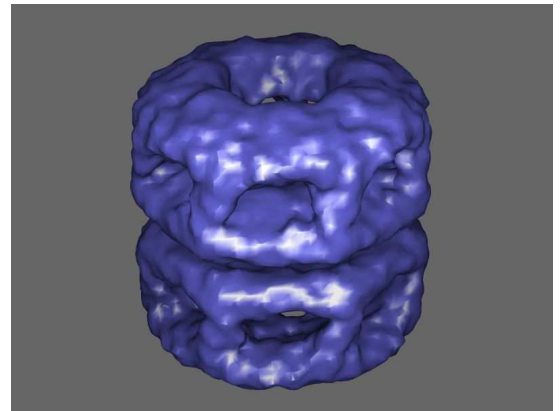
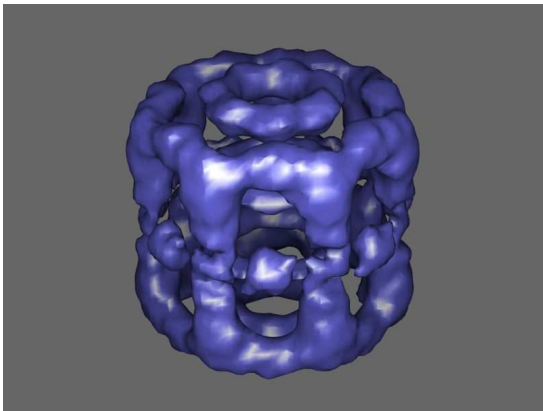
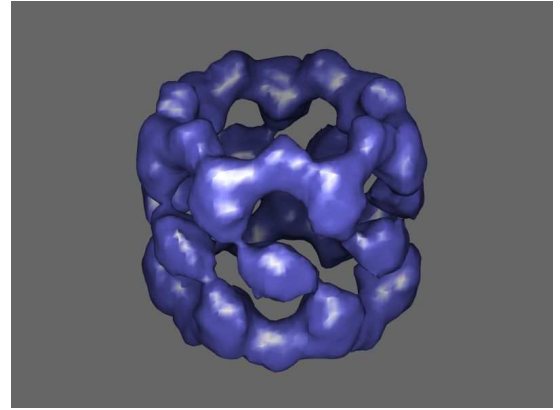
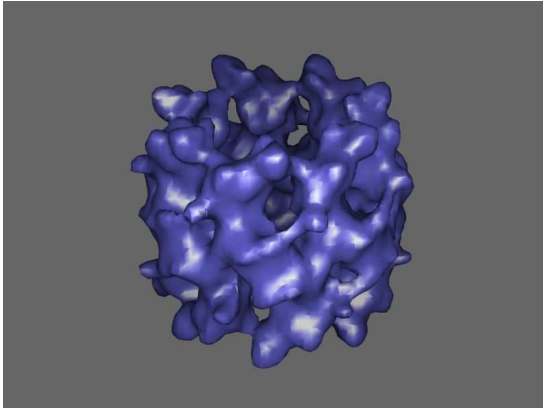
Filter
Hinge
M9
M10



CCD vs. Film







How do we get to Higher Resolutions?

- Get a better microscope
- Find a better microscopist
- Algorithm Improvements

Contrast Transfer Function

$$\overline{M}(s, \theta) = \overline{F}(s, \theta) C(s) E(s) + \overline{N}(s, \theta)$$

$$C(s) = \sqrt{1 - Q^2} \sin \gamma + Q \cos \gamma$$

$$\gamma = -\pi \left(\frac{1}{2} C_s \lambda^3 s^4 - \Delta Z \lambda s^2 \right)$$

