



## ZitR purification [↗](#)

PLOS One

[PALOMA VARELA](#)<sup>1</sup>

<sup>1</sup>Institute for Integrative Biology of the Cell, CEA, CNRS, Université Paris-Saclay, Gif-sur-Yvette, France

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[PALOMA VARELA](#)

### ABSTRACT

### EXTERNAL LINK

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### THIS PROTOCOL ACCOMPANIES THE FOLLOWING PUBLICATION

Varela PF, Velours C, Aumont-Niçaise M, Pineau B, Legrand P, Poquet I (2019) Biophysical and structural characterization of a zinc-responsive repressor of the MarR superfamily. PLoS ONE 14(2): e0210123. doi: [10.1371/journal.pone.0210123](https://doi.org/10.1371/journal.pone.0210123)

### PROTOCOL STATUS

**Working**

### MATERIALS TEXT

An untagged, recombinant (S2A, A4R, D8E) form of ZitR protein (145 amino acids, full length) from *L. lactis* subsp. *cremoris* strain MG1363 (UniProtKB/Swiss-Prot A2RNS2) was produced and purified as previously described (Lull et al, 2011). Recombinant ZitRMG protein was first over-produced in *E. coli* strain (BL21(DE3) (pVE8073) at 25 °C. It was then purified in the presence of ZnCl<sub>2</sub> in 2-steps by anion exchange chromatography followed by heparin-affinity chromatography. SDS-PAGE analysis revealed the presence of a few high molar mass contaminating *E. coli* proteins (not shown), which could be eliminated by gel filtration chromatography in 20 mM Tris-HCl (pH 7.0), 200 mM NaCl and 100 μM ZnSO<sub>4</sub>. At the end of this purification process, we observed a major dimeric form, and a minor tetrameric form (data not shown). The major dimeric form was subsequently studied.

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