

I. CancerGenomicTestStdzFormat Details Correction

The corrections to the released details.

※It does not describe the version that is no longer supported.

| No. | Target version | Correction point. (Before) | Correction |
|-----|----------------|-------------------------------|--|
| 1 | v1.0.2 | P.27-28 | Corrected the description of the genePairs key as follows. |
| | | | <div>Before:</div> <p>* If this tag is omitted (and there is no mention in matePieceLocation), any pair of geneSymbols will be searched for unordered pairs of geneSymbols, but if you want to control the search in detail, please describe it as ["A-B", "B-A", "C-B"].</p> <div>After:</div> <p>* If this tag is omitted any pair of geneSymbols will be searched for unordered pairs of geneSymbols, but if you want to control the search in detail, please describe it as ["A-B", "B-A", "C-B"].</p> |
| 2 | v1.1.1 | P.28 | Same as above. |

II. For inquires

Please contact the C-CAT Help Desk.

E-Mail: helpdesk_c-cat@ml.res.ncc.go.jp