



## Life Sciences Seminar

# Single-Particle Cryo-EM: A rising star in structural biology

**Doreen Matthies**

National Institutes of Health (NIH)

Host:

Cryo-EM is currently revolutionizing the field of structural biology. Due to recent advances in detector technologies, hardware & software improvements, this method is becoming more and more mainstream in solving high-resolution structures. It complements methods such as X-ray crystallography & NMR spectroscopy with the advantage of not having to form well-ordered crystals or having an upper size limit. Progress in the field of single-particle cryo-EM will be discussed with examples of soluble & membrane protein complexes yielding resolutions of better than 3 Å. A study on 200 kDa Mg<sup>2+</sup> channel CorA will be used as a detailed example. CorA is a homo-pentamer and is gated by intracellular Mg<sup>2+</sup>. We determined the structures of CorA in the presence and absence of Mg<sup>2+</sup> using single-particle cryo-EM. The five-fold symmetric Mg<sup>2+</sup>-bound closed state of CorA at an overall 3.8 Å resolution reveals side-chain densities as well as densities for several magnesium ions. Cryo-EM of Mg<sup>2+</sup>-free open state CorA revealed dramatic differences in the conformation of the channel with large asymmetric changes in the soluble domain of the pentameric complex suggesting an unprecedented gating mechanism.

**Monday, January 16, 2017 08:45am - 09:45am**

Mondi Seminar Room 2, Central Building



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