

Supporting Information

Semi-rational directed evolution of a Deepsea-derived P450_{S18} for Phenazines Construction

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1. Supplementary Figures

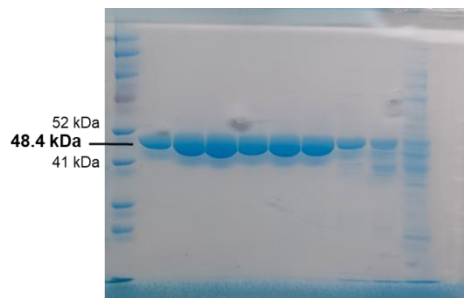


Figure S1. SDS-PAGE analysis of purified P450_{S18} (48.4 kDa). Separation was performed using a 12% acrylamide gel and was stained with Coomassie Brilliant Blue.

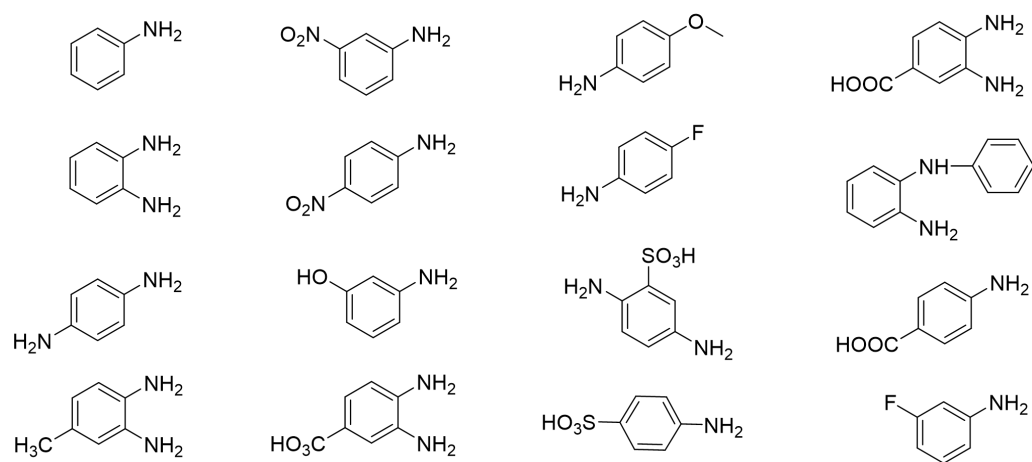


Figure S2. Substrates tested for P450_{S18}.

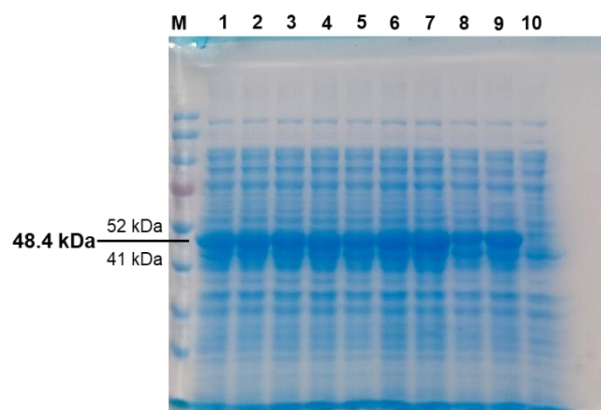


Figure S3. SDS-PAGE analysis of the crude enzyme of P450_{S18} and its mutants. M: protein marker; lane 1-8: F82A, Q88A, F176A, P246A, R245A, F295A, T296A and F292A. lane 9: wild-type-P450_{S18}. lane 10: empty vector without the P450_{S18} gene. Separation was performed using a 12% acrylamide gel and was stained with Coomassie Brilliant Blue

