

CORRECTION

Correction: DNA microarray analysis of rheumatoid arthritis susceptibility genes identified by genome-wide association studies

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See related research by Sugino *et al.*, <http://arthritis-research.com/content/12/2/401>

After publication of our recent article [1], we noticed an error in Table 1. Gene expression of STAT4 should be listed as “down” and not “up”. The description of the table in the article is unchanged and the corrected table is provided here.

Competing interests

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References

1. Sugino H, Lee HM, Nishimoto N: DNA microarray of rheumatoid arthritis susceptibility genes identified by genome-wide association studies. *Arthritis Research & Therapy* 2010, **12**:401.

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Table 1. Candidate genes identified from rheumatoid arthritis genome-wide association studies

Gene	GeneID	PMID	Gene expression (up or down)	Microarray P values ^a
CD244	605554	18794858	Up	1.0 x 10 ⁻¹⁶
PADI4	605347	12833157	Up	2.32 x 10 ⁻⁵
SLC22A2	602608	14608356	Up	1.94 x 10 ⁻⁶
PTPN22	600716	15208781	Up	9.66 x 10 ⁻⁸
CTLA4	123890	16380915	No change	0.767
STAT4	600558	17804842	Down	1.38 x 10 ⁻⁸

^aP values determined by comparison between 112 rheumatoid arthritis patients and 45 healthy individuals.

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