

**Supplemental Information**  
**for**  
**Leveraging the Structure of DNAJA1 to Discover Potentially Novel Pancreatic Cancer Therapies**

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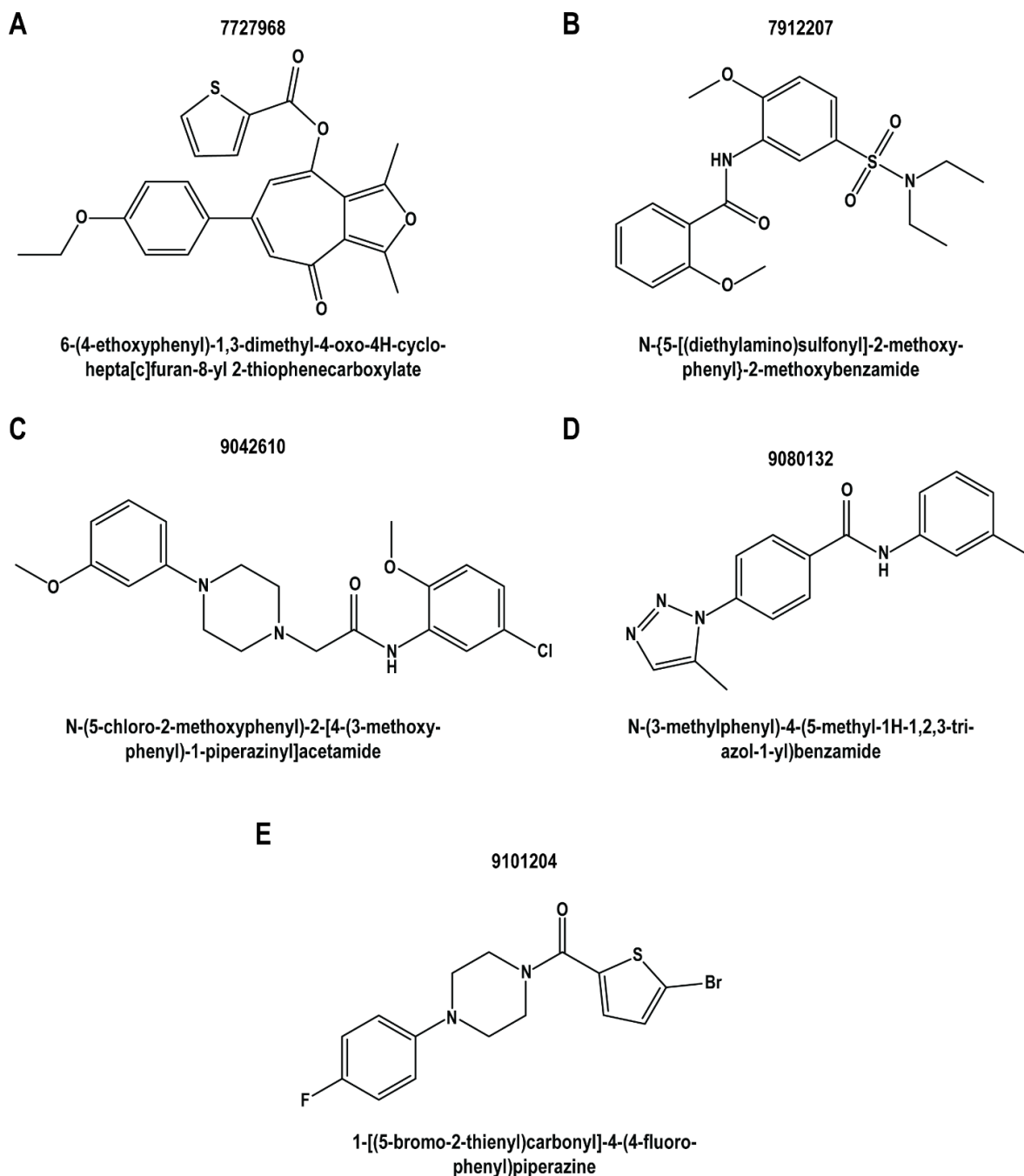
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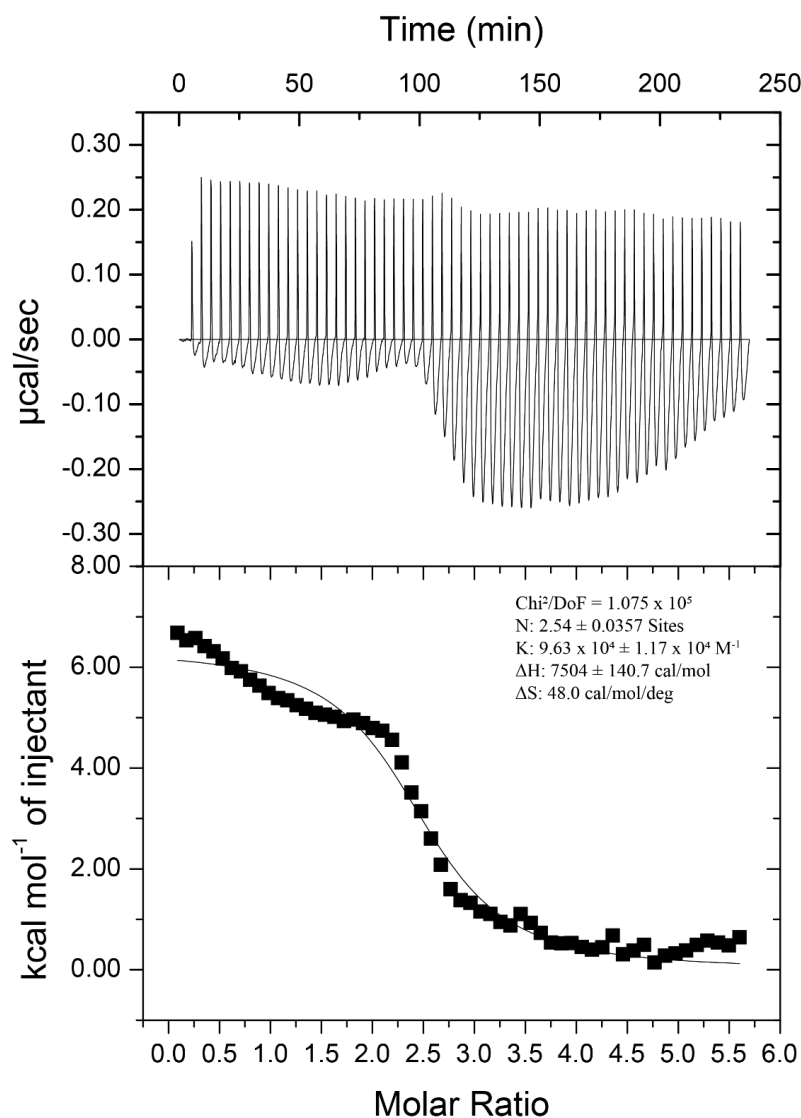
**Table S2. Pairwise Tanimoto coefficients for lead compounds.**

