RMSD(solved, model) vs. similarity to top template

Legend:
- TCR-pMHC RMSD, PCC: -0.702, p-val: 0.000
- pMHC RMSD, PCC = 0.047, p-val: 0.786
- peptide RMSD, PCC = -0.187, p-val: 0.276

X-axis: Sequence similarity to top template (%)
Y-axis: RMSD