Erratum to: A novel missense mutation of *CYLD* gene in a Chinese family with multiple familial trichoepithelioma

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Unfortunately, the original version of this article contains some mistakes in “Results” section (the correction also reflects some changes in Figure caption 3 and Table 1), which we correct here.

Results

Direct DNA sequencing showed a novel heterozygous missense mutation (2335G>A) in exon 17 of *CYLD* gene.

This mutation results in replacement of Asp (GAT) by Asn (AAT) at amino acid Asp779Asn (Fig. 3). It was not found in any of the healthy controls showing that it is a novel pathogenic mutation, not a common polymorphism.

Figure caption 3

Mutant sequence and wild-type sequence show missense mutation in exon 17 causing a Asp779Asn change in *CYLD* gene.

The online version of the original article can be found under doi:[10.1007/s00403-009-1003-1](https://doi.org/10.1007/s00403-009-1003-1).

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Table 1 Summary of *CYLD* gene mutations in patients with multiple familial trichoepithelioma

Family no.	Mutation	Amino-acid change	Mutation location	Mutation type	Reference
1	1364_1365delAA	454fsX476	Exon 10	Nonsense	Liang [9]
2	1462delA	Ile488fsX497	Exon 10	Frameshift	Zheng [18]
3	1787G>A	Gly596Asp	Exon 11	Missense	Zuo [19]
4	1862 + 2T>G	–	Exon 12	Splice	Liang [10]
5	IVS12 + 1G>A	–	Exon 12	Splice	Huang [5]
6	2104-2105insA	702fsX723	Exon 15	Frameshift	Salhi [13]
7	2241-2242delAG	Glu747fsX763	Exon 16	Nonsense	Liang [10]
8	2128C>T	Gln710X	Exon 17	Nonsense	Zheng [18]
9	2335G>A	Asp779Asn	Exon 17	Missense	This study
10	2339T>G	Leu780X	Exon 17	Nonsense	Bowen [2]
12	2355-2358delCAGA	Pro785fsX917	Exon 18	Nonsense	Zhang [17]
13	2409C>G	Tyr803X	Exon 18	Nonsense	Liang [9]
14	2687G>C	Gly896Ala	Exon 20	Missense	España [3]
15	2711C>T	Pro904Leu	Exon 20	Missense	Lv [7]
16	2822A>T	Asp941Val	Exon 20	Missense	Zheng [18]