

Erratum to: Impact of New Genomic Technologies on Understanding Adverse Drug Reactions

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In the online published article **Section 3, para 2, lines 4–5**, which previously read:

...Gordon et al. applied targeted NGS of multiple genes from over 14,000 subjects, to illustrate the extent of potentially...

Should read:

...Gordon et al. applied targeted NGS of multiple genes from 6503 subjects, to illustrate the extent of potentially...

The online version of the original article can be found under doi:[10.1007/s40262-015-0324-9](https://doi.org/10.1007/s40262-015-0324-9).

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