

Supplementary Figures

Figure S1 Simulated Structure positions.	3
Figure S2 Synthetic Contact maps	4
Figure S3 iPSC 4DMax agreement with 3DMax.	5
Figure S4 Synthetic Hi-C from 4D Structures	6
Figure S5 Heat Maps Comparing iPSC reconstruction to iPSC Real Data	11
Figure S7 Heat Maps Comparing Cardio Reconstruction	13
Figure S8 Interpolated Hi-C from iPSC 4D Structure	16
Figure S9 Interpolated Hi-C from Cardiomyocyte 4D Structures	20
Figure S10 iPSC Contact Map Similarity	21
Figure S11 Reconstructed iPSC Trajectory Curves	23
Figure S12 iPSC AB Interpolation Scatter Plots	24
Figure S13 Pearson Mats	30
Figure S14 4D Model Similarity of Interpolated iPSC and Full iPSC Models	31
Figure S15 4D Model Similarity of Interpolated Cardio and Cull Cardio	32
Figure S16 Run Time	34
Figure S17 Cardio Schematic	34
Figure S18 AB Vec Pearson Matrices	40
Figure S19 Hyper Parameter Tuning	42
Figure S20 Performance Comparison to Numerical Interpolation	44
Videos	45
Video S1 Collection 1 Synthetic Videos	45
Video S2 Collection 2 iPSC Videos	45
Video S3 Collection 3 Cardio Videos	45
Video S4 Collection 4 iPSC Missing Chros	45
Video S5 Collection 5 Cardio Missing Chros	45
Video S6 Collection 6 Changing Resolution	45
Video S7 Collection 7 Changing Granularity	45

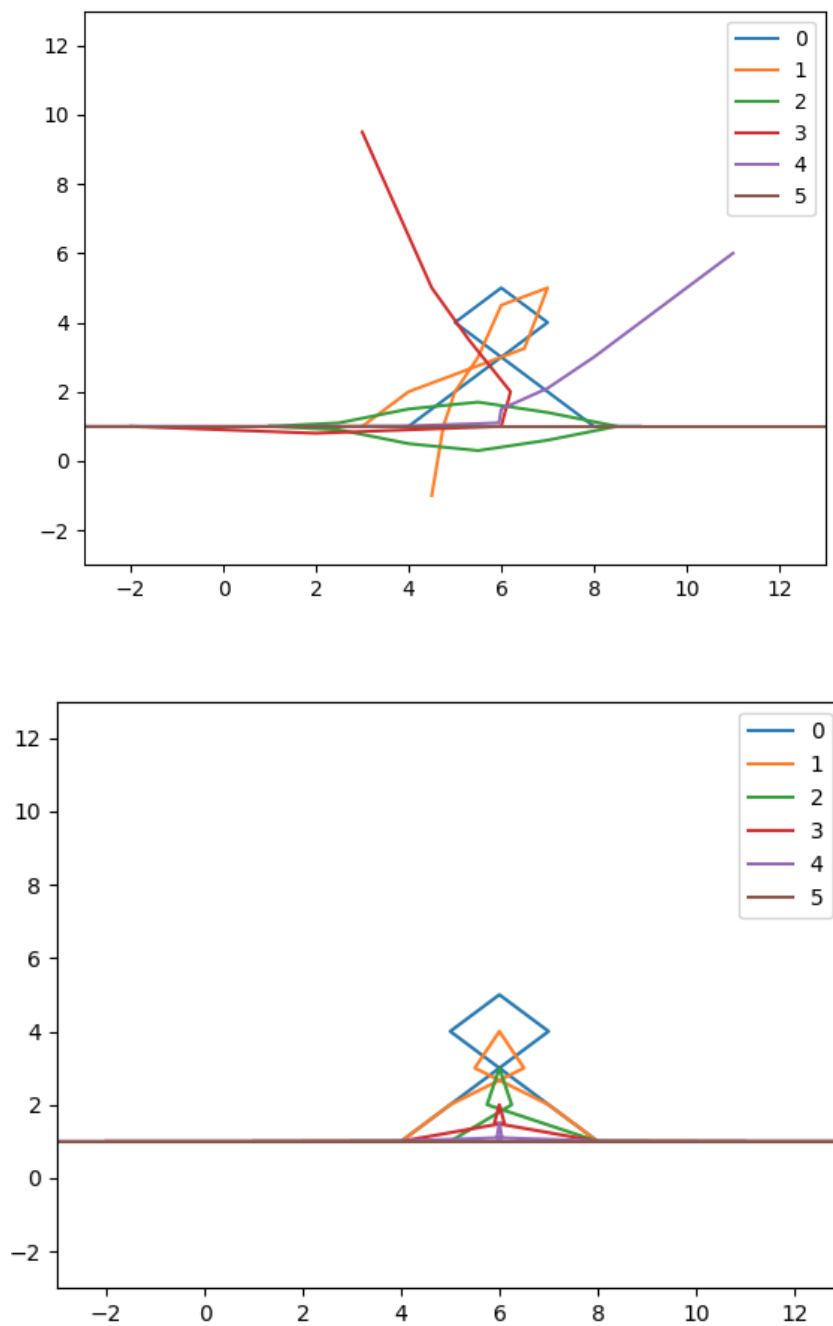


Figure S1 Simulated Structure positions.

Two dimensional projections of positions of synthetic chromosomes a. Structure 1. b. Structure 2. Each structure begins and ends with the same confirmation, but structure 1 is opened in a swinging motion while structure 2 is unraveled by both ends extending outward.

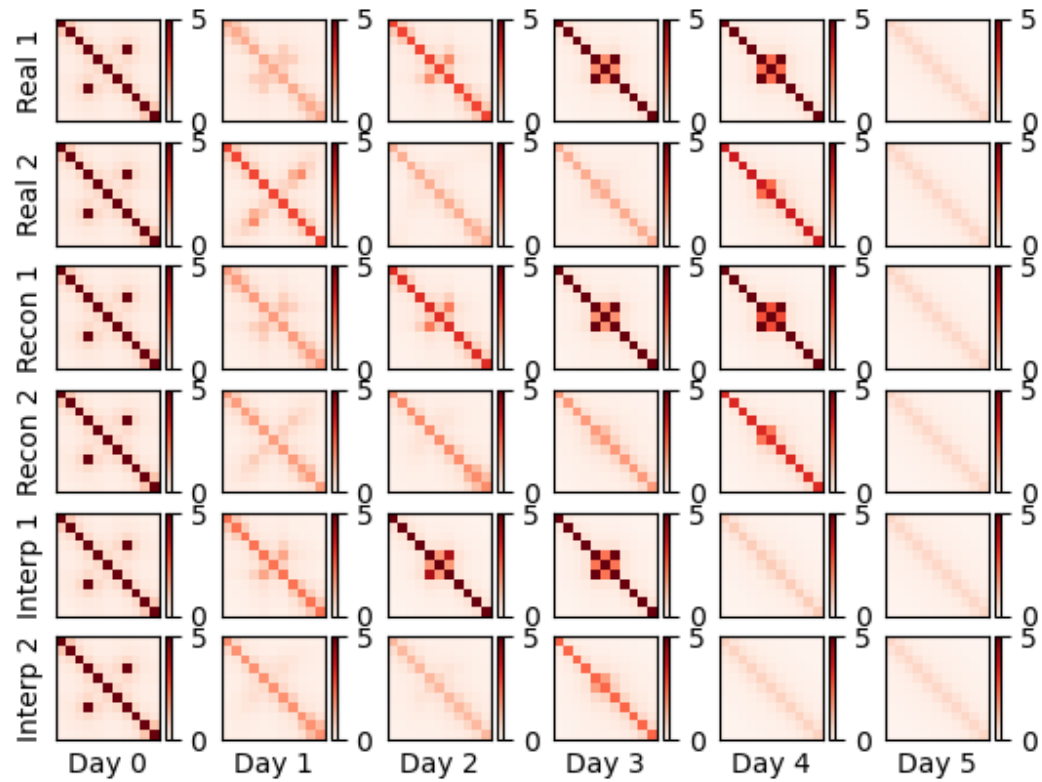


Figure S2 Synthetic Contact maps

Heatmaps of each day in each synthetic chromosome. The top 2 rows show heat maps of original synthetic data, middle two rows show heatmaps of 4DMax reconstructed structures and bottom two show interpolations if the algorithm 4DMax algorithm is not shown heat maps on given day.

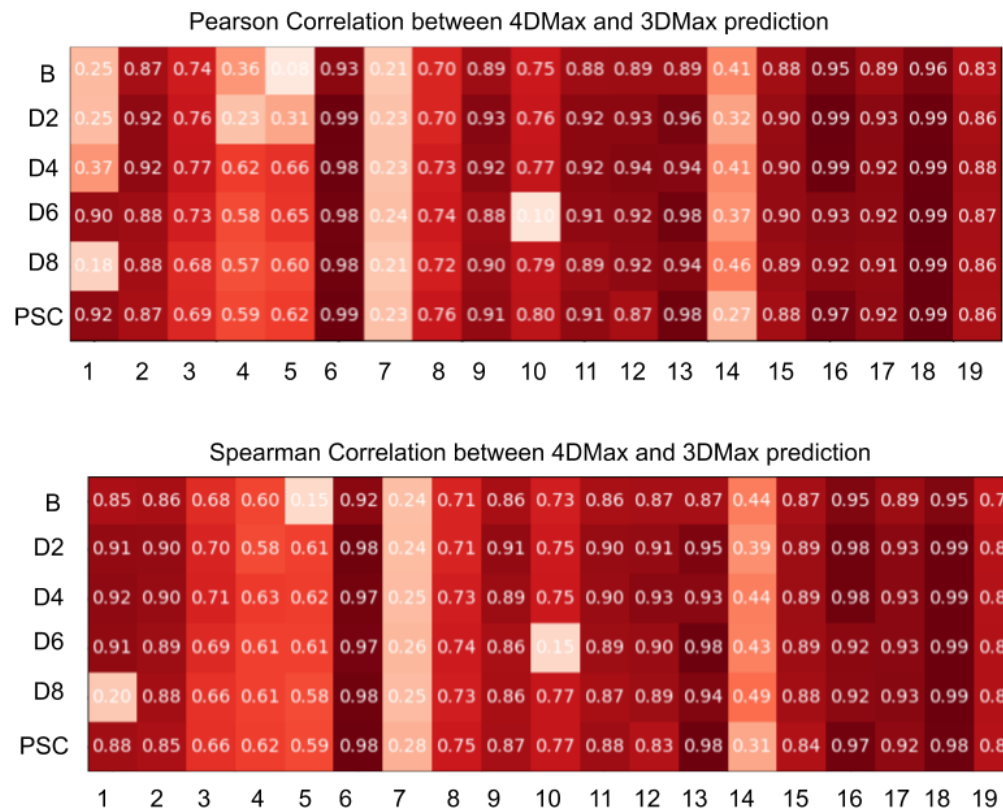


Figure S3 iPSC 4DMax agreement with 3DMax.

(a) pearson and (b) spearman correlation between 4DMax structures and 3D structure predicted using 3DMax algorithm.

