

Table 4.5 Genetic and related effects of DDT in non-human mammals in vitro

| Species | Tissue, cell line | End-point | Test | Results | | Dose | Comments | Reference |
|-----------------|------------------------|------------|--------------------------------|------------------------------|---------------------------|--------------------|----------|--------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| Rat | Primary hepatocytes | DNA damage | Alkaline elution | – | NT | 106 µg/mL | DDT | Sina et al. (1983) |
| Rat | Primary hepatocytes | DNA damage | UDS assay | – | NT | 35 µg/mL | DDT | Probst et al. (1981) |
| Rat | Primary hepatocytes | DNA damage | UDS assay | – | NT | 35 µg/mL | DDT | Maslansky & Williams (1981) |
| Mouse | Primary hepatocytes | DNA damage | UDS assay | – | NT | 35 µg/mL | DDT | Maslansky & Williams (1981) |
| Mouse, Balb/c | Primary hepatocytes | DNA damage | UDS assay | – | NT | 35 µg/mL | p,p'-DDT | Klaunig et al. (1984) |
| Chinese hamster | Primary hepatocytes | DNA damage | UDS assay | – | NT | 35 µg/mL | DDT | Maslansky & Williams (1981) |
| Chinese hamster | V79 cells | DNA damage | Alkaline elution | – | – | 354 µg/mL | DDT | Swenberg et al. (1976) |
| Chinese hamster | V79 cells | DNA damage | Alkaline elution | – | – | 1060 µg/mL | DDT | Swenberg (1981) |
| Rat | Liver epithelial cells | Mutation | ARL/HGPRT assay | – | NT | NR | DDT | Telang et al. (1981) |
| Rat, Fisher | Primary hepatocytes | Mutation | Host-mediated assay (in vitro) | – | NT | 10 ⁻⁴ M | DDT | Tong et al. (1981) |
| Mouse | Balb/c 3T3 fibroblasts | Mutation | Cell transformation assay | + | + | 10 µg/mL | DDT | Fitzgerald et al. (1989) |
| Mouse | Embryo cells | Mutation | Cell transformation assay | – | NT | 15 µg/mL | p,p'-DDT | Langenbach & Gingell (1975) |
| Chinese hamster | V79 cells | Mutation | Hprt mutation assay | – | NT | 14.2 µg/mL | p,p'-DDT | Tsushimoto et al. (1983) |
| Chinese hamster | V79 cells | Mutation | Hprt mutation assay | – | NT | 35 µg/mL | p,p'-DDT | Kelly-Garvert & Legator (1973) |

Table 4.5 Genetic and related effects of DDT in non-human mammals in vitro

| Species | Tissue, cell line | End-point | Test | Results | | Dose (LEC or HIC) | Comments | Reference |
|---|---------------------------------|--------------------|---|------------------------------|---------------------------|----------------------|------------------|--------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| Chinese hamster | V79 cells | Mutation | 6-thioguanine (6-TG) resistance assay | – | NT | 14.2 µg/mL | <i>p,p'</i> -DDT | Tsushima et al. (1983) |
| | | | Diphtheria toxin (<i>DT</i>) resistance assay | – | NT | 14.2 µg/mL | <i>p,p'</i> -DDT | |
| Rat, kangaroo <i>Potorous tridactylis apicalis</i> | Aneuploid established cell line | Chromosomal damage | Chromosomal aberrations | + | NT | 10 µg/mL | <i>p,p'</i> -DDT | Palmer et al. (1972) |
| | | | | + | NT | 10 µg/mL | <i>o,p'</i> -DDT | |
| Chinese hamster | V79 cells | Chromosomal damage | Chromosomal aberrations | – | NT | 45 µg/mL | <i>p,p'</i> -DDT | Kelly-Garvert & Legator (1973) |
| Chinese hamster | B14 F28 cells | Chromosomal damage | Chromosomal aberrations | + | NT | 49 µg/mL × 4h | <i>p,p'</i> -DDT | Mahr & Miltenburger (1976) |
| Rabbit, New Zealand White | Lymphocytes | Chromosomal damage | Chromosomal aberrations | – | NT | 100 µg/mL | <i>p,p'</i> -DDT | Hart et al. (1972) |
| Artic beluga whale <i>Delphinapterus leucas</i> | Skin fibroblasts | Chromosomal damage | Micronucleus formation | + | – | 10 µg/mL | <i>p,p'</i> -DDT | Gauthier et al. (1999) |

+, positive; –, negative; DDT, dichlorodiphenyltrichloroethane; HIC, highest ineffective concentration; LEC, lowest effective concentration; NT, not tested; UDS, unscheduled DNA synthesis

