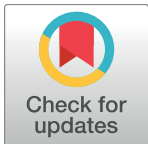


CORRECTION

# Correction: Thermal stability and structural changes in bacterial toxins responsible for food poisoning

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There is an error with the Turn content of the SEE pH 6.0 Zn<sup>2+</sup> sample. Please see the corrected [Table 1](#) here.



## OPEN ACCESS

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**Table 1. Theoretical and experimental content of secondary structure (%) determined for SEA, SEE and SEH at pH 6.0 in presence of EDTA or Zn<sup>2+</sup>.**

SE protein	$\alpha$ -helix	$\beta$ -sheet	Turn	Unordered	RMSD**
Theoretical SEA (1ESF)*	17.6	32.1	12.0	38.3	–
SEA pH 6.0 EDTA	10.5	39.4	16.5	33.6	0.119
SEA pH 6.0 Zn <sup>2+</sup>	12.2	41.7	9.3	36.8	0.101
Theoretical SEE (5FK9)*	15.5	26.5	8.2	49.8	–
SEE pH 6.0 EDTA	6.6	46.2	14.4	32.8	0.151
SEE pH 6.0 Zn <sup>2+</sup>	8.6	32.6	18.9	39.9	0.178
Theoretical SEH (1ENF)*	18.3	41.0	9.0	31.7	–
SEH pH 6.0 EDTA	13.2	37.4	11.2	38.2	0.093
SEH pH 6.0 Zn <sup>2+</sup>	12.1	31.1	12.5	44.3	0.084

\*PDB accession code,

\*\*RMSD between experimental and fitted data.

<https://doi.org/10.1371/journal.pone.0175989.t001>

## Reference

1. Regenthal P, Hansen JS, André I, Lindkvist-Petersson K (2017) Thermal stability and structural changes in bacterial toxins responsible for food poisoning. PLoS ONE 12(2): e0172445. <https://doi.org/10.1371/journal.pone.0172445> PMID: 28207867