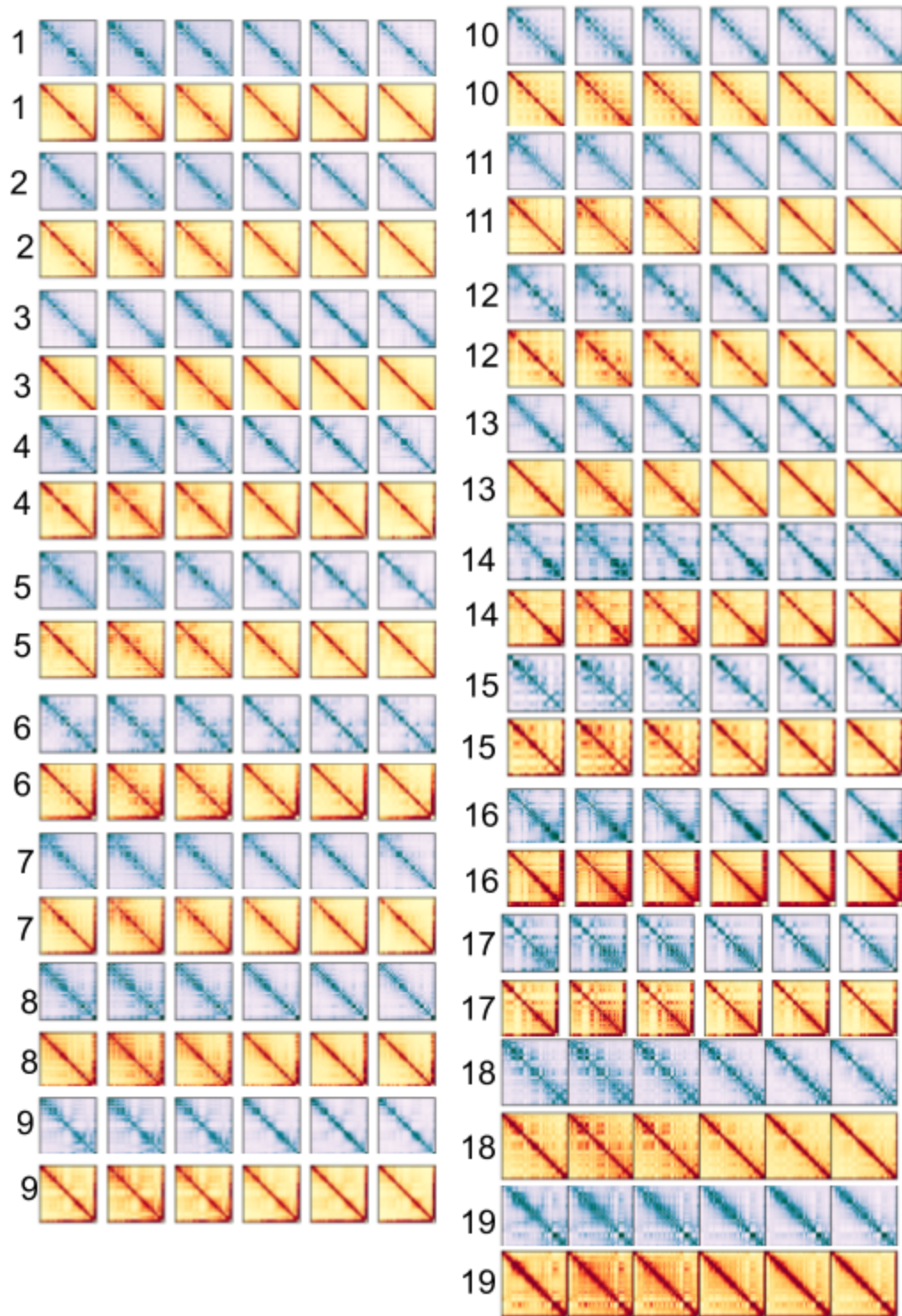
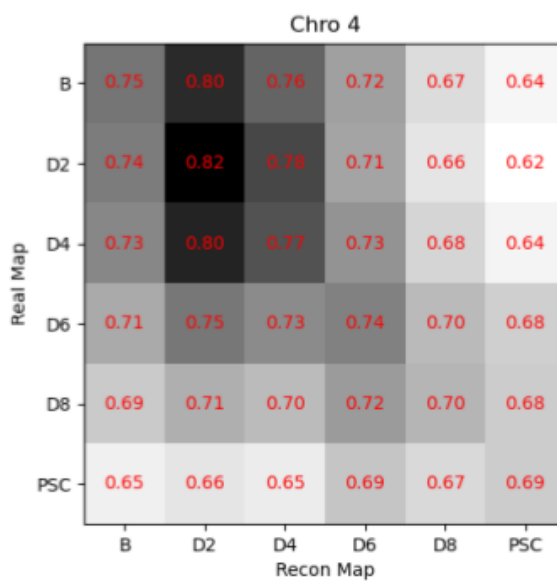
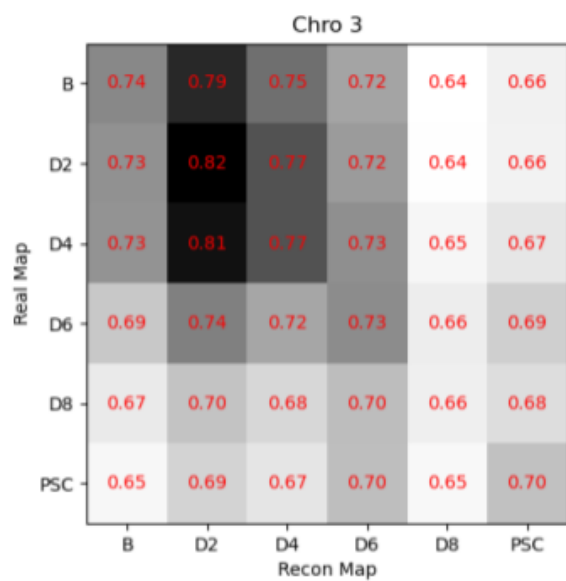
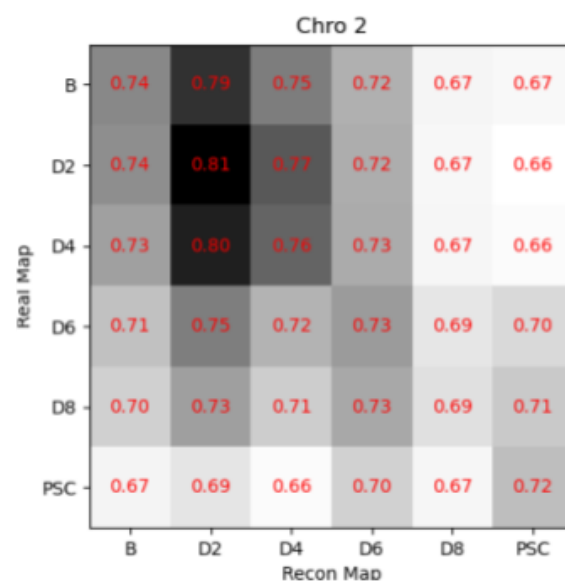
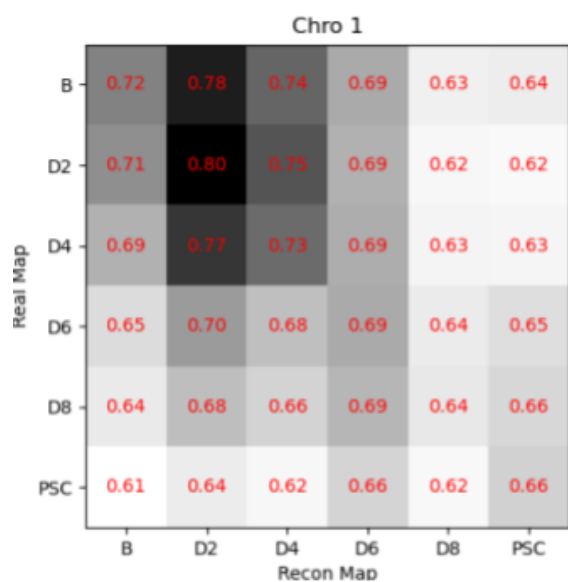
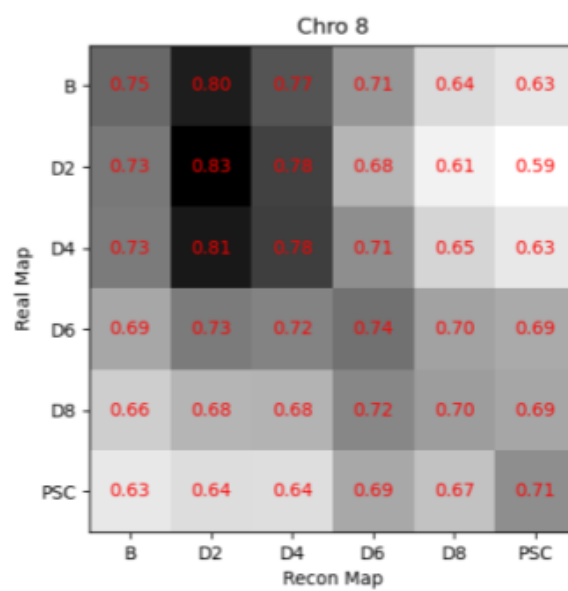
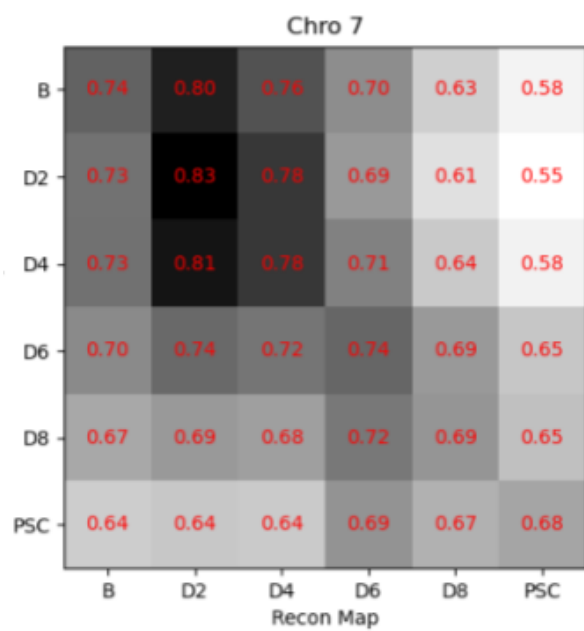
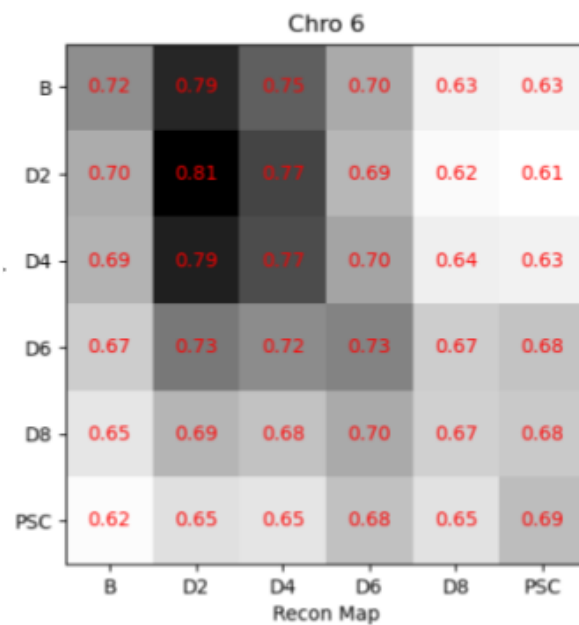
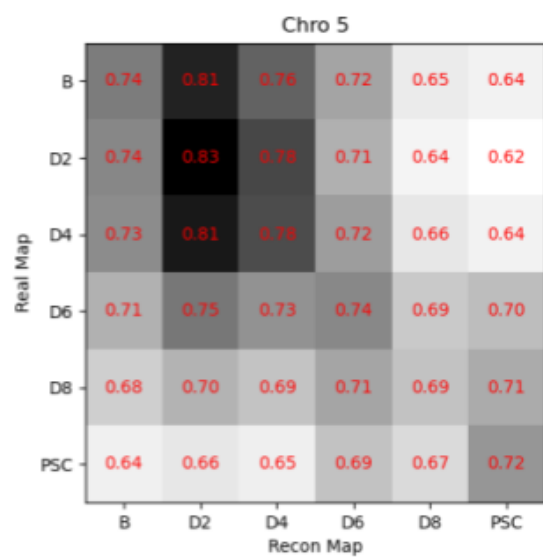
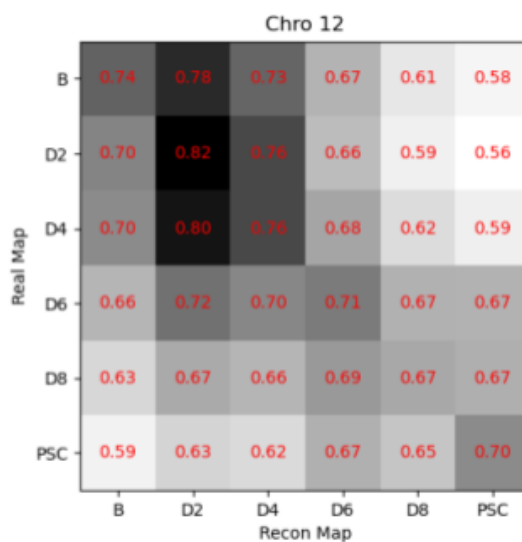
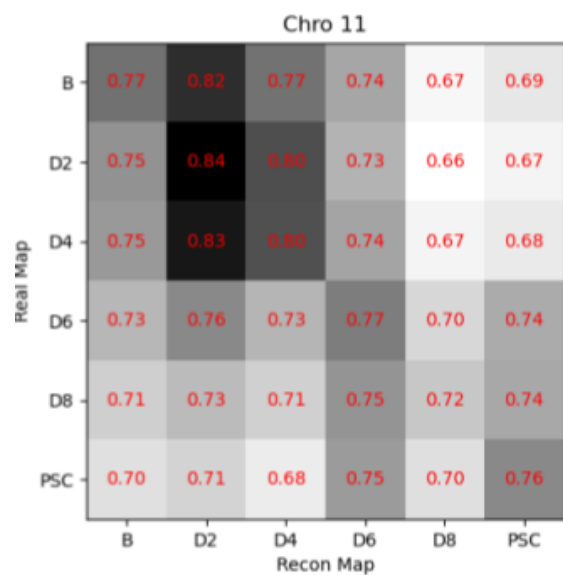
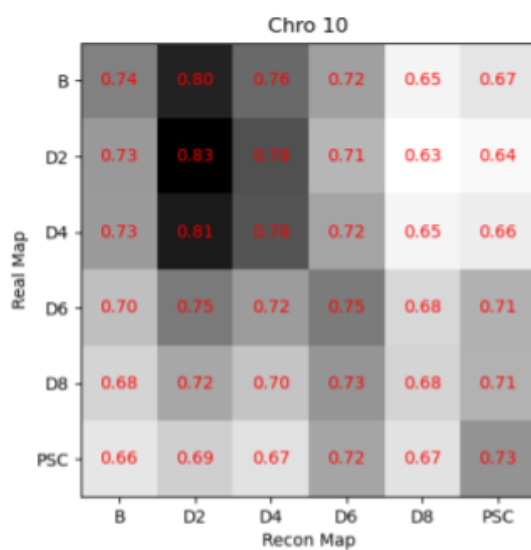
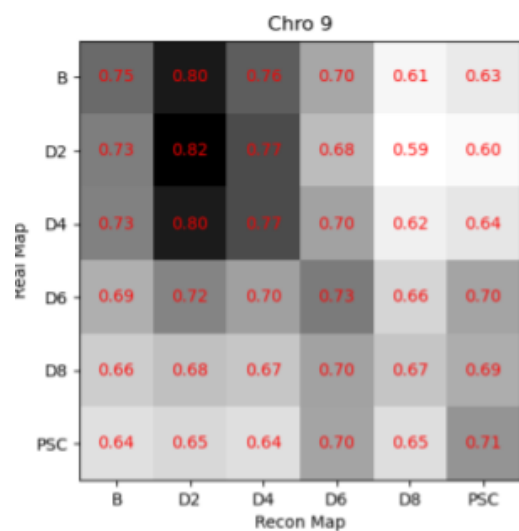


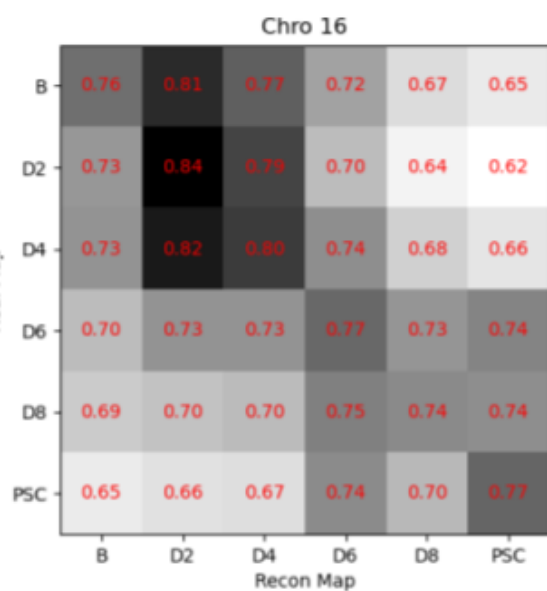
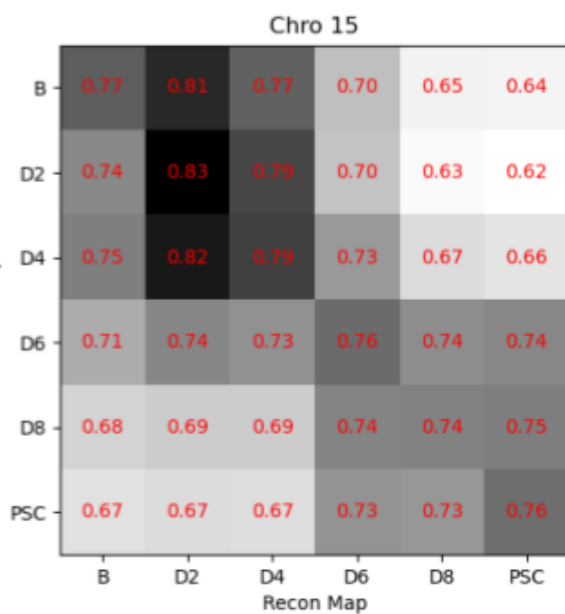
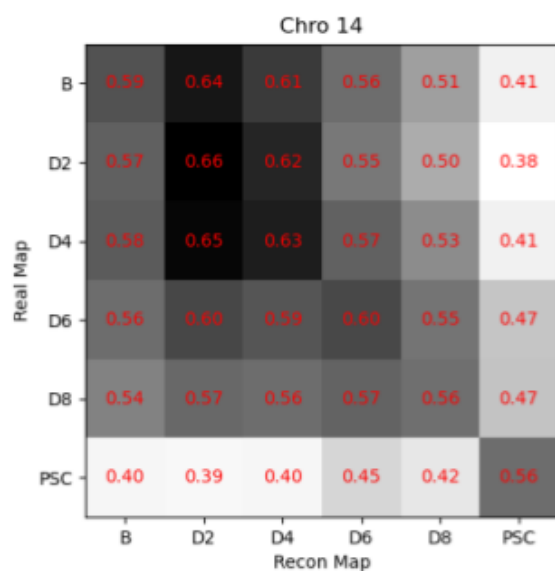
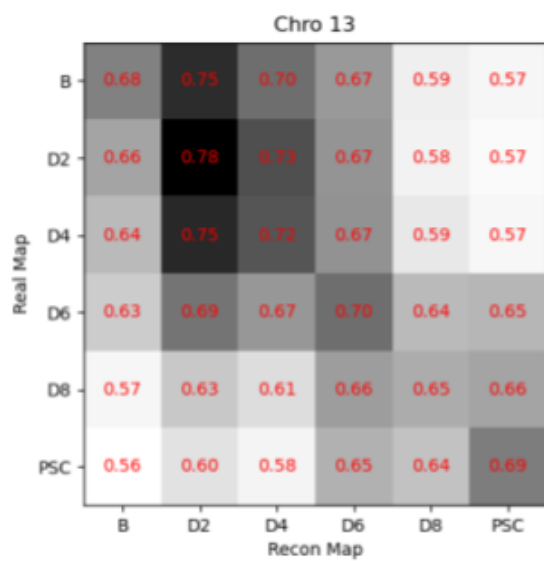
Figure S4 Synthetic Hi-C from 4D Structures

