Figure S4. 3D models of selected domains in the *Drosophila* genome.
Superimposed 3D structures for selected models in cluster #1 for each of the 50 modeled domains. Models are colored by their particle chromatin type as previously defined [1]. They can be directly visualized using TADkit by visiting the Web site http://www.3DGenomes.org/datasets/serra_etal.

2L:58,763-1,067,636	2L:17,502,364-18,533,147	3L:128,671-1,169,578	3R:1,540,632-2,547,798	X:26,253-1,033,981
2L:604,451-1,610,765	2L:21,395,984-22,412,832	3L:218,525-1,232,505	3R:4,627,784-5,632,660	X:1,234,758-2,243,328
2L:2,890,809-3,902,511	2R:510,323-1,538,959	3L:4,420,797-5,552,000	3R :4,935,081-5,939,573	X:2,348,243-3,377,271
2L:3,874,187-4,878,079	2R:1,851,157-2,937,500	3L:8,062,026-9,067,732	3R:10,108,917-11,125,089	X:2,831,084-3,850,088
2L:6,045,705-7,078,729	2R:3,129,889-4,144,094	3L:10,619,399-11,633,875	3R:11,830,184-12,834,808	X:11,803,883-12,972,782
2L:7,882,539-8,889,439	2R:6,628,590-7,641,170	3L:16,030,279-17,045,046	3R:16,737,550-17,758,875	X:13,910,272-14,912,855
2L:7,954,116-8,962,062	2R:9,877,764-10,888,792	3L:22,793,448-23,817,780	3R:18,922,749-19,929,202	X:15,340,984-16,350,132
2L:8,533,966-9,545,865	2R:15,391,841-16,436,911	3R:241-1,012,283	3R:19,748,864-20,776,661	X:15,590,302-16,600,049
2L:9,516,384-10,525,431	2R:19,557,610-20,564,425	3R:68,298-1,076,568	3R:23,123,438-24,126,051	X:19,389,629-20,397,756
2L:15,185,753-16,246,568	2R:20,132,958-21,145,985	3R:1,189,408-2,192,628	4:60,451-1,083,771	X:21,107,582-22,132,184