

Supplementary Materials: Detoxification of Mycotoxins through Biotransformation

Peng Li, Ruixue Su, Ruya Yin, Daowan Lai, Mingan Wang, Yang Liu and Ligang Zhou

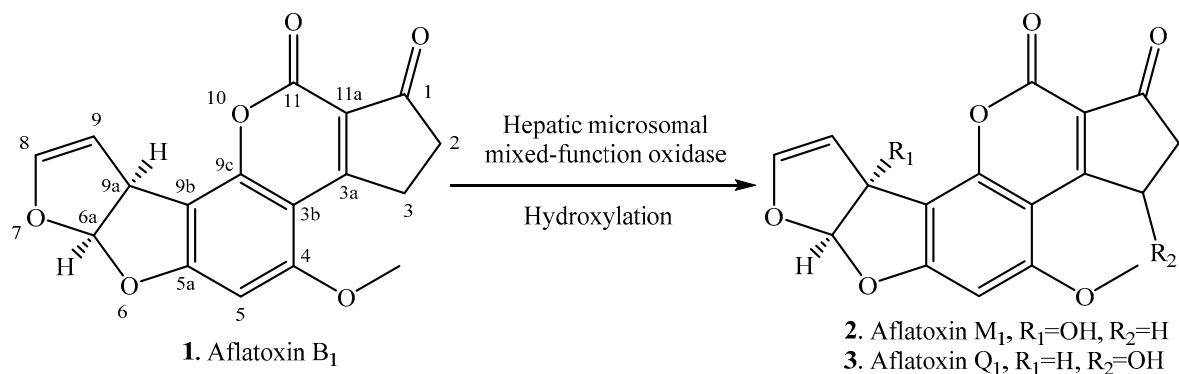


Figure S1. Transformation of aflatoxin B₁ (1) by hepatic microsomal mixed-function oxidase of rhesus monkey [1].

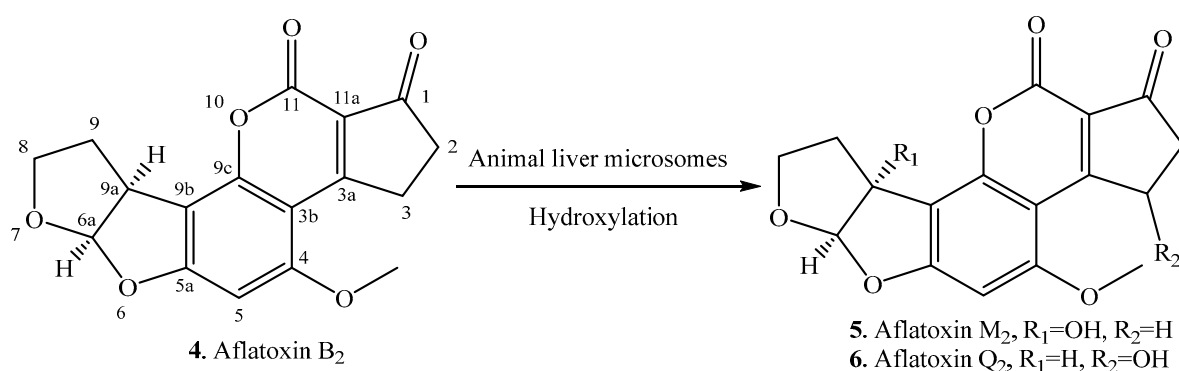


Figure S2. Transformation of aflatoxin B₂ (4) with hydroxylation by animal liver microsomes [2].

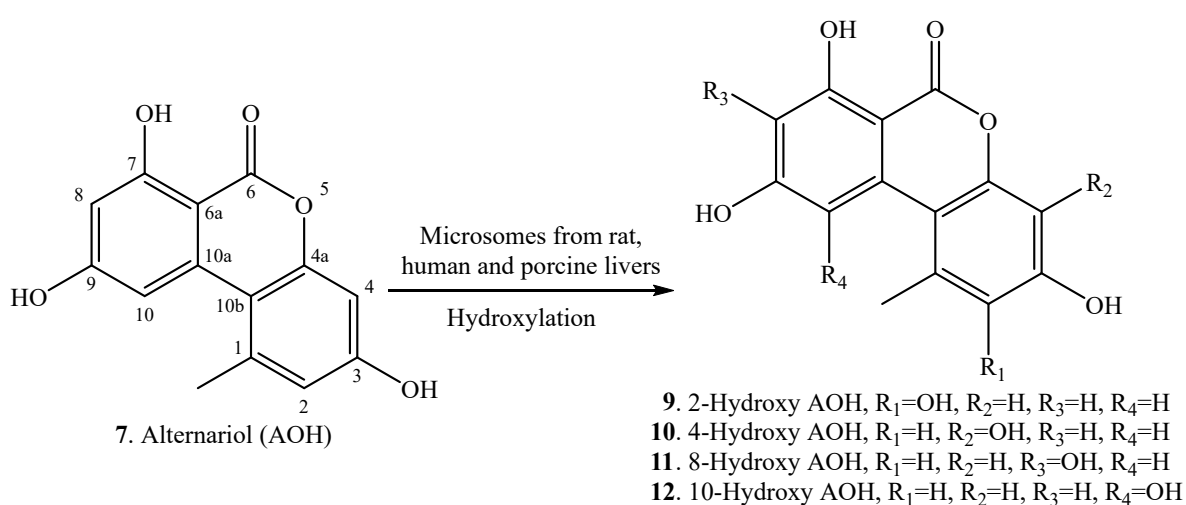


Figure S3. Transformation of alternariol (7) with hydroxylation by the microsomes from rat, human and porcine livers [3].

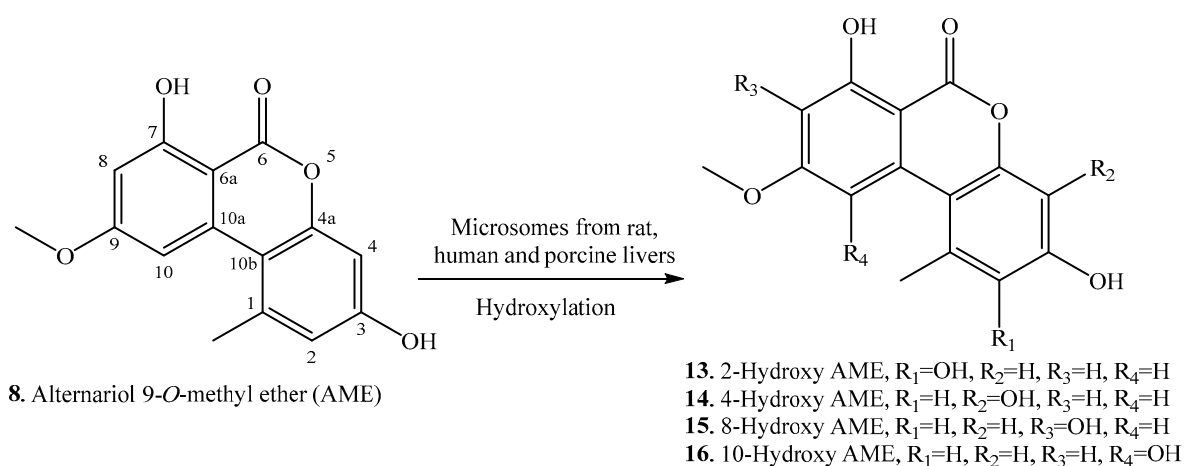


Figure S4. Transformation of alternariol 9-O-methyl ether (8) with hydroxylation by the microsomes from rat, human and porcine livers [3].

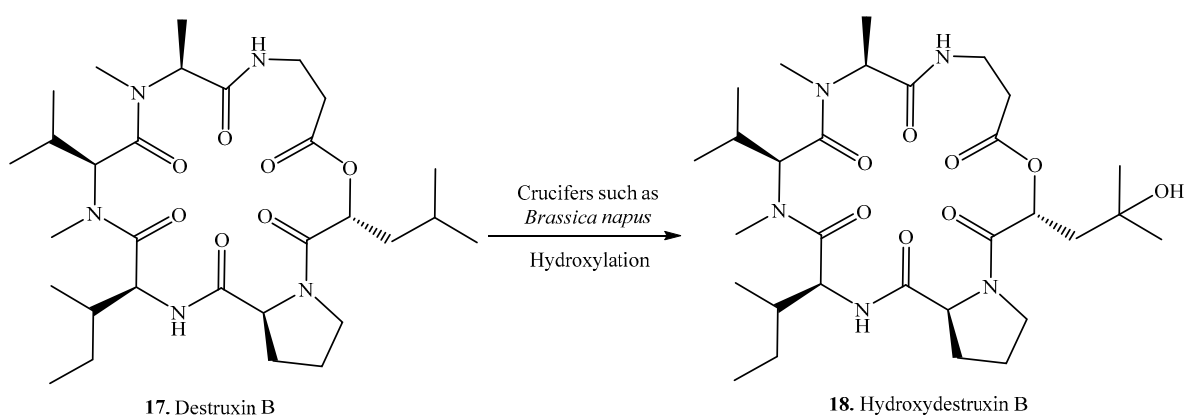


Figure S5. Transformation of destruxin B (17) with hydroxylation by crucifers such as *Brassica napus* [4].

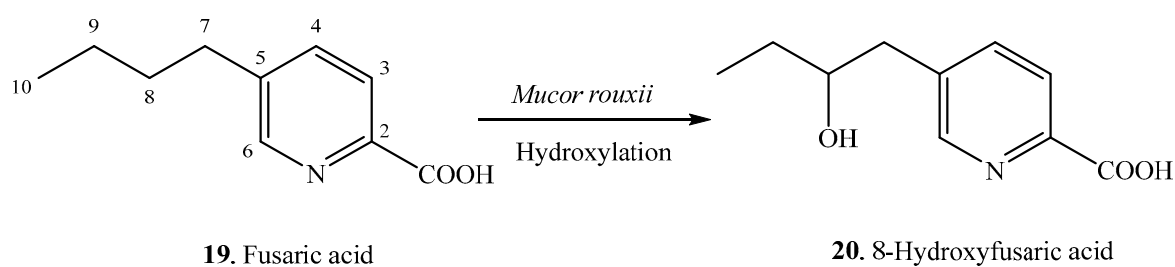


Figure S6. Transformation of fusaric acid (19) with hydroxylation by *Mucor rouxii* [5].

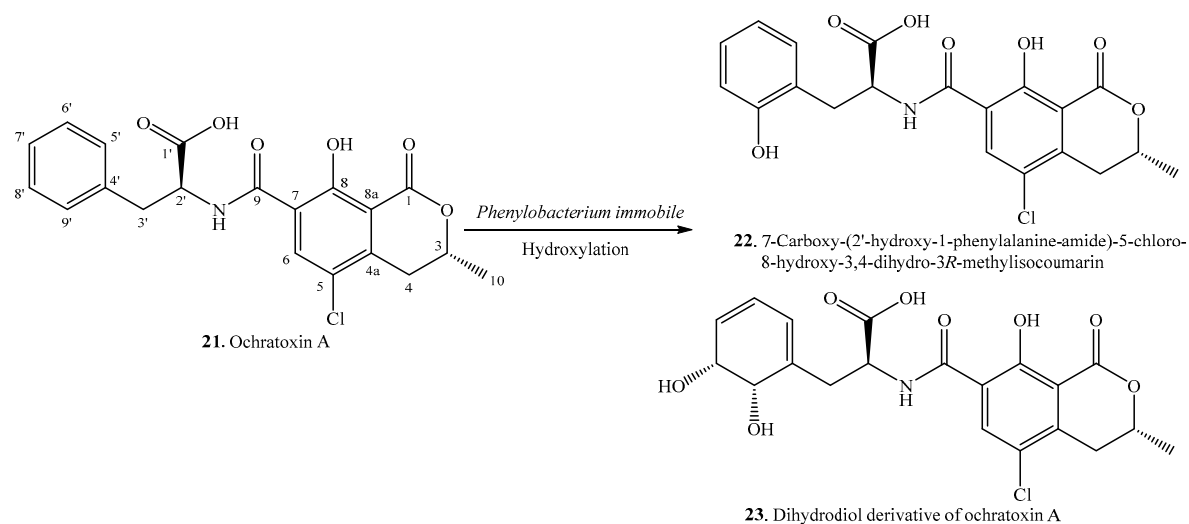


Figure S7. Transformation of ochratoxin A (21) with hydroxylation by *Phenyllobacterium immobile* [6].

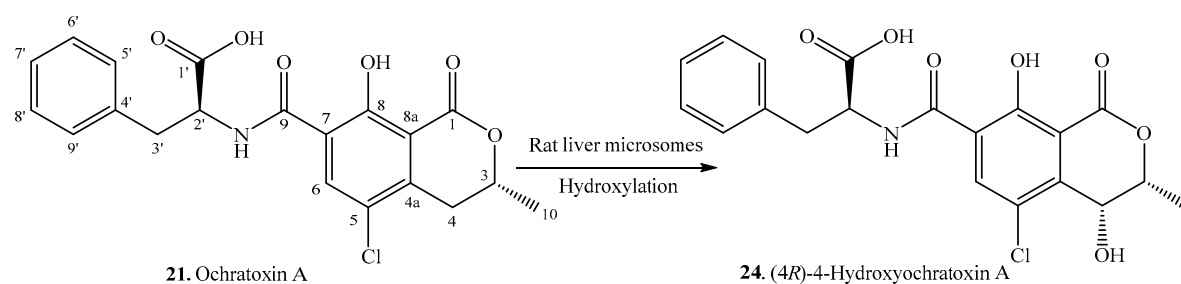


Figure S8. Transformation of ochratoxin A (21) with hydroxylation by rat liver microsomes [7].

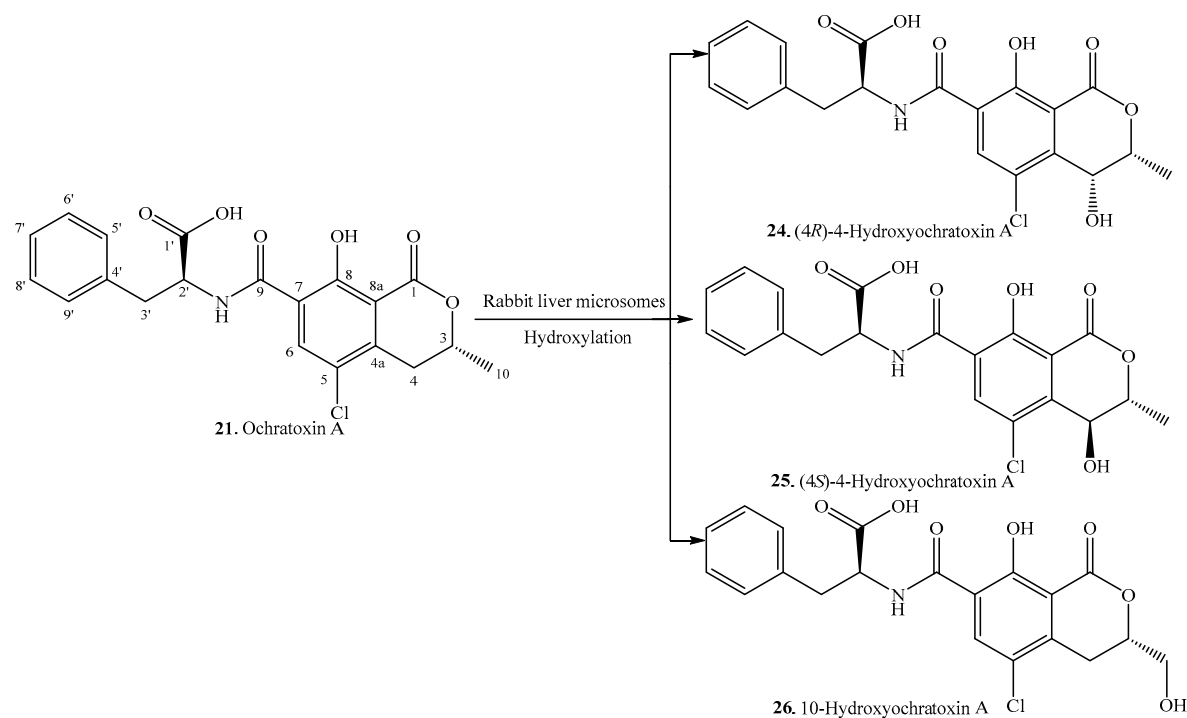


Figure S9. Transformation of ochratoxin A (21) with hydroxylation by rabbit liver microsomes [8].

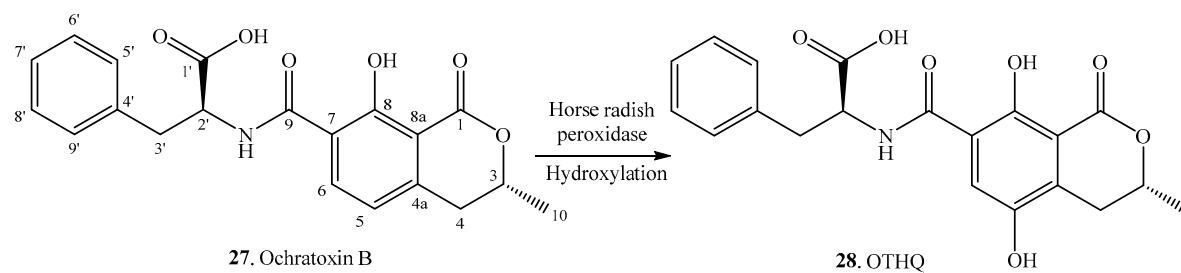


Figure S10. Transformation of ochratoxin B (27) with hydroxylation by horse radish peroxidase [9].

