

Supporting Table S1. Autodock binding energy evaluation of 5AR homology model in complex with different substrates
(kcal/mol)

| SUBSTRATES | ENERGY |
|---|--------|
| <i>NADP⁺</i> | -9 |
| <i>NADP⁺ + testosterone</i> | -8 |
| <i>NADP⁺ + finasteride</i> | -9 |
| <i>NADP⁺ + beta-sitosterol</i> | -11 |
| <i>NADP⁺ + stigmasterol</i> | -9 |
| <i>NADP⁺ + campesterol</i> | -10 |
| <i>NADP⁺ + daucosterol*</i> | -10 |
| <i>NADP⁺ + oleic acid</i> | -8 |
| <i>NADP⁺ + lauric acid</i> | -8 |
| <i>NADP⁺ + linoleic acid</i> | -9 |
| <i>NADP⁺ + palmitic acid</i> | -8 |

*Bound in different site