



Correction for Volume 15, p. 376

Genome Res. 2005 15: 601

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Errata**Genome Research 15: 231–240 (2005)****Divergent V1R repertoires in five species: Amplification in rodents, decimation in primates, and a surprisingly small repertoire in dogs**

Janet M. Young, Marijo Kambere, Barbara J. Trask, and Robert P. Lane

The line designating intact V1R's was inadvertently omitted from Table 1. The complete Table 1 is reprinted below.

Table 1. Number of V1R-like sequences found in the genomes of five species

Mouse	Rat	Dog	Chimpanzee	Human	
364	220	65	116	117	Total V1R-like sequences
165 (50%)	106 (49%)	8 (13%)	0 (0%)	2 (2%)	Intact V1Rs
165 (50%)	110 (51%)	54 (87%)	102 (100%)	115 (98%)	Pseudogenes
34	4	3	14	0	Partial sequences
95%	90%	98%	94%	100%	Assembly coverage ^a
190	120	9	0	2	Intact V1Rs extrapolated to complete genome

Partial sequences are interrupted by assembly gaps and could be intact or pseudogenes. Complete sequence information is needed before their status can be determined. The percentages shown are intact and pseudogene proportions of all V1Rs with full-length sequence available.

^aSee Methods.

Genome Research 15: 376–384 (2005)**Protein interaction mapping: A *Drosophila* case study**

Etienne Formstecher, Sandra Aresta, Vincent Colluar, Alexandre Hamburger, Alain Meil, Alexandra Trehin, Céline Reverdy, Virginie Betin, Sophie Maire, Christine Brun, Bernard Jacq, Monique Arpin, Yohanns Bellaiche, Saverio Bellusci, Philippe Benaroch, Michel Bornens, Roland Chanet, Philippe Chavrier, Olivier Delattre, Valérie Doye, Richard Fehon, Gérard Faye, Thierry Galli, Jean-Antoine Girault, Bruno Goud, Jean de Gunzburg, Ludger Johannes, Marie-Pierre Junier, Vincent Mirouse, Ashim Mukherjee, Dora Papadopoulo, Franck Perez, Anne Plessis, Carine Rossé, Simon Saule, Dominique Stoppa-Lyonnet, Alain Vincent, Michael White, Pierre Legrain, Jérôme Wojcik, Jacques Camonis, and Laurent Daviet

The link, <http://imex.mbi.ucla.edu:60606/imex/IMEX.jsp> provided in the second column on page 382 in this article and in the Web site references is dysfunctional. However, the *Drosophila* protein interaction data are also available from BIND (<http://bind.ca>), Flybase (<http://flybase.bio.indiana.edu/>), and the authors' Web site (<http://pim.hybrigenics.com>). The authors apologize for any inconvenience this may have caused other investigators in the field.