

Table S1. Contribution of the top individual residues to the binding of indomethacin to HSA in the absence and in presence of quercetin obtained with the MM–GBSA approach. All values are in kcal mol⁻¹.

| HSA–Indomethacin Complex | | | HSA–Indomethacin–Quercetin Complex | | |
|---|-----------|------------------------------------|---|-----------|------------------------------------|
| Residue Name | Residue # | Contribution to ΔG _{bind} | Residue Name | Residue # | Contribution to ΔG _{bind} |
| Arg | 218 | -7.46 | Lys | 199 | -4.07 |
| Trp | 214 | -4.57 | Trp | 214 | -2.65 |
| Lys | 199 | -2.76 | Leu | 198 | -2.10 |
| Phe | 211 | -1.79 | Phe | 211 | -1.62 |
| Lys | 195 | -1.48 | Arg | 218 | -1.59 |
| Ser | 202 | -1.31 | Lys | 195 | -1.59 |
| Ala | 215 | -1.01 | His | 242 | -1.16 |
| Leu | 198 | -0.99 | Ser | 202 | -1.15 |
| Leu | 203 | -0.87 | Ala | 215 | -0.97 |
| Ala | 210 | -0.72 | Leu | 481 | -0.84 |
| Total ΔG_{bind} (Indomethacin) | | -33.5 | Total ΔG_{bind} (Indomethacin) | | -38.7 |