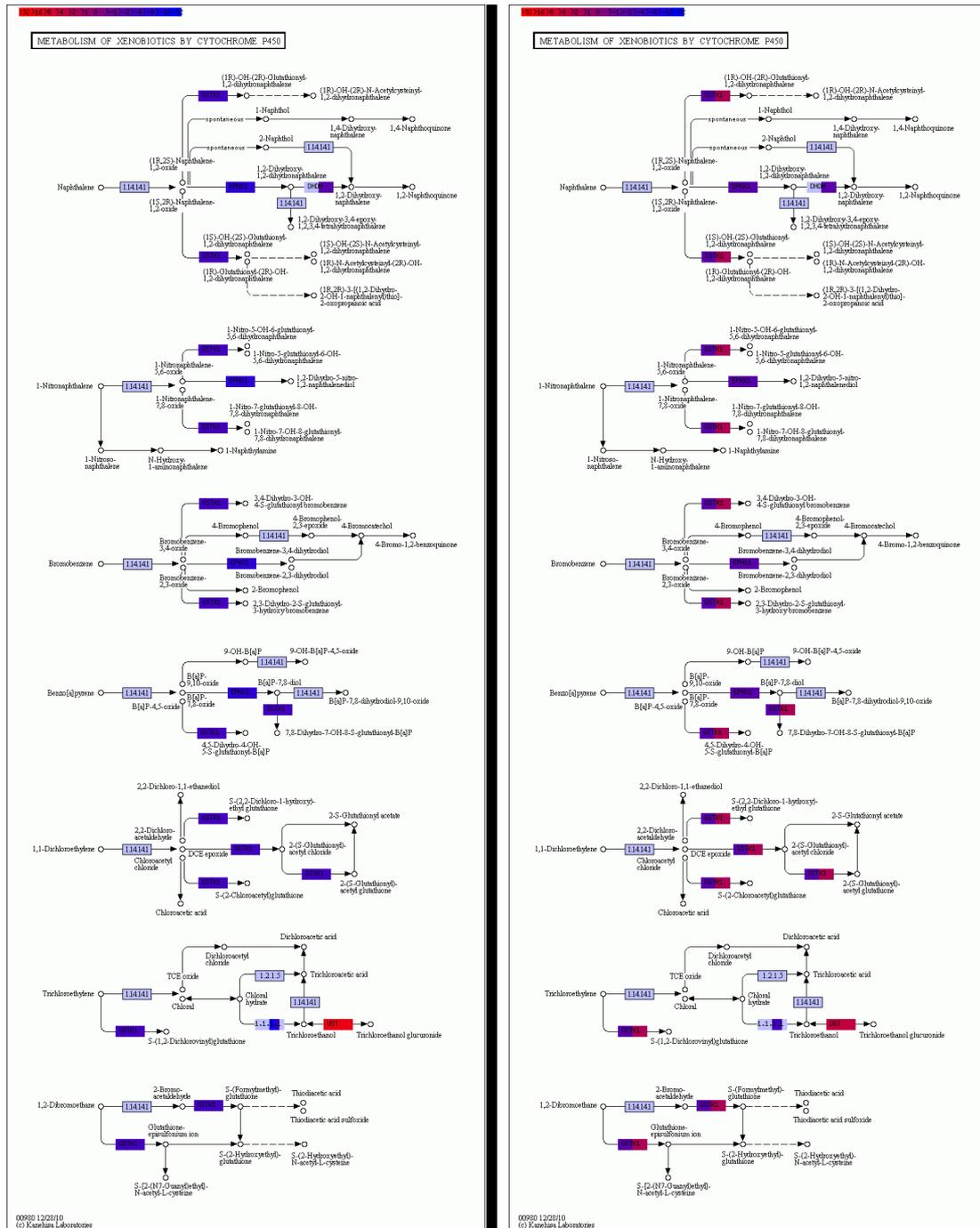


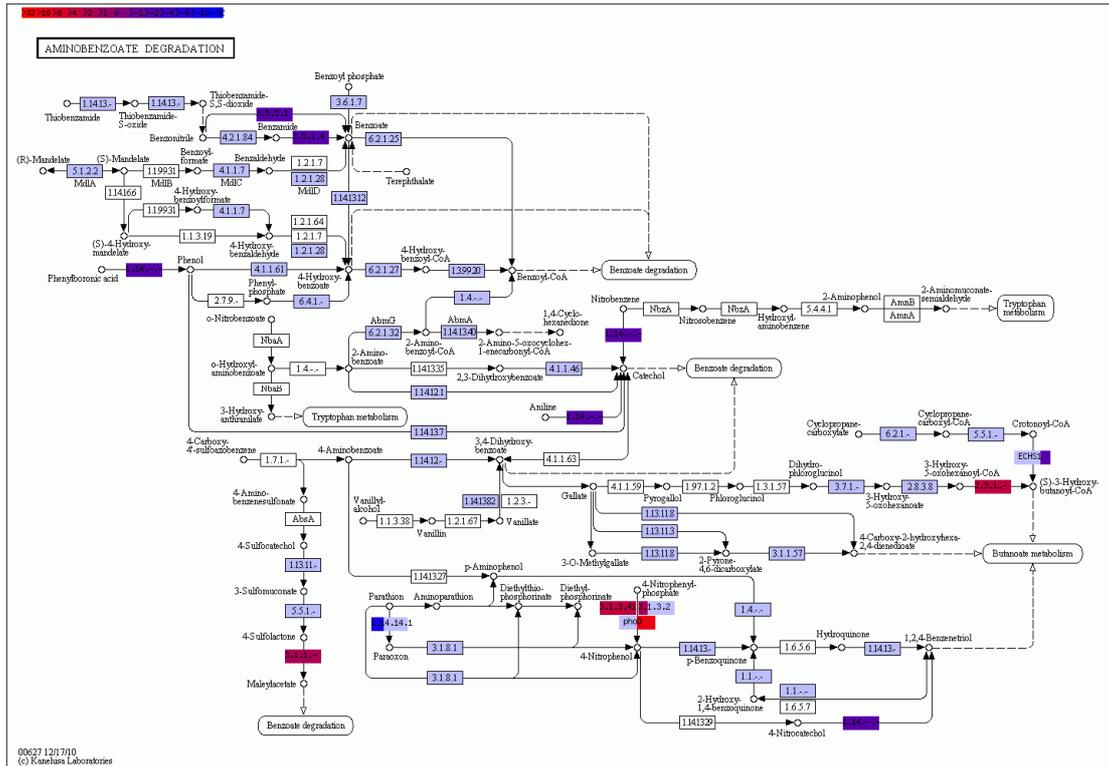
Supplementary Materials