Table 4.6 Genetic and related effects of metabolites of DDT in non-human mammalian cells in vitro

| Species | Tissue, cell line | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|-----------------|--|------------|---|------------------------------------|---------------------------------|-------------------------------|------------------|-----------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| Rat | Primary hepatocytes | DNA damage | Alkaline elution | + | NT | 95 µg/mL | <i>p,p'</i> -DDE | Sina et al. (1983) |
| Rat | Primary hepatocytes | DNA damage | UDS assay | - | NT | 31 µg/mL | DDE | Maslansky & Williams (1981) |
| Rat | Primary hepatocytes | DNA damage | UDS assay | - | NT | 2000 µg/mL | <i>p,p'</i> -DDE | Williams et al. (1982) |
| Mouse | Primary hepatocytes | DNA damage | UDS assay | - | NT | 31 µg/mL | DDE | Maslansky & Williams (1981) |
| Mouse | Embryo cells | Mutation | Cell transformation assay | - | NT | 15 µg/mL | <i>p,p'</i> -DDE | Langenbach & Gingell (1975) |
| Mouse | L5178Y mouse lymphoma cells | Mutation | TK mutation assay | + | + | 30 µg/mL | <i>p,p'</i> -DDE | Clive et al. (1979) |
| Mouse | L5178Y mouse lymphoma cells | Mutation | TK mutation assay | + | NT | 25 µg/mL | <i>p,p'</i> -DDE | McGregor et al. (1988) |
| Mouse | L5178Y mouse lymphoma cells | Mutation | <i>Hprt</i> mutation assay | + | NT | 16 µg/mL | <i>p,p'</i> -DDE | Amacher & Zelljadt (1984) |
| Chinese hamster | Wild-type V79 cells (6TG ^s) and a 6TG-resistant clone, T2–14 cells (6TG ^r) | Mutation | Inhibition of metabolic cooperation assay | + | NT | 10 µg/mL | DDE | Kurata et al. (1982) |
| | | | | + | NT | 2.5 µg/mL | TDE | |

Table 4.6 Genetic and related effects of metabolites of DDT in non-human mammalian cells in vitro

| Species | Tissue, cell line | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|---|---------------------------------|--------------------|----------------------------|------------------------------------|---------------------------------|-------------------------------|------------------|--------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| Chinese hamster | V79 cells | Mutation | <i>Hprt</i> mutation assay | + | NT | 25 µg/mL | <i>p,p'</i> -DDE | Kelly-Garvert & Legator (1973) |
| Rat, kangaroo <i>Potorous tridactylis apicalis</i> | Aneuploid established cell line | Chromosomal damage | Chromosomal aberrations | + | NT | 10 µg/mL | <i>p,p'</i> -DDE | Palmer et al. (1972) |
| | | | | + | NT | 10 µg/mL | <i>o,p'</i> -DDE | |
| | | | | + | NT | 10 µg/mL | <i>p,p'</i> -DDD | |
| | | | | + | NT | 10 µg/mL | <i>o,p'</i> -DDD | |
| | | | | - | NT | 100 µg/mL | <i>p,p'</i> -DDA | |
| Chinese hamster | V79 cells | Chromosomal damage | Chromosomal aberrations | + | NT | 35 µg/mL | <i>p,p'</i> -DDE | Kelly-Garvert & Legator (1973) |
| Chinese hamster | CHO cells | Chromosomal damage | Chromosomal aberrations | - | - | 60 µg/mL | DDE | Galloway et al. (1987) |
| | | | Sister-chromatid exchanges | - | (+) | 5 µg/mL | DDE | |
| Chinese hamster | B14 F28 cells | Chromosomal damage | Chromosomal aberrations | + | NT | 44 µg/mL × 4 h | <i>p,p'</i> -DDE | Mahr & Miltenburger (1976) |
| | | | | + | NT | 41 µg/mL × 4 h | <i>p,p'</i> -DDA | |
| | | | | - | NT | 75 µg/mL × 4 h | <i>p,p'</i> -DDD | |

+, positive; -, negative; DDA, 2,2-bis(4-chlorophenyl)-acetic acid; DDD, dichlorodiphenyldichloroethane; DDE, dichlorodiphenylchloroethylene; DDT, dichlorodiphenyltrichloroethane; HIC, highest ineffective concentration; LEC, lowest effective concentration, NT, not tested; TDE, 1,1-dichloro-2,2-bis(*p*-chlorophenyl)ethane; UDS, unscheduled DNA synthesis

