

Supplementary Data

**ORPHAN MACRODOMAIN (HUMAN C6ORF130) IS AN *O*-ACYL-ADP-RIBOSE
DEACYLASE: SOLUTION STRUCTURE AND CATALYTIC PROPERTIES**
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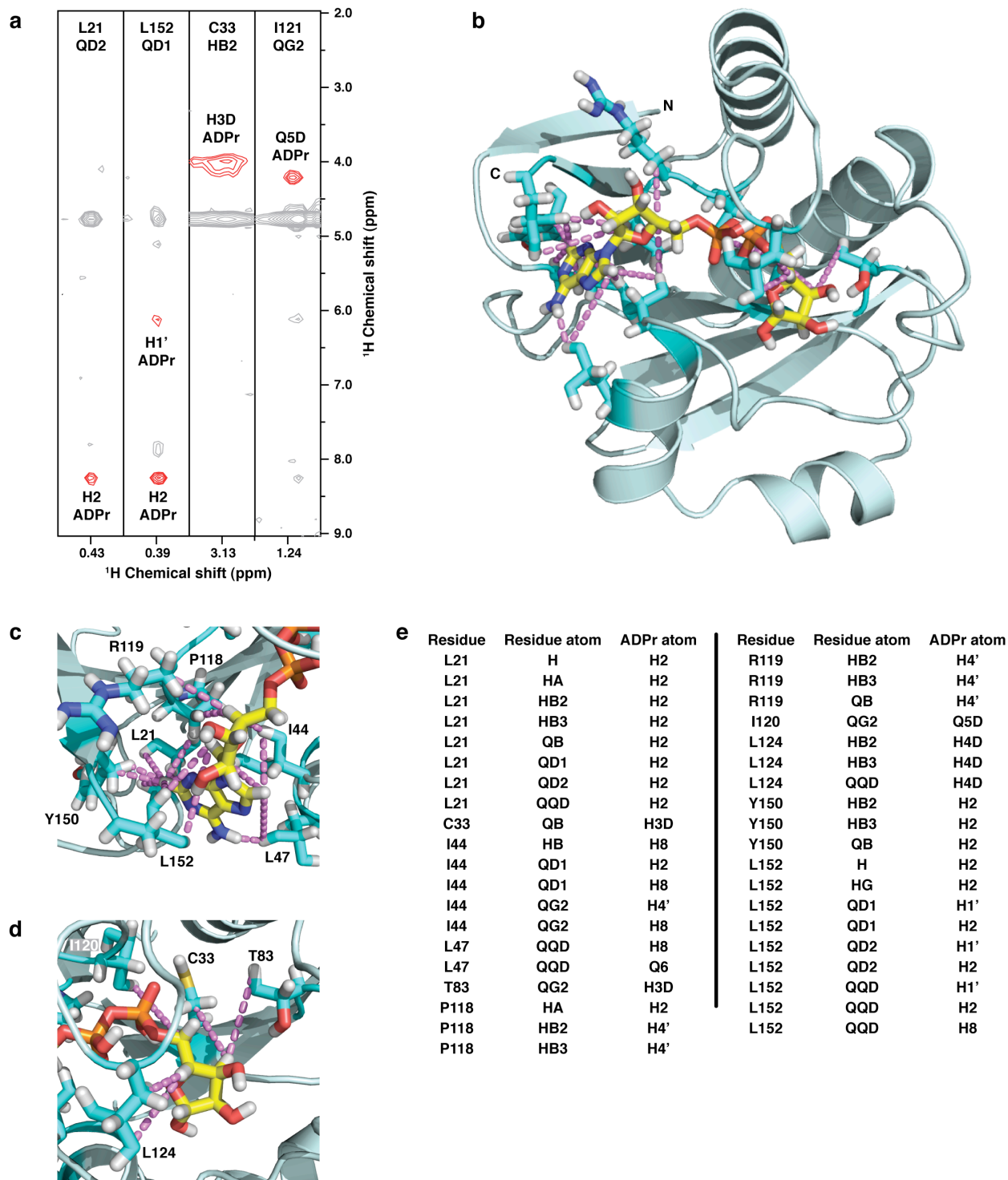


Figure S1. Intermolecular NOE contacts defining the C6orf130:ADPr complex. (a) Representative strips from the 3D ^{13}C -edited/ ^{13}C -filtered NOESY spectrum containing intermolecular NOE crosspeaks. (b) Overview of the bound ADPr ligand with NOE distance constraints. Close-up views of the adenosine and proximal ribose (c) and distal ribose (d) with intermolecular NOE constraints. (e) All intermolecular NOEs used to solve the structure are listed.

Table S1. Sequence comparison of C6orf130 protein with macro family proteins

Proteins	C6orf130 ^a	MacroD1 ^b	MacroD2	<i>E. coli</i> YmdB	<i>S. aureus</i>
SAV0325					
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C6orf130		19.7 ^c	19.1	17.8	21.7
macroD1			61.1	47.2	38.5
macroD2				49.7	32.3
<i>E. coli</i> YmdB					40.7
<i>S. aureus</i> SAV0325					

^aMungall, A. J. et al. (2003) *Nature* **425**, 805-811.

^bChen, D. et al. (2011) *J. Biol. Chem.* **286**, 13261-13271

^cNumbers indicate percentage of sequence identity between the overlapping region of the two sequences compared.