



Erratum

## Correction: Construction of a multiplex mutation hot spot PCR panel: the first step towards colorectal cancer genotyping on the GS Junior platform

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The Acknowledgments section in our paper [1] was incomplete and should instead read as follows: The results published here are in whole or part based upon data generated by The Cancer Genome Atlas managed by the NCI and NHGRI (http://cancergenome.nih.gov). The TCGA data analyzed here are available through dbGAP, accession phs000178.v9.p8 (http://www.ncbi.nlm.nih.gov/projects/gap/cgi-bin/study.cgi?study\_id=phs000178.v9.p8).

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## References

1. Péterfia B, Kalmár A, Patai ÁV, Csabai I, Bodor A, Micsik T, Wichmann B, Egedi K, Hollósi P, Kovalszky I, Tulassay Z, Molnár B. Construction of a multiplex mutation hot spot PCR panel: the first step towards colorectal cancer genotyping on the GS Junior platform. J Cancer 2017; 8(2):162-173. doi:10.7150/jca.16037.