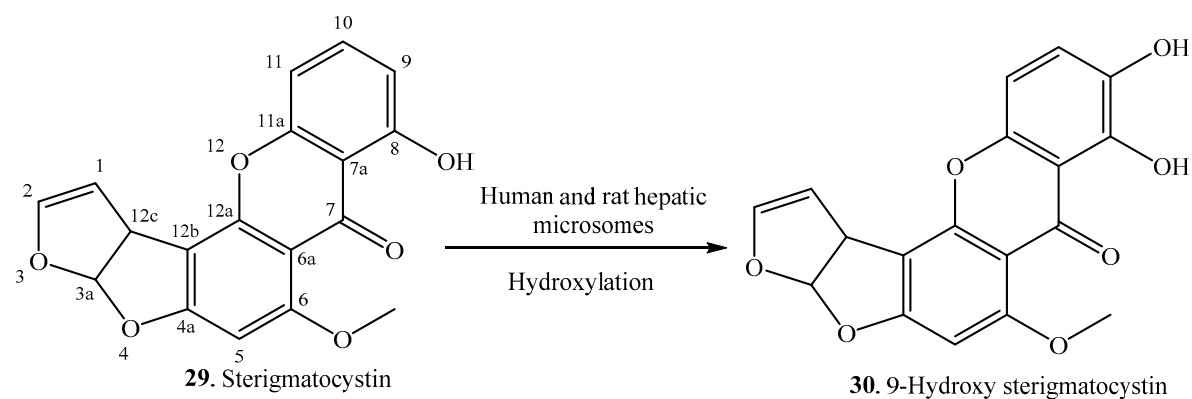


Figure S11. Transformation of sterigmatocystin (29) with hydroxylation by human and rat hepatic microsomes [10].

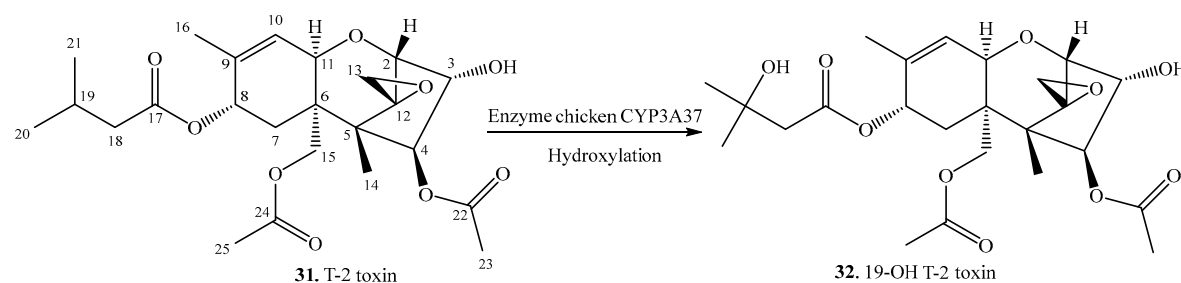


Figure S12. Transformation of T-2 toxin (31) with hydroxylation by chicken CYP3A37 [11].

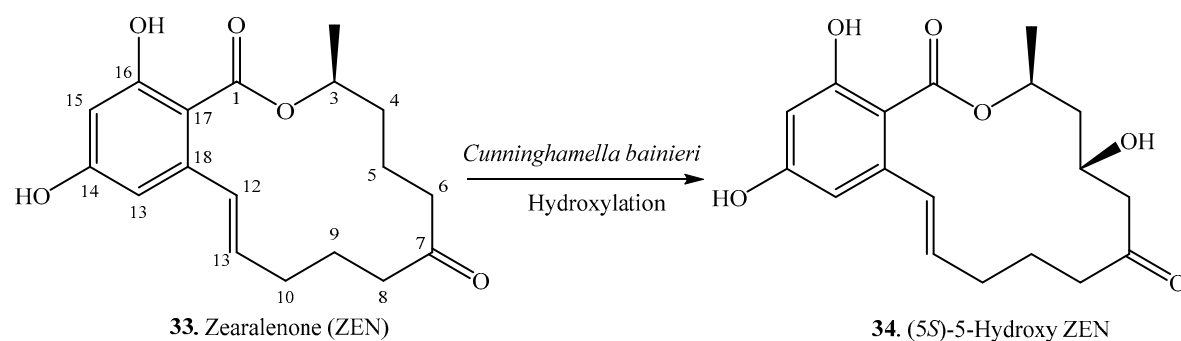


Figure S13. Transformation of zearalenone (33) with hydroxylation by *Cunninghamella bainieri* [12].

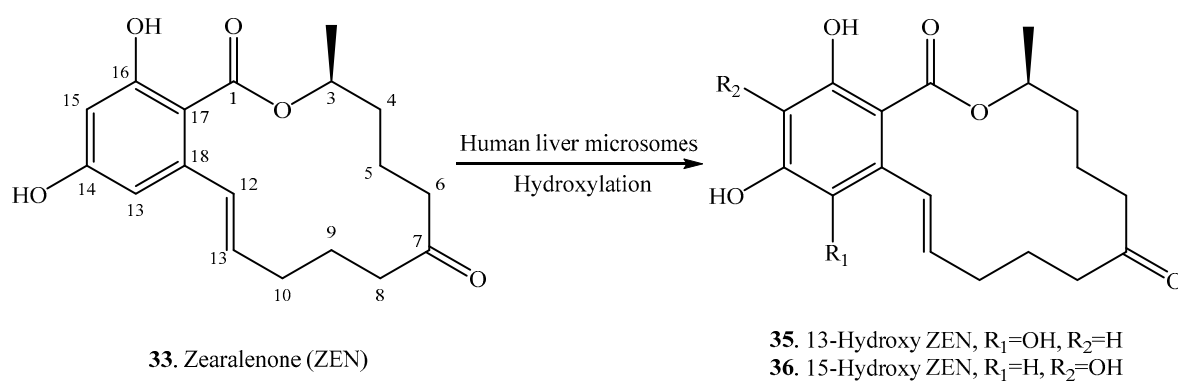


Figure S14. Transformation of zearalenone (33) with hydroxylation by human liver microsomes [13].

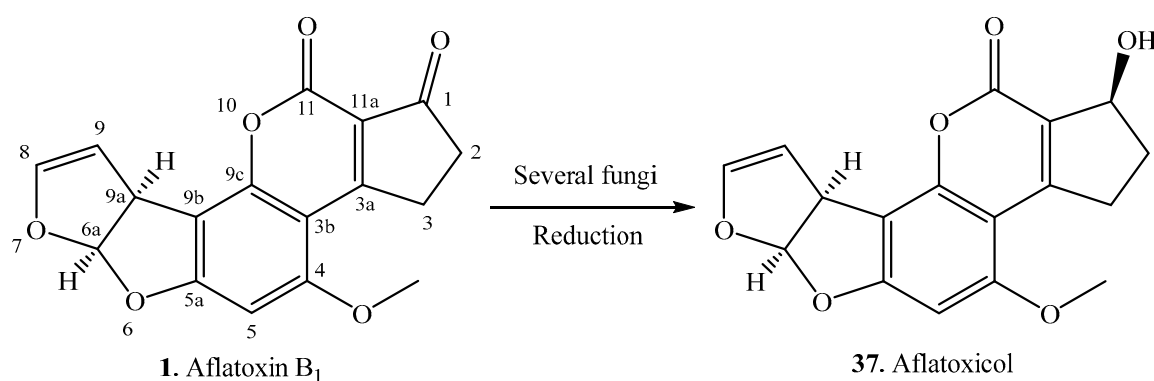


Figure S15. Transformation of aflatoxin B₁ (1) with reduction by several fungi [14].

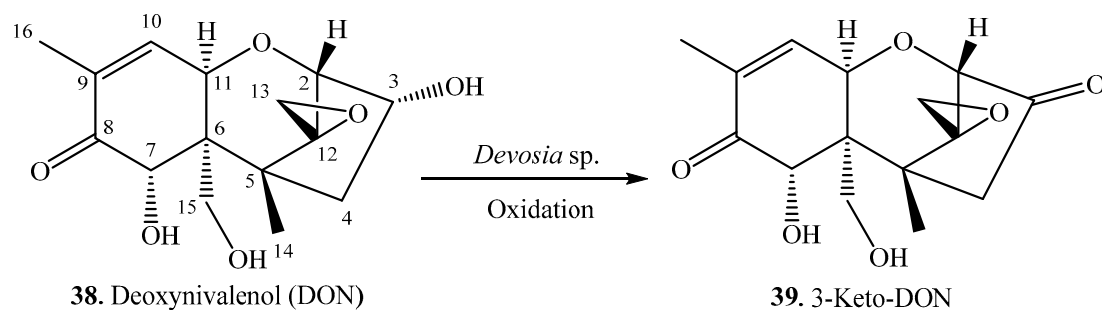


Figure S16. Transformation of deoxynivalenol (38) with oxidation by *Devosia* sp. [15].

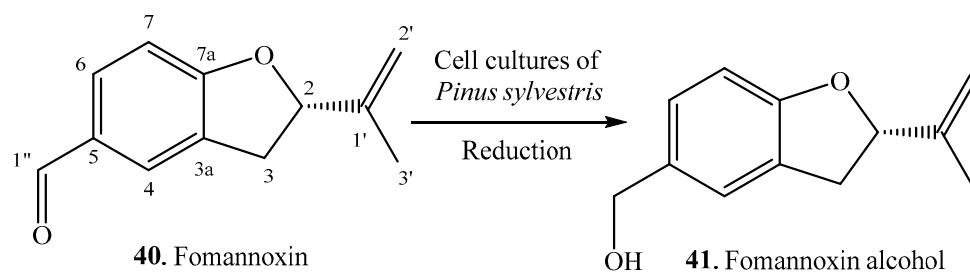


Figure S17. Transformation of fomannoxin (40) with reduction by cell cultures of *Pinus sylvestris* [16].

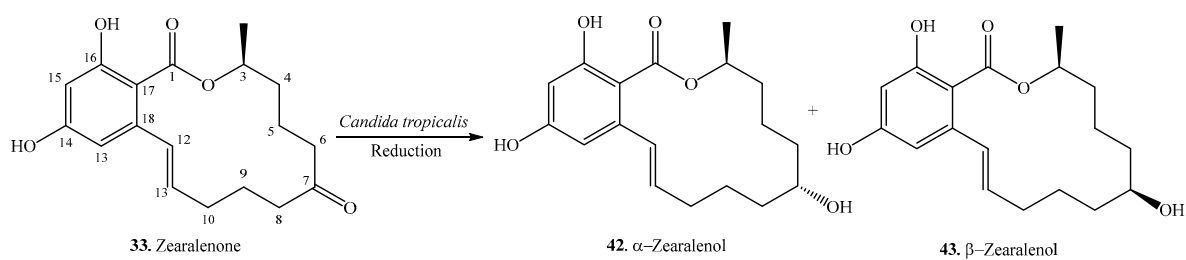


Figure S18. Transformation of zearalenone (33) with reduction by *Candida tropicalis* [17].

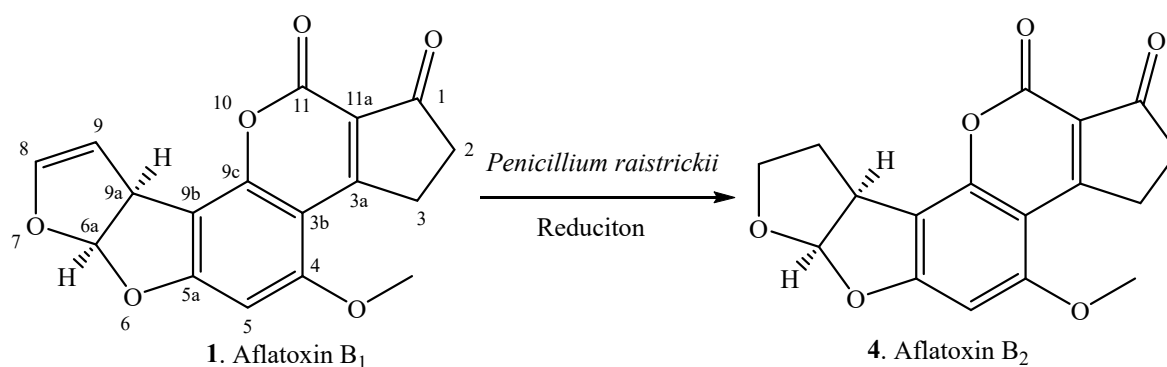


Figure S19. Transformation of aflatoxin B₁ (1) with reduction by the fungus *Penicillium raistrickii* [2].

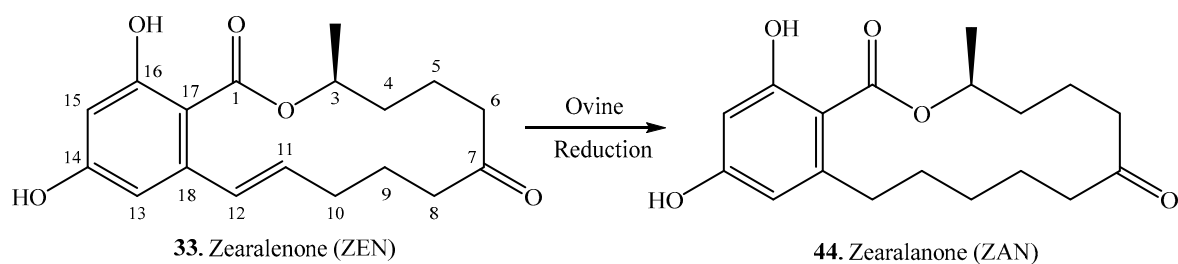


Figure S20. Transformation of zearalenone (33) with reduction in ovine [18].

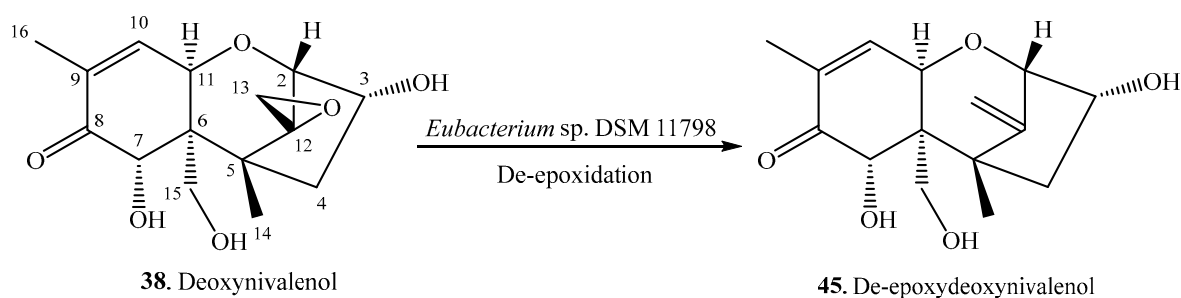


Figure S21. Transformation of deoxynivalenol (38) with de-epoxidation by *Eubacterium* sp. DSM 11798 [19].

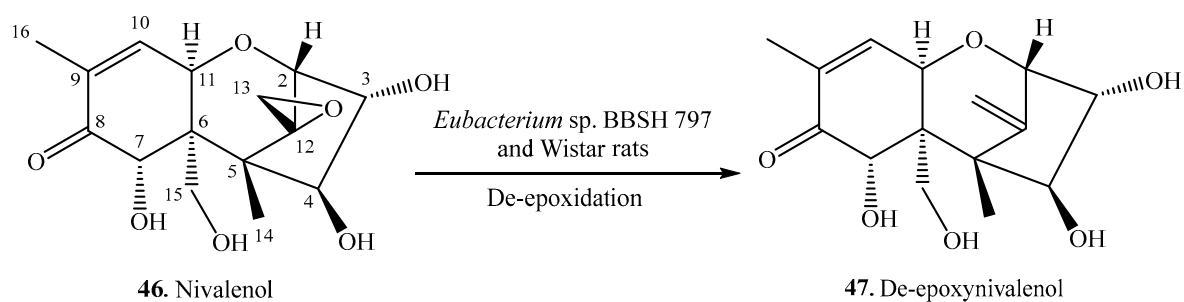


Figure S22. Transformation of nivalenol (**46**) with de-epoxidation by the bacterium *Eubacterium* sp. BBSH 797 [20] and Wistar rats [21].

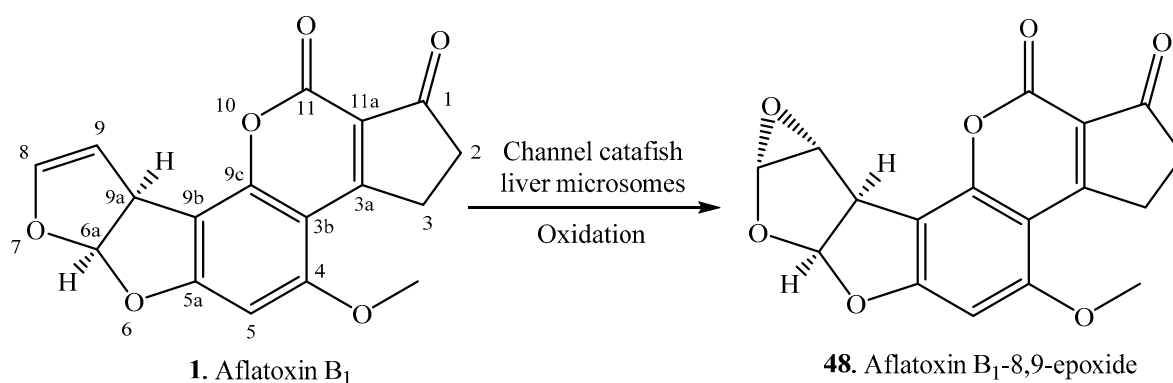


Figure S23. Transformation of aflatoxin B₁ (**1**) with oxidation by channel catfish microsomes [22].

