

ERRATUM

doi:10.1038/nature08491

Structural basis for leucine-rich nuclear export signal recognition by CRM1

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Nature 458, 1136–1141 (2009)

In the first paragraph of this Article, Leu was incorrectly listed as Lys. The correct sentence is: “Several mutagenesis and computational studies have led to a consensus sequence ϕ -X(2–3)- ϕ -X(2–3)- ϕ -X- ϕ (in which ϕ is Leu, Val, Ile, Phe or Met, X is any amino acid, and the numbers in parentheses denote the number of repeats) that matches most known LR-NESs, but it is so broad that it is found in most helix-containing proteins^{17,18,19,20,21}.”