Chromosomal contact maps on each day for each chromosome in iPSC









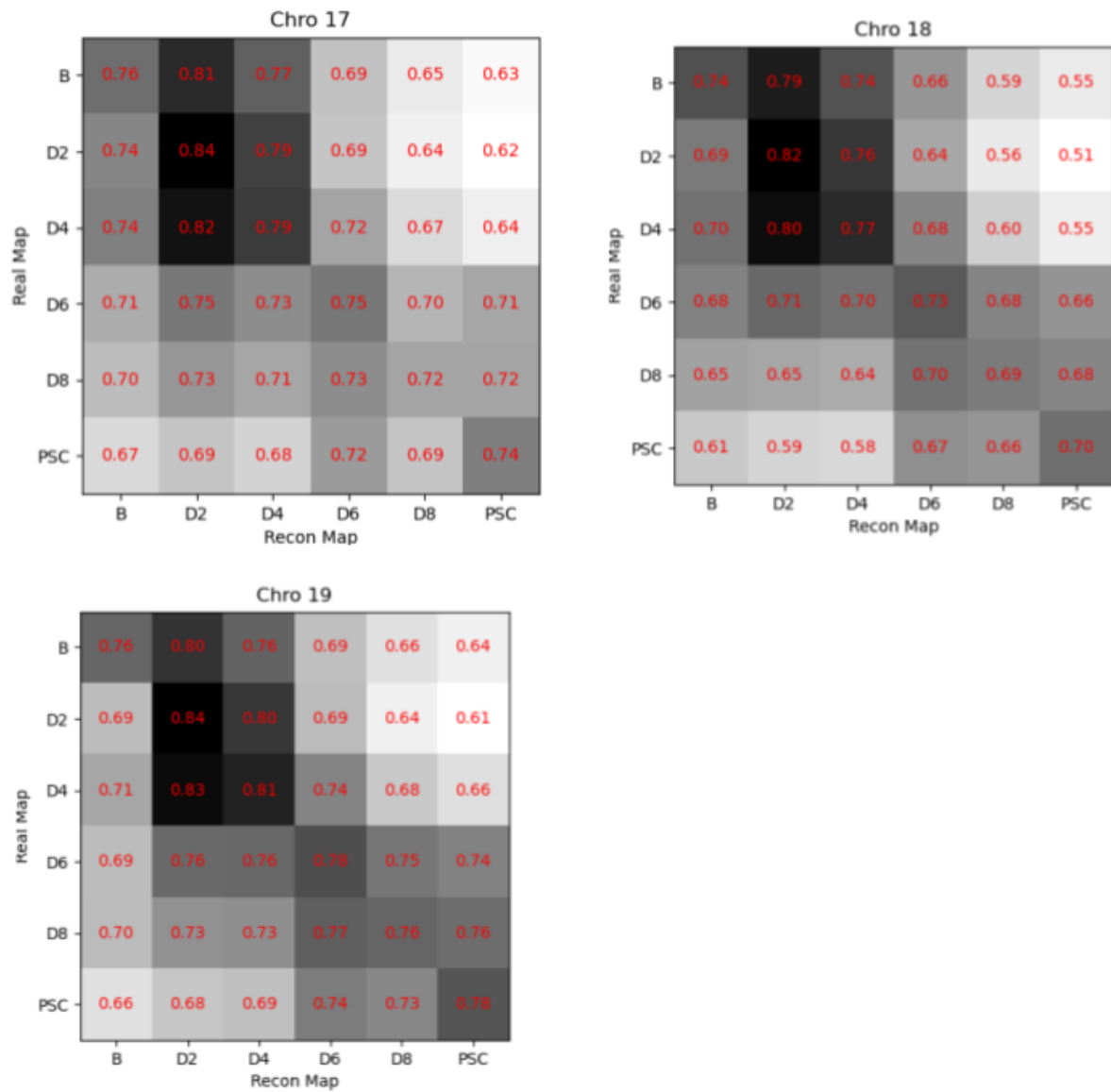
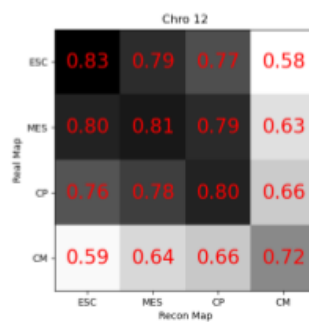
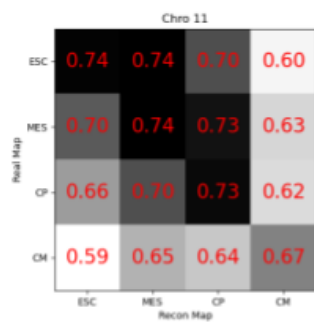
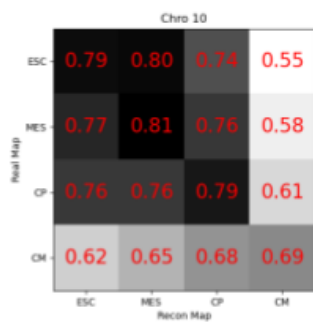
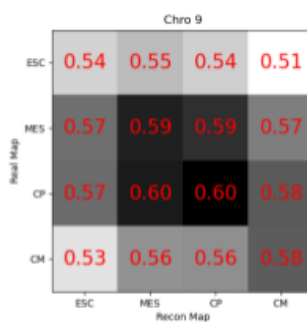
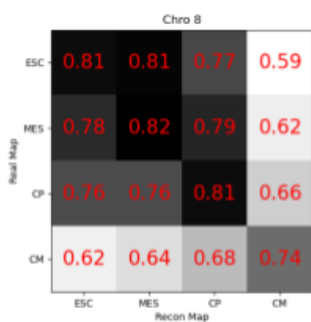
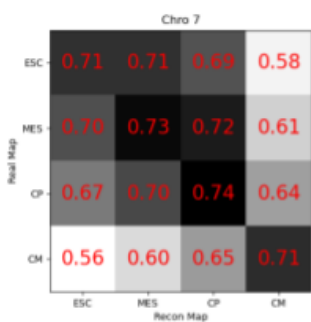
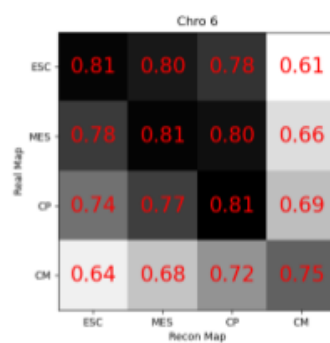
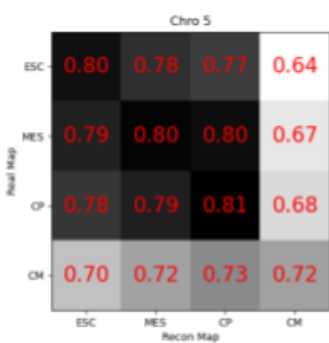
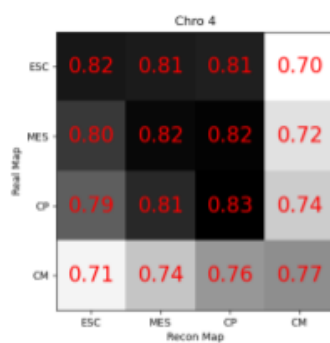
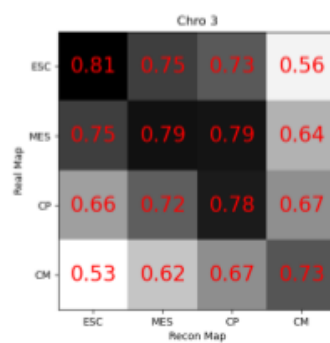
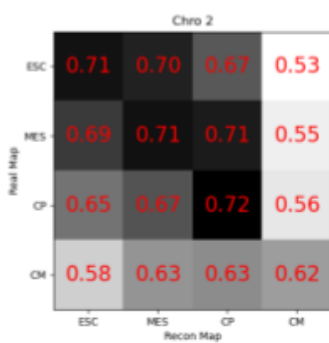
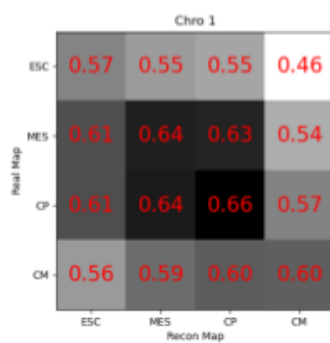


Figure S5 Heat Maps Comparing iPSC reconstruction to iPSC Real Data

Pearson correlation between real Hi-C contact maps and 4d reconstructed maps.



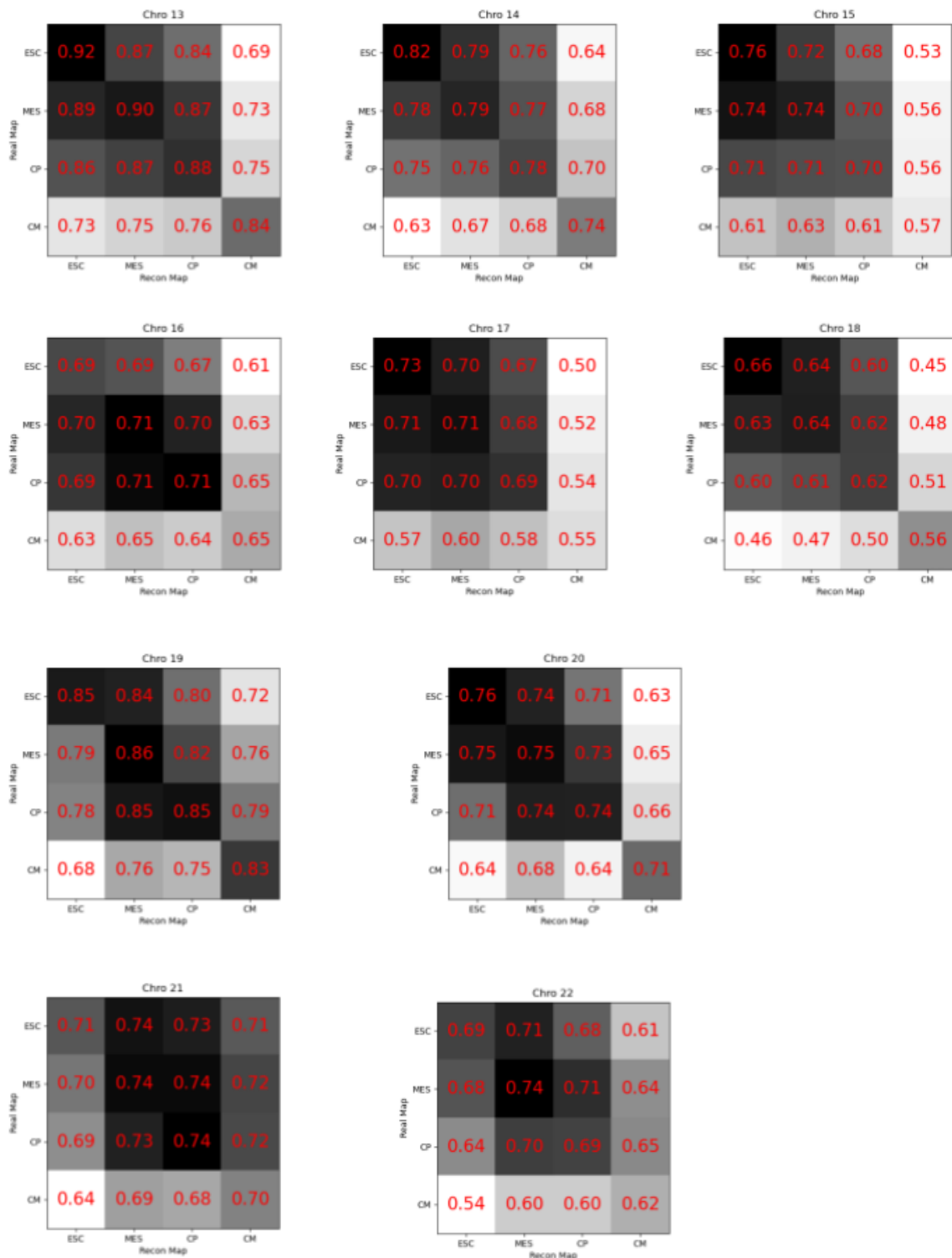
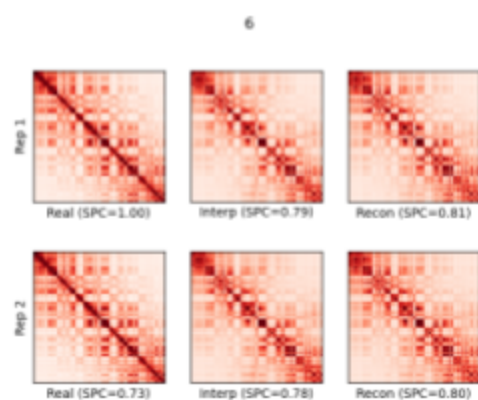
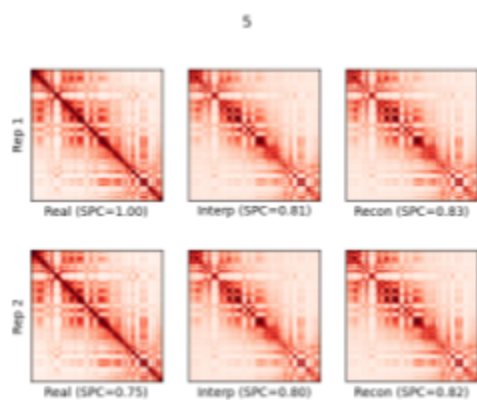
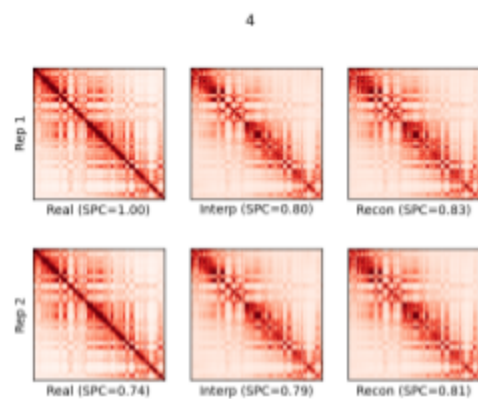
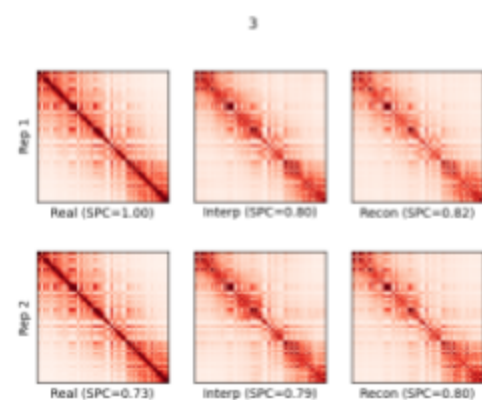
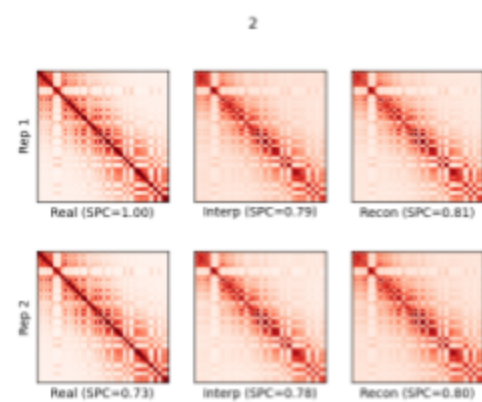
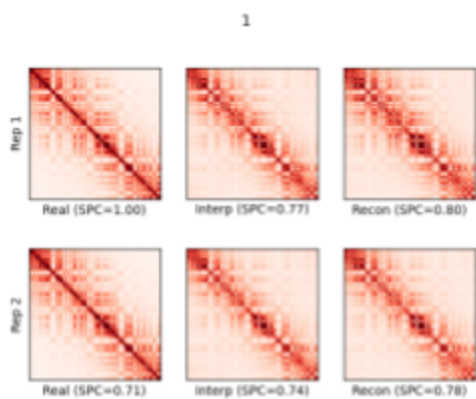
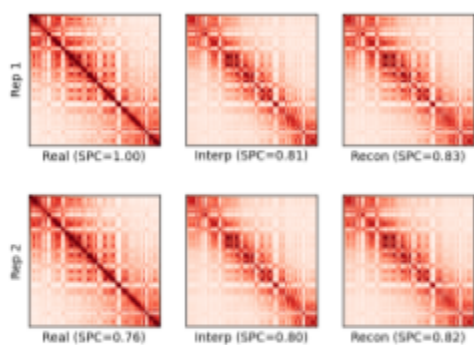


Figure S7 Heat Maps Comparing Cardio Reconstruction

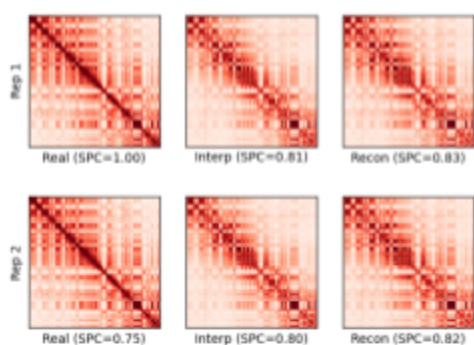
Pearson correlation between real Hi-C contact maps and 4d reconstructed maps.