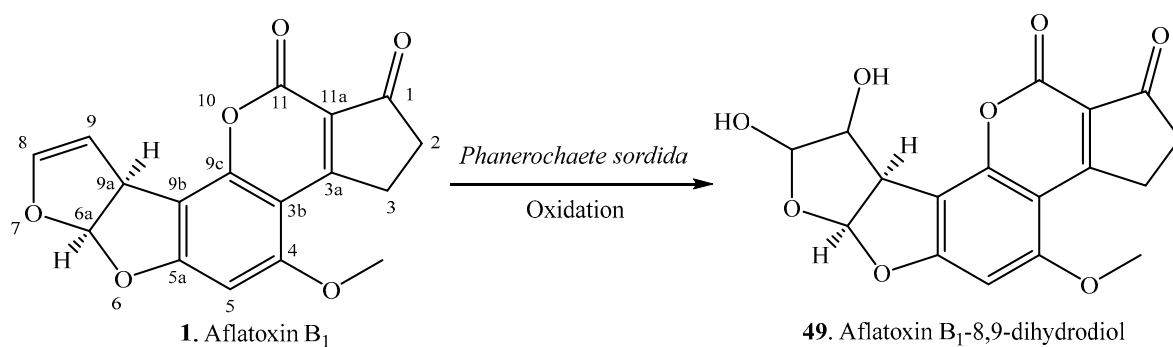


Figure S24. Transformation of aflatoxin B₁ (**1**) with oxidation by *Phanerochaete sordida* YK-624 [23].

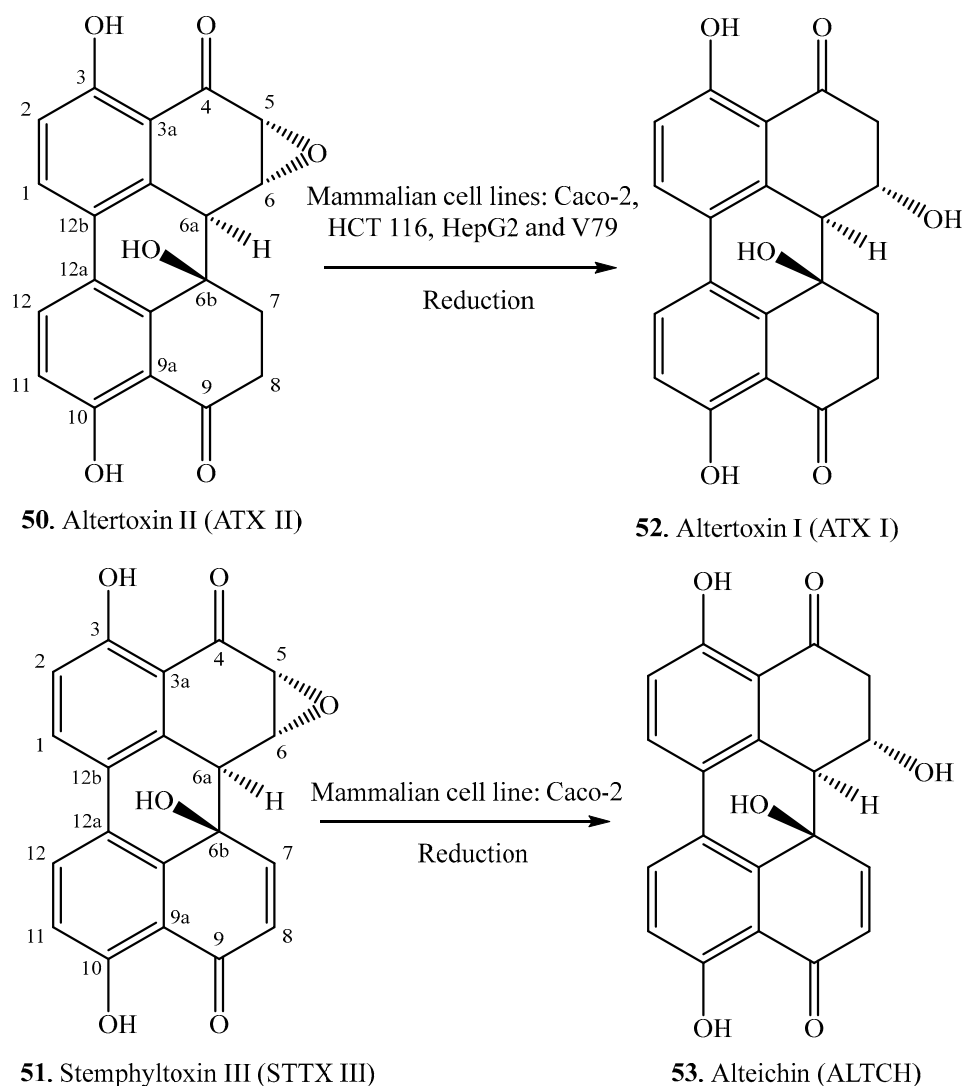


Figure S25. Transformation of altertoxin II (50) and stemphytoxin III (51) with reduction by mammalian cells [24].

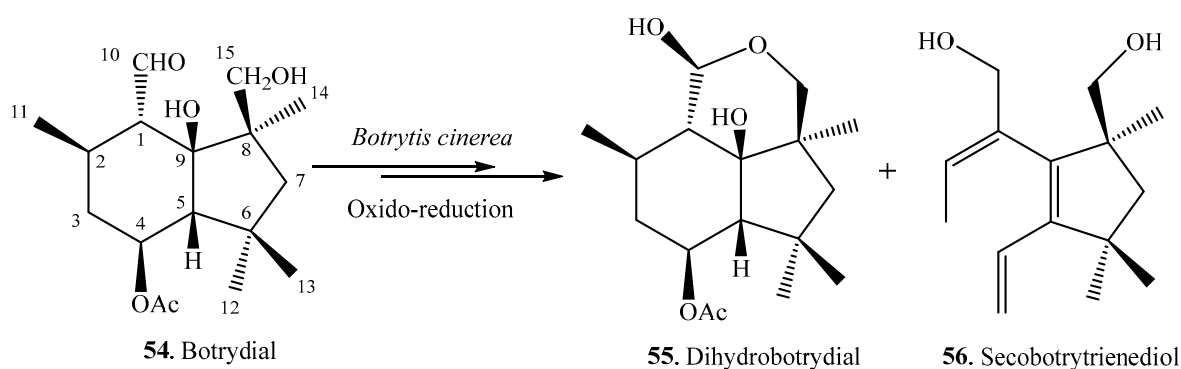


Figure S26. Transformation of botrydial (54) with oxido-reductions by *Botrytis cinerea* [25].

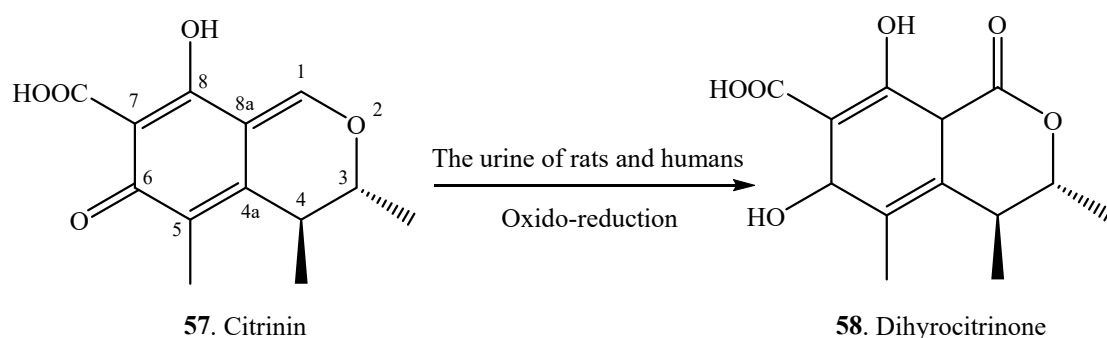


Figure S27. Transformation of citrinin (57) with oxido-reduction in the urine of rats and humans [26,27].

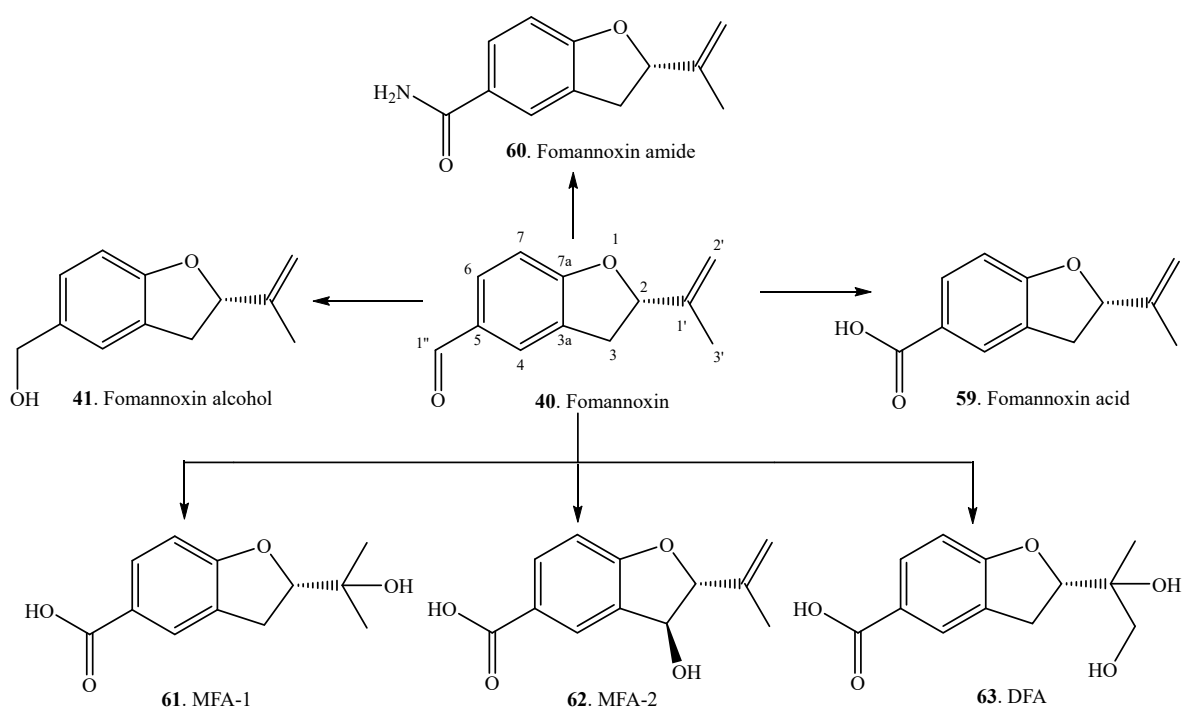


Figure S28. Transformation of fomannoxin (40) with oxido-reduction by rhizosphere-associated *Streptomyces* sp. AcH 505 [28].

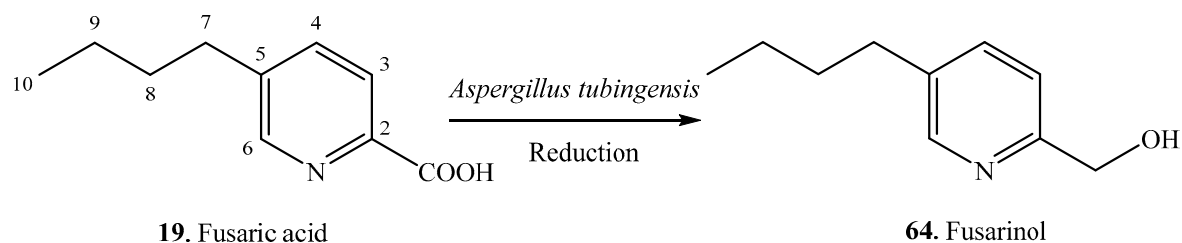


Figure S29. Transformation of fusaric acid (19) with reduction by *Aspergillus tubingensis* [29].

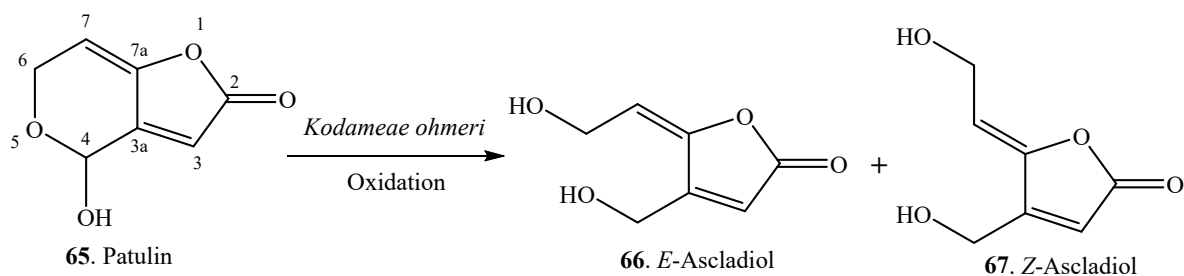


Figure S30. Transformation of patulin (65) with reduction by the yeast *Kodamea ohmeri* [30].

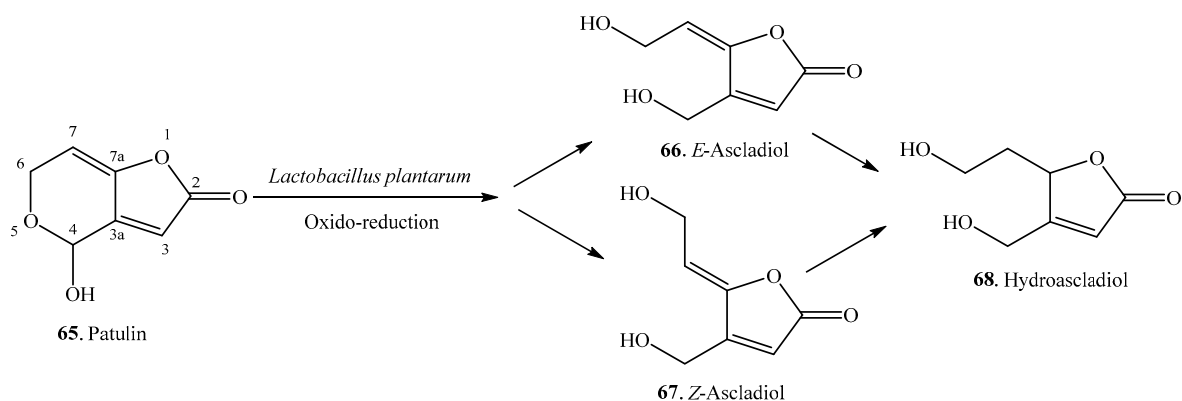


Figure S31. Transformation of patulin (65) with reduction by the bacterium *Lactobacillus plantarum* [31].

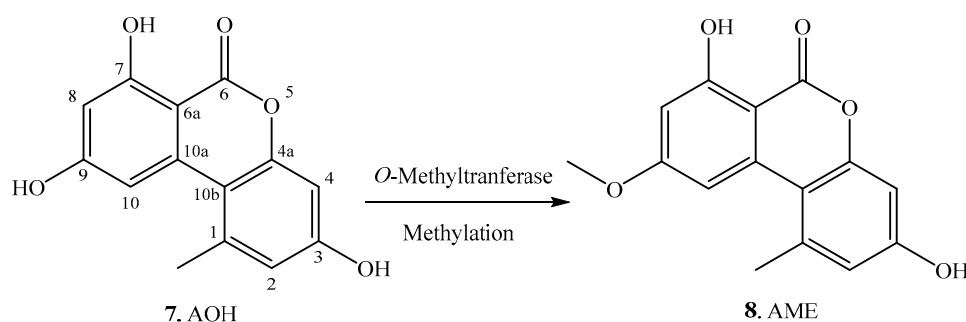


Figure S32. Transformation of alternariol (7) with methylation by *O*-methyltransferase [32].

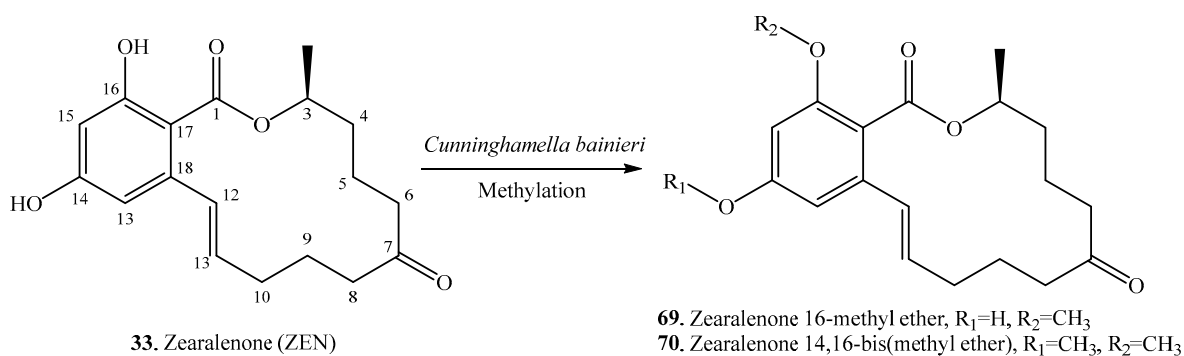


Figure S33. Transformation of zearalenone (33) with methylation by *Cunninghamella bainieri* [12].