$$E(s) = e^{-B s^2}$$

$$|N|^2 = n_1 e^{n_2 s + n_3 s^2 + n_4 \sqrt{s}}$$

$$M(s)^2 = F(s)^2 C(s)^2 E(s)^2 + N(s)^2$$

8 Parameters

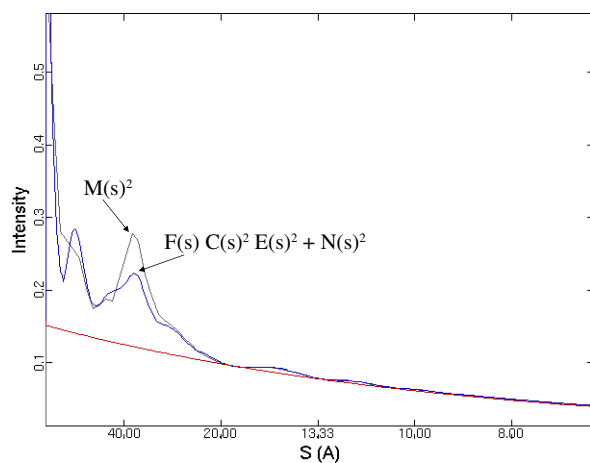
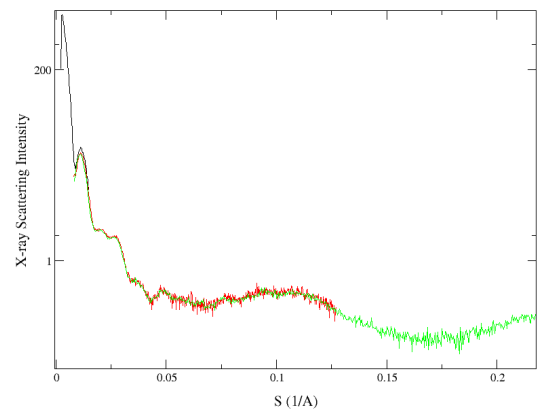
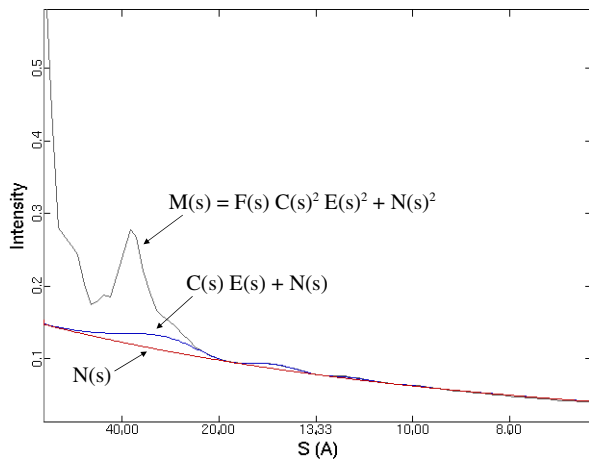
ΔZ - Defocus

Q - Amplitude Contrast

B - Gaussian Envelope Width

k - Signal Amplitude

n_{1-4} - Noise Parameters



CTF Correction

$$\overline{T}(s, \theta) = \sum_i k_i \overline{M}_i(s, \theta)$$

$$k_i = ?$$

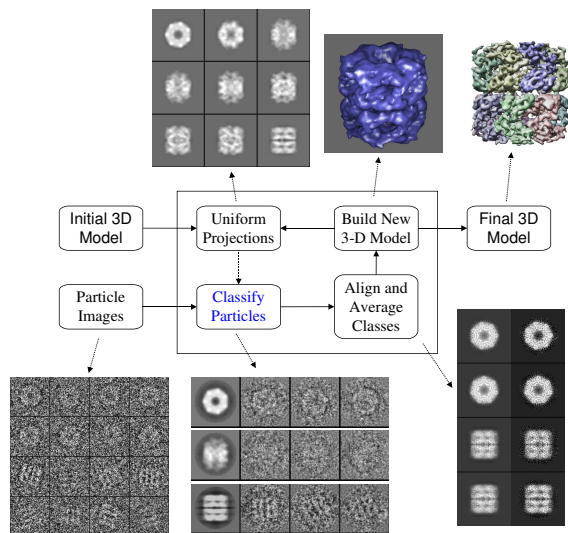
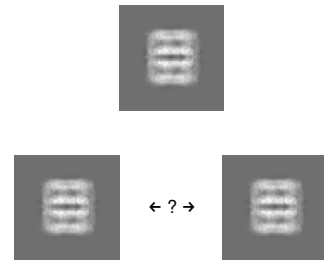
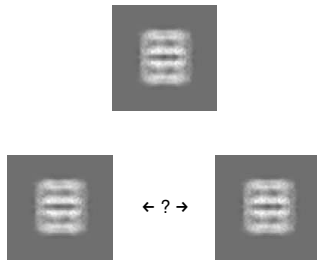
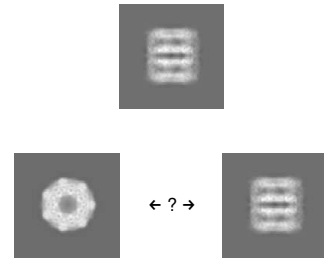
- Maximize SNR of $T(s, q)$
- Minimize variance between $T(s, q)$ and $F(s, q)$

