Cell Chemical Biology, Volume 28

## **Supplemental information**

Evolution of kinase polypharmacology across HSP90 drug discovery

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**Figure S1. Protein-ligand interactions schemes for selected HSP90 inhibitors, related to Figure 5.** a Docking pose with the top MOE score (S = -8.20) for SNX-2112 in ABL1 kinase. **b** Docking pose with the top MOE score (S = -8.02) for Debio-0932 in ABL1 kinase. **c** Docking pose with the top MOE score (S = -8.73) for geldanamycin in ABL1 kinase. **d** Docking pose with the top MOE score (S = -6.94) for radicicol in ABL1 kinase. The MOE ligand interaction tool was used to generate the schematic diagrams of protein-ligand interactions.