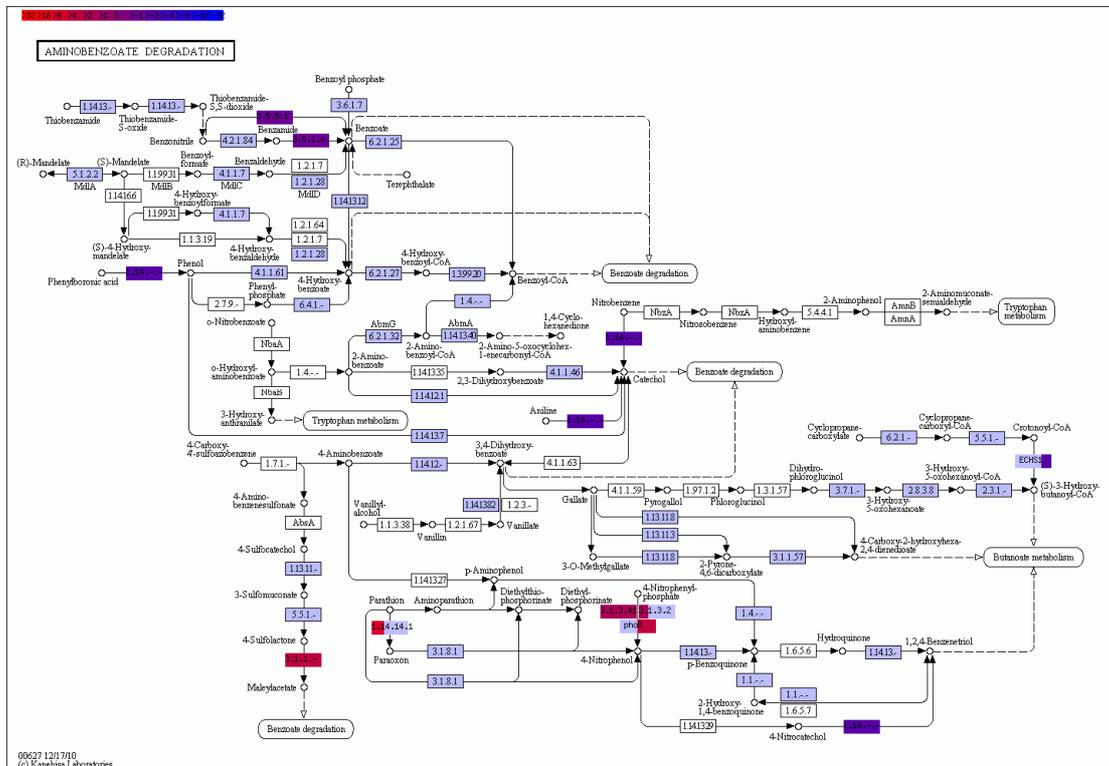
AD

AG

Figure S1. KEGG pathways for metabolism of xenobiotics by cytochrome P450.

