Recovering Ensembles of Chromatin Conformations from Contact Probabilities – Supplementary Data Dario Meluzzi and Gaurav Arya

Table S1. Harmonic restraints used when simulating test systems and when performing the ensemble recovery procedure.

| | Test system ^a | | | | Ensemble recovery | | | |
|--|--------------------------|---------|------------------|-------------|-------------------|--------|---------------------------|--------------------|
| N^{b} | Loops ^c | n^* d | i,j ^e | $k_{i,j}$ f | n^{g} | i,j h | $k_{i,j}(2)^{\mathrm{i}}$ | $k_{i,j}(n+1)^{j}$ |
| 25 | 2 free | 3 | 13, 23 | 2.5 | 3 | 13, 23 | 2.462 | 2.506 |
| | | | 3, 13 | 2.5 | | 3, 13 | 2.552 | 2.544 |
| <u></u> | | | 3, 23 | 2.5 | | 3, 23 | 2.558 | 2.544 |
| 25 | 2 tied | 4 | 13, 23 | 2.5 | 4 | 13, 23 | 2.564 | 2.518 |
| | | | 3, 13 | 2.5 | | 3, 13 | 2.558 | 2.591 |
| 7. ; У. | <u> </u> | | 3, 23 | 2.5 | | 3, 23 | 2.406 | 2.521 |
| | | | 8, 18 | 2.5 | | 8, 18 | 2.489 | 2.411 |
| 35 | 3 free | 4 | 13, 23 | 2.5 | 6 | 13, 23 | 2.539 | 2.634 |
| | | | 23, 33 | 2.5 | | 23, 33 | 2.509 | 2.485 |
| | | | 3, 13 | 2.5 | | 3, 13 | 2.512 | 2.494 |
| | | | 3, 33 | 2.5 | | 3, 33 | 2.500 | 2.531 |
| | | | | | | 13, 33 | 0.000 | 0.000 |
| | | | | | | 3, 23 | 0.000 | 0.000 |
| 35 | 3 tied | 6 | 13, 23 | 2.5 | 9 | 13, 23 | 2.539 | 2.645 |
| | | | 18, 28 | 2.5 | | 18, 28 | 2.610 | 2.706 |
| | | | 23, 33 | 2.5 | | 23, 33 | 2.387 | 2.284 |
| | iiii | | 3, 13 | 2.5 | | 3, 13 | 2.413 | 2.446 |
| | | | 3, 33 | 2.5 | | 3, 33 | 2.505 | 2.597 |
| | | | 8, 18 | 2.5 | | 8, 18 | 2.665 | 2.607 |
| | | | | | | 11, 27 | 0.000 | 0.000 |
| | | | | | | 13, 33 | 0.000 | 0.000 |
| | | | | | | 3, 23 | 0.162 | 0.019 |
| 45 | 4 free | 5 | 13, 23 | 2.5 | 10 | 13, 23 | 2.520 | 2.456 |
| | | | 23, 33 | 2.5 | | 23, 33 | 2.549 | 2.501 |
| \sim | $\sim \sim \sim 1$ | | 3, 13 | 2.5 | | 3, 13 | 2.463 | 2.496 |
| / Y | <u> </u> | | 3, 43 | 2.5 | | 3, 43 | 2.524 | 2.527 |
| | | | 33, 43 | 2.5 | | 33, 43 | 2.521 | 2.546 |
| | | | | | | 13, 33 | 0.000 | 0.000 |
| | | | | | | 13, 43 | 0.000 | 0.000 |
| | | | | | | 23, 43 | 0.002 | 0.000 |
| | | | | | | 3, 23 | 0.000 | 0.000 |
| | | | | | | 3, 33 | 0.000 | 0.000 |
| 45 | 4 tied | 8 | 13, 23 | 2.5 | 13 | 13, 23 | 2.297 | 2.367 |
| | | | 18, 28 | 2.5 | | 18, 28 | 2.555 | 2.654 |
| 60 | | | 23, 33 | 2.5 | | 23, 33 | 2.305 | 2.664 |
| <u> / </u> | <u> </u> | | 28, 38 | 2.5 | | 28, 38 | 2.670 | 2.516 |
| | | | 3, 13 | 2.5 | | 3, 13 | 2.411 | 2.402 |
| | | | 3, 43 | 2.5 | | 3, 43 | 2.385 | 2.480 |
| | | | 33, 43 | 2.5 | | 33, 43 | 2.190 | 2.255 |
| | | | 8, 18 | 2.5 | | 8, 18 | 2.484 | 2.591 |
| | | | | | | 13, 43 | 0.317 | 0.081 |
| | | | | | | 14, 32 | 0.089 | 0.000 |
| | | | | | | 24, 42 | 0.243 | 0.328 |
| | | | | | | 3, 23 | 0.000 | 0.141 |
| | | | | | | 4, 32 | 0.000 | 0.000 |

^a Each test system is shown schematically as a straight chain of black beads with additional harmonic restraints represented by gray arcs connecting the restrained beads. ^b Number of beads in the chain. ^c Number and type of induced loops. Free loops result from connecting loop end-beads with harmonic restraints, shown as gray arcs in the schematics, while tied loops result from connecting middle beads across free loops, shown as dotted gray arcs in the schematics. ^d Number of harmonic restraints used to induce the loops in the test system.

^e Indexes of beads connected by harmonic restraints in the test systems. The first bead has index 1.
^f Spring constant used for harmonic restraints in simulations of test systems.
^g Number of harmonic restraints used during the ensemble recovery procedure.
^h Indexes of beads connected by harmonic restraints during the ensemble recovery procedure.
ⁱ Restraint spring constants recovered for optimal conformation ensemble when using two parameters in the general linear model to predict spring constants from contact probabilities (CPs).

^j Same as $k_{i,j}(2)$, but using n+1 parameters in the general linear model.