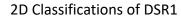
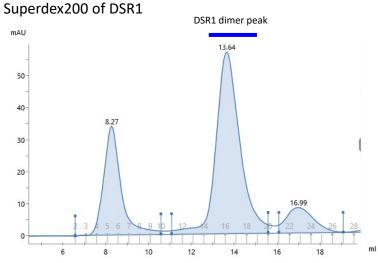
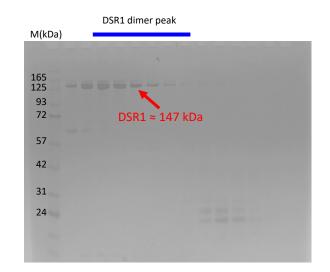
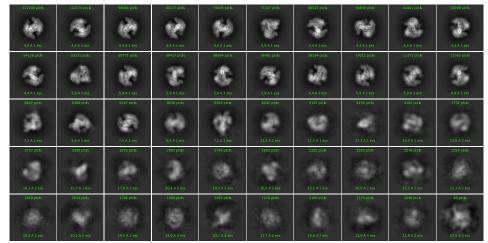
The purification and cryo-EM structure of DSR1 (E. Coli) at 3.16 Å

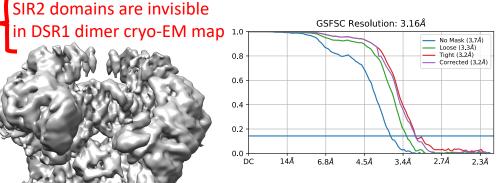




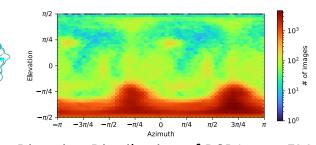
Buffer: 25mM Tris, pH=7.5, 250mM NaCl, 2mM betaMe, 0.002% DDM





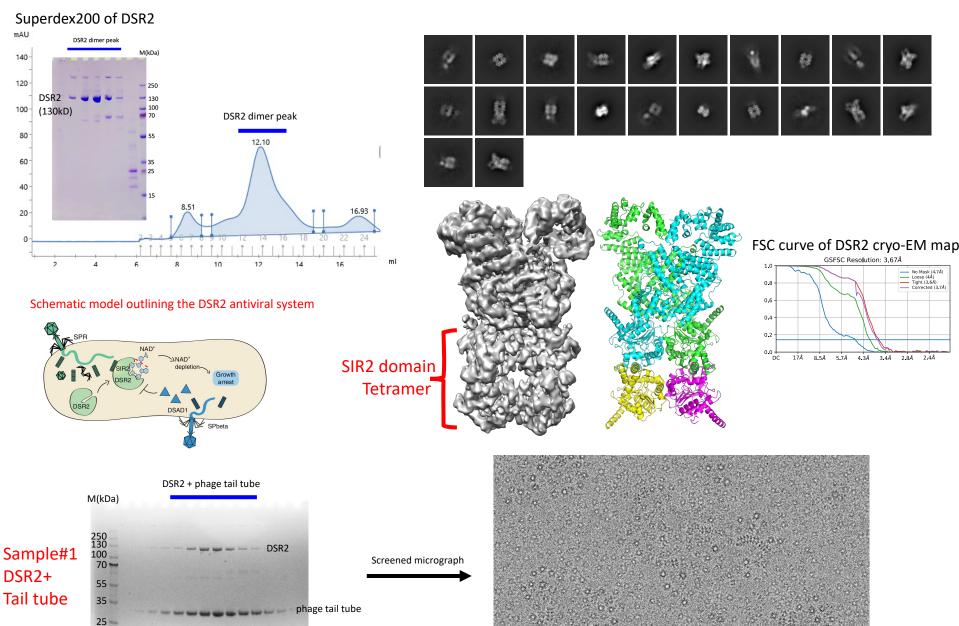


FSC curve of DSR1 cryo-EM map



Direction Distribution of DSR1 cryo-EM map

The purification and cryo-EM structure of DSR2 (B. Subt) at 3.67 Å and our sample of DSR1+tail tube



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