CTF Correction

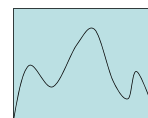
$$\bar{T}(s, \theta) = \frac{\overset{\text{Wiener Filter}}{F^2(s)R(s)}}{1 + F^2(s)R(s)} \sum_i \frac{\overset{\text{CTF Correction}}{1}}{C_i(s)E_i(s)} \frac{\overset{\text{SNR Weight}}{R_i(s)}}{R(s)} \bar{M}_i(s, \theta)$$

$$R_i(s) = \frac{C_i^2(s)E_i^2(s)}{N_i^2(s)} \quad R(s) = \sum_i \frac{C_i^2(s)E_i^2(s)}{N_i^2(s)}$$

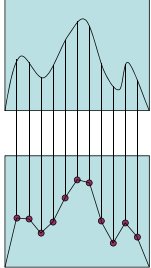
Image Classification



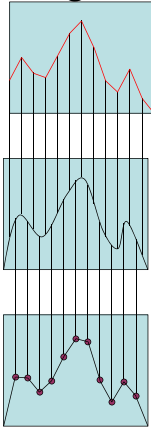
Alignment/Registration



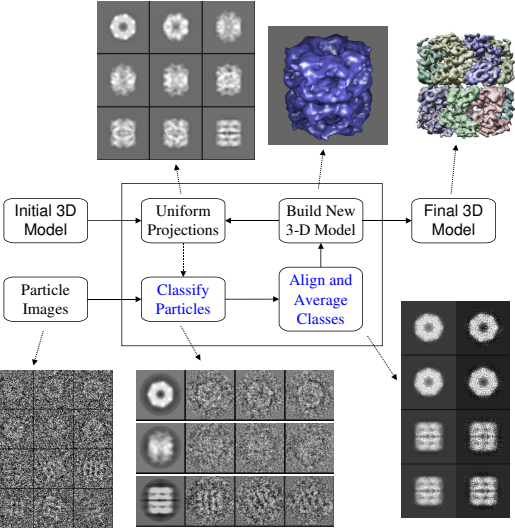
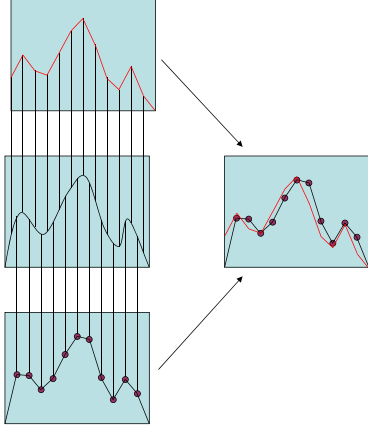
Alignment/Registration



Alignment/Registration

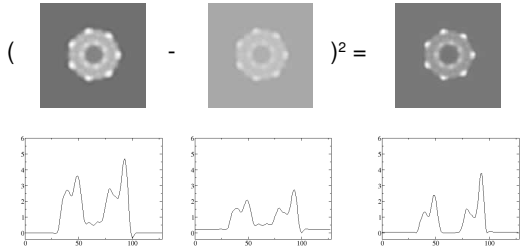


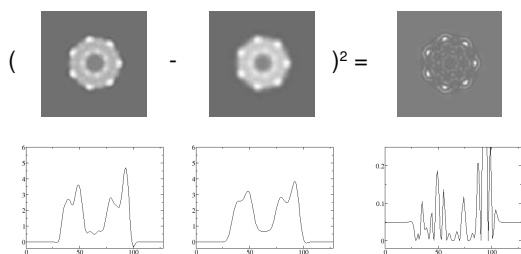
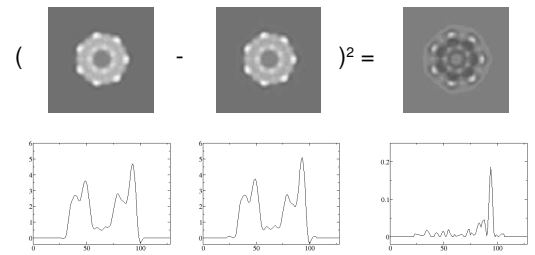
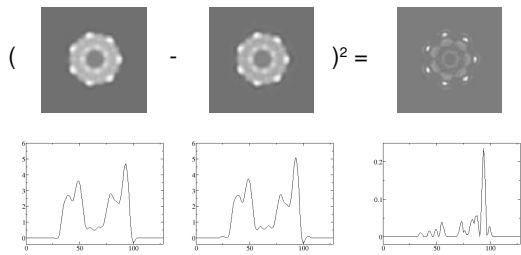
Alignment/Registration



