

## Correction: Transcription Factors Bind Thousands of Active and Inactive Regions in the *Drosophila* Blastoderm

Xiao-yong Li, Stewart MacArthur, Richard Bourgon, David Nix, Daniel A. Pollard, Venky N. Iyer, Aaron Hechmer, Lisa Simirenko, Mark Stapleton, Cris L. Luengo Hendriks, Hou Cheng Chu, Nobuo Ogawa, William Inwood, Victor Sementchenko, Amy Beaton, Richard Weiszmann, Susan E. Celniker, David W. Knowles, Tom Gingeras, Terence P. Speed, Michael B. Eisen, Mark D. Biggin

Correction for:

Li Xy, MacArthur S, Bourgon R, Nix D, Pollard DA, et al. (2008) Transcription factors bind thousands of active and inactive regions in the *Drosophila* blastoderm. PLoS Biol 6(2): e27. doi:10.1371/journal.pbio.0060027

The information in Table 1 for RNA polymerase II was incorrectly given for the form of the enzyme unphosphorylated at the C-terminal tail, which is recognized by the 8WG16 monoclonal antibody. The corrected version of the Table below gives the intended information for the enzyme phosphorylated at the C terminus, which is recognized by the H14 monoclonal antibody.

**Table 1.** Number of Regions Bound by Transcription Factors

Transcription Factors	Type	Antibody	Symmetric-Null Test				IgG Control Test			
			Number of Bound Regions		Overlap Between Bound Regions <sup>a</sup>		Number of Bound Regions		Overlap Between Bound Regions <sup>a</sup>	
			1%FDR	25%FDR	1% - 1%	1% - 25%	1%FDR	25%FDR	1% - 1%	1% - 25%
BCD	Maternal	Anti-BCD1	692	5,312	72%	97%	514	1,130	81%	94%
		Anti-BCD2	578	3,833	86%	99%	524	1,039	80%	96%
CAD	Maternal	Anti-CAD	1,392	6,484	NA	N/A	1,013	5,109	NA	N/A
HB	Gap	Anti-HB1	1,789	5,899	75%	94%	1,650	3,592	62%	84%
		Anti-HB2	1,726	8,865	80%	96%	1,174	2,400	91%	98%
KR	Gap	Anti-KR1	3,072	11,140	89%	99%	2,255	4,775	88%	99%
		Anti-KR2	3,415	10,774	80%	98%	2,274	4,440	87%	99%
GT	Gap	Anti-GT	966	3,958	NA	N/A	898	3,973	NA	N/A
KNI	Gap	Anti-KNI1	37	338	100	100%	117	530	90	98%
		Anti-KNI2	199	4,998	18%	87%	199	792	52%	94%
RNA Pol II	General	H14	1,822	6,927	NA	N/A	1,299	3,587	NA	N/A

<sup>a</sup>Percentage of 1% FDR bound regions for one antisera that overlap at least 500 bp with a 1% or 25% FDR bound region for the other antisera for the same factor. doi:10.1371/journal.pbio.0060190.t001

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