

Mourão AF, Caetano-Lopes J, Costa P, et al. Tumor necrosis factor- α -308 genotypes influence inflammatory activity and tumor necrosis factor- α serum concentrations in children with juvenile idiopathic arthritis. *J Rheumatol* 2009;36:837-42. In the Materials and Methods section, the details of DNA analysis and TNF- α assessment should have been given as follows:

“TNF- α gene -308 polymorphisms (G/A) were analyzed by RFLP using the forward primer 5'-AAT AGG TTT TGA GGG CCA TG-3' and the reverse primer 5'-ATC TGG AGG AAG CGG TAG TG-3'. The forward primer contained one nucleotide mismatch (underlined above), which allowed use of the restriction enzyme *NcoI* (New England Biolabs, Hitchin, UK) for detection of -308G/A polymorphisms.”

The following Table 1 should have been included. We regret the errors.

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Table 1. TNF- α promoter -308 genotype frequency in JIA subgroups and controls.

Group	Total No.	-308 GG N (%)	-308 GA/AA N (%)	p*
Control group	117	91 (77.8)	26 (22.2)	
JIA patients	114	87 (76.3)	27 (23.7)	0.876
Oligoarticular persistent	48	43 (89.6)	5 (10.4)	0.080
Oligoarticular extended	17	13 (76.5)	4 (23.5)	1.000
Polyarticular	24	15 (62.5)	9 (37.5)	0.123
Enthesitis-related arthritis	10	7 (70.0)	3 (30.0)	0.694
Systemic	9	6 (66.7)	3 (33.3)	0.684
Psoriatic arthritis	6	3 (50.0)	3 (50.0)	0.141

* Patient's genotype in comparison with the control group.