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FIG S1 Determination of the quaternary structure of AldR by gel filtration chromatography.

The purified AldR protein in 20 mM Tris-HCl buffer (pH 8.0) was incubated with 10 mM L
alanine (+Ala) or 10 mM L-cysteine (+Cys) for 1 h and subjected to gel filtration

chromatography on Superose 12 10/300 GL (24 ml, GE Healthcare) equilibrated with 20 mM

Tris-HCl buffer (pH 8.0) containing the corresponding amino acids. As the control, the

purified AldR protein in 20 mM Tris-HCl buffer without the amino acid was included in the experiment (control). (A) Elution profiles of the purified AldR protein. The elution volumes are indicated above the peaks representing the eluted AldR proteins. The left peaks are the aggregated AldR proteins that were eluted at the void volume of the column. The purified AldR protein was resolved in SDS-PAGE and stained with Coomassie brilliant blue. The theoretical molecular mass of His₆-tagged AldR is 19.4 kDa. (B) The plot of log molecular mass of each protein versus the ratios of elution volume to void volume (Ve/Vo) of the protein. The calculated molecular masses of native AldR in the absence of the amino acid (control) and presence of L-alanine (+Ala) or L-cysteine (+Cys) are given in the plot. Abbreviations: ADH, alcohol dehydrogenase; BSA, bovine serum albumin; CA, carbonic anhydrase.