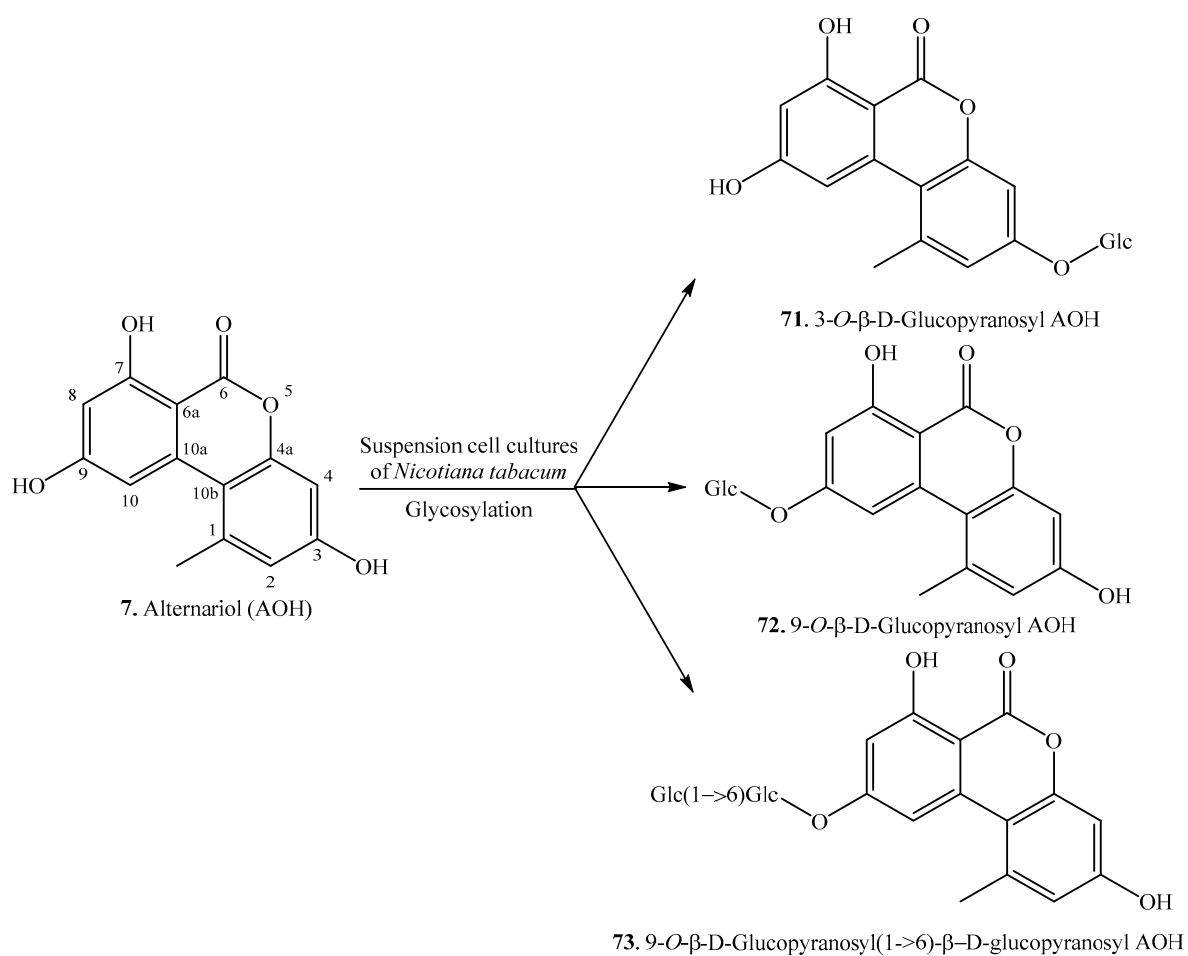


Figure S34. Transformation of alternariol (7) with glycosylation by suspension cell cultures of *Nicotiana batatum* [33].

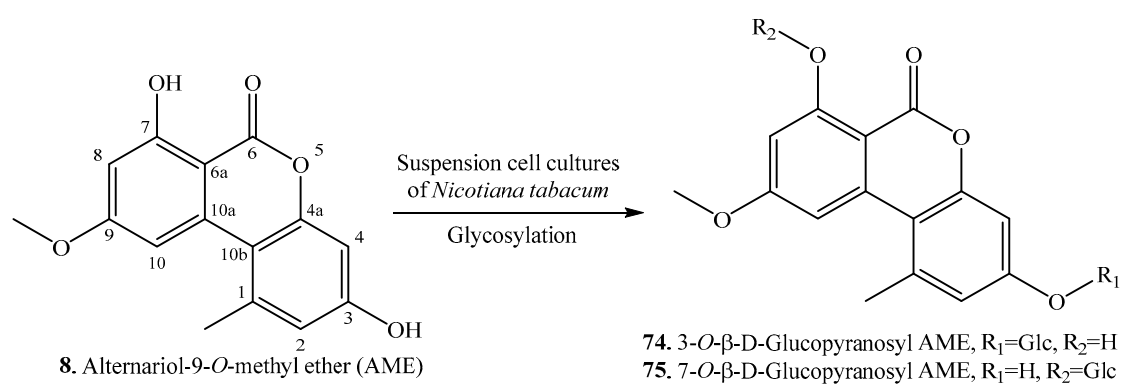


Figure S35. Transformation of alternariol 9-*O*-methyl ether (8) with glycosylation by suspension cell cultures of *Nicotiana batatum* [33].

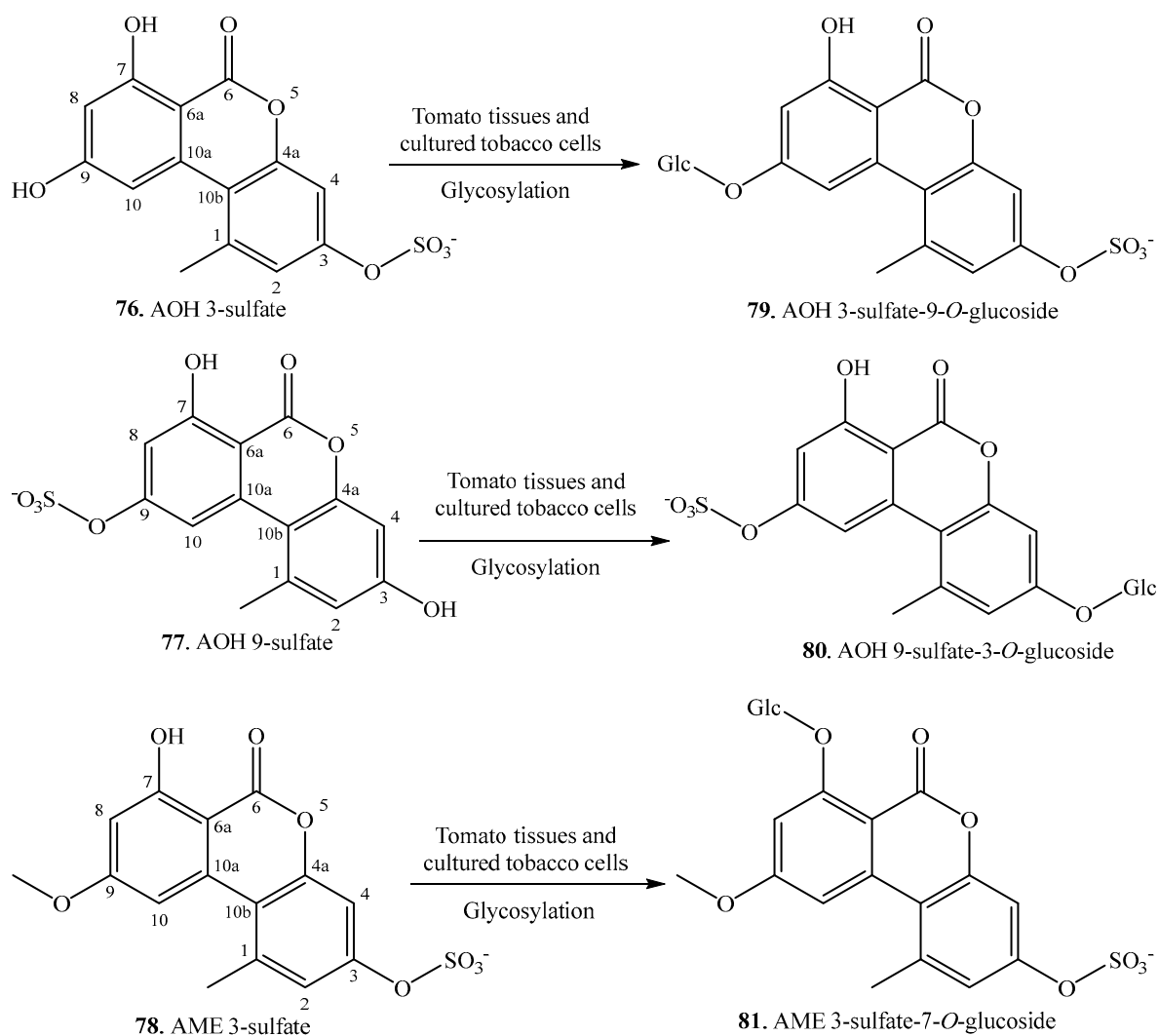


Figure S36. Transformation of alternariol 3-sulfate (76), alternariol 9-sulfate (77), and alternariol 9-O-methyl ether 3-sulfate (78) by tomato tissues and cultured tobacco cells [34].

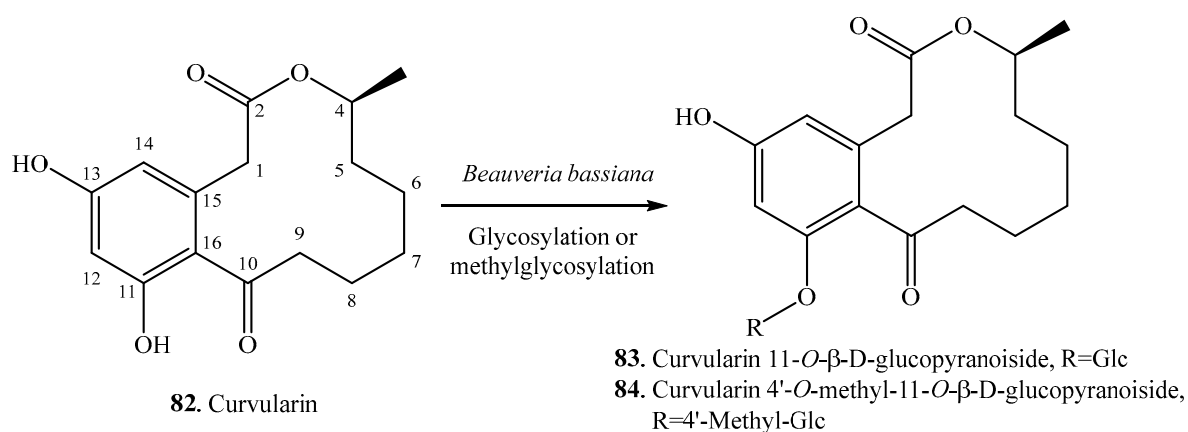


Figure S37. Transformation of curvularin (82) with glycosylation or methylglycosylation by *Beauveria bassiana* [35].

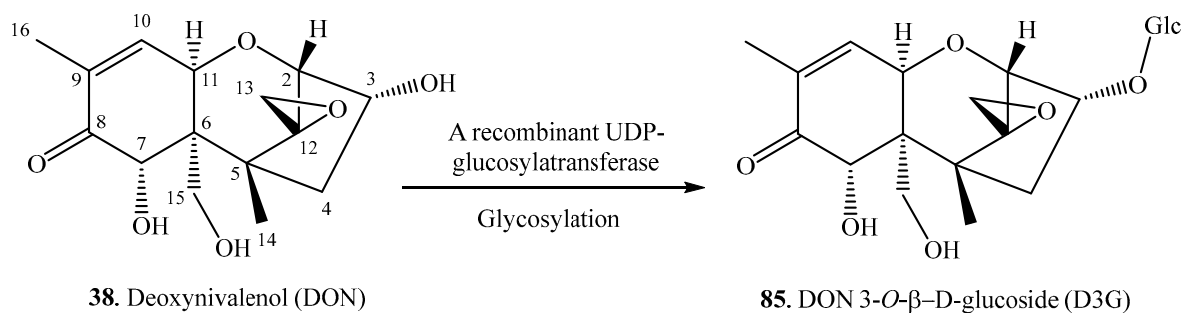


Figure S38. Transformation of deoxynivalenol (38) with glycosylation by a recombinant UDP-glucosyltransferase from rice [36].

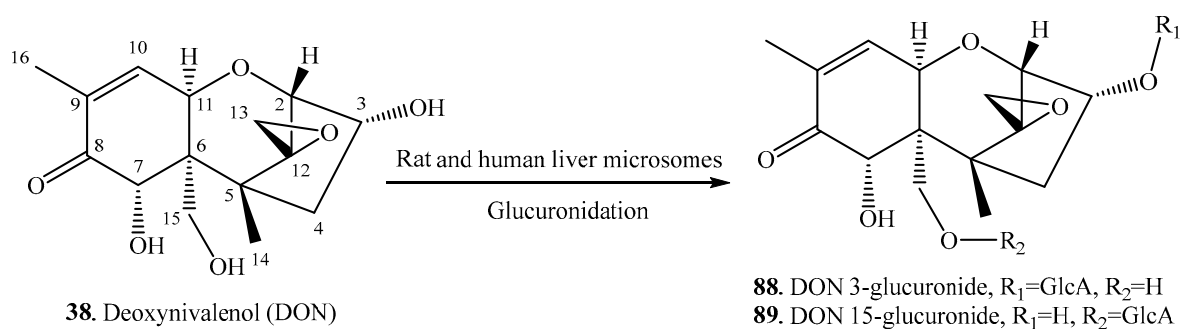


Figure S39. Transformation of deoxynivalenol (DON, 38) with glucuronidation by rat and human liver microsomes [37,38].

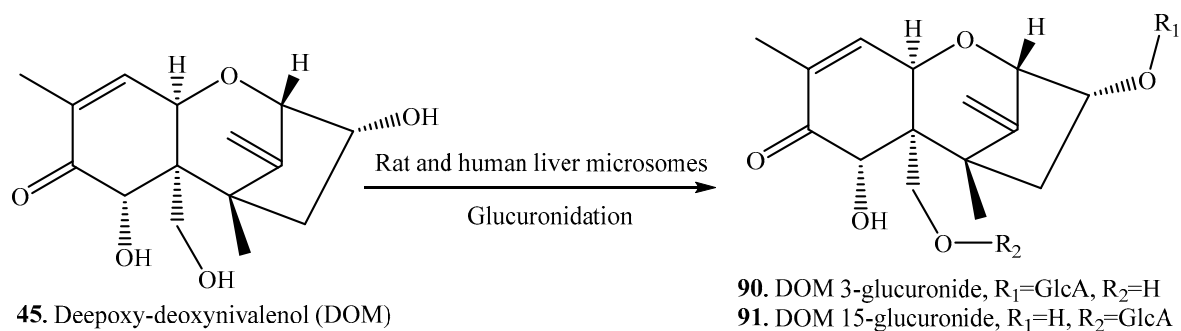


Figure S40. Transformation of deepoxy-deoxynivalenol (DOM, 45) with glucuronidation by rat and human liver microsomes [37,38].

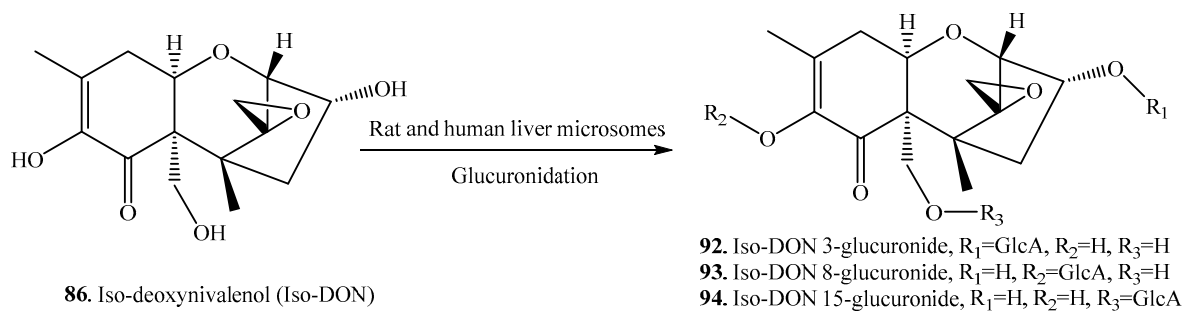


Figure S41. Transformation of iso-deoxynivalenol (iso-DON, 86) with glucuronidation by rat and human liver microsomes [37,38].

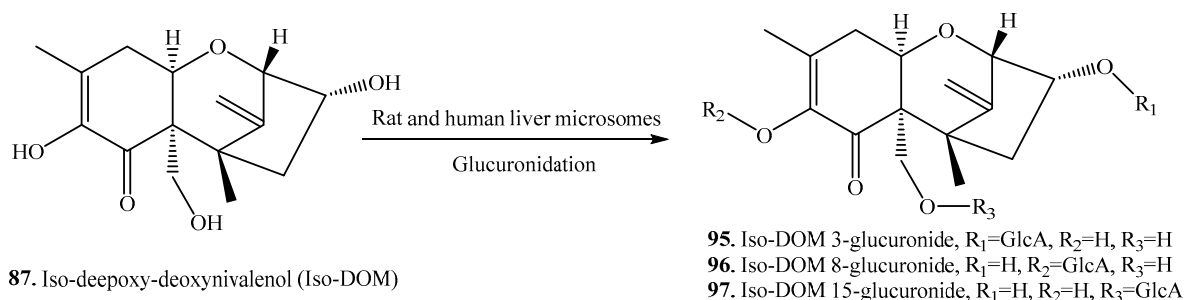


Figure S42. Transformation of iso-deepoxy-deoxynivalenol (iso-DOM, 87) with glucuronidation by rat and human liver microsomes [37,38].