Table 4.7 Genetic and related effects of DDT in non-mammalian systems

| Phylogenetic class | Species, strain, tissue | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------|---|--------------------|-----------------------------|------------------------------|---------------------------|-----------------------------|---|-----------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| Invertebrates | Zebra mussel, <i>Dreissena polymorpha</i> | Chromosomal damage | Micronucleus formation | + | NA | 0.1 µg/L for 168 h | <i>p,p'</i> -DDT | Binelli et al. (2008b) |
| | Shrimp, <i>Litopenaeus stylirostris</i> | DNA damage | DNA adducts | + | NA | Sublethal concentration NR | Exposed to DDT for 4 days; DDT replaced daily and dead larvae removed | Galindo Reyes et al. (2002) |
| | Shrimp, <i>Litopenaeus stylirostris</i> | DNA damage | DNA strand breaks | + | NA | Sublethal concentration NR | Exposed to DDT for 4 days; DDT replaced daily and dead larvae removed | Galindo Reyes et al. (2002) |
| Insects | <i>Drosophila melanogaster</i> y/R(1)2, vf/B ⁵ Yy ⁺ , male, germ-line cells | Chromosomal damage | Chromosomal aberrations | + | NA | 1 µg, in feeding media | Formulation (<i>p,p'</i> -DDT, 80%; <i>o,p'</i> -DDT, 18%; <i>p,p'</i> -DDE, 2%) | Clark (1974) |
| | <i>Drosophila melanogaster</i> y/R(1)2, vf/B ⁵ Yy ⁺ , male, germ-line cells | Chromosomal damage | Chromosomal aberrations | - | NA | 25 ppm, in feeding media | Formulated product of DDT not specified | Woodruff et al. (1983) |
| | <i>Drosophila melanogaster</i> , male, somatic cells | Mutation | Accord insertion | + | NA | NT | DDT | Catania et al. (2004) |
| | <i>Drosophila melanogaster</i> Canton-S, male, germ-line cells | Mutation | Dominant lethal | + | NA | 1 µg, in feeding media | Formulation (<i>p,p'</i> -DDT, 80%; <i>o,p'</i> -DDT, 18%; <i>p,p'</i> -DDE, 2%) | Clark (1974) |
| | <i>Drosophila melanogaster</i> , male, germ-line cells | Mutation | Sex linked recessive lethal | ± | NA | 0.14 mM, , in feeding media | DDT | Vogel (1972) |
| | <i>Drosophila melanogaster</i> , male, germ-line cells, germ-line cells | Mutation | Sex linked recessive lethal | - | NA | 20 µg, in feeding media | <i>p,p'</i> -DDT | Pielou (1952) |

Table 4.7 Genetic and related effects of DDT in non-mammalian systems

| Phylogenetic class | Species, strain, tissue | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|---------------------------------------|--|--------------------|--------------------------------|------------------------------|---------------------------|---|---|-------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| Lower eukaryote (yeast, mould, fungi) | <i>Drosophila melanogaster</i> Canton-S, male, germ-line cells | Mutation | Sex linked recessive lethal | – | NA | 20 µg/week, in feeding media up to 8 mo | Formulation (<i>p,p'</i> -DDT, 80%; <i>o,p'</i> -DDT, 18%; <i>p,p'</i> -DDE, 2%) | Clark (1974) |
| | <i>Saccharomyces cerevisiae</i> | Mutation | Mitotic gene conversion | – | NT | NR | DDT | Fahrig (1974) |
| | <i>Saccharomyces cerevisiae</i> D61.M | Mutation | Colony formation assay | – | – | 0.140 mM | <i>p,p'</i> -DDT | Albertini et al. (1988) |
| | <i>Saccharomyces cerevisiae</i> D61.M | Chromosomal damage | Chromosomal loss assay | – | – | 0.140 mM | <i>p,p'</i> -DDT | Albertini et al. (1988) |
| | <i>Aspergillus nidulans</i> haploid strain 35 | Mutation | Forward mutation | – | NT | 2.8 mM | <i>p,p'</i> -DDT | Crebelli et al. (1986) |
| | <i>Aspergillus nidulans</i> diploid strain P1 | Chromosomal damage | Aneuploidy | – | NT | 2.8 mM | <i>p,p'</i> -DDT | Crebelli et al. (1986) |
| | <i>Neurospora crassa</i> heterokaryon 12 | Mutation | Host-mediated assay (in vivo) | – | NT | 75 mg/mL | Formulation (<i>p,p'</i> -DDT, 80%; <i>o,p'</i> -DDT, 18%; <i>p,p'</i> -DDE, 2%) | Clark (1974) |
| | <i>Neurospora crassa</i> heterokaryon 12 | Mutation | Host-mediated assay (in vitro) | ± | NT | 75 mg/mL | Formulation (<i>p,p'</i> -DDT, 80%; <i>o,p'</i> -DDT, 18%; <i>p,p'</i> -DDE, 2%) | Clark (1974) |
| | | | | | | | Inconclusive results | |
| | <i>Tetrahymena thermophila</i> Cu428 | Toxicogenomics | Gene expression | + | NA | 4×10^{-6} mol/L per 24 h | DDT | Chang et al. (2011) |