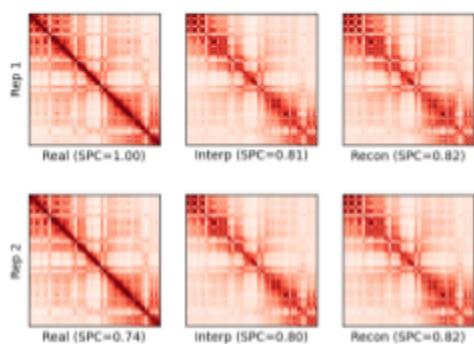
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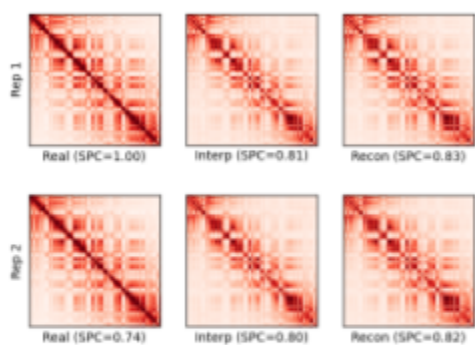
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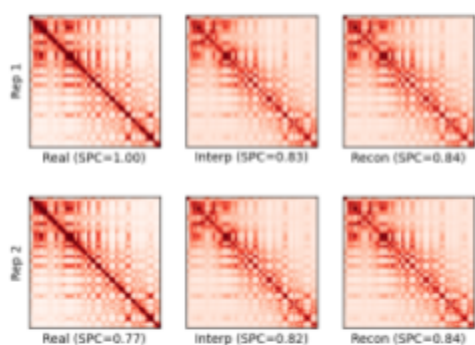
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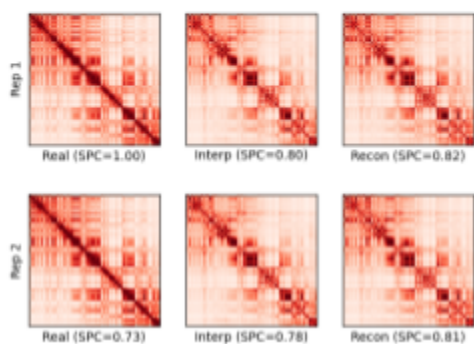
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11



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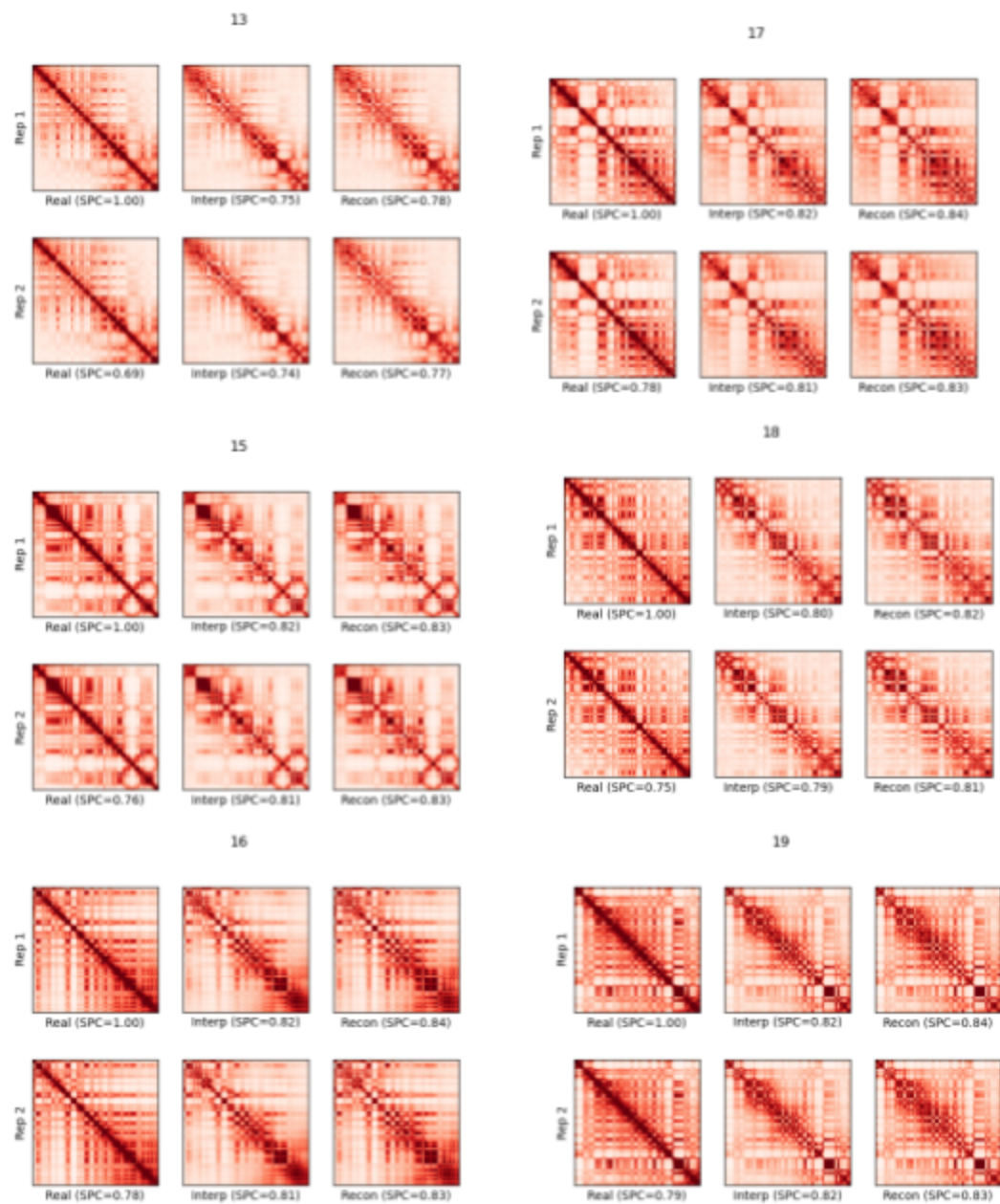
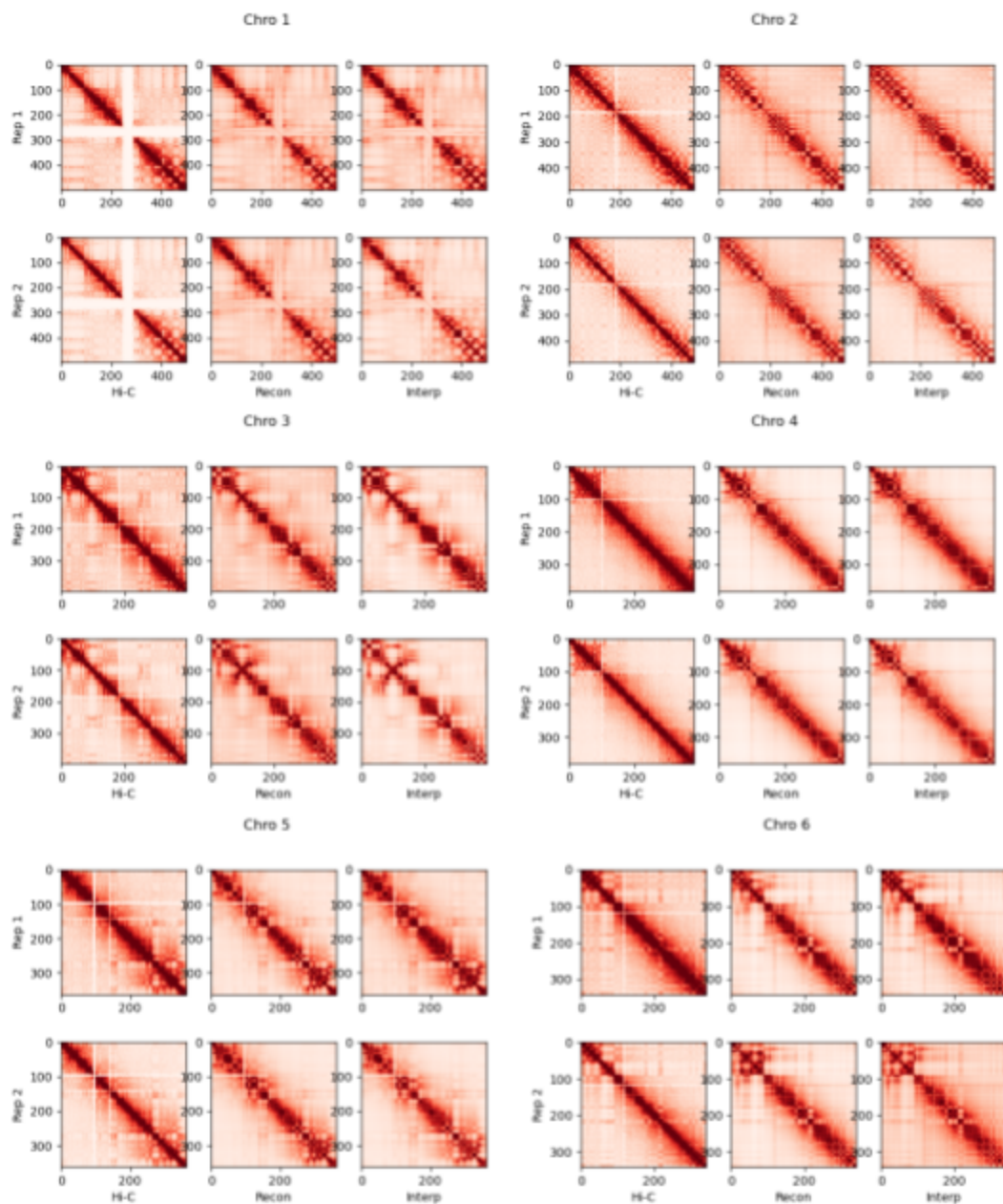
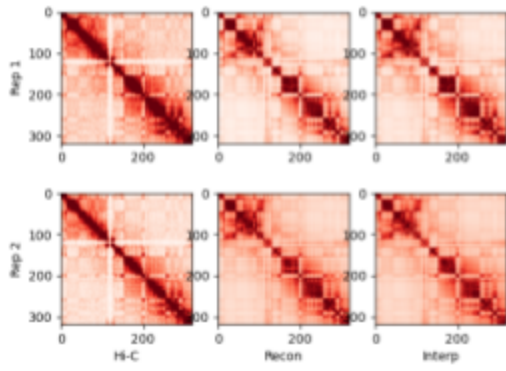


Figure S8 Interpolated Hi-C from iPSC 4D Structure

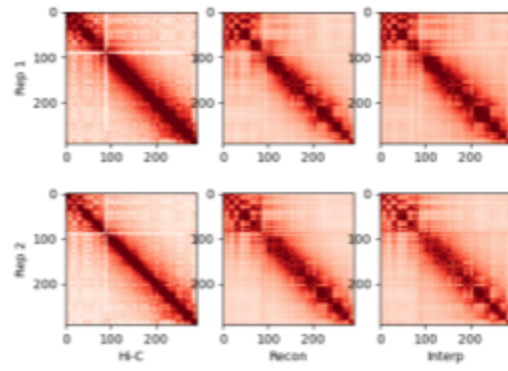
(left) Real Hi-C (middle) reconstructed Hi-C contact maps (right) interpolated contact maps.