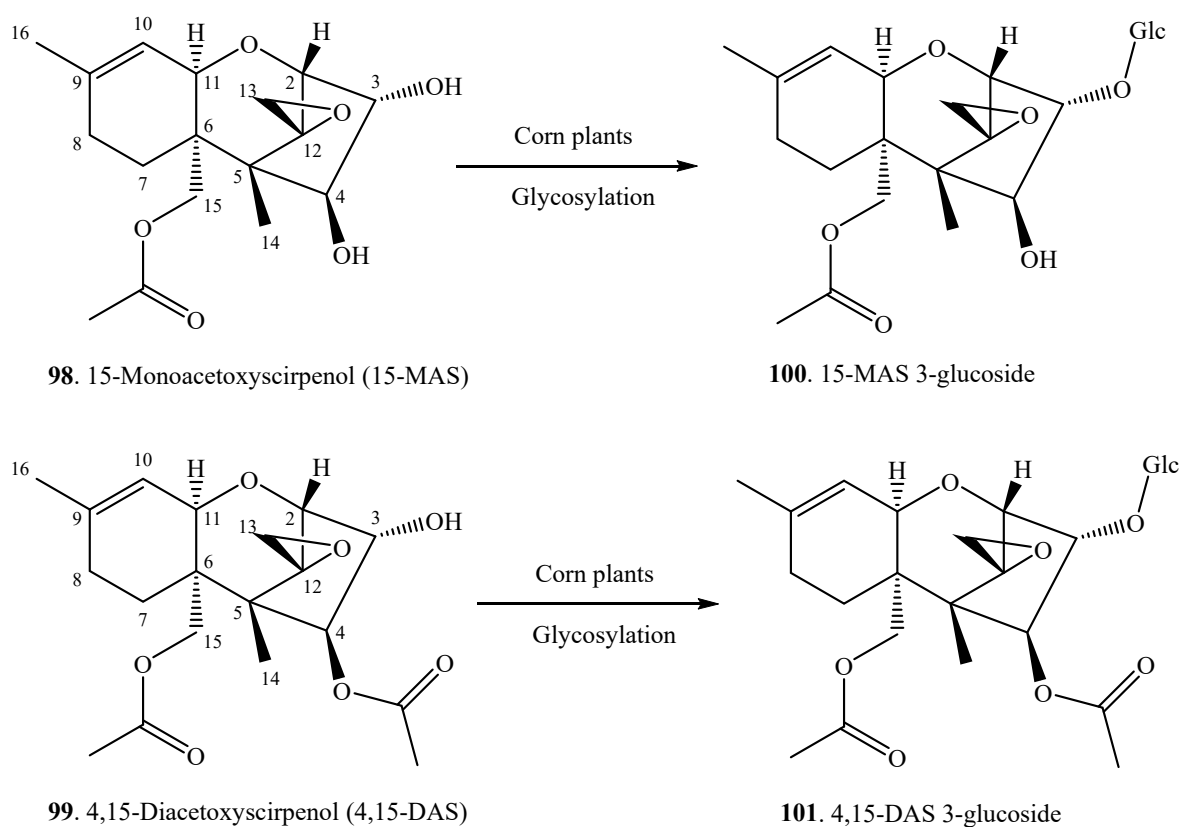


Figure S43. Transformation of 15-monoacetoxyscirpenol (98) and 4,15-diacetoxyscirpenol (99) with glycosylation respectively by corn plants [39].

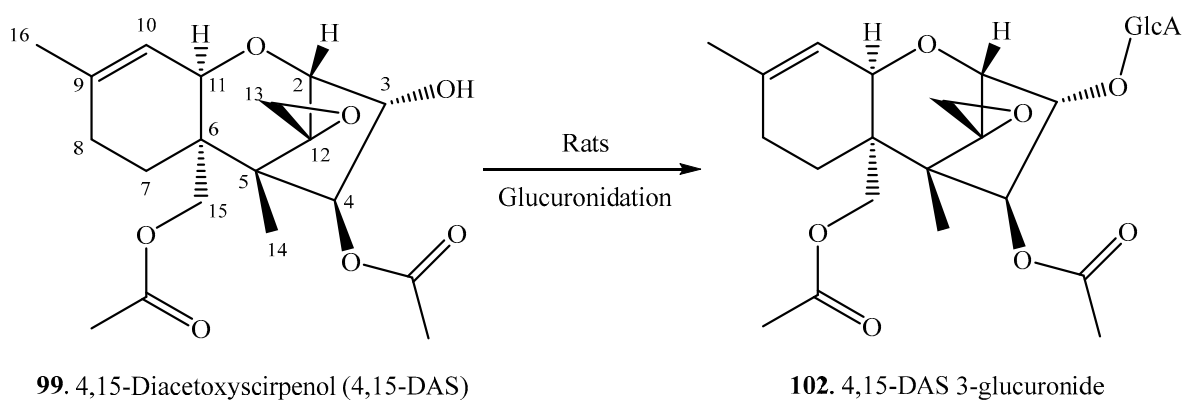


Figure S44. Transformation of 4,15-diacetoxyscirpenol (99) with glucuronidation in rats [40].

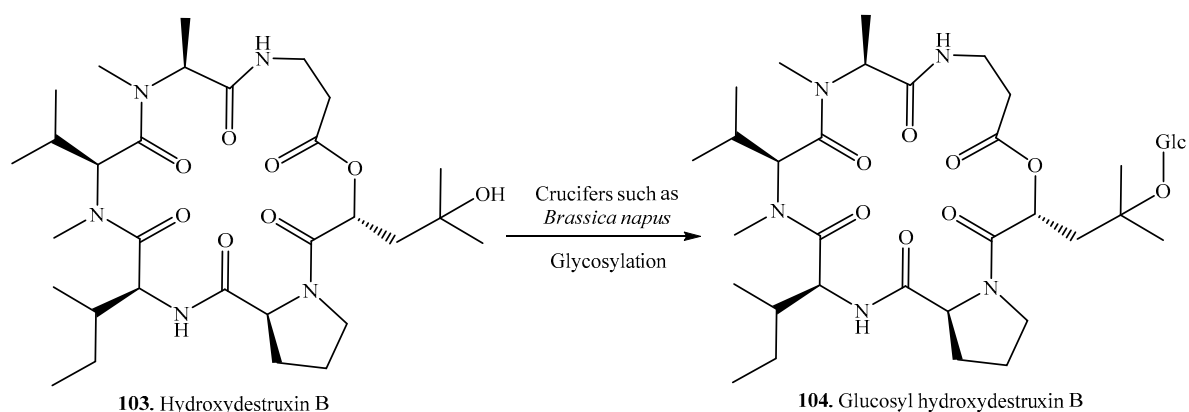


Figure S45. Transformation of hydroxydestruxin B (103) with glycosylation by crucifers such as *Brassica napus* [4].

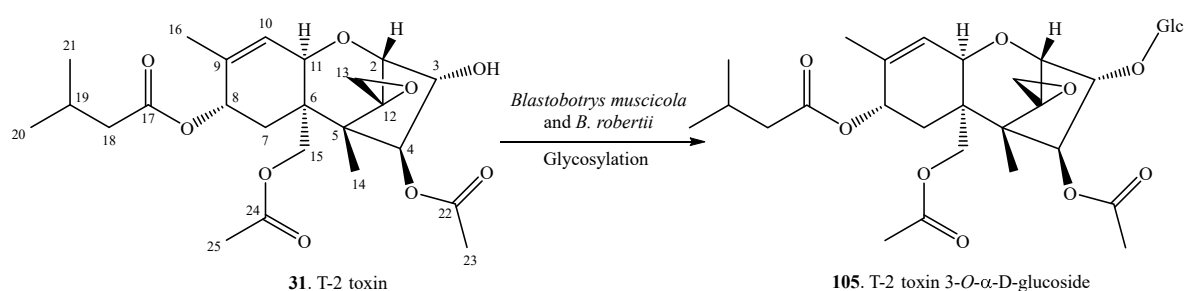


Figure S46. Transformation of T-2 toxin (31) with glycosylation by *Blastobotrys muscicola* and *B. robertii* [41].

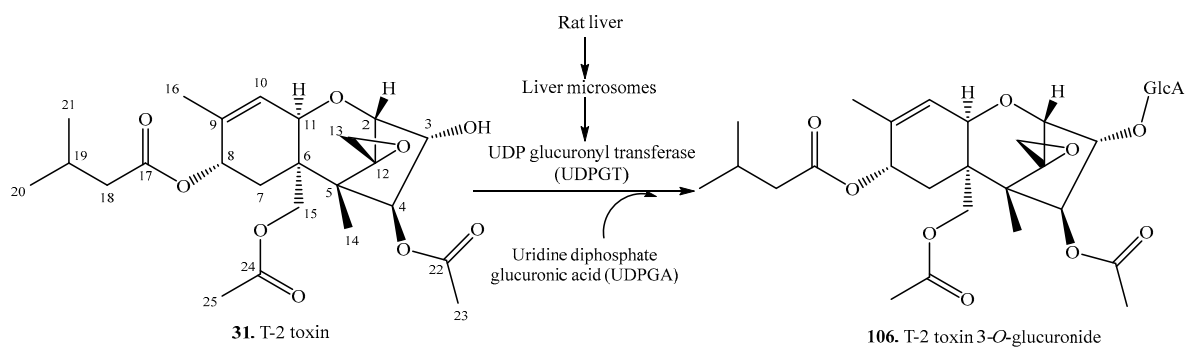


Figure S47. Transformation of T-2 toxin (31) with glucuronidation by rat liver microsomes [42].

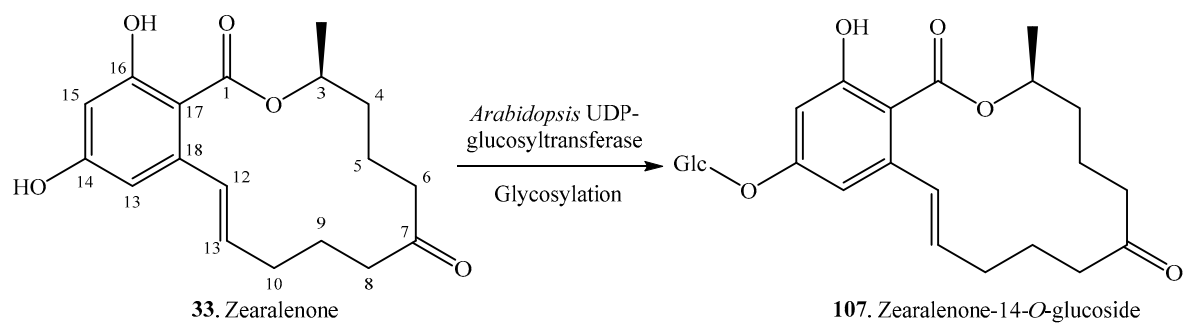


Figure S48. Transformation of zearalenone (33) by *Arabidopsis* UDP-glucosyltransferases expressed in *Saccharomyces cerevisiae* [43].

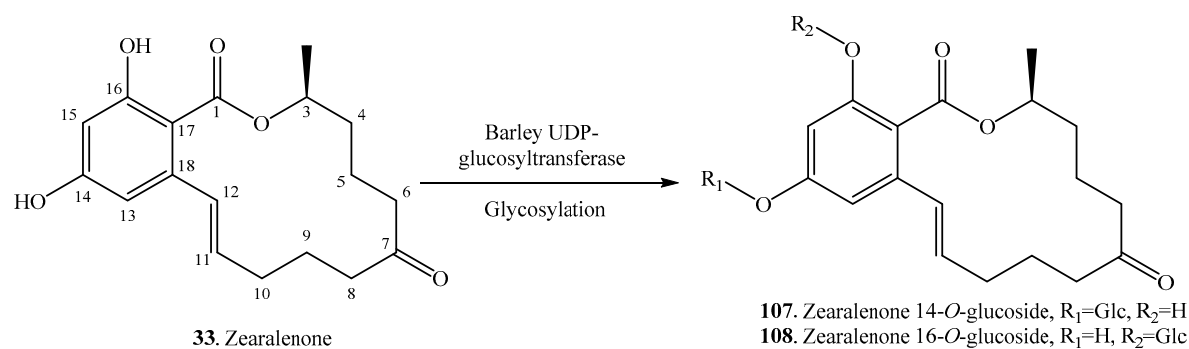


Figure S49. Transformation of zearalenone (**33**) by barley UDP-glucosyltransferases expressed in *Saccharomyces cerevisiae* [44].

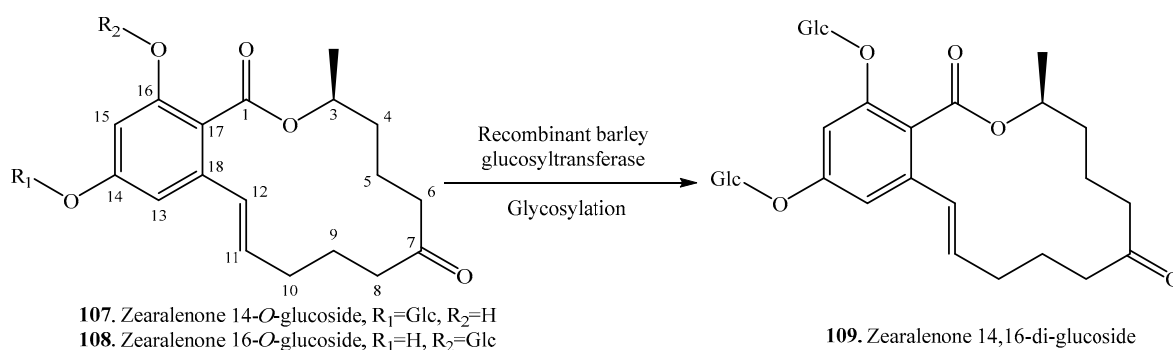


Figure S50. Transformation of zearalenone 14-O-glucoside (**107**) and zearalenone 16-O-glucoside (**108**) by the recombinant barley glucosyltransferases [45].

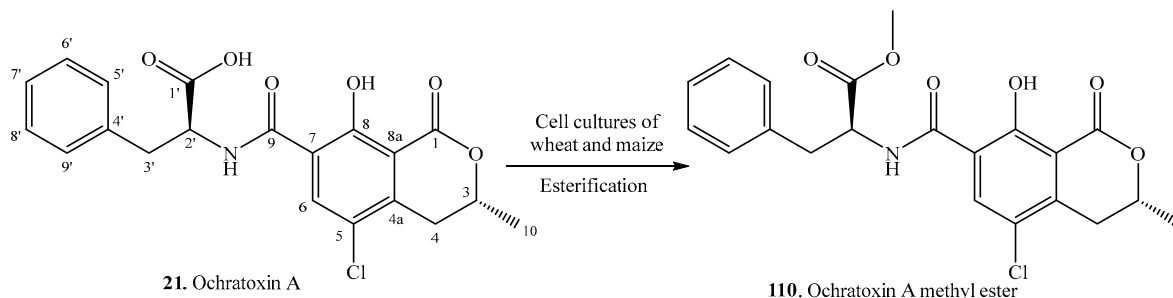


Figure S51. Transformation of ochratoxin A (**21**) with esterification by the cell cultures of wheat and maize [46].

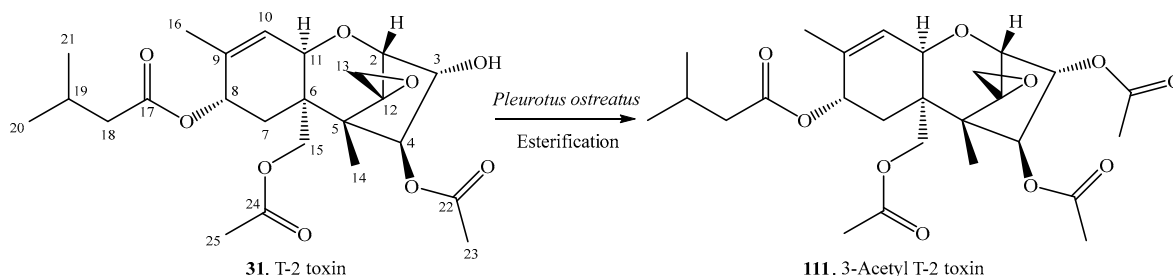


Figure S52. Transformation of T-2 toxin (**31**) with esterification (acetylation) by *Pleurotus ostreatus* [47].

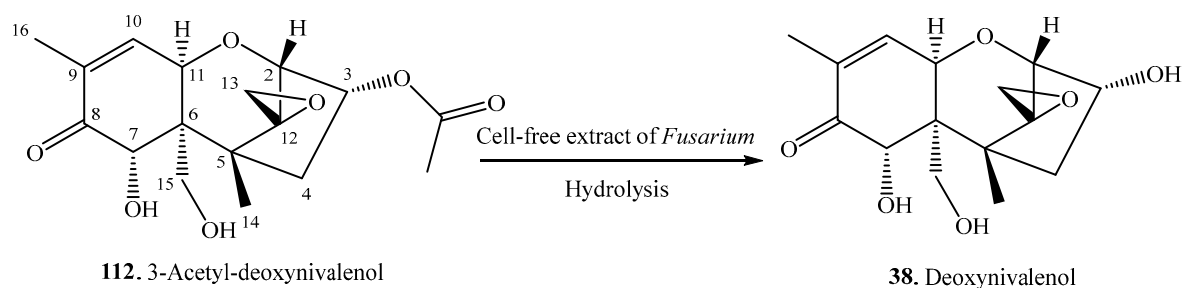


Figure S53. Transformation of 3-acetyl-deoxynivalenol (112) with hydrolysis (deacetylation) by cell-free extract of *Fusarium* sp. [48].