Measures of Similarity

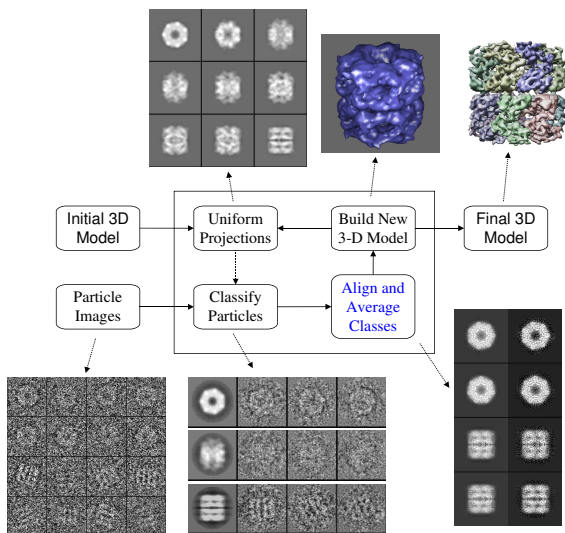
- Correlation Coefficient
- Variance (transformed density)
- Variance (matched filter)
- Phase Residual
- Mutual Information
- etc.



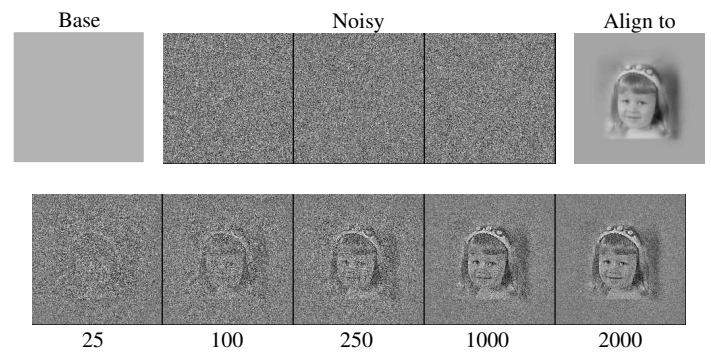


And the Answer is...

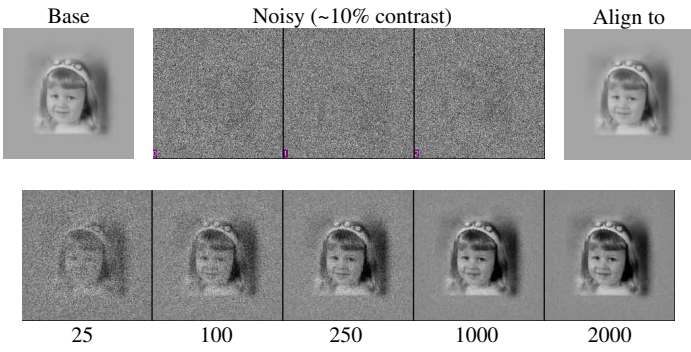
- Wiener filter particle
- Filter reference to match
- Normalize reference density to particle
- Calculate variance



