
dataset	Eco RNAP-Sal- Br
beamline	CHESS-F1
space group	P2 ₁ 2 ₁ 2 ₁
resolution range	50.00-4.70 Å (4.78-4.70 Å)
cell parameters (Å, °)	a=186.2, b=208.2, c=308.2 α=90.0, β=90.0, γ=90.0
completeness	1.000 (1.000)
multiplicity	16.6 (16.9)
mean I/σ	17.3 (3.4)
Rmerge	0.200 (0.962)

Supplementary file 3. Crystal structure of *E. coli* RNAP holoenzyme in complex with a bromine-containing Sal derivative: crystallization and refinement statistics.