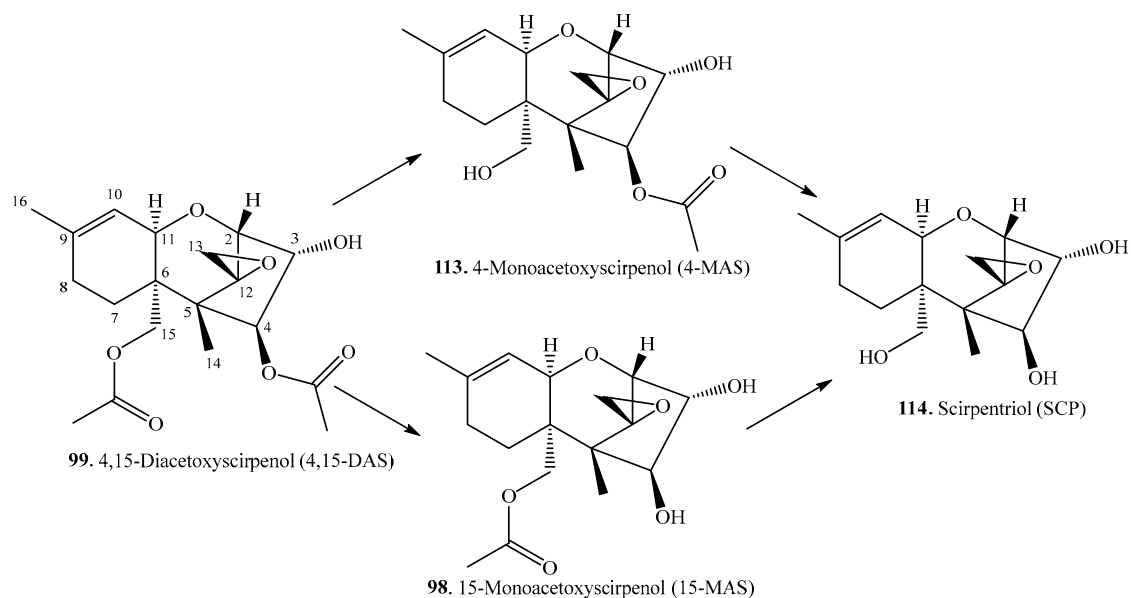


Figure S54. Transformation of 4,15-diacetoxyscirpenol (99) with hydrolysis (deacetylation) in rats [40].

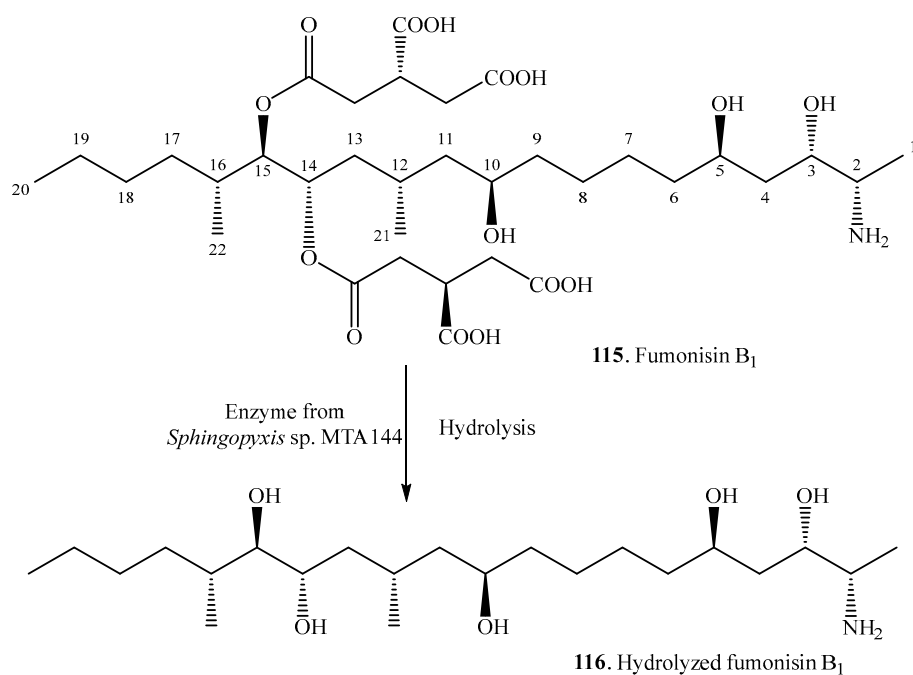


Figure S55. Transformation of fumonisin B₁ (115) with hydrolysis by the enzyme from *Sphingopyxis* sp. MTA144 [49].

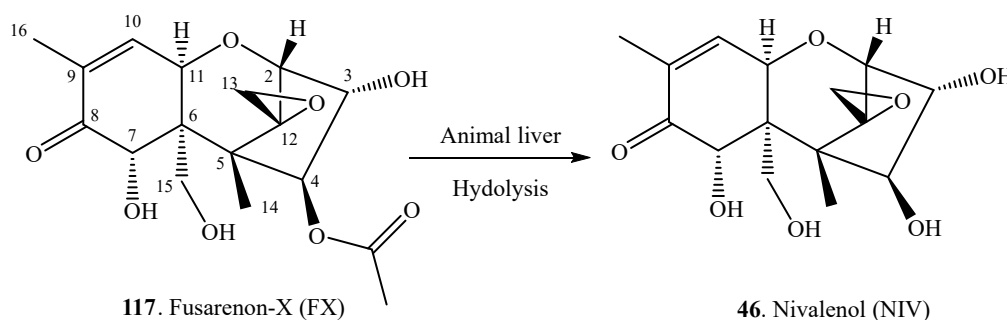


Figure S56. Transformation of fusarenon-X (117) with hydrolysis (deacetylation) in animal liver [50,51].

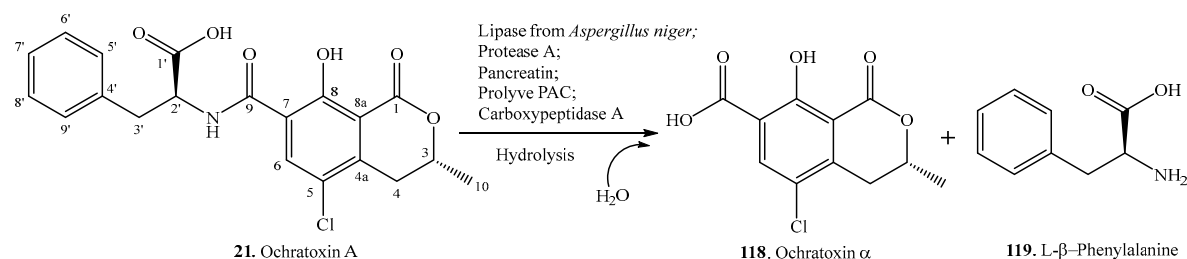


Figure S57. Transformation of ochratoxin A (21) with hydrolysis by enzymes [52–54].

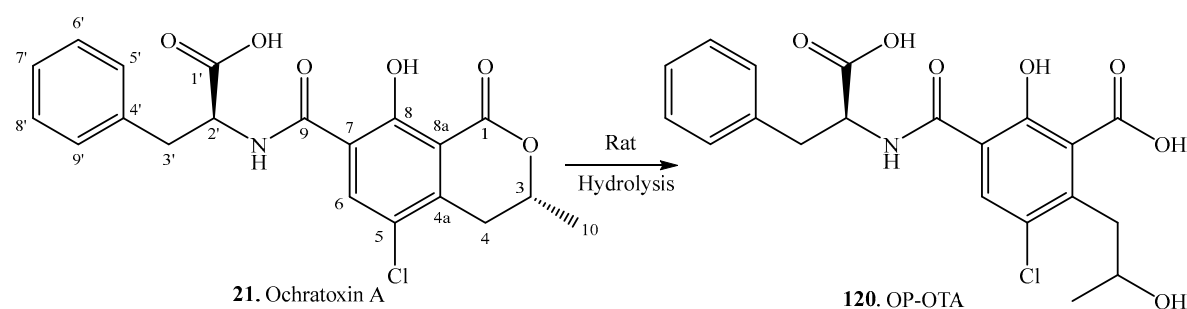


Figure S58. Transformation of ochratoxin A (21) with hydrolysis in rats [55].

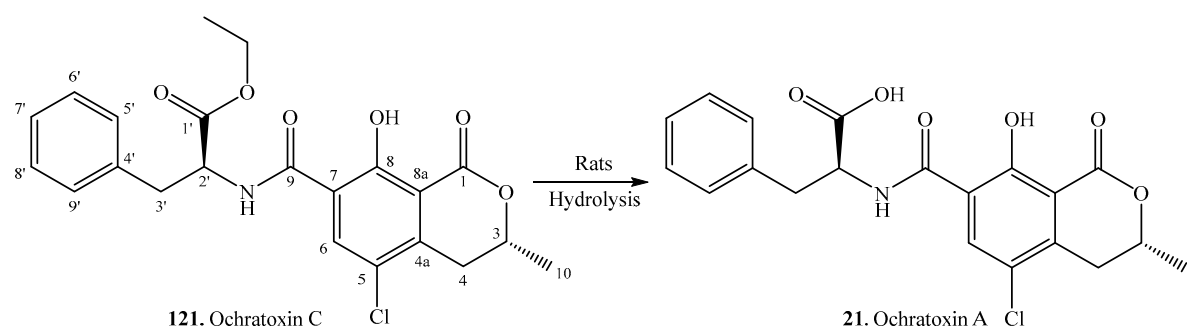


Figure S59. Transformation of ochratoxin C (121) with hydrolysis in rats [56].

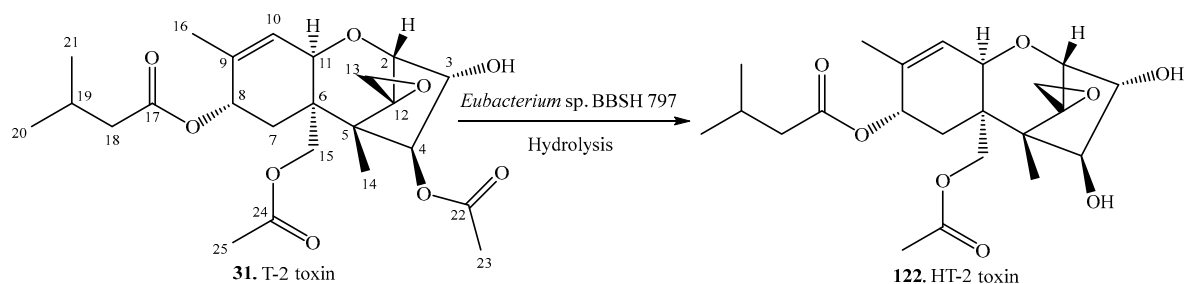


Figure S60. Transformation of T-2 toxin (31) with deacetylation by *Eubacterium* sp. 797 [57].

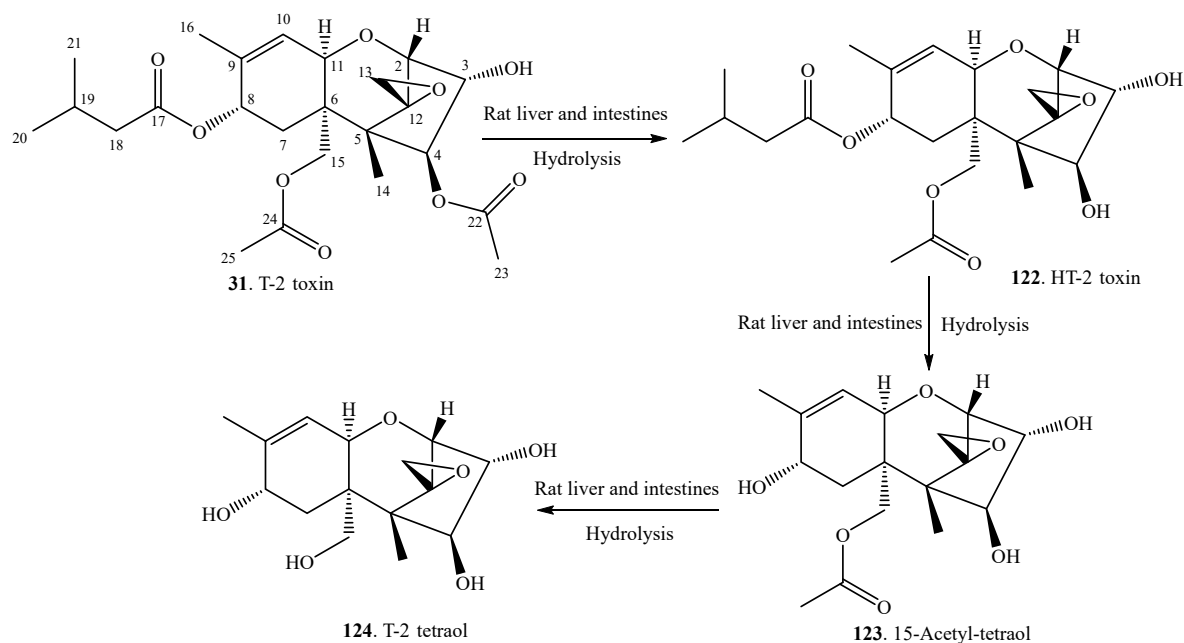


Figure S61. Transformation of T-2 toxin (31) with multi-step hydrolysis in rat liver and intestines [58].

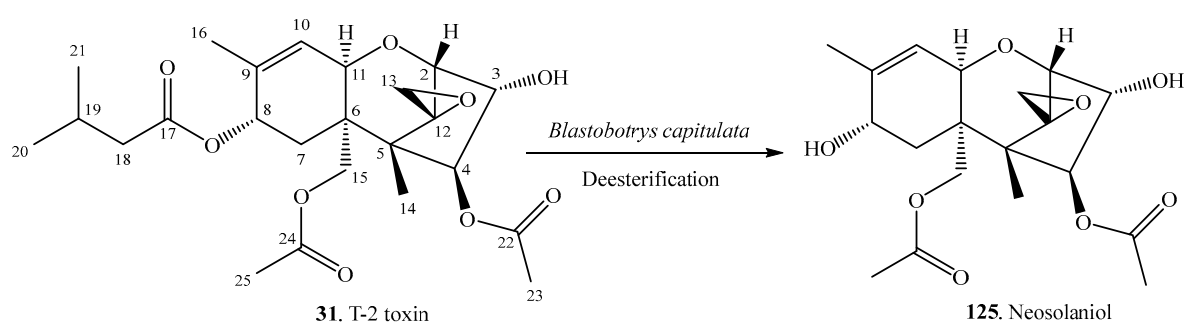


Figure S62. Transformation of T-2 toxin (31) with deesterification by the fungus *Blastobotrys capitulata* [41].

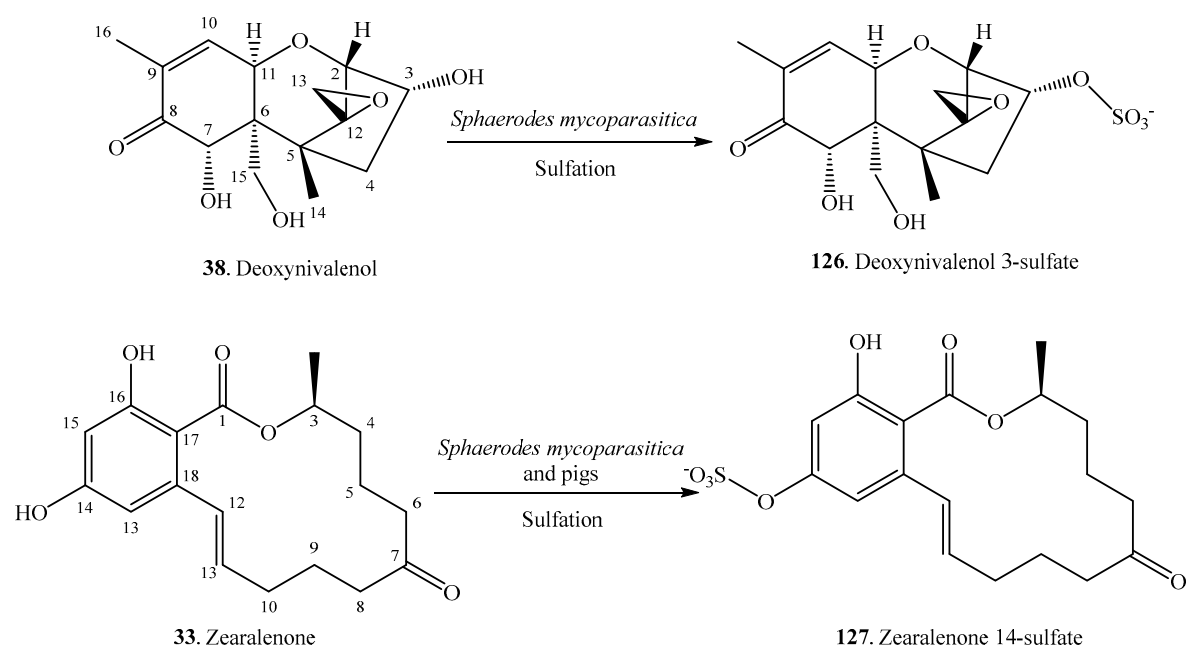


Figure S63. Both deoxynivalenol (DON, **38**) and zearalenone (ZEN, **33**) were converted to their corresponding sulfates DON 3-sulfate (**126**) and ZEN 14-sulfate (**127**) by the fungus *Sphaerodes mycoparasitica* [59]. ZEN (**33**) was converted to ZEN 14-sulfate (**127**) by pigs [60].

