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Erratum: Context-dependent protein stabilization by methionine-to-leucine substitution shown in T4 lysozyme (Protein Science (November 3, 1998) 7:3 (772))

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ERRATUM

Crystal structures of the psychrophilic α -amylase from *Alteromonas haloplanctis* in its native form and complexed with an inhibitor

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On page 565 in Volume 7, Number 3, 1998, under *Molecular fold and calcium binding site*, the domain numbering was wrongly reported. The correct numbering is domain A: residues 1 to 86 and 147 to 356, and domain B: residues 87 to 146. Domain C was correctly cited and remains as reported.

ERRATUM

Context-dependent protein stabilization by methionine-to-leucine substitution shown in T4 lysozyme

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On page 772 in Volume 7, Number 3, 1998, Table 5 of the above-referenced article should read as follows:

Table 5. *Contacts between introduced leucine side chains and surrounding atoms based on model building^a*

Model-built side chain		Contact atom in WT* lysozyme	Distance (Å)
Residue	Atom		
Leu 6	C ^{δ2}	Asp 10 O ^{δ1}	2.9
	C ^{δ2}	Asn 101 N ^{δ2}	2.9
	C ^{δ2}	Tyr 161 C ^{ε1}	3.0
	C ^{δ2}	Tyr 161 C ^ζ	3.0
	C ^{δ2}	Tyr 161 OH	2.9
Leu 102	C ^{δ2}	Trp 138 ^{ζ2}	3.1
	C ^{δ2}	Val 149 C ^{δ1}	3.1
Leu 106	None	None	—
Leu 120	None	None	—

^aThe table shows all interatomic approaches less than 3.2 Å between the model-built leucine side chain and the surrounding atoms in the model of WT* lysozyme.

Reprints of the article, with corrections, are available from B.W.M. (brian@uoxray.uoregon.edu).