



Size Exclusion Chromatography coupled to Multi Angle Light Scattering (SECMALS) [↗](#)

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[PALOMA VARELA](#)¹

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

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[PALOMA VARELA](#)

EXTERNAL LINK

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THIS PROTOCOL ACCOMPANIES THE FOLLOWING PUBLICATION

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PROTOCOL STATUS

Working

MATERIALS TEXT

To determine the oligomeric state of ZitR_{MG}, 50 ml of fully purified protein sample at approximately 1-4 mg.mL⁻¹ were loaded on a KW-803 column (Shodex) equilibrated at 0.5 mL.min⁻¹ flow rate (Shimadzu HPLC system) and supplemented with either 1 mM ZnSO₄ or 1 mM EDTA (Fig. S1). Detection was performed using a MiniDAWN TREOS multi angle light scattering detector and an Optilab T-rEX differential refractometer (Wyatt Technology) (biophysics platform, LEBS/IMAGIF). Molar mass and hydrodynamic radius were calculated with the Astra 6.1.1.17 software using a differential index of refraction (dn/dc) value of 0.183 mL.g⁻¹.



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