## Monolayer Purification for Single Particle EM



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## Historical Background

- Uzgiris and Kornberg (1983) 2D xtals of Ab on lipid Antigen
- Darst et al., 1988 RNA polymerase
- Avila-Sakar et al., 1994 50s ribosome subunit
- Kubalek et al., (1994)
   His-tagged HIV1 RT



Dietrich and Venien-Bryan (20

#### Monolayer Structures

290 projection maps

#### 15 Cryo-EM

#### Only 7 Cryo-EM

3D Reconstructions

## Current Specimen Limitations

- Screening conditions
- Fragile transfer step
- Specimen Flatness
- Alternative approach = Single particle EM

A combinatorial approach for protein purification / EM structural studies

- Develop the Ni-NTA monolayer technique as a novel purification method
- Single Particle Cryo-EM for 3D reconstruction
- Apply the methodology to a real system

## Monolayer Purification Setup

1) Add protein sample well Cell lysate w/ target (25 µl) 50mM Hepes+150 mM NaCl+imidazole

#### 2) Cast monolayer

Filler / Ni-NTA lipid (1mg/ml
in CHCl<sub>3</sub>)
Apply 1µl Mix w/ Hamilton
syringe
Incubate 15 - 30 min., 4°C
3) Sample with EM
grid

Apply clean EM grid Grid bar side on monolayer



## Basic Considerations

- Biological sample preparation
- Grid preparation
- Transfer Step
- Neg. stain screening
- Vitrification
- Low-dose Imaging

## Preparation of Cell Extracts





QuickTime™ and a TIFF (Uncompressed) decompressor are needed to see this picture.

Insect cells (Sf9)

Bacteria Mammalian cells (E.coli, BL21) (293T)

- Grow His-tagged construct
- Lyse cells with lysozyme, sonication
- Obtained cleared lysate; ML input

## Quantifoil Grid Preparation



Whatman #1 paper Saturated w/ Ethyl Acetate o/n in hood

Bake for 1 hr At least 100°C

## Transfer Methods



## Direct Transfer vs. Loop transfer



# 7Å Projection Map of SbpA



Norville et al., JSB (in

# Transfer of 2D crystal vs Single particles

- Crystals = Loop transfer / 5% Trehalose embedding /ethane
- Particles = Direct transfer /ethane

## Basic Considerations

- Biological sample preparation
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  Neg. stain screening
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# Negative Stain Screening Test specimen: Tf-TfR complex





#### Transferrin-Transferr Receptor Complex



## Negative Stain Screening



## Screening Monolayer Purified Tf-TfR complex from Sf9 extracts



In Soln.

2% ML

2% ML + 50 mM Imid.

# Neg. stain reconstruction using RCT



## Vitrification of ML specimens

- Place "Grid bar" on top of ML
- Remove grid w/ forceps
- Blot 3µl subphase



 Plunge into ethane

#### Manual vs. Automated Freezing

• Manual

- Vitrobot (F
- Uncontrolled Environment (22
  - Environment (22°C, 65% rh)

QuickTime™ and a TIFF (Uncompressed) decompressor are needed to see this picture.

- Calibrate blotting
- Blotting time  $\propto$  volume (µl)
- Consistent ice over entire grid
  - Gradient of vitreous ice

## Low-dose imaging

- Tecnai F-20 operating at 200 kV
- Quantifoil 2/1 (2 $\mu m$  holes, 1 $\mu m$  spacing)
- Magnification = 50,000x
- Defocus = -2 to -5  $\mu$ m
- 10 e<sup>-</sup> /  $Å^2$ , 1 sec exposure
- Images on Film, scanned w/ 7μm step (1.4 Å / pixel, 3 x 3 sub-sampling)

# Imaging Monolayer Purified Tf-TfR from Sf9 extracts



In Soln.

2% ML

20% ML

## Leginon for Screening ML Cryo-EM specimen



Taylor et al., JSB (in press)

## Hole selection based on radial density function



## High-throughput Potential



#### Current interests

• Adapt ML method for use with membrane proteins and other tags

 3D reconstructions of native complexes
 large data sets and automated routines

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