

Erratum

Erratum to “High incidence of the CFTR mutations 3272-26A → G and L927P in Belgian cystic fibrosis patients, and identification of three new CFTR mutations (186-2A → G, E588V, and 1671insTATCA)”
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The publisher regrets that a typing error has occurred in the above mentioned article relating to the name of one of the three new mutations identified in this study.

Instead of 1671insTATCA it must be:

- according to the old nomenclature: 1651insTATCA or 1651_1652insTATCA (standard numbering scheme for CF).
- according to the new nomenclature (nomenclature system as proposed by www.hgvs.org): c.1515_1519dupTATCA (p.Phe508SerfsX21) (with a stop at codon 528) (numbering according to GenBank NM_000492 with A of the start codon as position +1).

The mutant sequence is:

ATT AAA GAA AAT ATC ^A|**ta tca**| TCT TTG GTG TTT
CCT ATG ATG AAT ATA GAT ACA GAA GCG TCA TCA
AAG CAT GCC AAC TAG

The caret marks the start of codon 507, the inserted bases are shown in bold and lower case, and the premature termination codon is underlined. At protein level the first affected amino acid is at codon 508.

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