**MVKETYYDVLGVKPNATQEELKKAYRKLALKYHPDKNPNEGEKFKQISQA**  
**YEVLSDKRLGGEQAIKEGGAGGGFGSPMDIFDMFFGGGGRMQRERRGKNV**

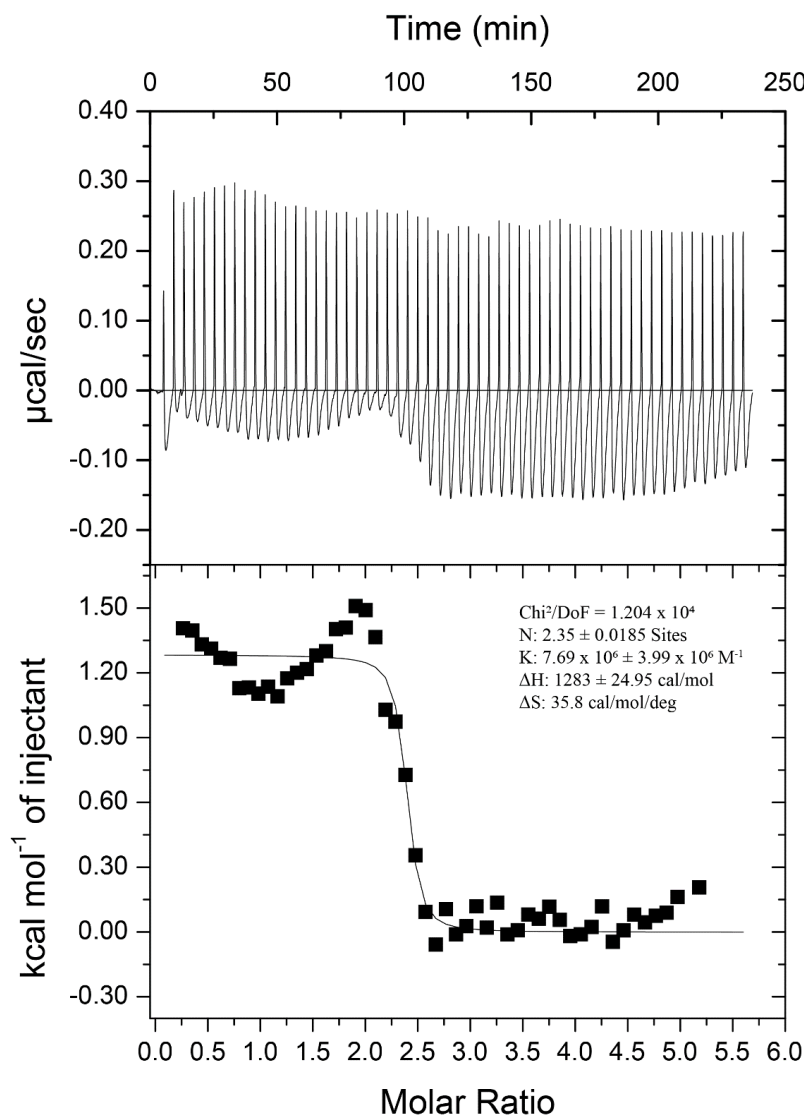
**Figure S1. DNAJA1-107 amino acid sequence.** One letter amino acid code for the first 107 residues of DNAJA1. Residues in red denote missing assignments in the NMR data.



