

CryoET of Single Particle CryoEM Grids Reveals Widespread Particle Adsorption to the Air-Water Interface, Which May be Reduced with New Plunging Techniques

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Single particle cryo-electron microscopy (cryoEM) has commonly been performed with the assumption that the protein particles were suspended safely between the air-water interfaces at the time of vitrification. Studies of dozens of single particle cryoEM samples on grids by cryo-electron tomography (cryoET) has revealed that the vast majority of particles are adsorbed to the air-water interfaces at the time of vitrification (Figure 1) [1].

Particle adsorption to the air-water interfaces may potentially cause particle preferred orientation, conformational changes, and particle degradation [1]. While particle adsorption to the air-water interfaces may be physically avoided using substrates [2, 3], an alternative method for reducing or possibly eliminating air-water interface interactions is to decrease the time between sample-to-grid application and plunge freezing sufficiently so as to limit the extent of particle diffusion to the air-water interfaces and reduce the amount of equilibration of particles adsorbed to the air-water interfaces.

Here we present three single particle specimen (apoferritin, hemagglutinin, and insulin receptor) prepared using Spotiton with varying sample application to plunge freezing times (spot-to-plunge times) (Figure 2). With a spot-to-plunge time of 400 ms, apoferritin, a protein complex with high symmetry, preferentially adsorbs to the air-water interfaces (Figure 2a), with few particles remaining non-adsorbed. This preferential adsorption is reduced significantly with a spot-to-plunge time of 100 ms (Figure 2b).

Both hemagglutinin and insulin receptor, at both long and short spot-to-plunge times, remain preferentially adsorbed to the air-water interfaces prior to vitrification. When hemagglutinin and insulin receptor are prepared with long spot-to-plunge times (800 and 600 ms, respectively), they both present a limited number of preferred orientations (Figure 3c). However, with spot-to-plunge times of 100 and 200 ms, respectively, a significant reduction in preferred orientations was seen (Figure 3d). With these three samples we show the potential for reducing and potentially eliminating air-water interactions.

References:

- [1] A Noble *et al*, BioRxiv (2017).
- [2] R Glaeser, Current Opinion in Colloid & Interface Science (2017).
- [3] TG Martin *et al*, PNAS **113** (2016), p. 7456.
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Sample Name	Example cross-sectional schematic diagram	Sample Name	Example cross-sectional schematic diagram	Sample Name	Example cross-sectional schematic diagram	Sample Name	Example cross-sectional schematic diagram
32 kDa Kinase		Neural Receptor		IDE		Apoferritin (0.5 mg/mL)	
Hemagglutinin		Protein with Bound Lipids (deglycosylated)		GDH		Apoferritin with 0.5 mM TCEP	
HIV-1 Trimer Complex 1		Protein with Bound Lipids (glycosylated)		GDH		Protein with Carbon Over Holes	
HIV-1 Trimer Complex 1		Lipo-protein		GDH + 0.001% DDM (2.5 mg/mL)		Protein and DNA Strands with Carbon Over Holes	
HIV-1 Trimer Complex 2		GPCR		DnaB Helicase-helicase Loader		T20S Proteasome	
Stick-like Protein 1		Rabbit Muscle Aldolase (1mg/mL)		Apoferritin		T20S Proteasome	
Stick-like Protein 2		Rabbit Muscle Aldolase (6mg/mL)		Apoferritin		T20S Proteasome	
Neural Receptor		Protein in Nanodisc (0.58 mg/mL)		Apoferritin		Mtb Proteasome	
				Apoferritin (1.25 mg/mL)		Protein on Streptavidin	

Figure 1. Single particle cryoEM grids plunged with a sample application to freeze time on the order of 1 second. The vast majority of particles are adsorbed to an air-water interface.

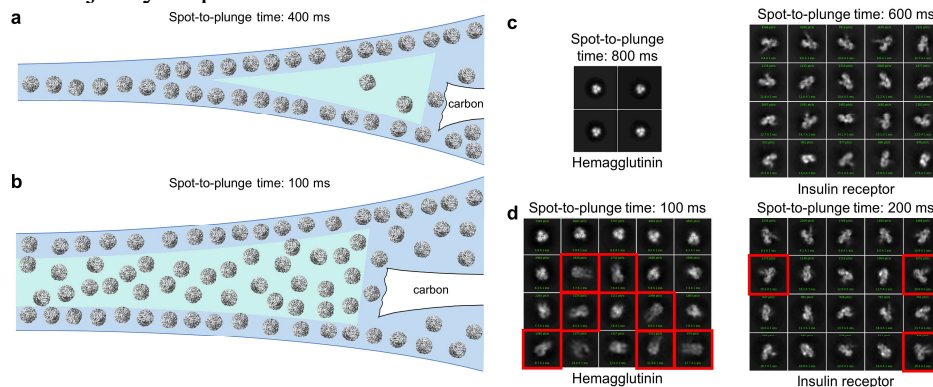


Figure 2. Effects of longer and shorter spot-to-plunge times using Spotiton. Cross-sections of tomograms of apoferritin grids prepared with a spot-to-plunge time of 400 ms (a) and 100 ms (b). The preparation in (b) resulted in more than an order of magnitude greater number of non-adsorbed particles than in (a). Long spot-to-plunge times of hemagglutinin and insulin receptor (c) resulted in limited number of preferred orientations, while shorter times (d) resulted in many more orientations (red boxes).