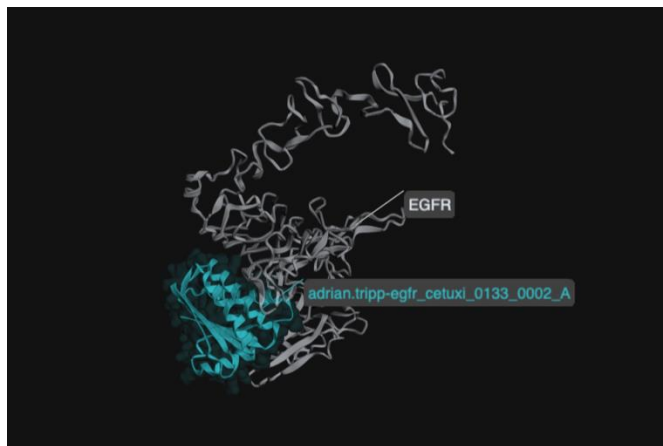
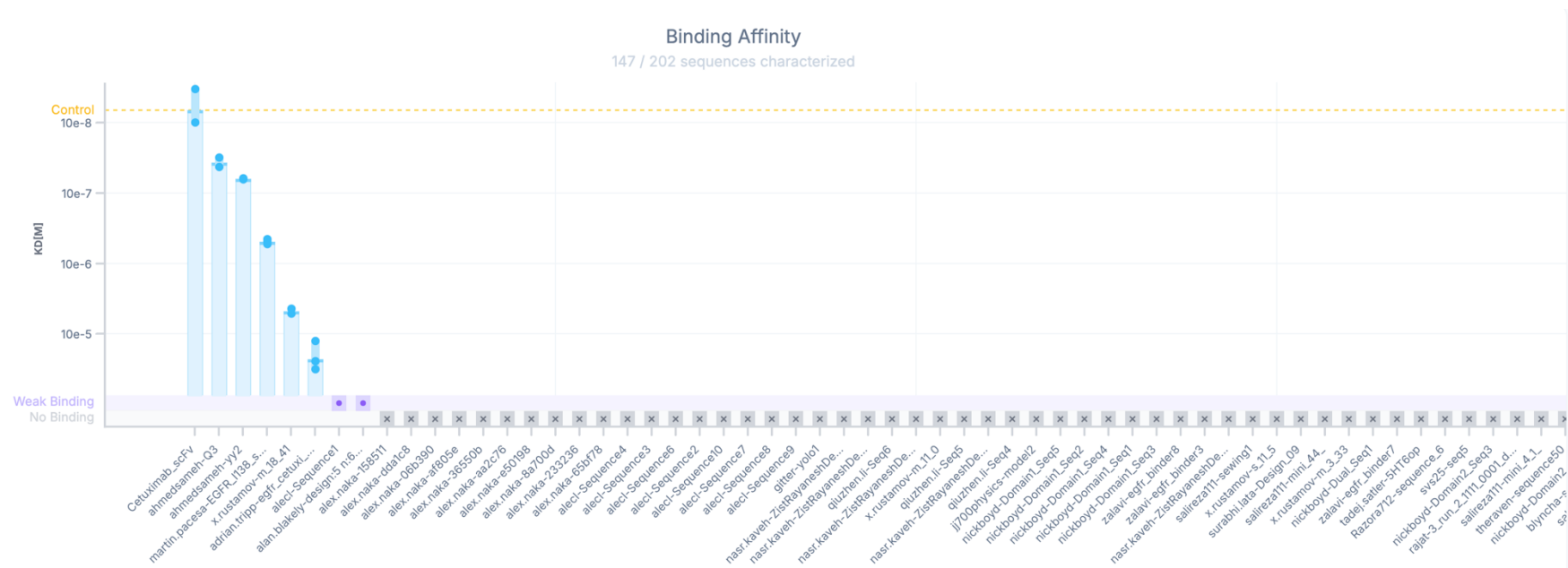


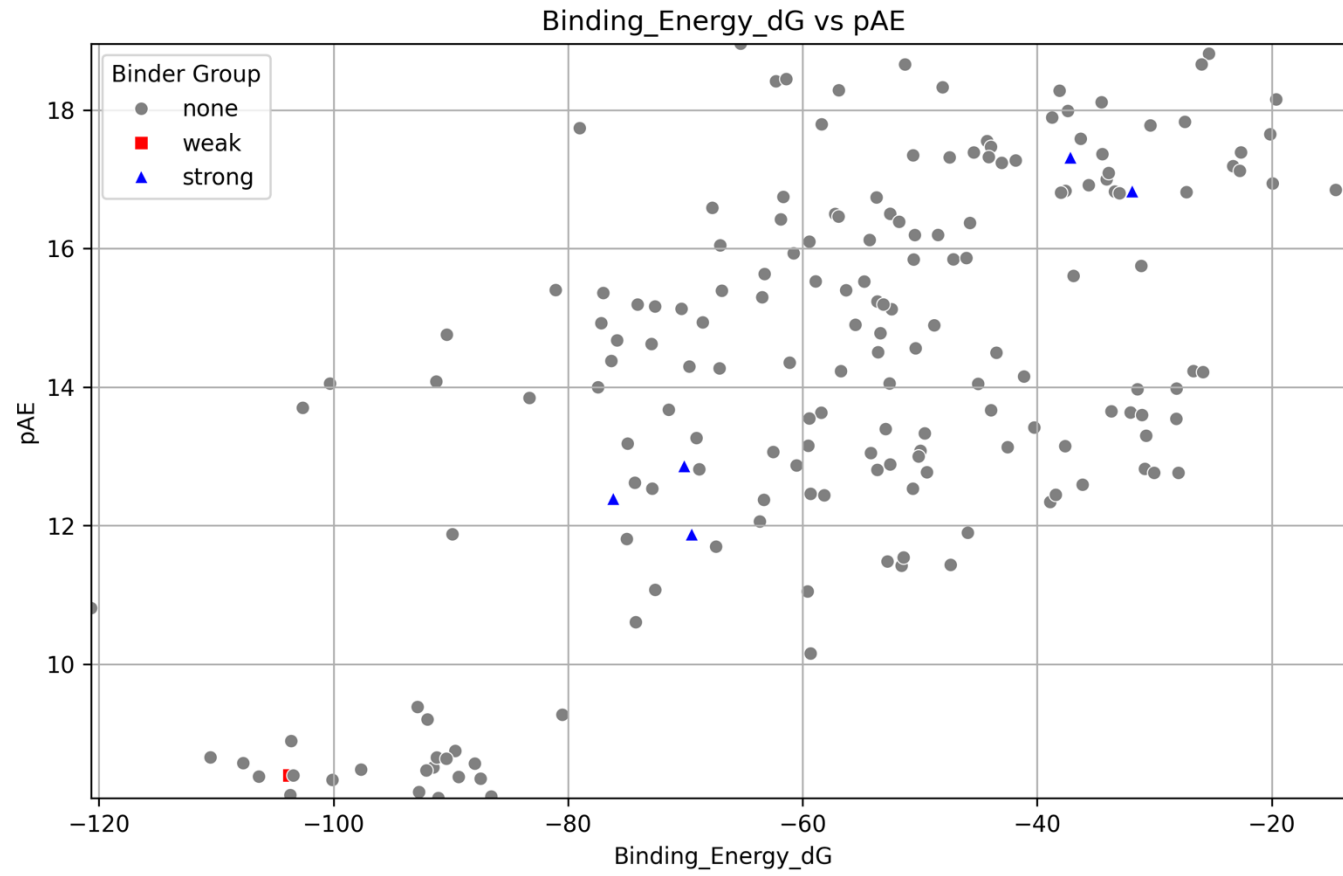
EGFR (Round1)



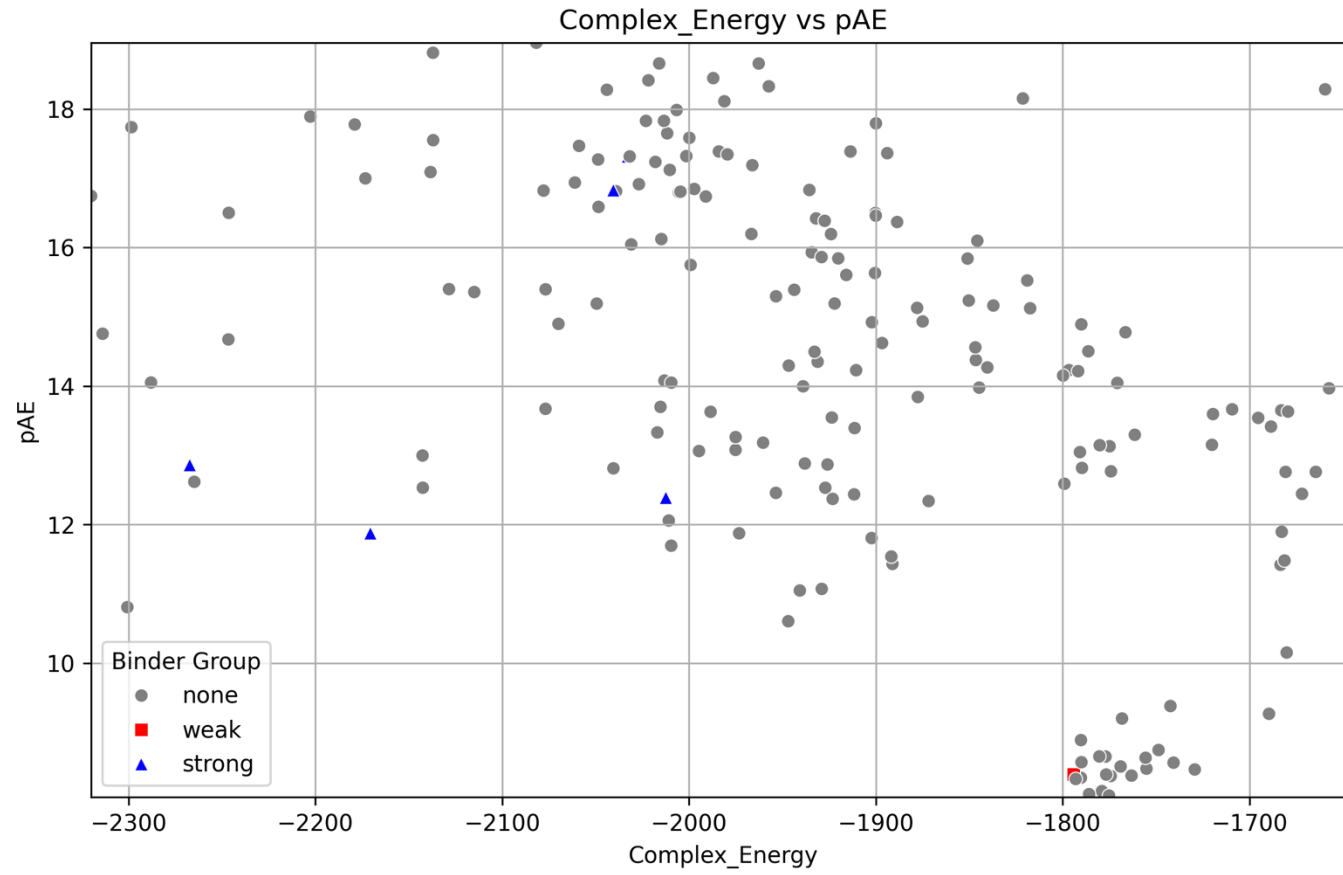
