|  |
| --- |
| ***Active*** |
| BRD4(1) | 0.83 |
| BRD3(2) | 0.84 |
| BRD3(1) | 0.95 |
| BRD4(2) | 0.70 |
| BRDT(1) | 0.65 |
|  |
| ***Inactive*** |
| TAF1L(2) | 0.57 |
| TIF1 | 0.64 |
| ATAD2 | 0.56 |
| SMARCA4 | 0.52 |
| BAZ2B | 0.55 |
| FALZ | 0.50 |
| PCAF | 0.55 |
| PB1(1) | 0.52 |
| TRIM28 | 0.48 |
| PB1(5) | 0.48 |

Table S8: Exemplar similarity of top (closest) pocket optimized structures to (+)-JQ1-BRD4(1) bound structure. This table shows the raw data from which the heatmap in Figure 8 was created.