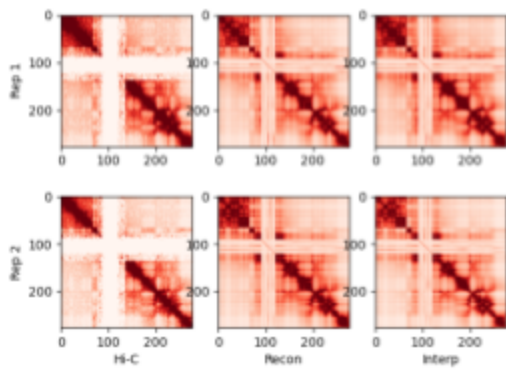
Chro 7



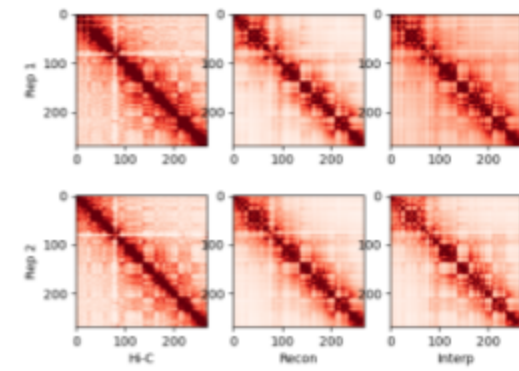
Chro 8



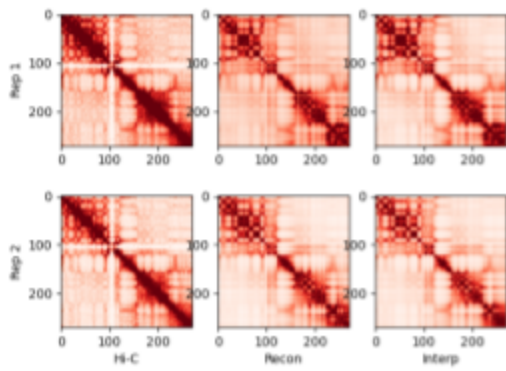
Chro 9



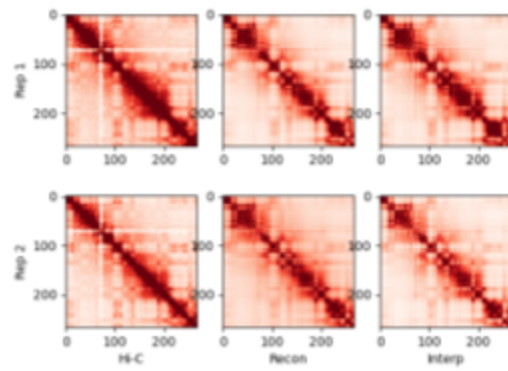
Chro 10



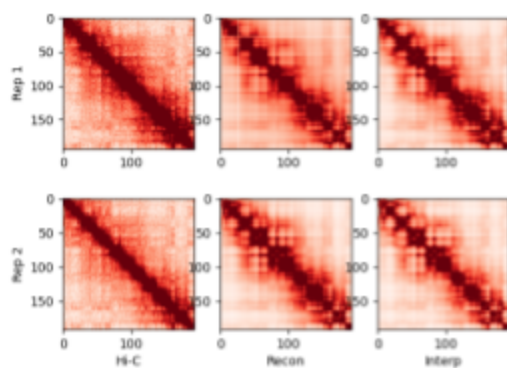
Chro 11



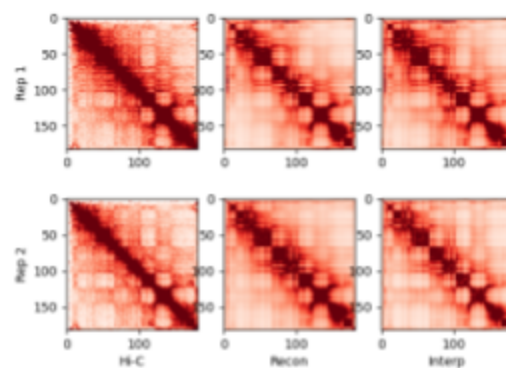
Chro 12



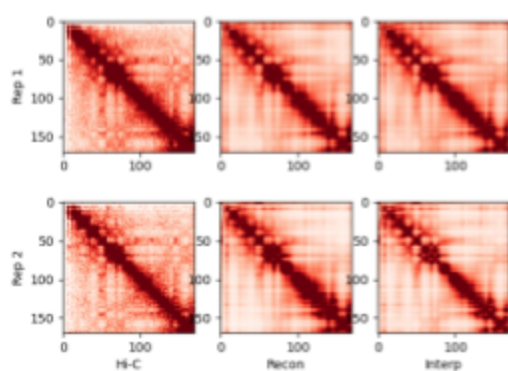
Chro 13



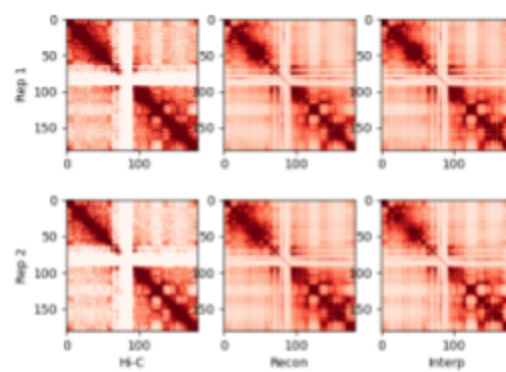
Chro 14



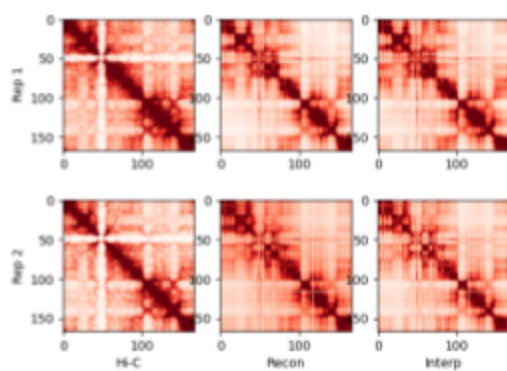
Chro 15



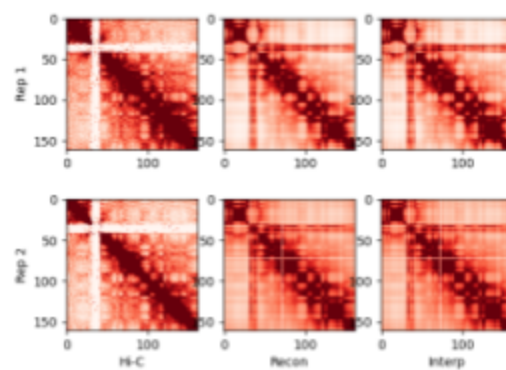
Chro 16



Chro 17



Chro 18



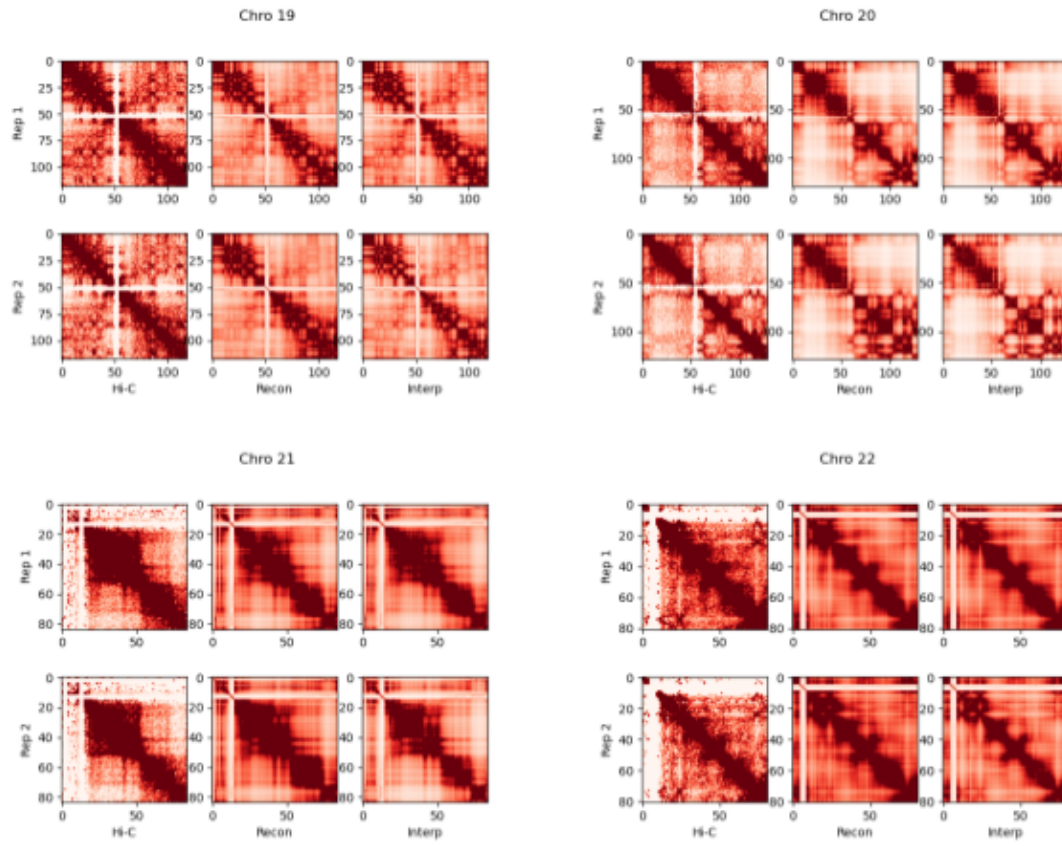


Figure S9 Interpolated Hi-C from Cardiomyocyte 4D Structures

(left) Real Hi-C (middle) reconstructed Hi-C contact maps (right) interpolated contact maps.

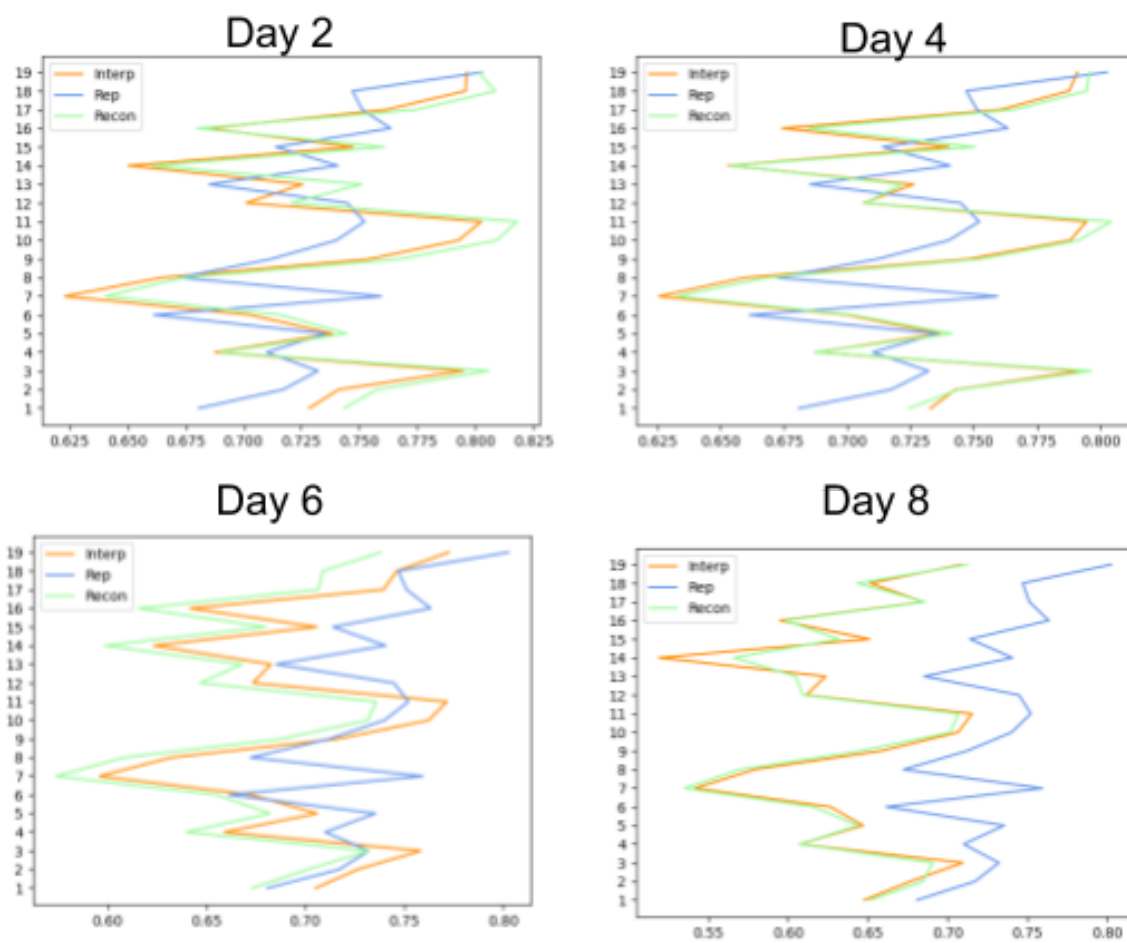
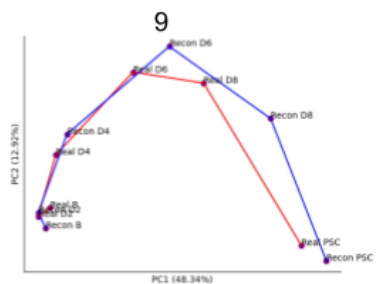
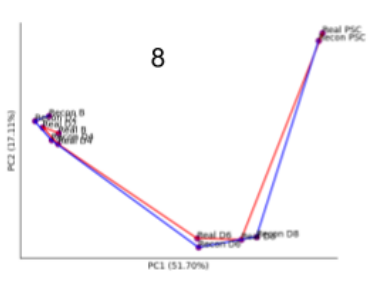
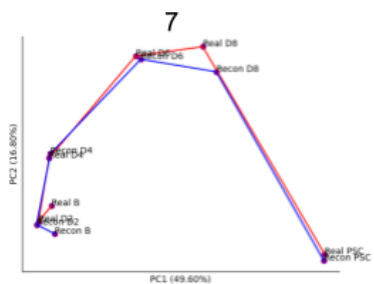
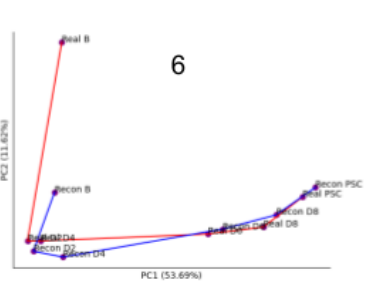
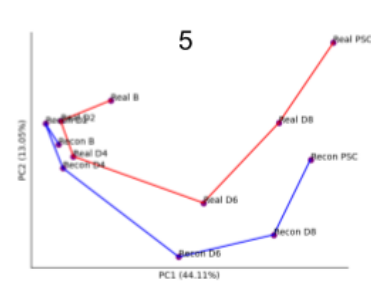
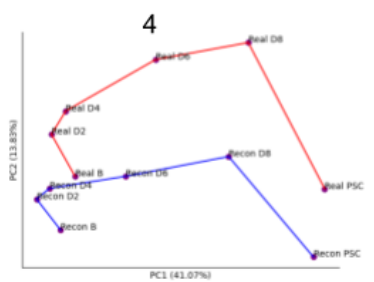
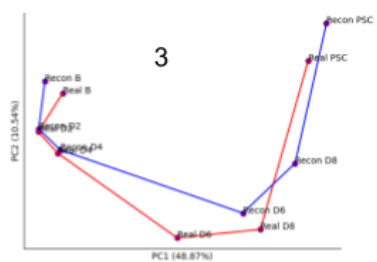
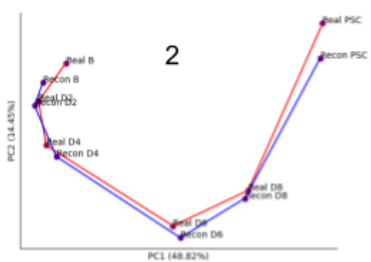
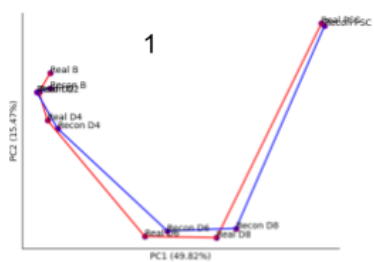


Figure S10 iPSC Contact Map Similarity

(orange) interpolation (green) reconstruction and (blue) biological replication.