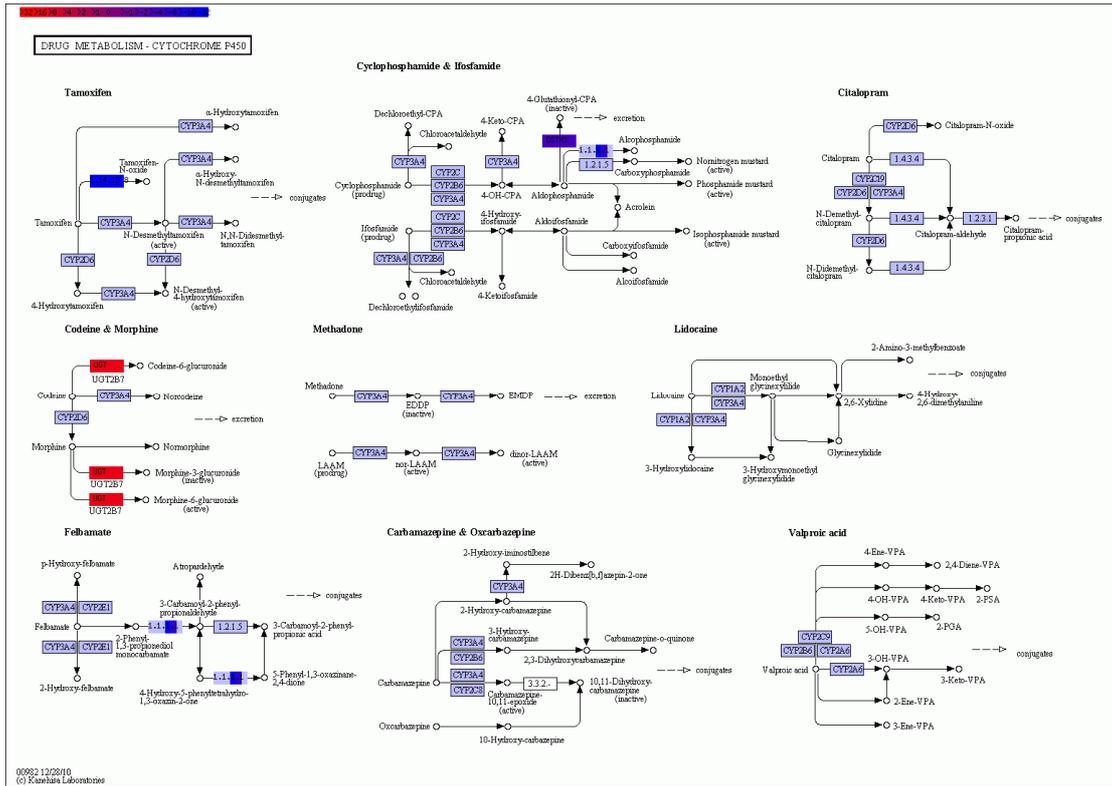


AD

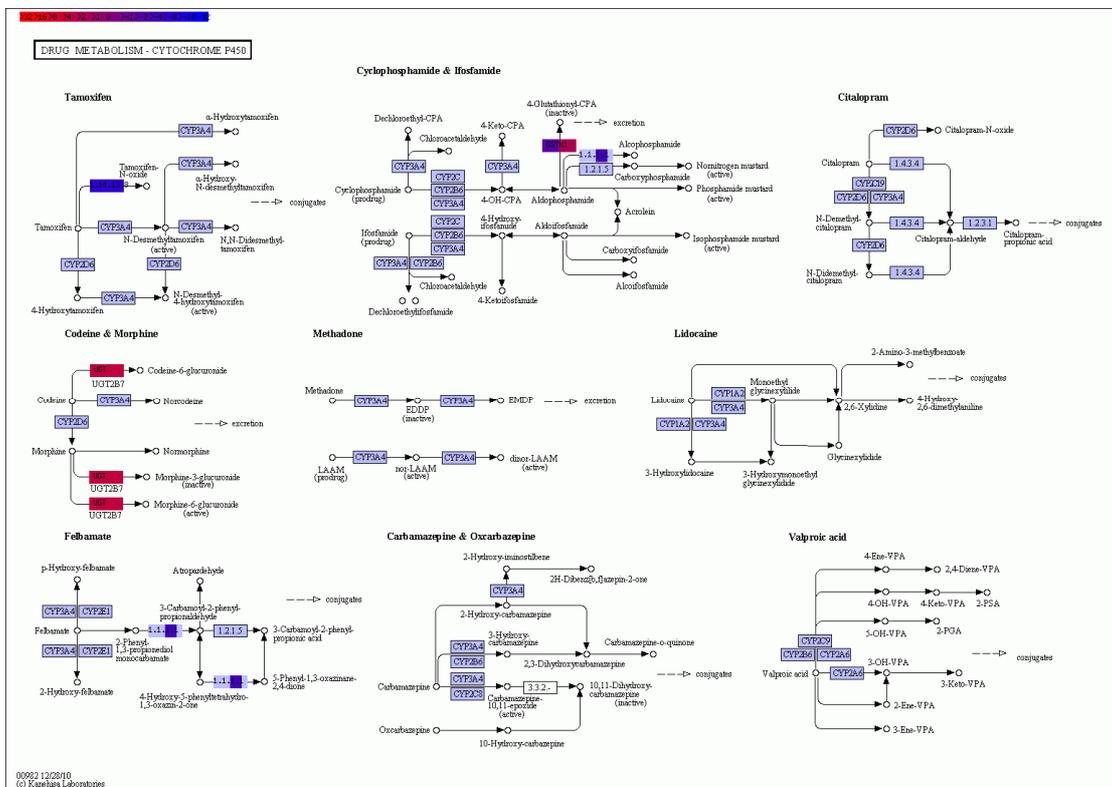


AG

Figure S2. KEGG pathways for aminobenzoate degradation.



AD



AG

Figure S3. KEGG pathways for drug metabolism-cytochrome P450.

Table S1 Primer sequence used for qRT-PCR in this study.

| Gene | Forward primer | Reverse primer |
|------------------|--------------------------|------------------------------|
| CYP-33C4 | GATACAGATACAGCAAGGAGAAT | AACAACCAGACGGAGGAA |
| CYP-33C2 | CAGTCTGGTTAATGAAGTT | ATAGCCGTTGAGATTGTA |
| CYP-33C9 | CTGATATTGTGGCTTGCTA | AGTCCTACGAGTTGTTGA |
| UGT-48 | GATGAGATGATGGACAATGGT | TTAGTTGAGATTCGCAAGCATA |
| DHS-2 | CGGTATTGCTGCTATTGG | AAGGAGAGGAATGAGAAGAT |
| SDR-3 | TATGAAGGTAATTGAAAGG | ATTATTGACCGAATGAAG |
| UNC-8 | GGAGATGTGGTCAACTTC | CAGAGGCACTAAGATTTCG |
| CBG09704 | GTGCTGAAGTTGTTATTC | TTGATGGTCTCGTTAATG |
| T08H10.1 | CCGCTATTATTGTTGTTCCA | TCCAGAGACCATCAGAT |
| Collagen protein | GACAGGCTAACAGAACTC | ATTGGATTGCTTCGTGAA |
| HSP-70 | CGAAGGAGAAGTTCAGATG | CTTAGGCGATGAGCAATC |
| PTR-12 | TTATCCAGAGGTCGTTGA | ATTCGCTCGTAGTTCTTG |
| Reference gene | TCCGTACCCTGAAGTTGGCTAACC | AAGTGGAGACGAGGGAATGGAA CC |