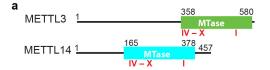
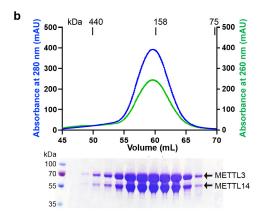
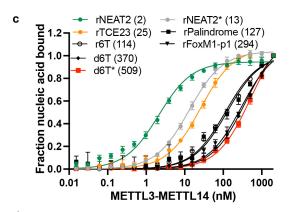
Project title: Structure of a human RNA methyltransferase complex

PI: Yogesh K Gupta, PhD

## **Preliminary data:**







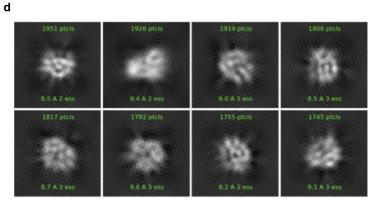


Figure 1. a. Domain architecture of human METTL3 and METTL14 methyltransferases. b. Chromatogram of final size exclusion chromatography step of purification (upper panel) human METTL3-METTL14 showing complexes (expressed in insect cells) co-eluted as single homogenous species. Blue, absorbance at 280 nm; green, absorbance at 260 nm (A260). Coomassie stained gels (lower panels) high purity of METTL3-METTL14 proteins in the SEC peak fractions. c. FP-based binding assay for DNA and RNA oligos showing highest affinity for rNEAT2 RNA (green), and lowest affinity for d6T\* oligo (red). Equilibrium dissociation constants (Kd) for each oligo are shown on the right side of the isotherms. The data were fit into one site specific binding model (Y=Bmax\*X/(Kd+X). d. Various 2D class)averages of METTL3-METTL14 complex (apo) calculated from KRIOS data measured at NCCAT, New York.