

1. The RIKEN Genome Exploration Research Group Phase I & II Team and FANTOM Consortium, Analysis of the mouse transcriptome based on functional annotation of 60,770 full-length cDNA, Nature, **420**, 563-573, 2002.
 2. Kawai J., Hayashizaki Y., DNABook, Genome Research, Special FANTOM Issue, **13**, Number 6b, June 2003.
 3. <http://geneome.riken.jp/>
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