

# Gene silencing of EREG mediated by DNA methylation and histone modification in human gastric cancers

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In this article, there is an error in Table 1. A revised Table 1 is provided below.

**Table 1 Methylation of EREG in normal and cancerous primary gastric tissue samples**

Normal tissue samples			Cancer tissue samples		
% methylation	Region 2	Region 4	% methylation	Region 2	Region 4
0 < % < 20	7/13 (54%)	13/13 (100%)	0 < % < 20	7/13 (54%)	11/13 (85%)
20 < % < 40	6/13 (46%)	0/13 (0%)	20 < % < 40	2/13 (15%)	2/13 (15%)
40 < % < 60	0/13 (0%)	0/13 (0%)	40 < % < 60	1/13 (8%)	0/13 (0%)
60 < % < 80	0/13 (0%)	0/13 (0%)	60 < % < 80	3/13 (23%)	0/13 (0%)
80 < % < 100	0/13 (0%)	0/13 (0%)	80 < % < 100	0/13 (0%)	0/13 (0%)