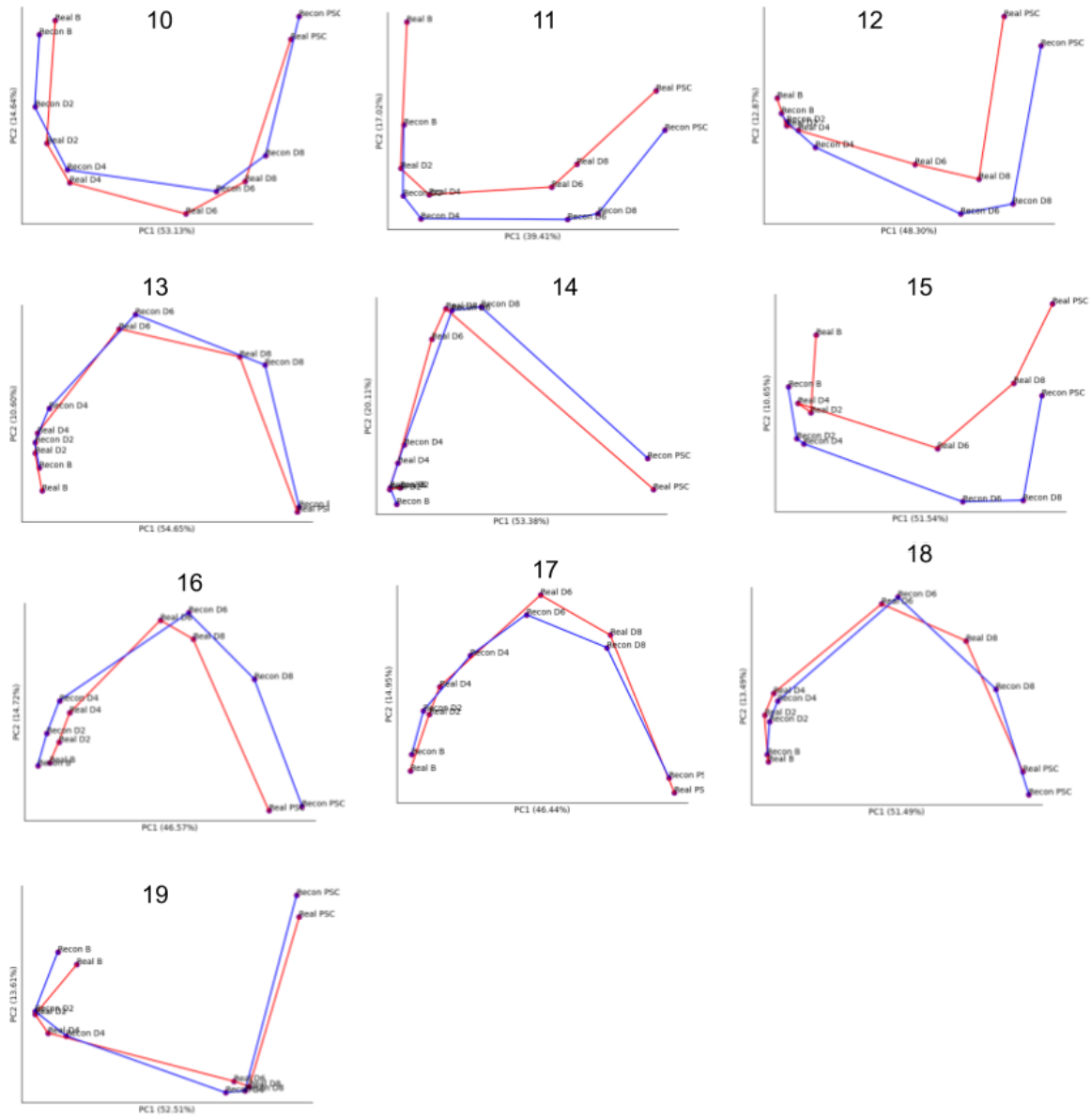


Figure S11 Reconstructed iPSC Trajectory Curves

(red) real Hi-C Trajectory (blue) Reconstructed Hi-C trajectory

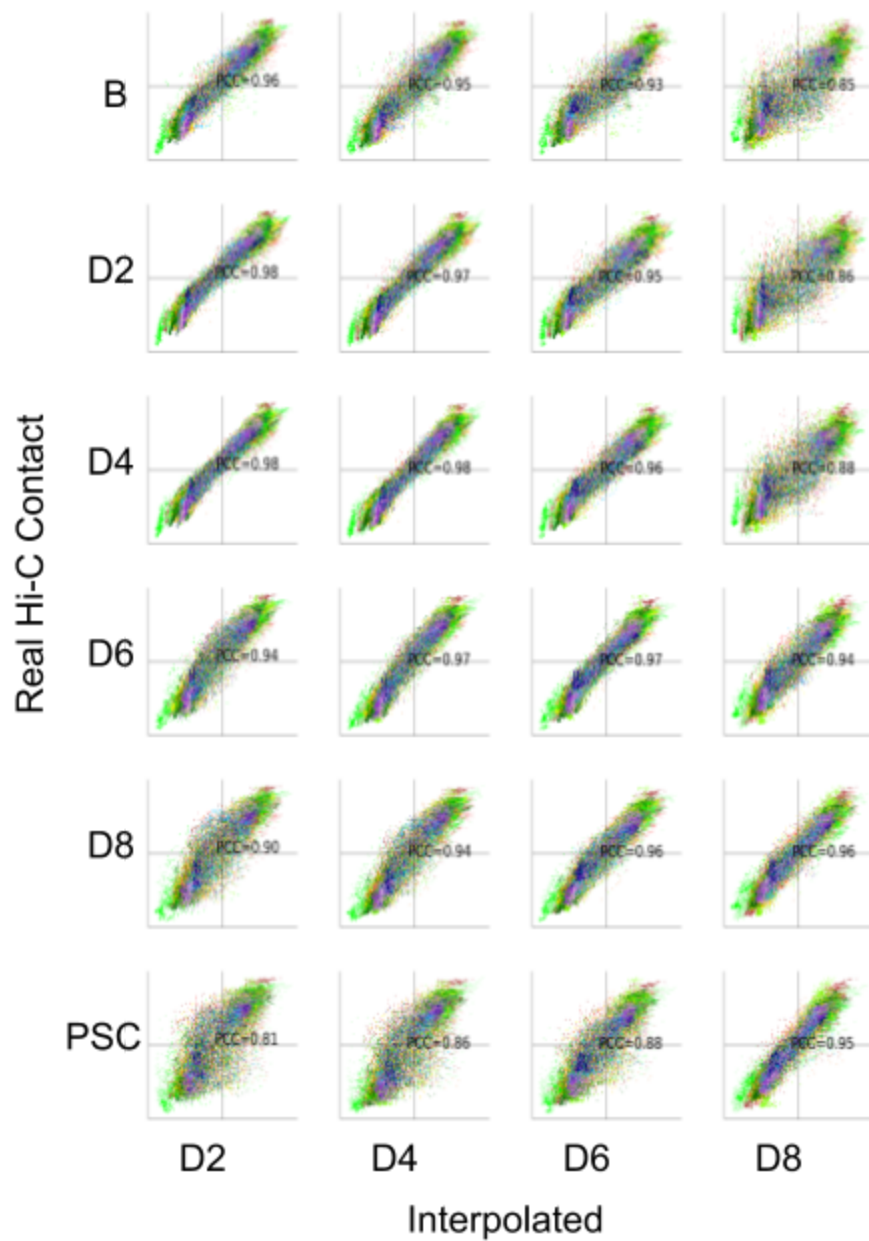
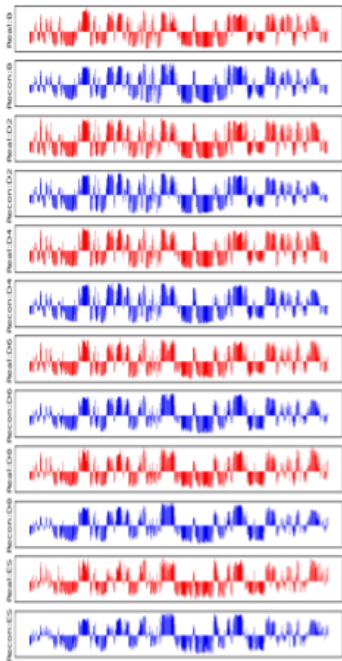


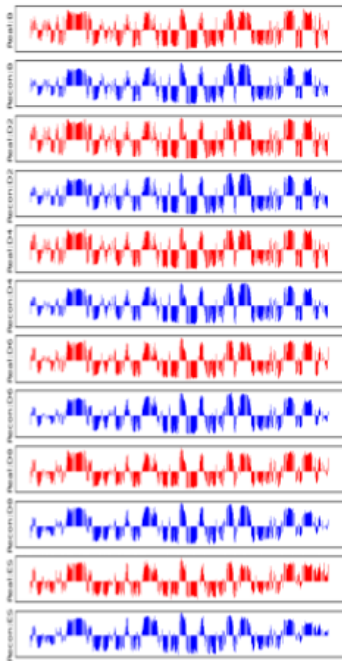
Figure S12 iPSC AB Interpolation Scatter Plots

Scatter plot of PC1 values from interpolated and biological real Hi-C contact maps.

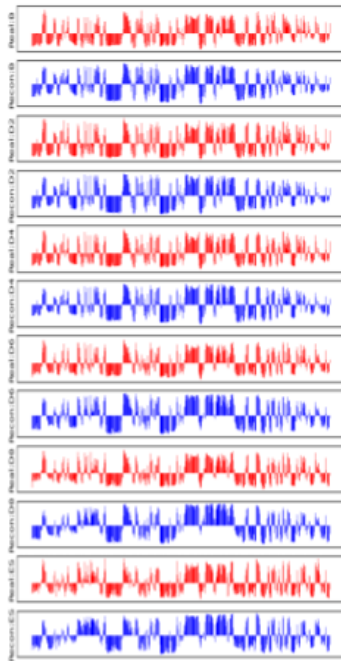
Chro 1



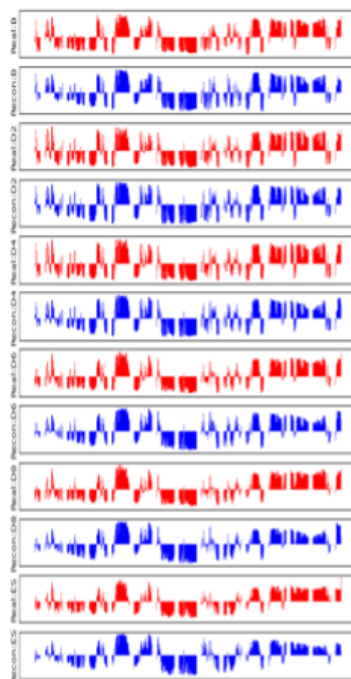
Chro 2



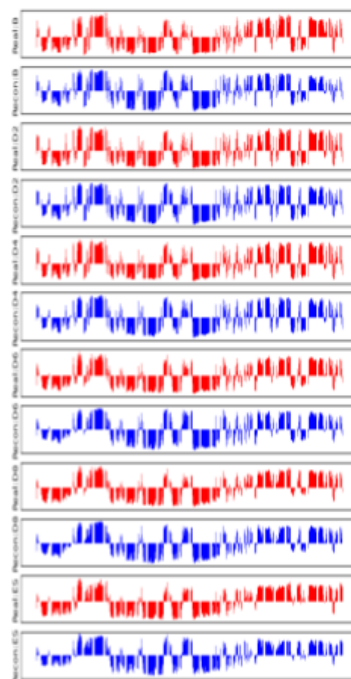
Chro 3



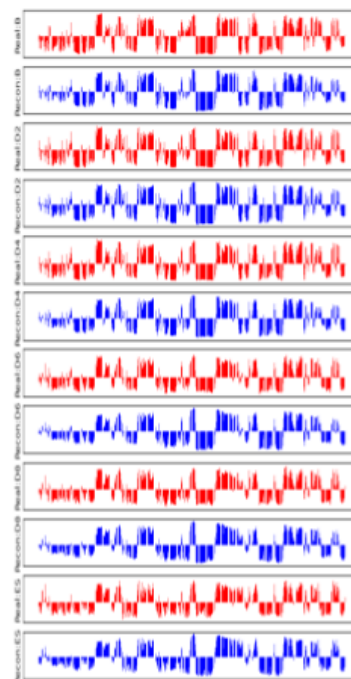
Chro 4



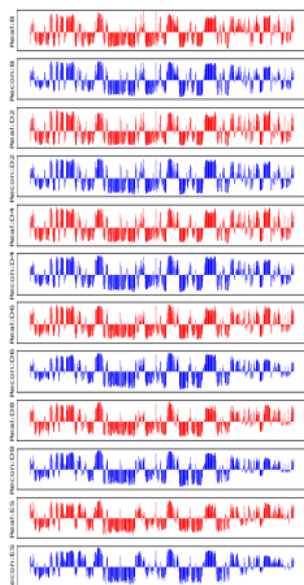
Chro 5



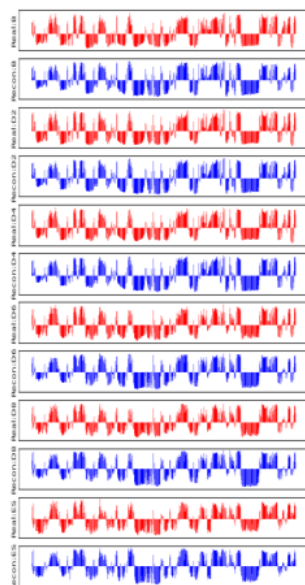
Chro 6



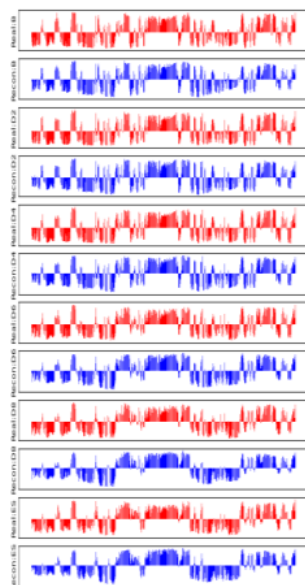
Chro 7



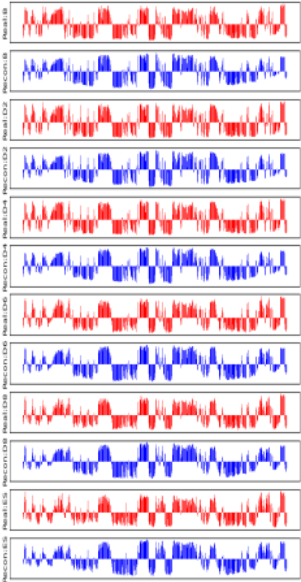
Chro 8



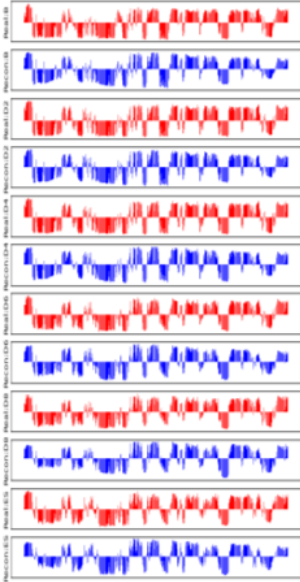
Chro 9



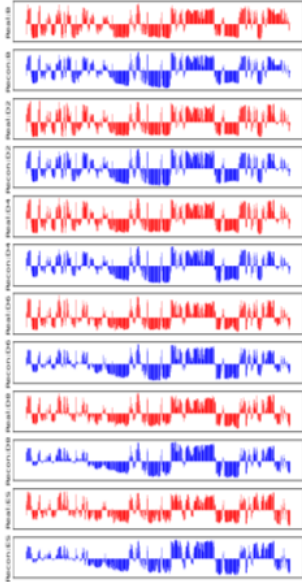
Chro 10



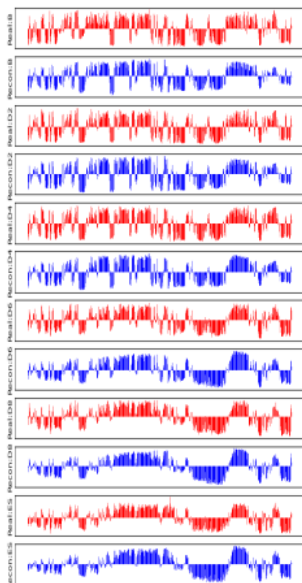
Chro 11



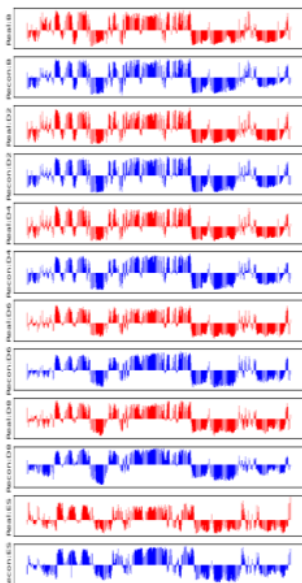
Chro 12



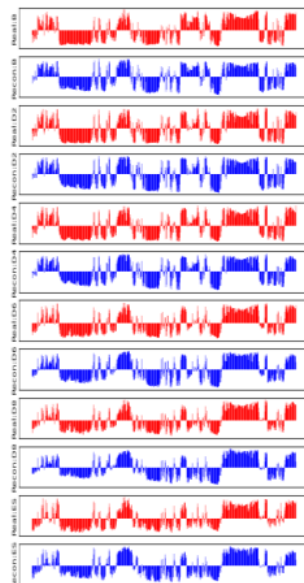
Chro 13



Chro 14



Chro 15



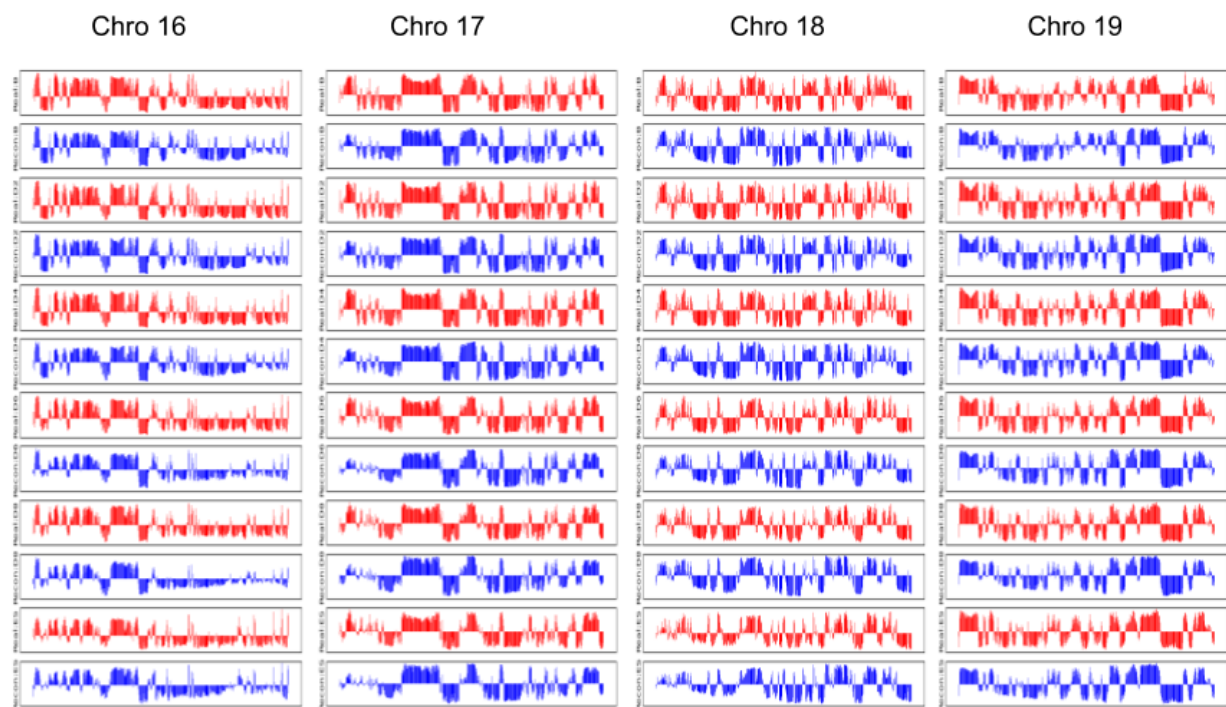


Figure S13 Pearson Mats

Plots of AB compartment assignments (red) biological Hi-C (blue) reconstructed Hi-C contacts.