201 Protein:
Expressed: 147
(B/U): 7/147



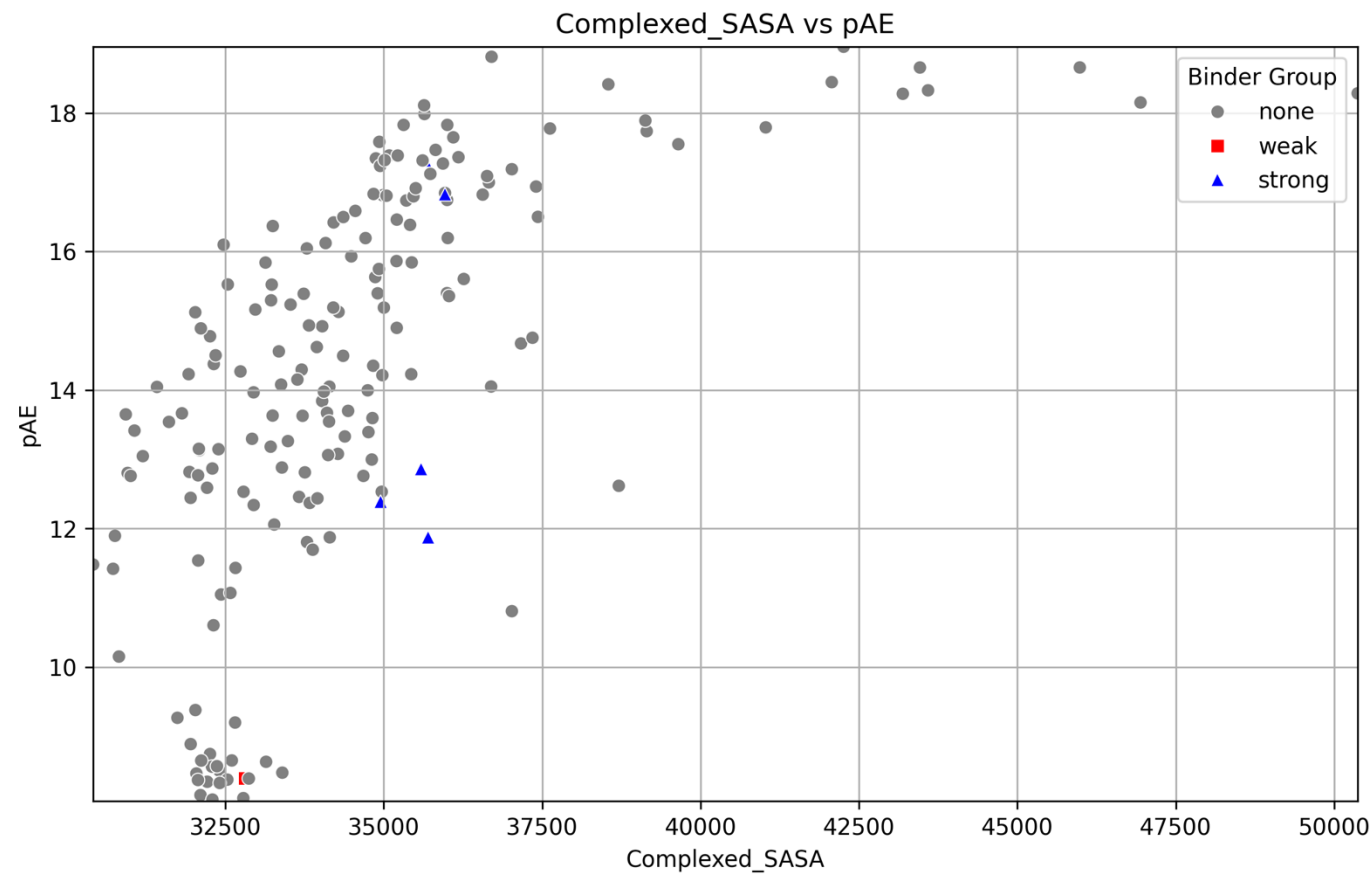
Binding Energy vs pAE



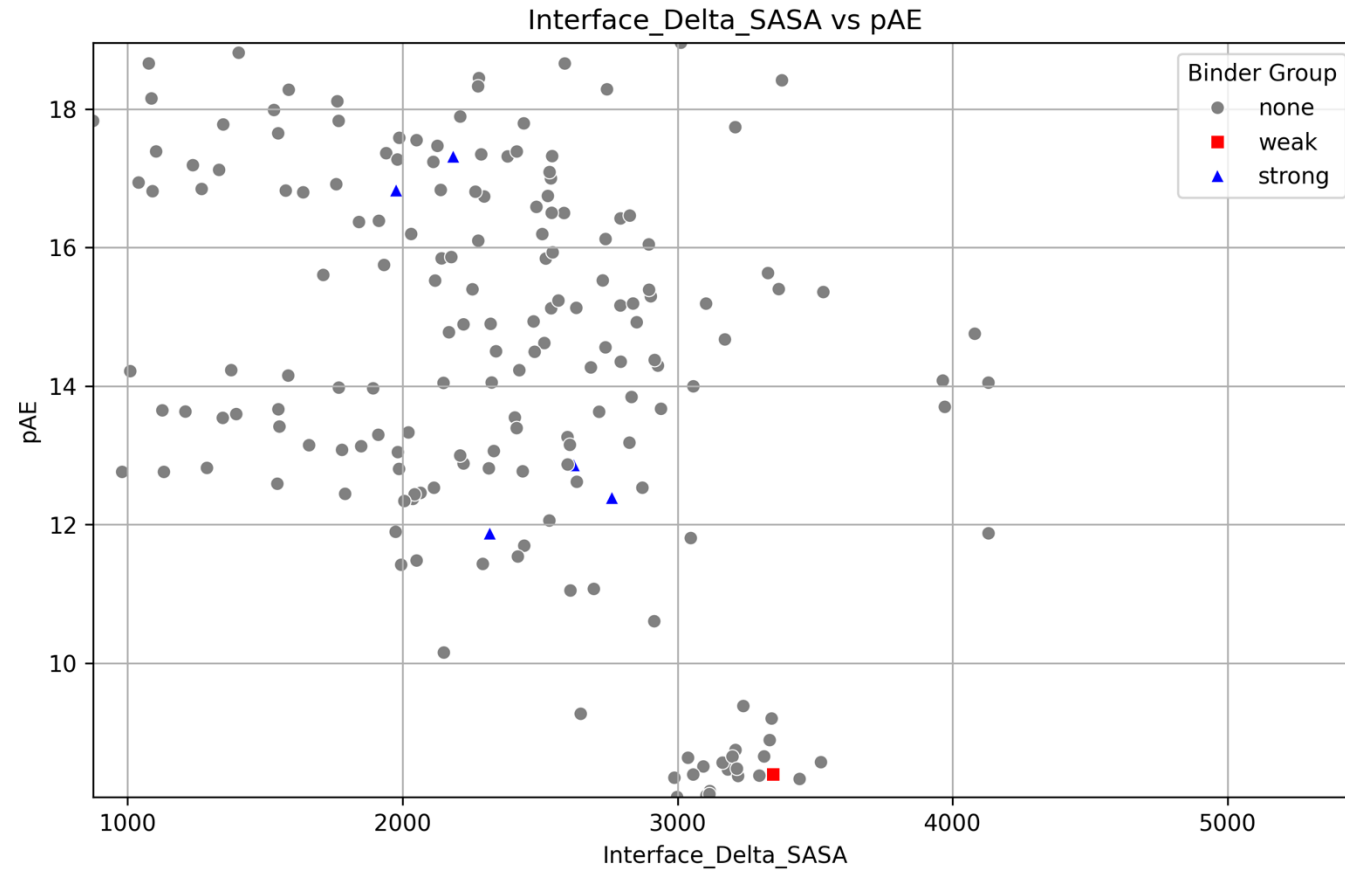
Complexed Energy vs pAE



