Correction

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Corrigendum

Oviz-Bio: a web-based platform for interactive cancer genomics data visualization

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In Figure 1, the protein layer of gene PIK3CA should start from the NO.2 exon. An updated figure is provided below, and the published article has been updated.

This correction does not affect the results and conclusions of the article.

REFERENCE


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†The authors wish it to be known that, in their opinion, the first two authors should be regarded as joint First Authors.

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Figure 1. Demo representation of the ‘Mut On Genes’ visualization and features. Mutations on gene PIK3CA are displayed in three layers (genome, cDNA and protein), which can be switched by clicking layer tags. Mutations are denoted by different icons according to their types and function changes. Icons have tooltips containing related sample list. Exons and mutations are linked among the three layers to support synchronous highlights. Function domains are denoted by colored areas in the protein layer. The mutation list is downloaded from the TumorPortal database (40).