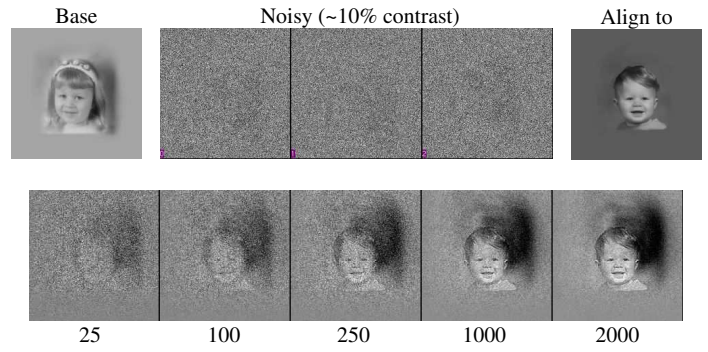
Model Bias



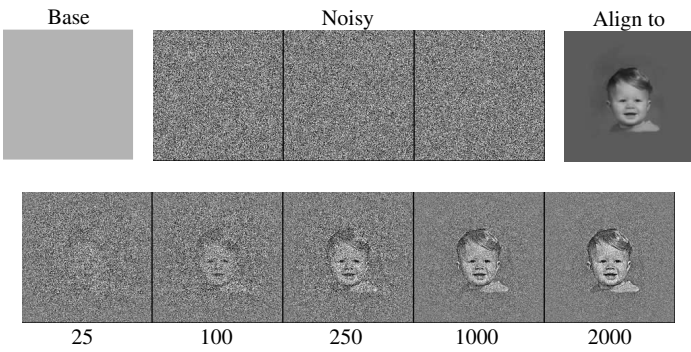
Model Bias



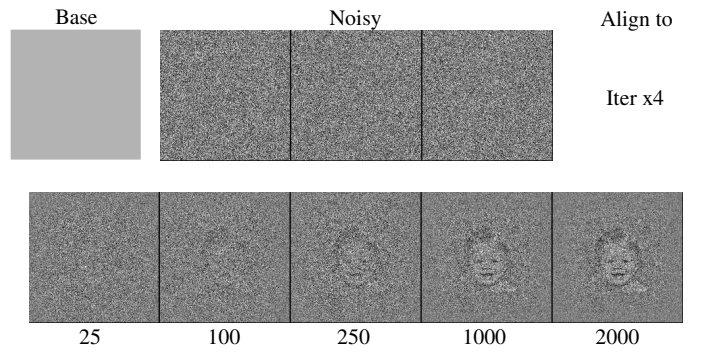
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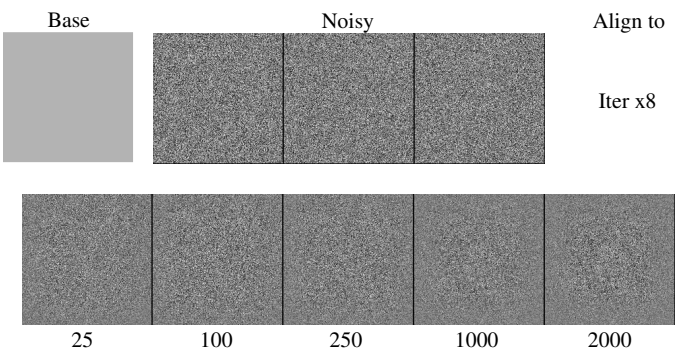
Model Bias



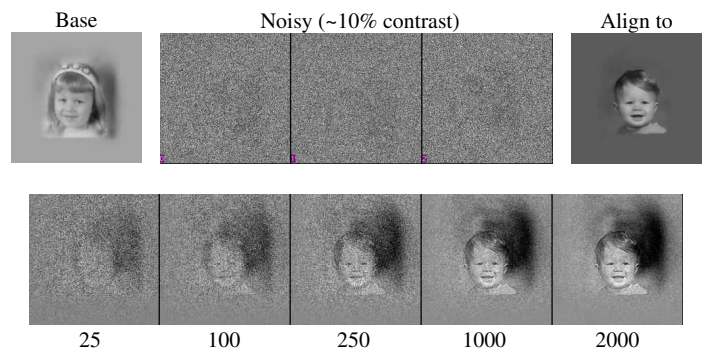
Model Bias



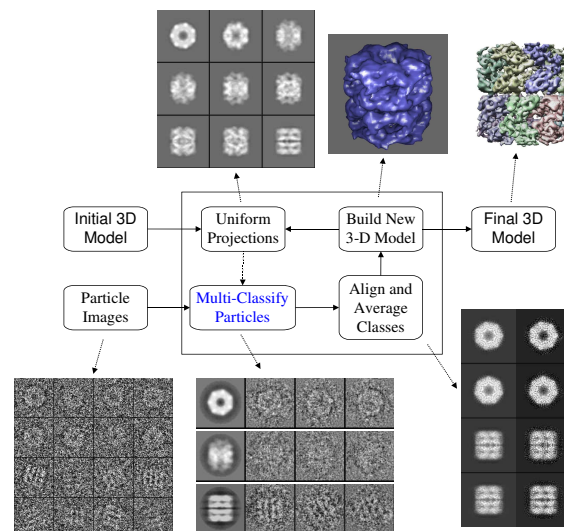
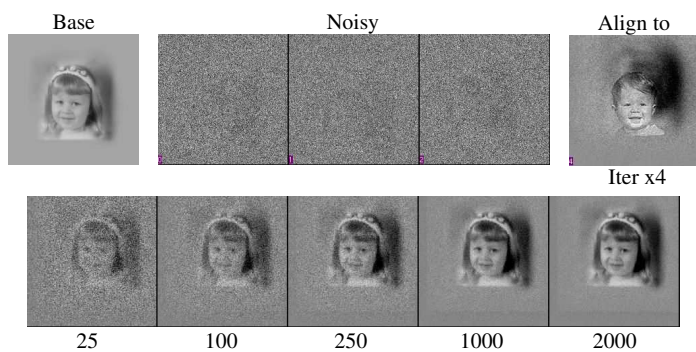
Model Bias



Model Bias



Model Bias



- Each particle -> best n classes
- More restrictive exclusion from class-avg

The Future

- Better similarity criteria
- Improved CTF model
- Per-particle CTF (at least defocus)
- Beam tilt
- Better 3-D reconstruction
- New refinement methodologies