**Figure S2. Structures and nomenclatures of lead compounds.** Compounds confirmed to bind DNAJA1-107 were ordered from Hit2Lead.com<sup>©</sup> and include **A.** 7727958, **B.** 7912207, **C.** 9042610, **D.** 9080132, and **E.** 9101204.



**Figure S3. Binding of 7727968 to DNAJA1-107.** Isothermal titration calorimetry (ITC) plot demonstrating the binding of 7727968 to DNAJA1-107. (*top*) Raw isothermal ITC data corresponding to the heat intensity versus injection number and plotted as equivalent ratios. (*bottom*) Plot of the integrated enthalpies normalized per mol of injectant and normalized against the equivalent ratios. The raw data was fit to a standard binding curve with stoichiometry set as a variable. The best-fit line is superimposed onto the experimental data and the resulting thermodynamic parameters are shown as an insert.



**Figure S4. Binding of 910120 to DNAJA1-107.** Isothermal titration calorimetry (ITC) plot demonstrating the binding of 910120 to DNAJA1-107. (*top*) Raw isothermal ITC data corresponding to the heat intensity versus injection number and plotted as equivalent ratios. (*bottom*) Plot of the integrated enthalpies normalized per mol of injectant and normalized against the equivalent ratios. The raw data was fit to a standard binding curve with stoichiometry set as a variable. The best-fit line is superimposed onto the experimental data and the resulting thermodynamic parameters are shown as an insert.

**Table S1. Phosphorous compounds used to calibrate ligand efficiency (LE)**

6 -Phosphogluconic acid trisodium salt  
 Adenosine 5'-diphosphate sodium salt  
 alpha-D-glucose 1,6-phosphate disodium salt hydrate  
 Dihydroxyacetone phosphate hemimagnesium salt hydrate  
 D-fructose 1,6-bisphosphate trisodium salt hydrate  
 D-fructose 6-phosphate disodium salt hydrate  
 D-Mannose 6-phosphate disodium salt hydrate  
 D-Ribose 5-phosphate disodium salt dihydrate  
 Glycerol 2-phosphate disodium salt hydrate  
 Lithium potassium acetyl phosphate  
 O-phospho-L-serine  
 O-phospho-L-Tyrosine  
 O-Sulfo-L-serine  
 rac -Glycerol 3 -phosphate disodium salt  
 Riboflavin 5'-phosphate  
 Thiamine pyrophosphate

**Table S2. Pairwise Tanimoto coefficients for lead compounds<sup>a,b</sup>**

	<b>7727968</b>	<b>7912207</b>	<b>9042610</b>	<b>9080132</b>	<b>9101204</b>
<b>7727968</b>	1	0.30 (0.16)	0.16(0.16)	0.22 (0.16)	0.14 (0.16)
<b>7912207</b>	0.30 (0.16)	1	0.23(0.29)	0.28 (0.44)	0.14 (0.23)
<b>9042610</b>	0.16(0.16)	0.23(0.29)	1	0.20(0.26)	0.25(0.41)
<b>9080132</b>	0.22 (0.16)	0.28 (0.44)	0.20(0.26)	1	0.15 (0.26)
<b>9101204</b>	0.14 (0.16)	0.14 (0.23)	0.25(0.41)	0.15 (0.26)	1

<sup>a</sup>Values outside parentheses represent atom pair (AP) coefficients. Values in parentheses represent maximum common substructure (MCS) coefficients.

<sup>b</sup>See **Figure S2** for the compound IDs and structures