

Correction

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**Genome-wide subcellular localization of putative outer membrane and extracellular proteins in *Leptospira interrogans* serovar Lai genome using bioinformatics approaches**Wasna Viratyosin\*<sup>1</sup>, Supawadee Ingsriswang<sup>1</sup>, Eakasit Pacharawongsakda<sup>1</sup> and Prasit Palittapongarnpim<sup>1,2</sup>

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**Correction**

After the publication of this work [1], we discovered a typographic error in Table 6 (56 Putative extracellular proteins (EX) derived from the 1 out of 3 predictions with significant DBSubloc or/and GTD prediction). For the entry of LA3370/LIC10793 (a conserved hypothetical protein/surface antigen) [1], the homologous structure predicted by GTD should read, "1l0q", a structural surface layer of protein binding containing tandem beta propeller. In the final print version, it appeared as 1loq (letter o replacing zero). This distinction is important as 1loq is the PDB structural code for a different protein, Orotidine 5'-phosphate decarboxylase, which is localized as cytoplasmic and not extracellular or secreted.

**References**

1. Viratyosin W, Ingsriswang S, Pacharawongsakda E, Palittapongarnpim P: **Genome-wide subcellular localization of putative outer membrane and extracellular proteins in *Leptospira interrogans* serovar Lai gen.** *BMC Genomics* 2008, **9**:181.