



## Erratum

*Genome Res.* 2009 19: 690

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## Errata

### Genome Research 18: 631–639 (2008)

#### High-throughput biochemical analysis of *in vivo* location data reveals novel distinct classes of POU5F1(Oct4)/DNA complexes

Dean Tantin, Matthew Gemberling, Catherine Callister, and William Fairbrother

In the author list for this paper, author name William Fairbrother should have appeared as William G. Fairbrother.

The authors apologize for any confusion this may have caused.

### Genome Research 19: 500–509 (2009)

#### Mapping DNA structural variation in dogs

Wei-Kang Chen, Joshua D. Swartz, Laura J. Rush, and Carlos E. Alvarez

The authors have discovered minor changes in the data, which affect one sentence in text and one species row in Table 2.

On page 501, the fourth sentence beneath the heading “aCGH of pedigreed dogs” should read:

We identified 152 CNVs—in 57 CNV regions (CNVRs)—at high confidence (Fig. 2; Table 1; Supplemental Table S2).

In addition, the row that displays the data for the Dog species is incorrect and has been replaced. The corrected Table 2 is reprinted below.

**Table 1B.** Comparison of dog CNV to that reported in other mammals

| Species  | Total no. of CNVs | Mean no. of CNVs per individual                                   | No. of CNVRs | No. of deletions, no. of amplifications | Amplified avg. (mean)  | Deletion avg. (mean)  | References           |
|--|-------------------|---|--------------|---|--|-----------------------|----------------------|
| Dog ( <i>n</i> = 9)  | 152               | 16.9 (385 k oligo aCGH)   | 57           | 85, 67                                  | 476.5 kb   | 272.2 kb              | This study           |
| Mouse ( <i>n</i> = 21)   | 80                | 22 (385 k oligo aCGH)   |              |   |  | 271.5 kb <sup>a</sup> | Graubert et al. 2007 |
| Rat ( <i>n</i> = 10, 5 M oligo <sup>b</sup> ; <i>n</i> = 3, 375 k) | 643               | 63 (5 M exon-specific oligo <sup>b</sup> ); 11 (385 k oligo aCGH) |              |   | 5.2 kb (5 M exon-specific oligo <sup>b</sup> ); 256 kb (375 k oligo aCGH) <sup>a</sup> | 102 kb <sup>a</sup>   | Guryev et al. 2008   |
| Macaque ( <i>n</i> = 9)  | 123               | 14 (385 k oligo aCGH)   |              |   |  |                       | Lee et al. 2008      |
| Human ( <i>n</i> = 270)  | 1447              | 70 (BAC aCGH); 24 (SNP array)                                     |              |   | 341 kb (BAC aCGH); 206 kb (SNP array) <sup>a</sup>                                     |                       | Redon et al. 2006    |

<sup>a</sup>Average CNV size, amplifications and deletions combined.

<sup>b</sup>5 M exon-specific, non-CGH oligo array.