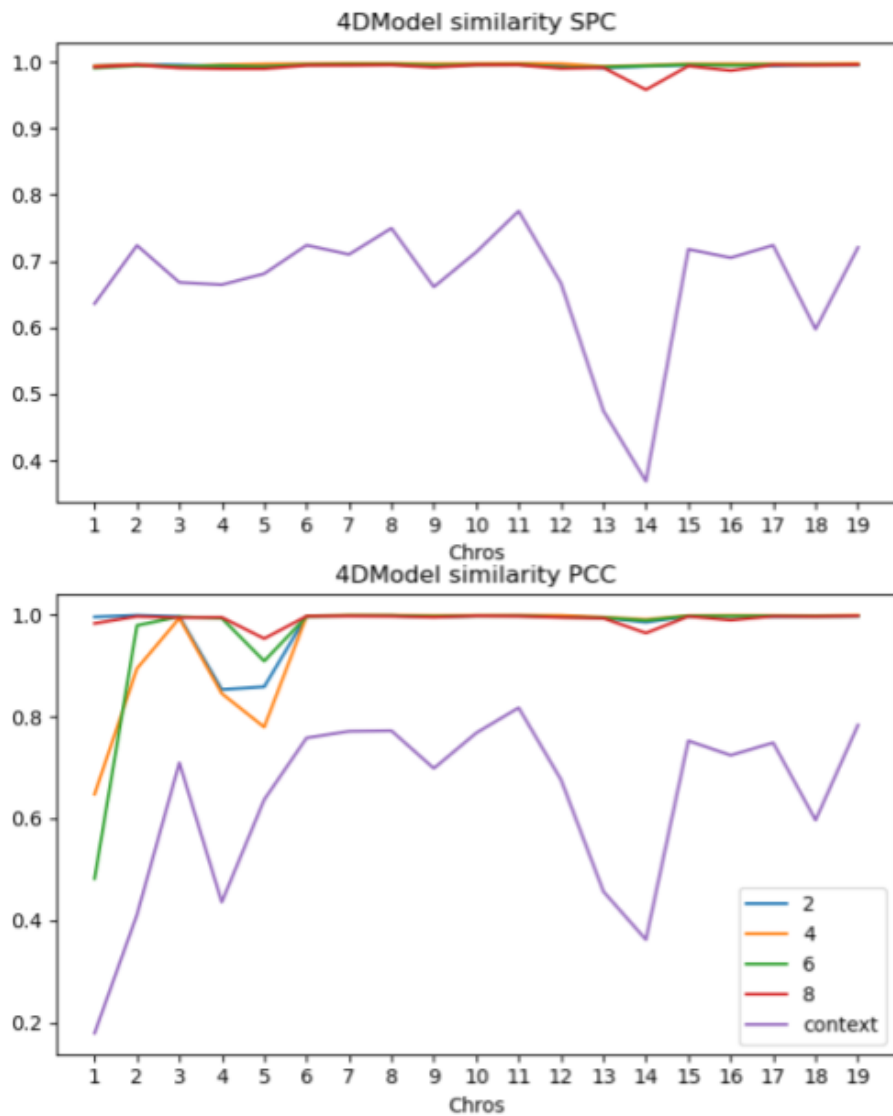


Figure S14 4D Model Similarity of Interpolated iPSC and Full iPSC Models

Avg spearman and pearson correlation between distance vectors of structural conformation at all time points of full cardiomyocyte structures and interpolation structures. The context metrics show the correlation between distance vectors for the structural conformation at the beginning and end of an interpolation structure. This indicated consistently higher similarity between two 4D structures at each time than between a 4D structures starting and ending conformation.

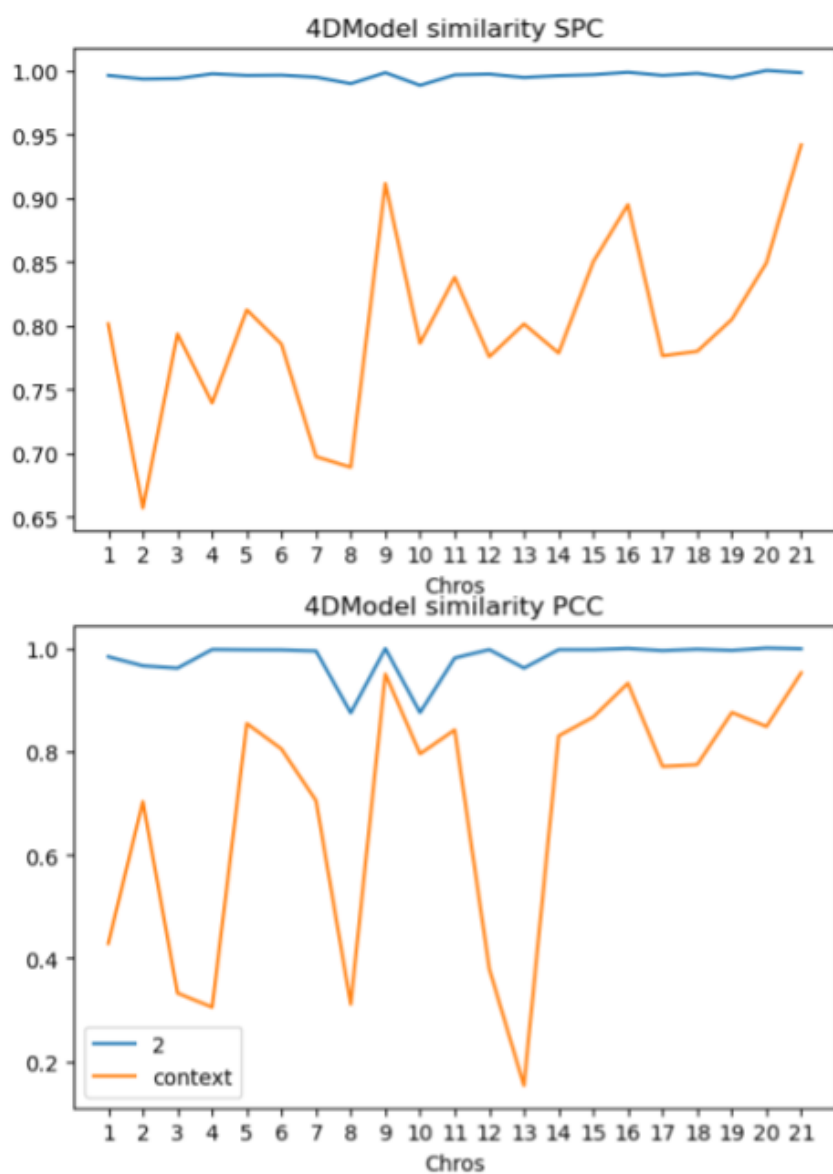


Figure S15 4D Model Similarity of Interpolated Cardio and Cull Cardio

Avg spearman and pearson correlation between distance vectors of structural conformation at all time points of full cardiomyocyte structures and interpolation structures. The context metrics show the correlation between distance vectors for the structural conformation at the beginning and end of an interpolation structure. This indicated consistently higher similarity between two 4D structures at each time than between a 4D structures starting and ending conformation.

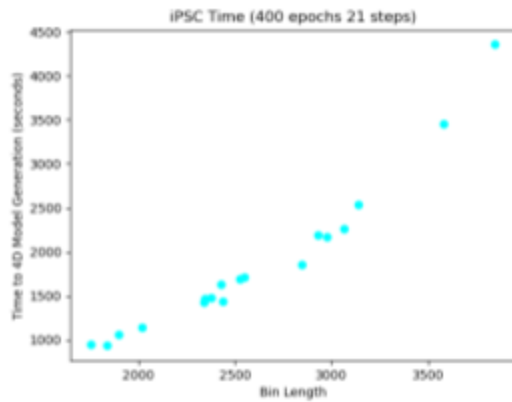
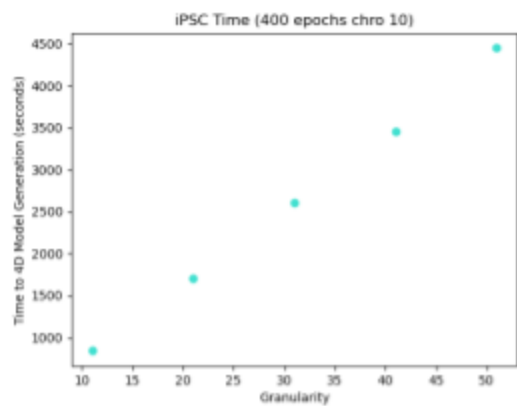
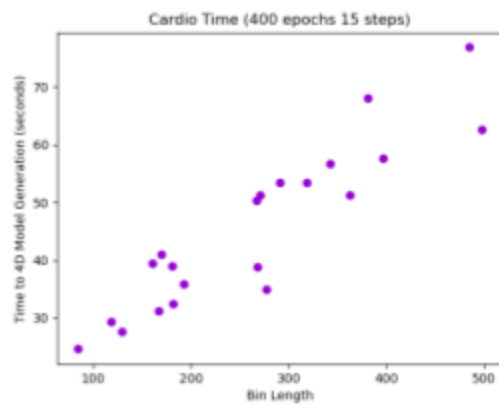
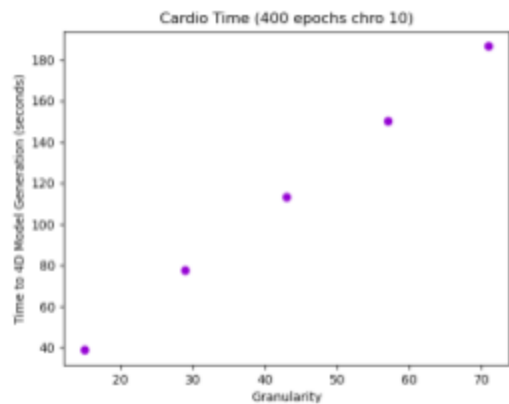


Figure S16 Run Time

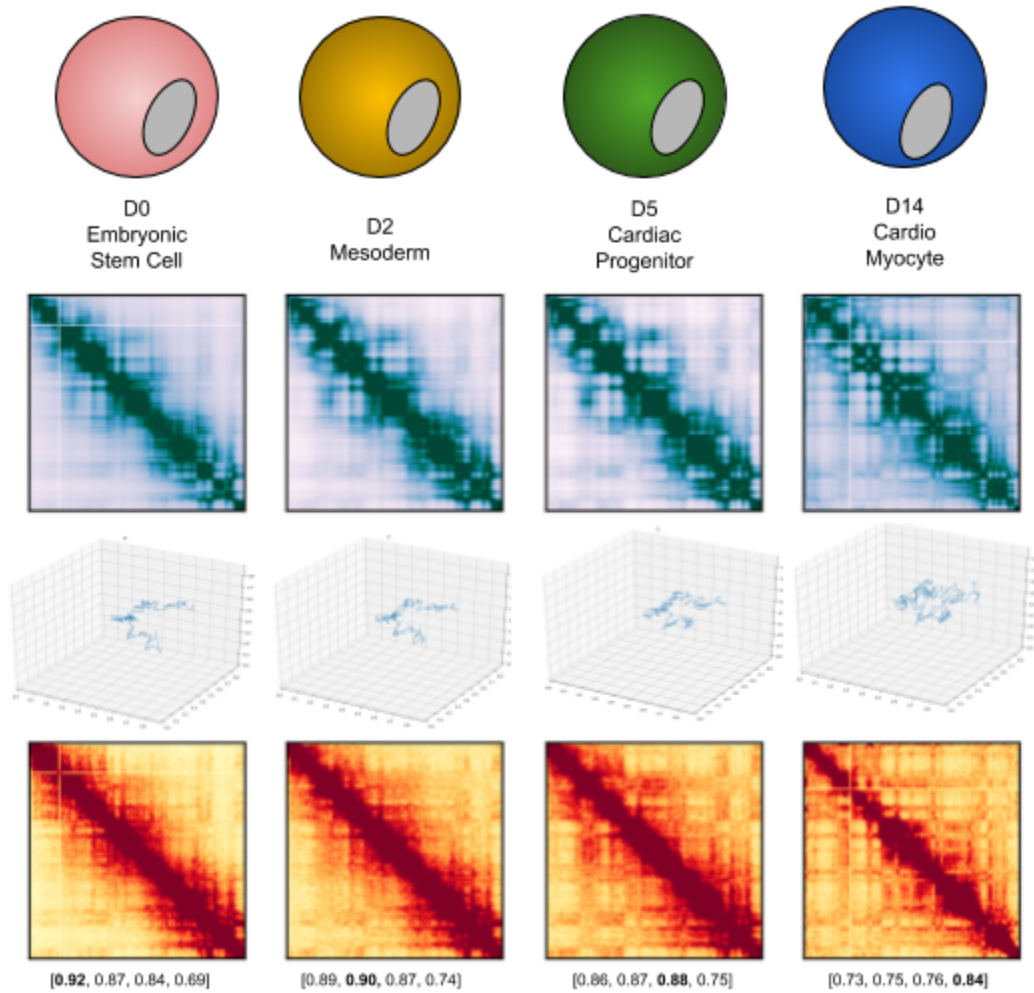
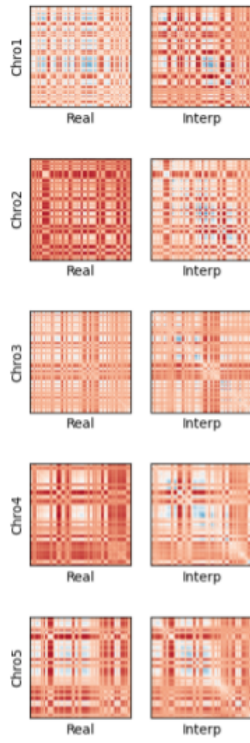


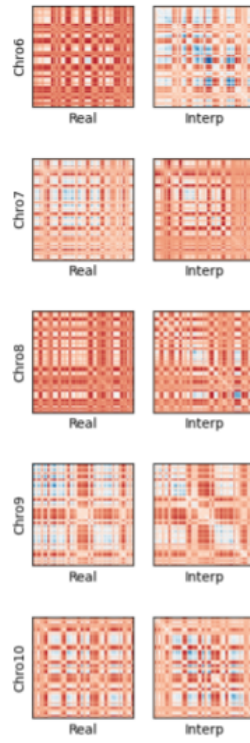
Figure S17 Cardio Schematic

Chromosome 13. (a) Days Hi-C experiments are taken. (b) Hi-C contact matrices © reconstructed structure (4) Reconstructed Contact maps

