**Table S2. Docking score of protein and SSH molecules.**

|  |  |  |  |
| --- | --- | --- | --- |
| **EGFR** | **MMP9** | **SRC** | **MTOR** |
| −46.26849 | −44.176273 | −42.484798 | −50.211506 |
| −42.776367 | −42.307373 | −38.409981 | −48.231388 |
| −42.662792 | −41.085983 | −37.829121 | −47.403488 |
| −41.209278 | −40.582821 | −37.187149 | −45.551258 |
| −41.011459 | −40.273621 | −36.803013 | −44.048946 |
| −40.690716 | −39.997375 | −30.912102 |  |
| −40.562599 | −39.424683 | −30.909056 |  |
| −39.341671 |  |  |  |
| −38.997875 |  |  |  |

The Vina score indicates the binding activity to the protein. The more negative the score, the more stable the interaction of the compound with the protein. EGFR, estimated glomerular filtration rate; MMP9, matrix metalloproteinase 9; mTOR, mammalian target of rapamycin; SRC, proto-oncogene tyrosine-protein kinase Src; SSH, Schisanhenol.