STRUCTURAL BIOLOGY

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Supporting information for article:

Single support serial isomorphous replacement phasing

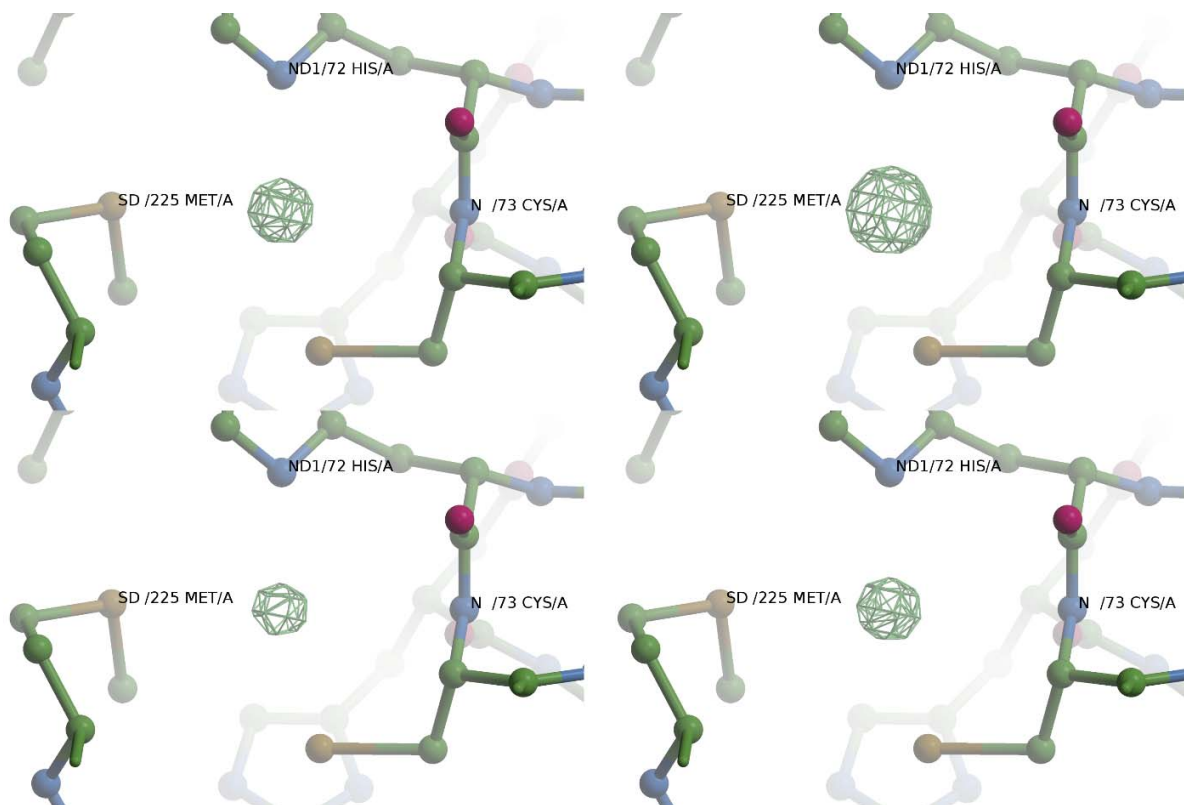Nicolas Foos, Mahmoud Rizk and Max H. Nanao

Figure S1 Fo-Fc difference maps of proteinase K for each CODGAS group (clockwise from upper left, group 1,2,3 and 4), contoured at 35 standard deviations above the mean difference density value. Maps were calculated using phenix.maps (Adams *et al.*, 2010) using default parameters. Density images were created in Coot (Emsley *et al.*, 2010).