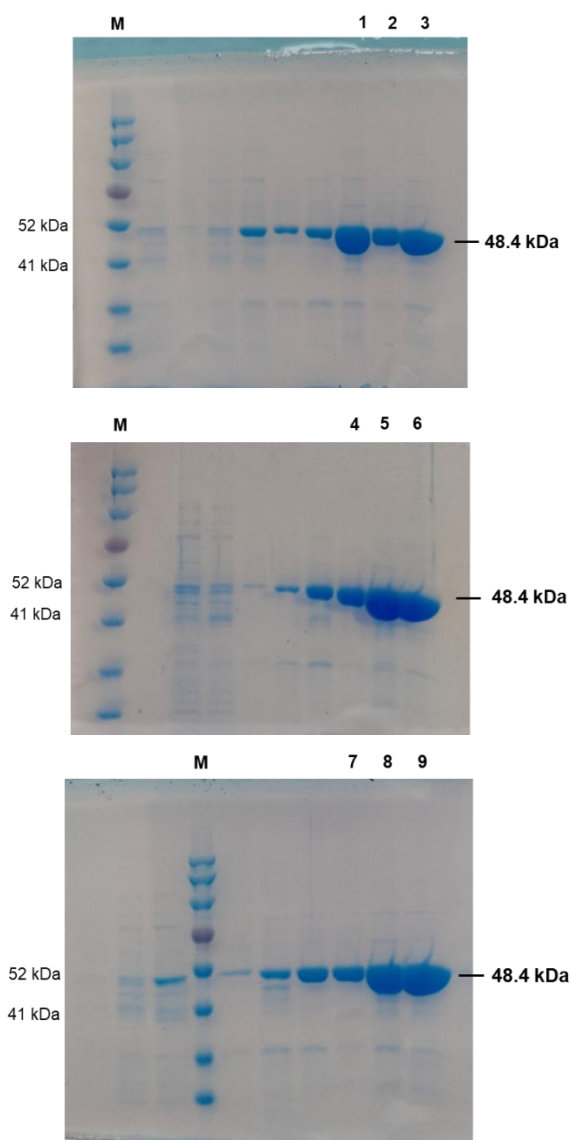


Figure S4. SDS-PAGE analysis of P450_{S18} and its mutants. M: protein marker; lane 1-8: F82A, Q88A, F176A, P246A, R245A, F295A, T296A and F292A. lane 9: wild-type-P450_{S18}. Separation was performed using a 12% acrylamide gel and was stained with Coomassie Brilliant Blue.

2. Supplementary Tables

Table S1. Bacteria and plasmids used in this study.

| Strains or plasmids | Description | Reference or source |
|-----------------------------------|---|---------------------|
| <i>E. coli</i> | | |
| <i>E. coli</i> DH5 α | <i>dam</i> ⁺ , <i>dcm</i> ⁺ , F ⁻ Φ 80/ <i>lacZ</i> Δ M15 Δ (<i>lacZ</i> YA- <i>argF</i>) U169, <i>recA1</i> , <i>endA1</i> , <i>hsdR17</i> (<i>r</i> _K ⁻ , <i>m</i> _K ⁺), <i>phoA</i> , <i>supE44</i> λ - <i>thi-1</i> <i>gyrA96</i> , <i>relA1</i> , for general cloning and preparing methylated DNA | Stratagene |
| <i>E. coli</i> BL21 (DE3) | F ⁻ , <i>ompT</i> , <i>hsdSB</i> (<i>r</i> _B ⁻ , <i>m</i> _B ⁻), <i>gal</i> , <i>dcm</i> (DE3) | Novagen |
| Plasmids | | |
| pET28a | Kan ^R , expression vector | Novagen |
| pET28a::P450 _{S18} | pET28a harboring P450 _{S18} | This study |
| pET28a::P450 _{S18} F176A | pET28a harboring P450 _{S18} F176A | This study |
| pET28a::P450 _{S18} P246A | pET28a harboring P450 _{S18} P246A | This study |
| pET28a::P450 _{S18} R245A | pET28a harboring P450 _{S18} R245A | This study |
| pET28a::P450 _{S18} F295A | pET28a harboring P450 _{S18} F295A | This study |
| pET28a::P450 _{S18} T296A | pET28a harboring P450 _{S18} T296A | This study |
| pET28a::P450 _{S18} F292A | pET28a harboring P450 _{S18} F292A | This study |
| pET28a::P450 _{S18} Q88A | pET28a harboring P450 _{S18} Q88A | This study |
| pET28a::P450 _{S18} F82A | pET28a harboring P450 _{S18} F82A | This study |

Table S2. The primer pairs used in this study.

| | | Primer pairs used for inactivation (5'-3') ^a | Restriction sites | Size (bp) |
|---------------------------|-------------------------|---|-------------------|-----------|
| Protein expression in | P450 _{S18} -FP | GGAATTC <u>CATATG</u> aattcaggtaagcaaatac | Nde I | 1281 |
| <i>E. coli</i> BL21 (DE3) | P450 _{S18} -RP | CCGCTCGAGttacttaacctttatattc | Xho I | |
| Site-directed mutagenesis | F176A-FP | gacatgattgatgcaGCAggcgcaacaggcccac | | |
| | F176A-RP | gtgggcctgttgccgTGctgcatcaatcatgtc | | |
| | P246A-FP | ctgaaccttttacggGCAattgtggcaattgcc | | |
| | P246A-RP | ggcaattgccacaatTGCccgtaaaagggtcag | | |
| | R245A-FP | attctgaacctttaGCAccgattgtggcaattg | | |
| | R245A-RP | caattgccacaatcggTGCtaaaagggtcagaat | | |
| | F295A-FP | ctatccgtttgcacccGCAacaggggcgtaacag | | |
| | F295A-RP | ctgttaacgcccctgtTGCgggtgcaaacggatag | | |
| | T296A-FP | ccgtttgcaccctttGCAggggcgtaacagcg | | |
| | T296A-RP | cgctgttaacgcccTGCAaagggtgcaaacgg | | |
| | F292A-FP | cgtaggtactatccgGCAgcaccctttacaggg | | |
| | F292A-RP | ccctgtaaagggtgcTGCcgatagtagctacg | | |
| | Q88A-FP | ggtgtaggtggtgtgGCAgggatggatggggaag | | |
| | Q88A-RP | cttcccatccatcccTGCcacaccacctacacc | | |
| | F82A-FP | gaataaaagaaagcttgGCAggtgtaggtggtgtgcaag | | |
| | F82A-RP | cttgacaccacctacaccTGCcaagcttctttattc | | |

^aUnderlined letters represent restriction site