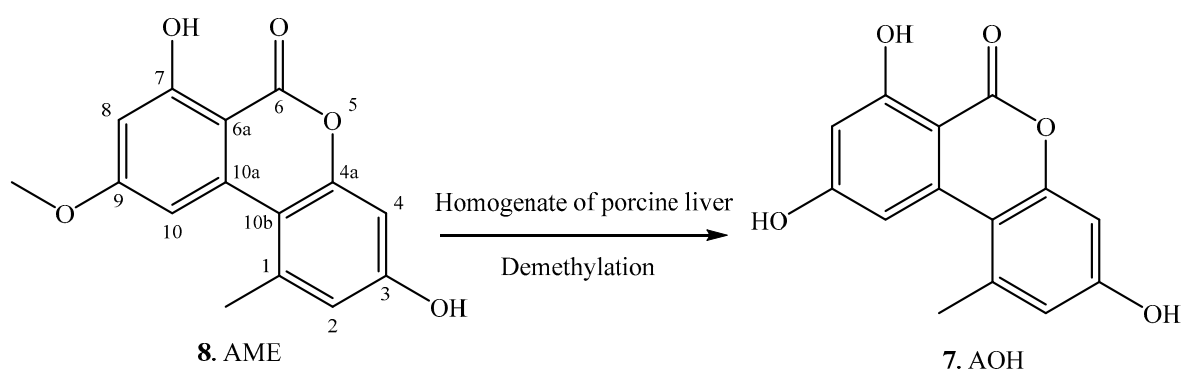


Figure S64. Transformation of alternariol 9-O-methylether (AME, **8**) with demethylation by the homogenate of porcine liver in the presence of NADPH [61].

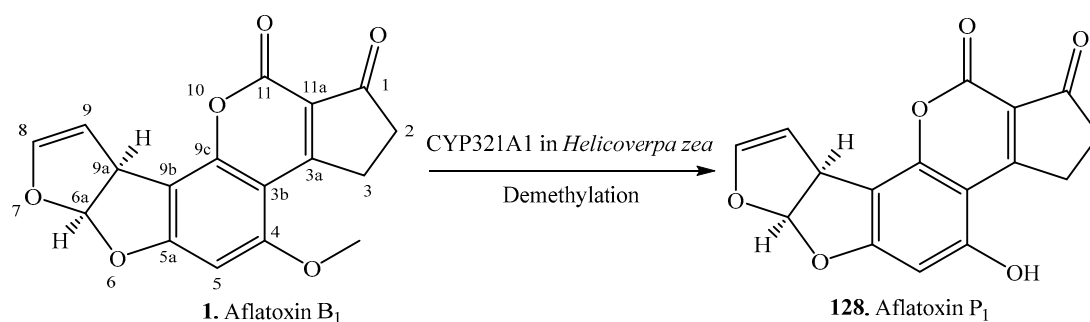


Figure S65. Transformation of aflatoxin B₁ (**1**) with demethylation by CYP21A1 in *Helicoverpa zea* [62].

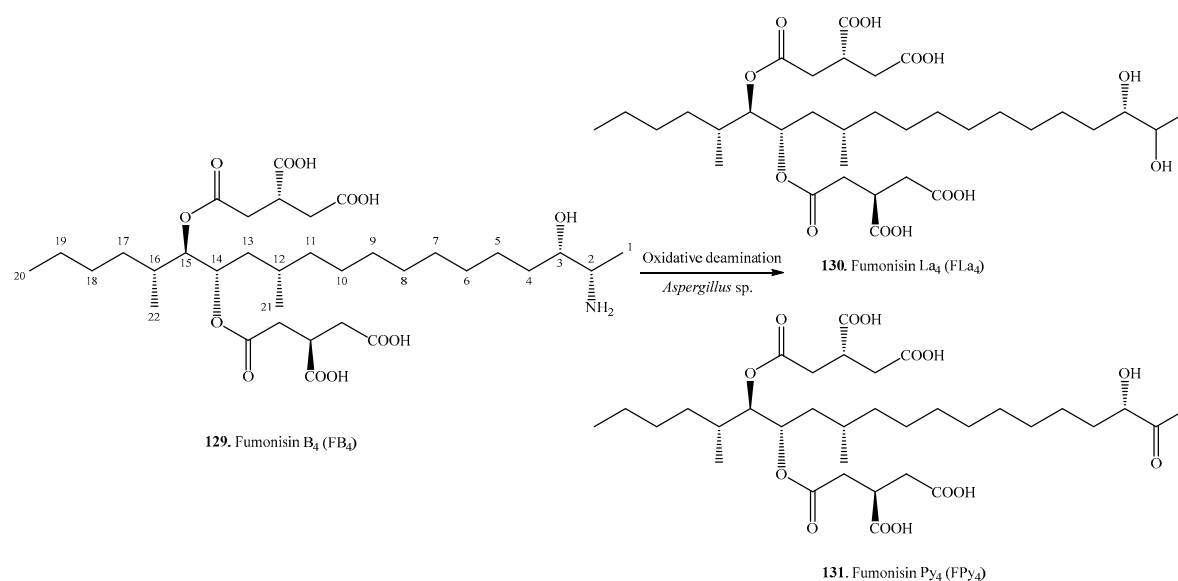


Figure S66. Transformation of fumonisin B₄ (129) with oxidative deamination by *Aspergillus* sp. [63].

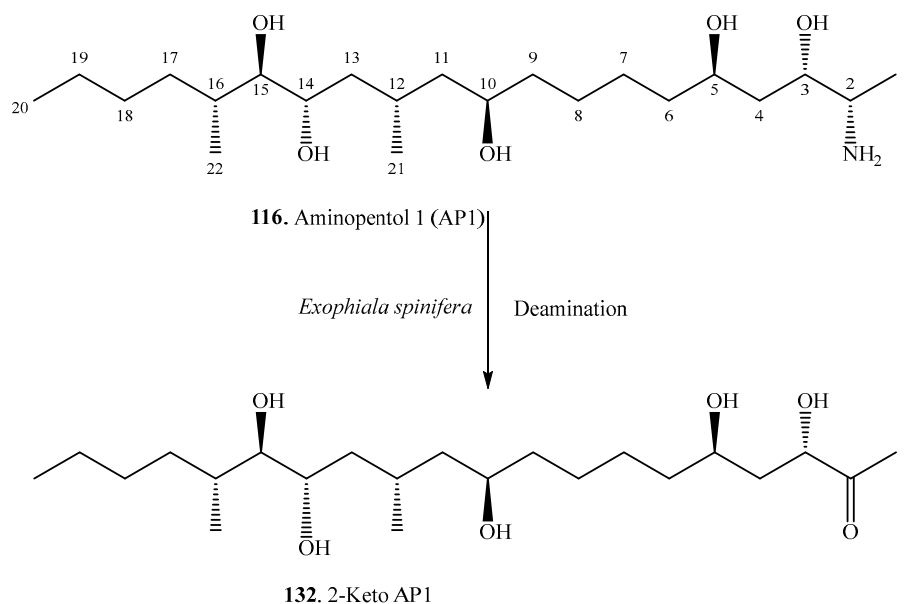


Figure S67. Transformation of aminopentol 1 (AP1, 116) with oxidative deamination by *Exophiala spinifera* [64].

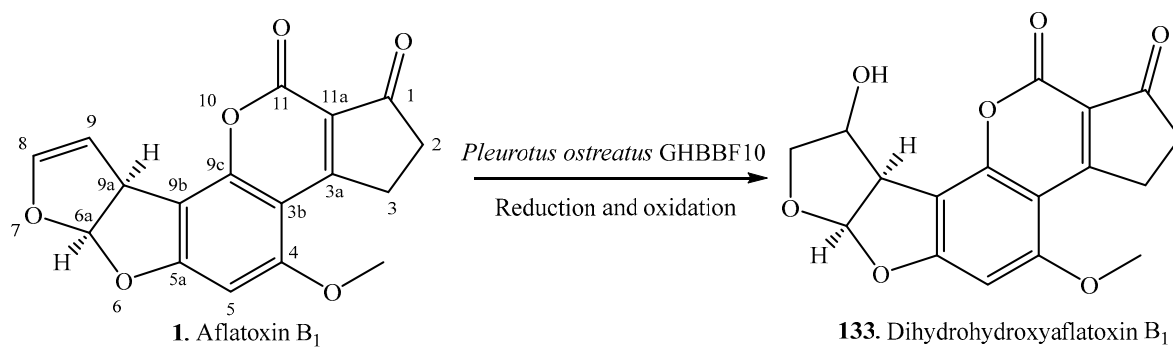


Figure S68. Transformation of aflatoxin B₁ (1) with reduction and oxidation by *Pleurotus ostreatus* GHBBF10 [65].

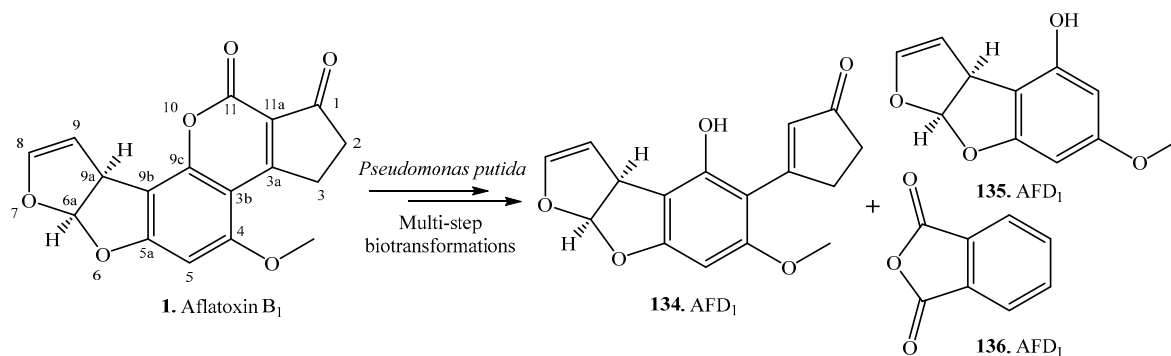


Figure S69. Transformation of aflatoxin B₁ (1) with hydrolysis, decarboxylation and oxidation-reduction by *Pseudomonas putida* [66].

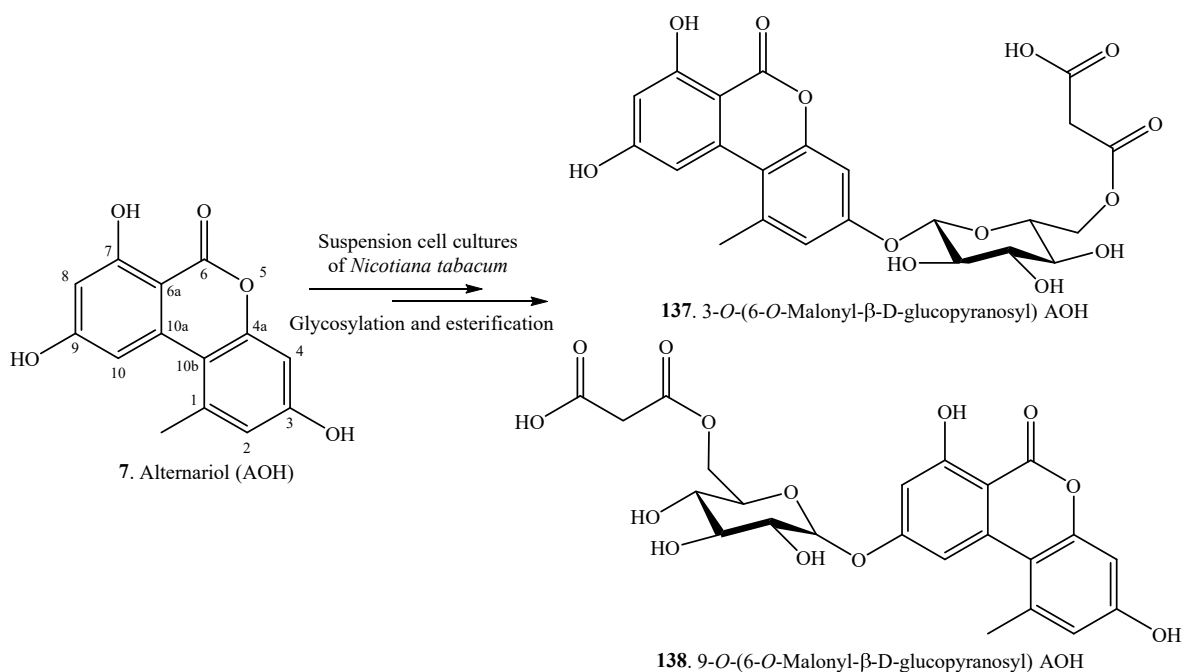


Figure S70. Transformation of alternariol (7) through glycosylation and esterification by suspension cell cultures of *Nicotiana batatum* [33].

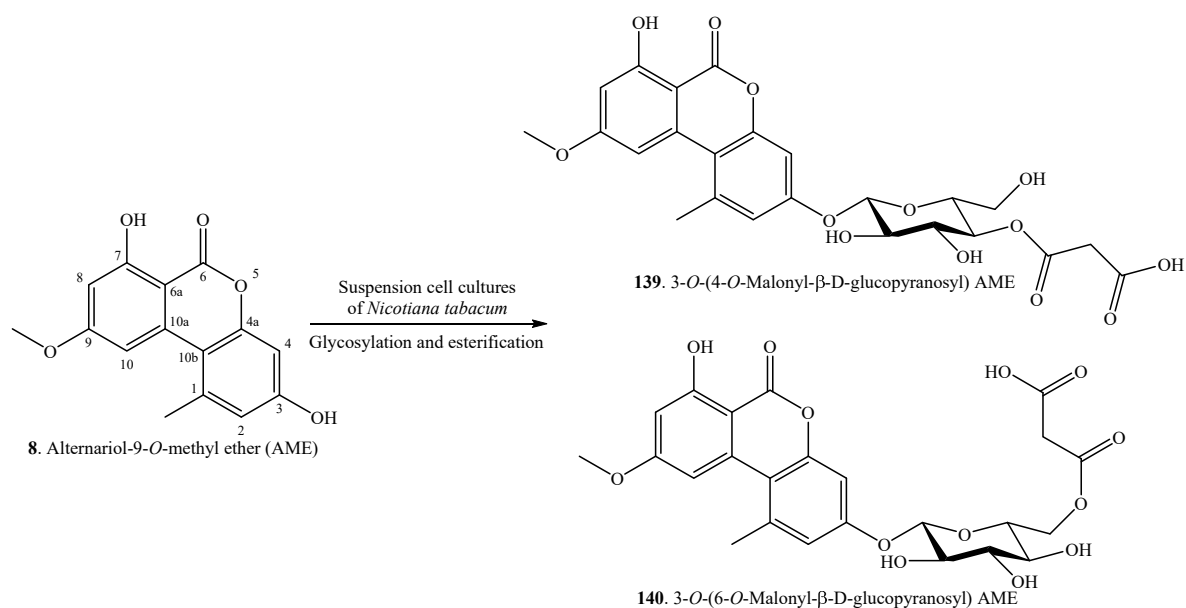


Figure S71. Transformation of alternariol 9-O-methyl ether (8) through glycosylation and esterification by suspension cell cultures of *Nicotiana batatum* [33].

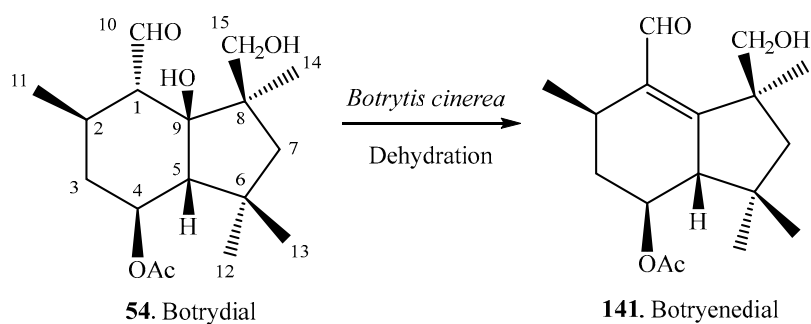


Figure S72. Transformation of botrydial (54) with dehydration by *Botrytis cinerea* [25].

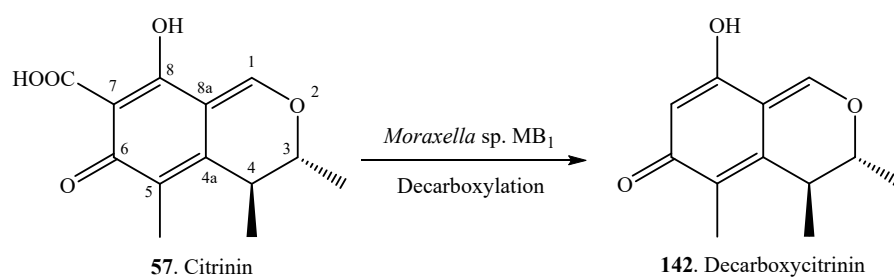


Figure S73. Transformation of citrinin (57) with decarboxylation by *Moraxella* sp. MB₁ [67].