Complexed SASA vs pAE

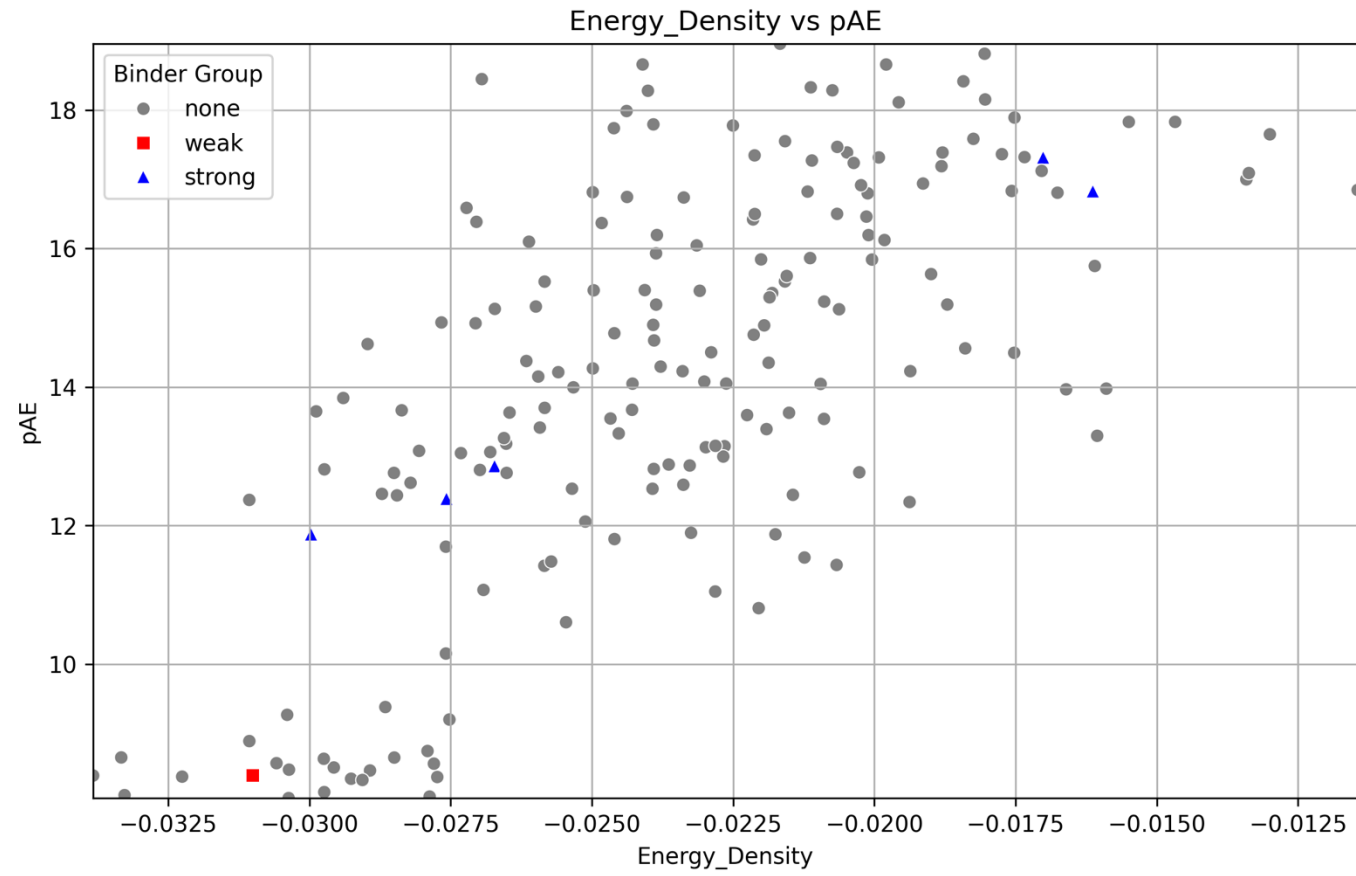


Interface deltaSASA vs pAE

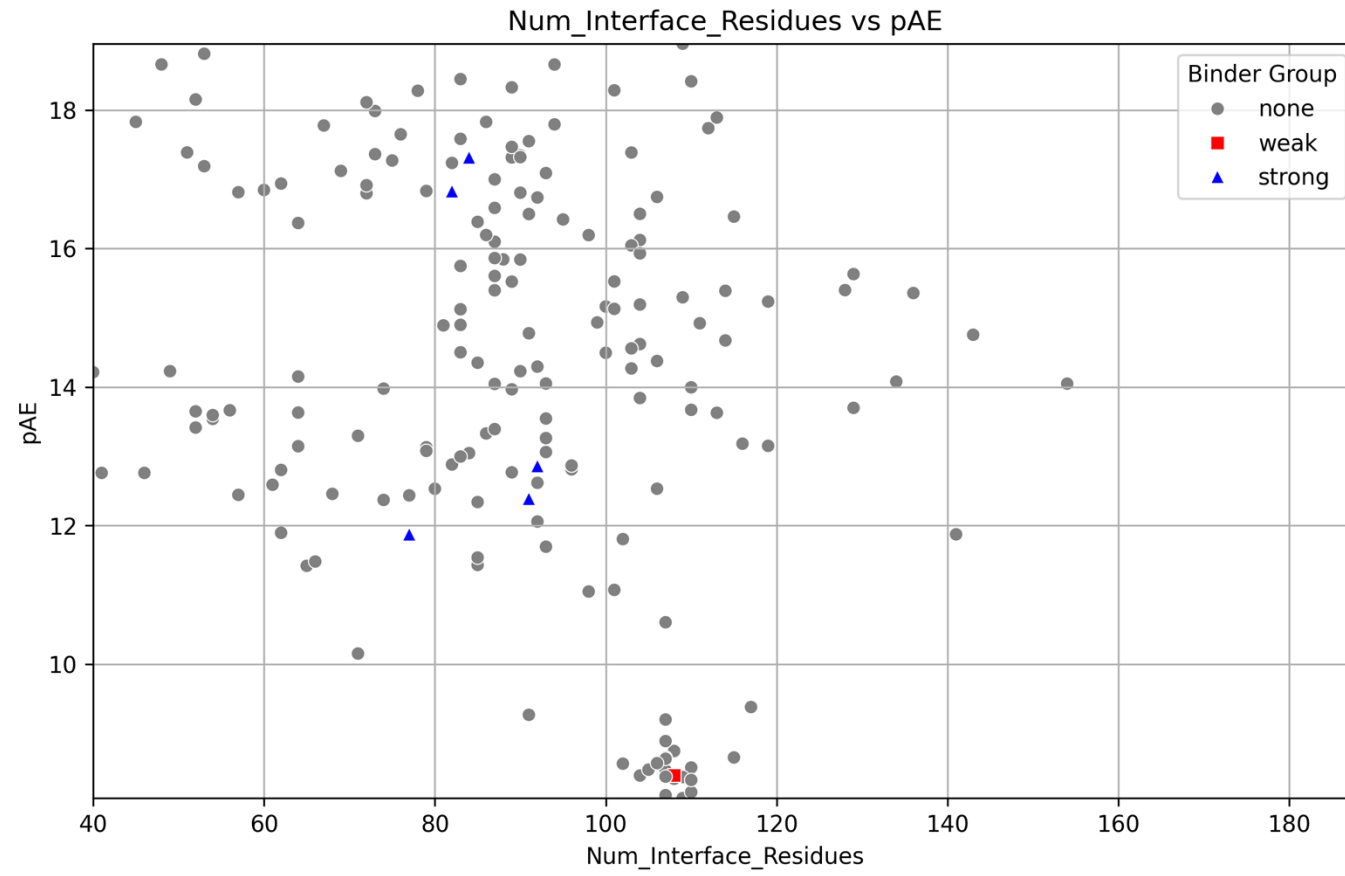


Energy Density (dg/dsasa) vs pAE

The scatter plot, titled "Energy_Density vs pAE", displays the relationship between Energy Density (dg/dsasa) on the x-axis and pAE on the y-axis. The x-axis ranges from -0.0325 to -0.0125, and the y-axis ranges from 10 to 18. A legend in the top-left corner identifies the data points by "Binder Group": grey circles for "none", a red square for "weak", and blue triangles for "strong". The plot shows a dense distribution of grey circles, with a few red squares and blue triangles scattered throughout. The data points are distributed across the entire range of both axes, with a higher density of points in the central region (Energy Density between -0.025 and -0.020, pAE between 12 and 16).



Interface res vs pAE



Total H-bond Energy vs pAE