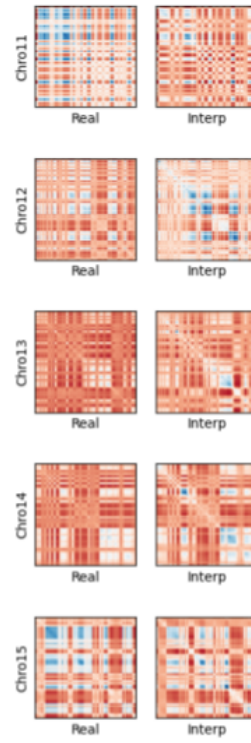
Day B



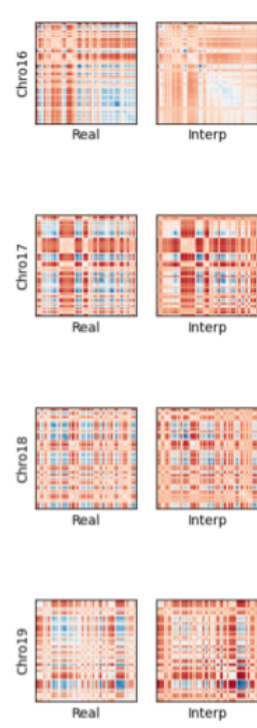
Day B



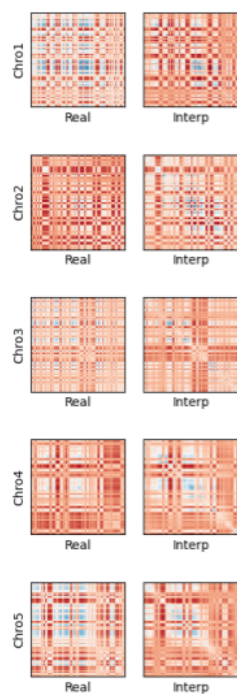
Day B



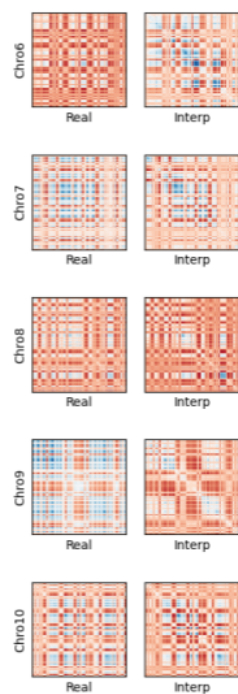
Day B



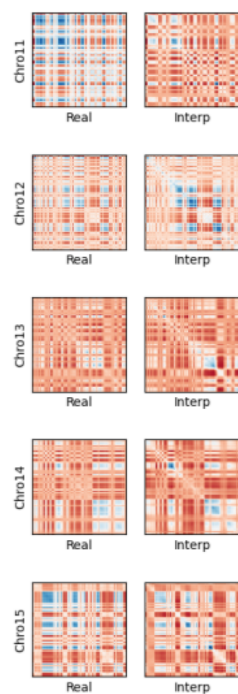
Day D2



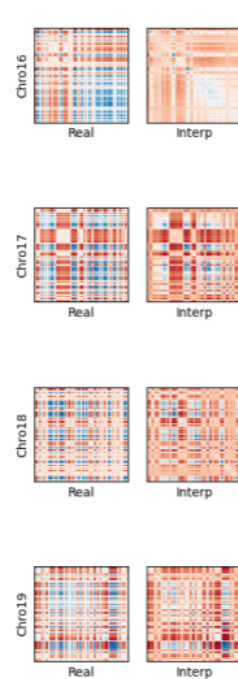
Day D2



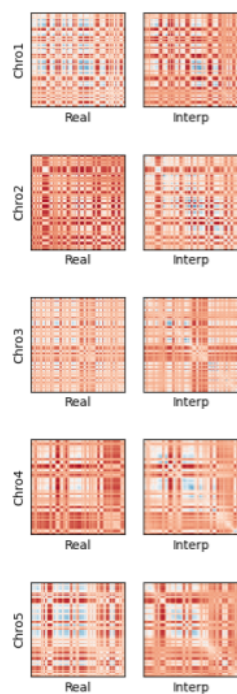
Day D2



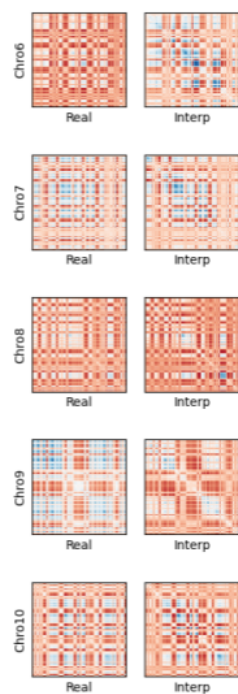
Day D2



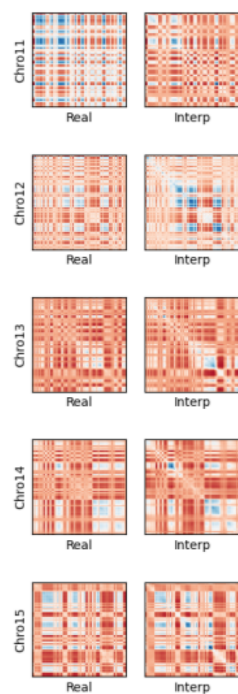
Day D4



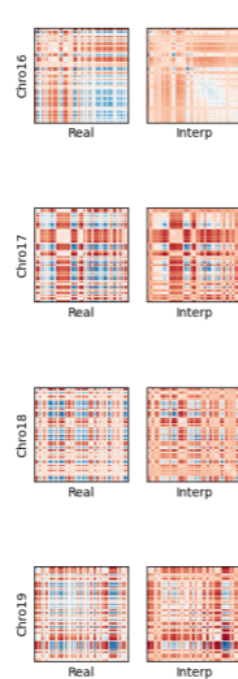
Day D4



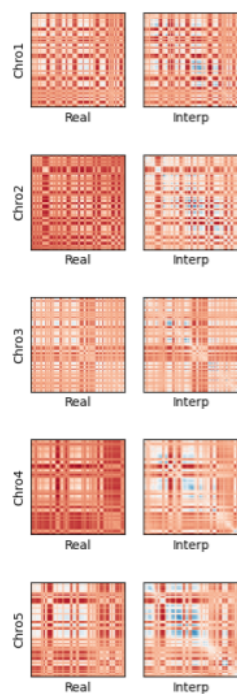
Day D4



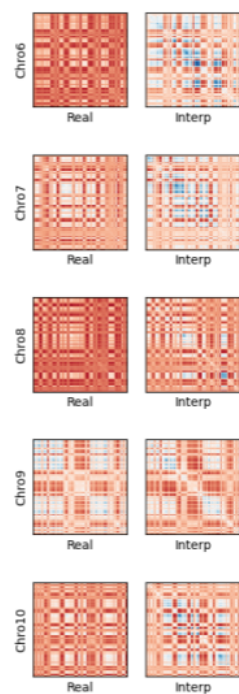
Day D4



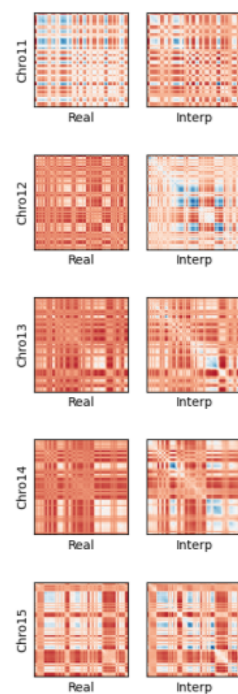
Day D6



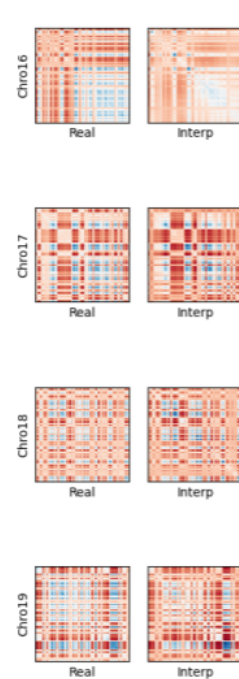
Day D6



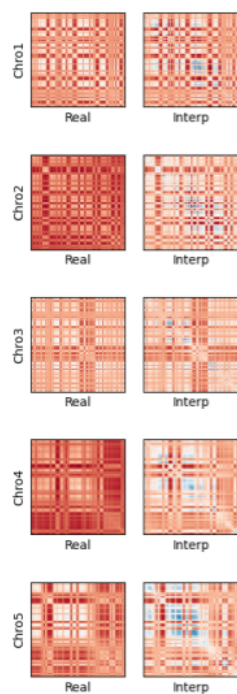
Day D6



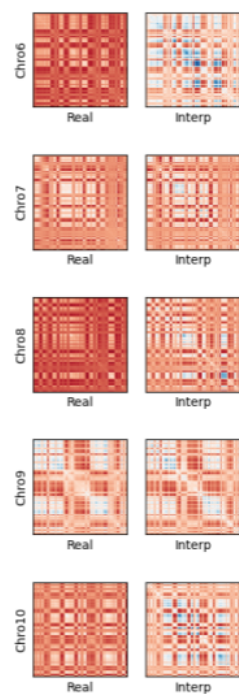
Day D6



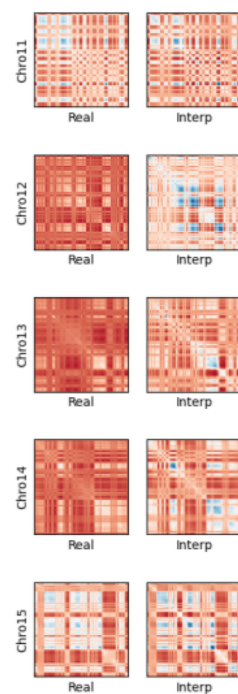
Day D8



Day D8



Day D8



Day D8

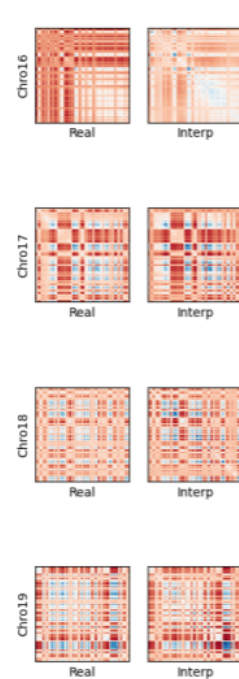




Figure S18 AB Vec Pearson Matrices

AB compartments Pearson correlation matrices for (left) real Hi-C and (right) reconstructed Hi-C data.

# eta	lr	alpha	avg_ pcc
1	1.00E-01	0.6	-0.0317
1	1.00E-01	0.8	-0.0358
1	1.00E-01	1	-0.038
1	1.00E-03	0.6	-0.0017
1	1.00E-03	0.8	-0.0035
1	1.00E-03	1	-0.0071
1	1.00E-04	0.6	0.7492
1	1.00E-04	0.8	0.7162
1	1.00E-04	1	0.6774
1	1.00E-05	0.6	0.354
1	1.00E-05	0.8	0.4005
1	1.00E-05	1	0.4323
1000	1.00E-01	0.6	nan
1000	1.00E-01	0.8	nan
1000	1.00E-01	1	nan
1000	1.00E-03	0.6	-0.2323
1000	1.00E-03	0.8	-0.1823
1000	1.00E-03	1	-0.1452
1000	1.00E-04	0.6	0.7493
1000	1.00E-04	0.8	0.7166
1000	1.00E-04	1	0.6769
1000	1.00E-05	0.6	0.3546
1000	1.00E-05	0.8	0.4209
1000	1.00E-05	1	0.4741
1000000	1.00E-01	0.6	nan
1000000	1.00E-01	0.8	nan
1000000	1.00E-01	1	nan
1000000	1.00E-03	0.6	nan
1000000	1.00E-03	0.8	nan
1000000	1.00E-03	1	nan
1000000	1.00E-04	0.6	nan
1000000	1.00E-04	0.8	nan

1000000	1.00E-04	1	nan
1000000	1.00E-05	0.6	0.252
1000000	1.00E-05	0.8	0.2787
1000000	1.00E-05	1	0.2924

Figure S19 Hyper Parameter Tuning

# Day	Chro	Numerical Interp	4DMax recon	4DMax interp
D2		1	0.734	0.777
D4		1	0.712	0.732
D6		1	0.663	0.693
D2		2	0.751	0.785
D4		2	0.727	0.758
D6		2	0.685	0.727
D2		3	0.756	0.798
D4		3	0.733	0.76
D6		3	0.685	0.726
D2		4	0.764	0.794
D4		4	0.742	0.769
D6		4	0.693	0.721
D2		5	0.772	0.804
D4		5	0.747	0.773
D6		5	0.693	0.712
D2		6	0.748	0.785
D4		6	0.729	0.761
D6		6	0.677	0.699
D2		7	0.775	0.8
D4		7	0.748	0.769
D6		7	0.698	0.703
D2		8	0.764	0.801
D4		8	0.741	0.772
D6		8	0.682	0.697
D2		9	0.763	0.797
D4		9	0.732	0.763
D6		9	0.667	0.692
D2		10	0.764	0.802
D4		10	0.739	0.77
D6		10	0.688	0.72
D2		11	0.791	0.813
D4		11	0.764	0.785

D6	11	0.716	0.767	0.741
D2	12	0.746	0.815	0.776
D4	12	0.715	0.756	0.743
D6	12	0.661	0.721	0.681
D2	13	0.716	0.779	0.742
D4	13	0.689	0.72	0.712
D6	13	0.639	0.7	0.674
D2	14	0.732	0.808	0.774
D4	14	0.707	0.752	0.744
D6	14	0.645	0.712	0.677
D2	15	0.781	0.834	0.802
D4	15	0.76	0.792	0.78
D6	15	0.709	0.762	0.704
D2	16	0.789	0.842	0.807
D4	16	0.773	0.8	0.786
D6	16	0.721	0.769	0.718
D2	17	0.79	0.837	0.803
D4	17	0.771	0.796	0.784
D6	17	0.712	0.752	0.702
D2	18	0.766	0.816	0.782
D4	18	0.737	0.772	0.751
D6	18	0.673	0.73	0.658
D2	19	0.802	0.835	0.794
D4	19	0.791	0.808	0.781
D6	19	0.746	0.781	0.702

Figure S20 Performance Comparison to Numerical Interpolation

Videos

Videos are stored as gif files and are available:

<http://sysbio.rnet.missouri.edu/3dgenome/4DMax/>

S1 Video Collection 1 Synthetic Videos

<http://sysbio.rnet.missouri.edu/3dgenome/4DMax/Movies%201%20Synthetic.zip>

S2 Video Collection 2 iPSC Videos

<http://sysbio.rnet.missouri.edu/3dgenome/4DMax/Movies%202%20iPSC%20Full.zip>

S3 Video Collection 3 Cardio Videos

<http://sysbio.rnet.missouri.edu/3dgenome/4DMax/Movies%203%20Cardio%20Videos.zip>

S4 Video Collection 4 iPSC Missing Chros

<http://sysbio.rnet.missouri.edu/3dgenome/4DMax/Movies%204%20iPSC%20Missing.zip>

S5 Video Collection 5 Cardio Missing Chros

<http://sysbio.rnet.missouri.edu/3dgenome/4DMax/Movies%205%20Cardio%20Missing%20Chros.zip>

S6 Video Collection 6 Changing Resolution

<http://sysbio.rnet.missouri.edu/3dgenome/4DMax/Movies%206%20Changing%20Resolution.zip>

S7 Video Collection 7 Changing Granularity

<http://sysbio.rnet.missouri.edu/3dgenome/4DMax/Movies%207%20Changin%20Granularity.zip>

S8 Video Collection 8 Changing Granularity

<http://sysbio.rnet.missouri.edu/3dgenome/4DMax/Movies%208%20Hyper%20Param.zip>