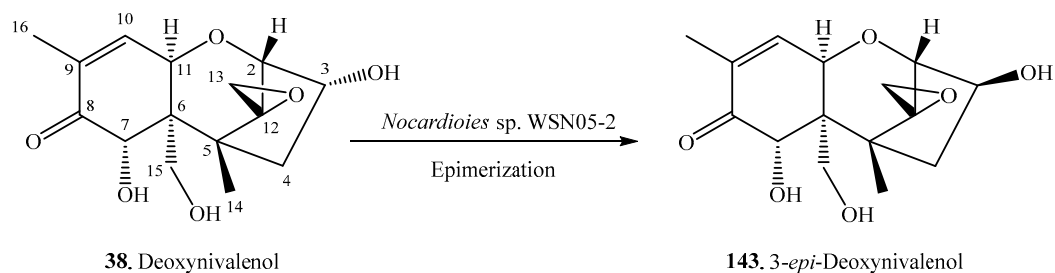


Figure S74. Transformation of deoxynivalenol (38) with epimerization by *Nocardioles* sp. WSN05-2 [68].

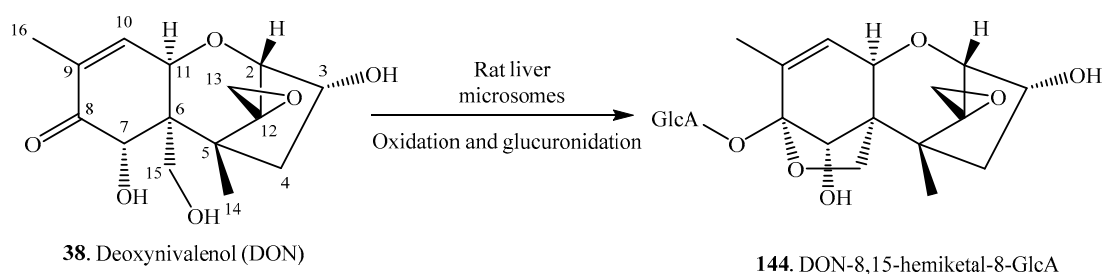


Figure S75. Transformation of deoxynivalenol (DON, 38) with oxidation and glucuronidation by rate liver microsomes [37,38].

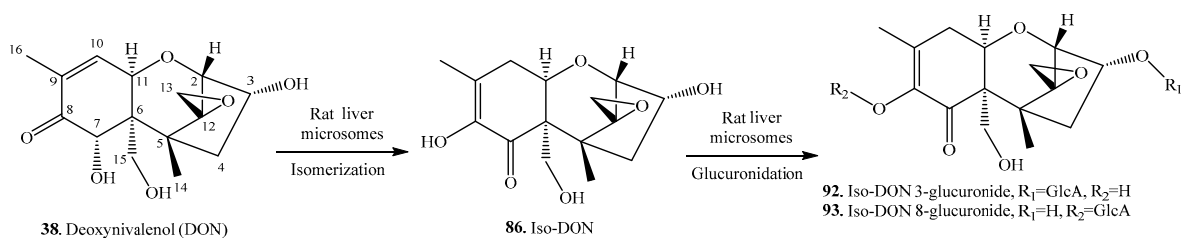


Figure S76. Transformation of deoxynivalenol (DON, 38) with isomerization and glucuronidation by rate liver microsomes [37,38].

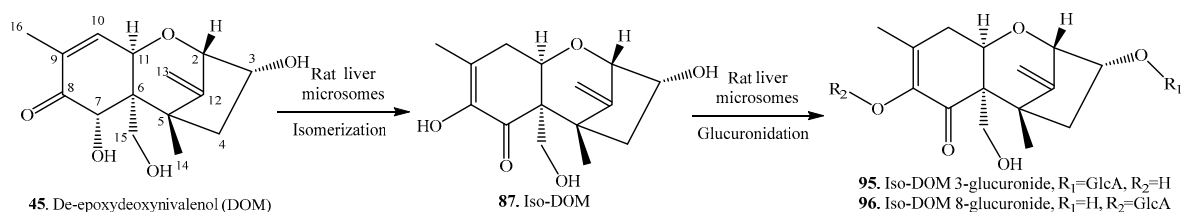


Figure S77. Transformation of deepoxy-deoxynivalenol (DOM, 45) with isomerization and glucuronidation by rate liver microsomes [37,38].

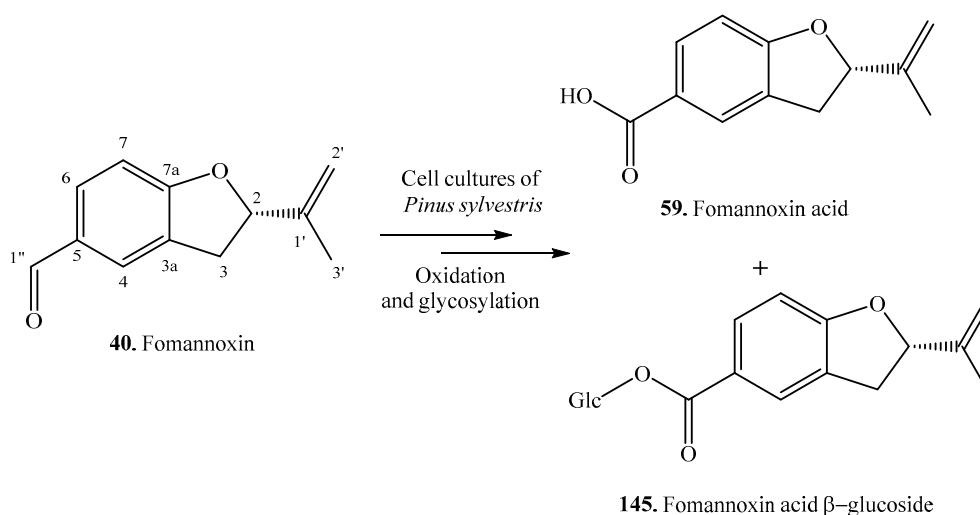


Figure S78. Transformation of fomannoxin (40) with oxidation and glycosylation by cell cultures of *Pinus sylvestris* [16].

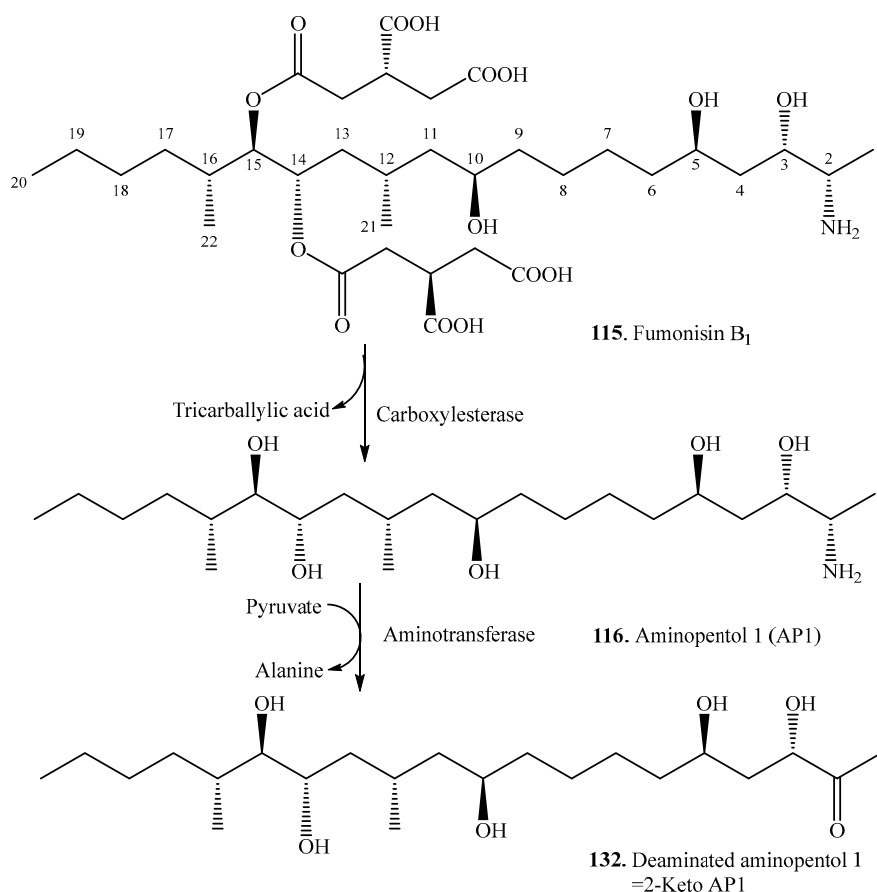


Figure S79. Transformation of fumonisin B₁ (115) with hydrolysis and deamination by the recombinant enzymes from the bacterium *Sphingopyxis* sp. MTA144 [69].

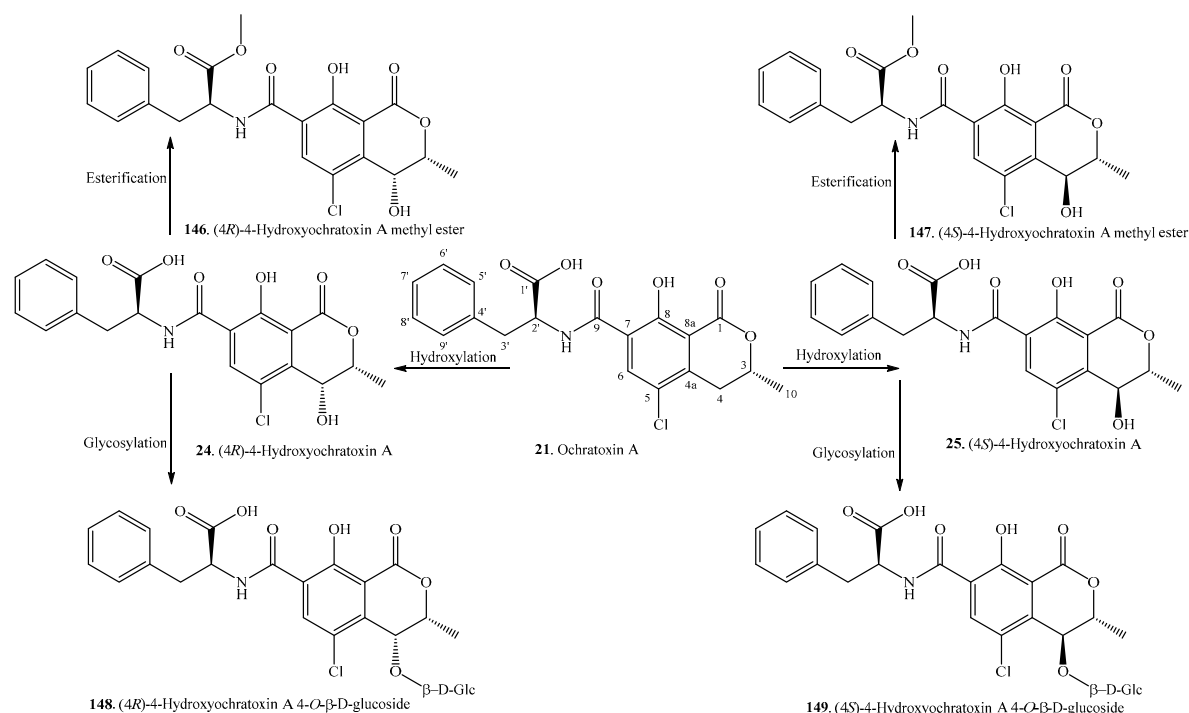


Figure S80. Transformation of ochratoxin A (21) with hydroxylation further esterification or glycosylation by the cell cultures of wheat and maize [46].

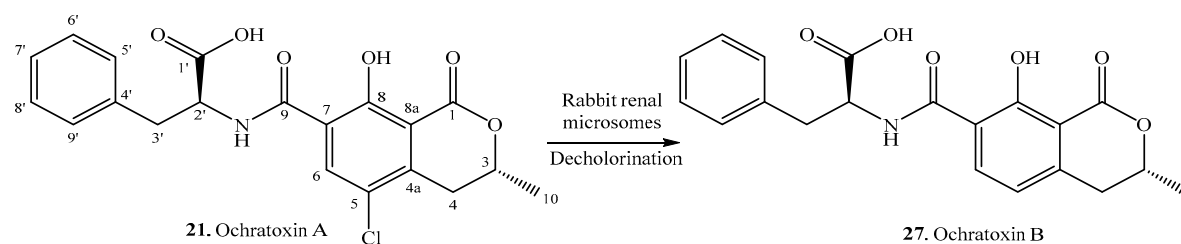


Figure S81. Transformation of ochratoxin A (21) with dechlorination in the renal microsomes [70].

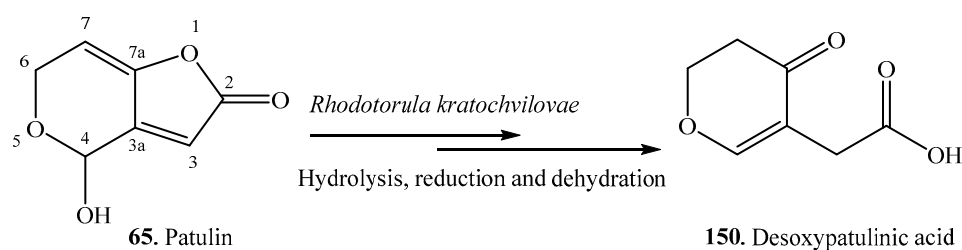


Figure S82. Transformation of patulin (65) with hydrolysis, reduction and dehydration by *Rhodotorula kratochvilovae* [71].

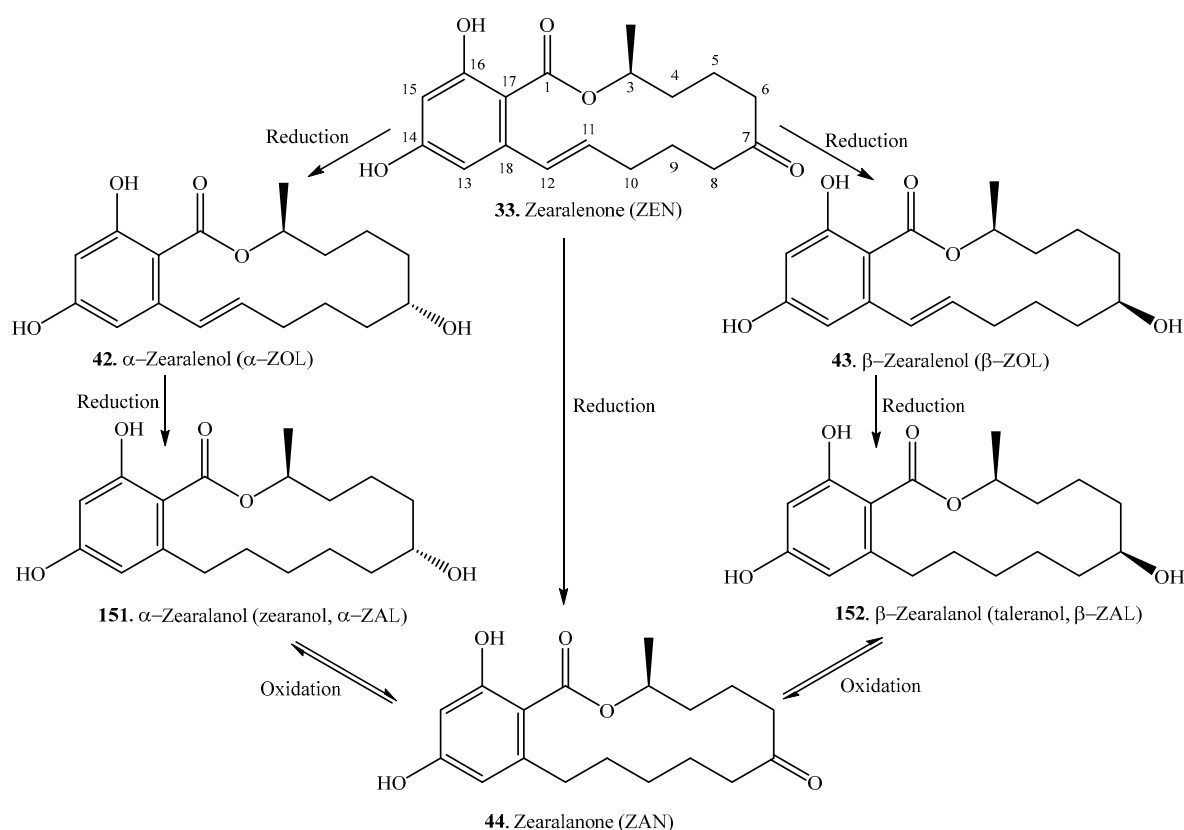


Figure S83. Transformation of zearalenone (33) with multi-step oxido-reductions in human body [72].

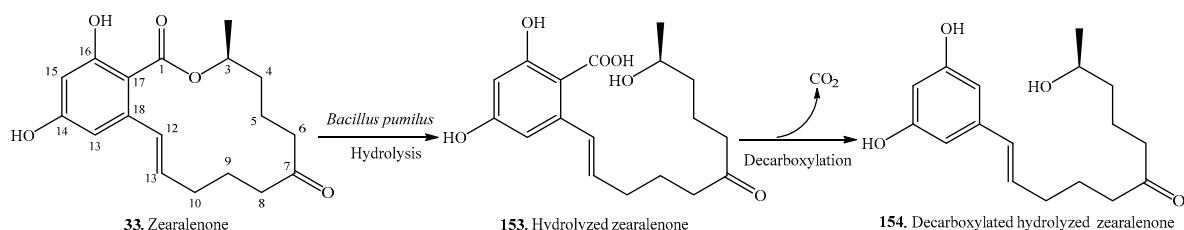


Figure S84. Transformation of zearalenone (33) with hydrolysis and decarboxylation by *Bacillus pumilus* [42].

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