Table 4.7 Genetic and related effects of DDT in non-mammalian systems

| Phylogenetic class | Species, strain, tissue | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|-----------------------|------------------------------------|-----------|------------------|------------------------------|---------------------------|----------------------------|----------|----------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| Prokaryote (bacteria) | <i>Salmonella typhimurium</i> TA92 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDT | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 500 µg/plate | DDT | De Flora et al. (1989) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 2500 µg/plate | DDT | Byeon et al. (1976) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 170 µg/plate | DDT | Bartsch et al. (1980) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 177 µg/plate | DDT | Nishimura et al. (1982) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 370 µg/plate | DDT | Van Dijck & Van de Voorde (1976) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 500 µg/plate | DDT | Bruce & Heddle (1979) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 35 µg/plate | DDT | Probst et al. (1981) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | NT | – | 2500 µg/plate | DDT | Simmon et al. (1977) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDT | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 5000 µg/plate | p,p'-DDT | Moriya et al. (1983) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 177 µg/plate | p,p'-DDT | Planche et al. (1979) |

Table 4.7 Genetic and related effects of DDT in non-mammalian systems

| Phylogenetic class | Species, strain, tissue | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------|--------------------------------------|-----------|------------------|------------------------------|---------------------------|----------------------------|----------|----------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 35 µg/plate | DDT | Probst et al. (1981) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 370 µg/plate | DDT | Van Dijck & Van de Voorde (1976) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 177 µg/plate | DDT | Nishimura et al. (1982) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 170 µg/plate | DDT | Bartsch et al. (1980) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 500 µg/plate | DDT | De Flora et al. (1989) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 2500 µg/plate | DDT | Byeon et al. (1976) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 500 µg/plate | DDT | Bruce & Heddle (1979) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDT | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | NT | – | 2500 µg/plate | DDT | Simmon et al. (1977) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 5000 µg/plate | p,p'-DDT | Moriya et al. (1983) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 177 µg/plate | p,p'-DDT | Planche et al. (1979) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 35 µg/plate | DDT | Probst et al. (1981) |

Table 4.7 Genetic and related effects of DDT in non-mammalian systems

| Phylogenetic class | Species, strain, tissue | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------|--------------------------------------|-----------|------------------|------------------------------|---------------------------|----------------------------|----------|------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 2500 µg/plate | DDT | Byeon et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 2500 µg/plate | DDT | Marshall et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 500 µg/plate | DDT | Bruce & Heddle (1979) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDT | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | NT | – | 2500 µg/plate | DDT | Simmon et al. (1977) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 5000 µg/plate | p,p'-DDT | Moriya et al. (1983) |
| | <i>Salmonella typhimurium</i> TA1536 | Mutation | Reverse mutation | – | – | 2500 µg/plate | DDT | Marshall et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | – | – | 35 µg/plate | DDT | Probst et al. (1981) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | – | – | 2500 µg/plate | DDT | Marshall et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | – | – | 500 µg/plate | DDT | Bruce & Heddle (1979) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDT | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | NT | – | 2500 µg/plate | DDT | Simmon et al. (1977) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | – | – | 5000 µg/plate | p,p'-DDT | Moriya et al. (1983) |

Table 4.7 Genetic and related effects of DDT in non-mammalian systems

| Phylogenetic class | Species, strain, tissue | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------|--------------------------------------|------------|------------------|------------------------------|---------------------------|----------------------------|----------|----------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | – | – | 2500 µg/plate | DDT | Byeon et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | – | – | 35 µg/plate | DDT | Probst et al. (1981) |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | – | – | 2500 µg/plate | DDT | Marshall et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | NT | – | 2500 µg/plate | DDT | Simmon et al. (1977) |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDT | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | – | – | 5000 µg/plate | p,p'-DDT | Moriya et al. (1983) |
| | <i>Salmonella typhimurium</i> TA1978 | Mutation | Reverse mutation | – | – | 370 µg/plate | DDT | Van Dijck & Van de Voorde (1976) |
| | <i>Salmonella typhimurium</i> C3076 | Mutation | Reverse mutation | – | – | 35 µg/plate | DDT | Probst et al. (1981) |
| | <i>Salmonella typhimurium</i> D3052 | Mutation | Reverse mutation | – | – | 35 µg/plate | DDT | Probst et al. (1981) |
| | <i>Salmonella typhimurium</i> G46 | Mutation | Reverse mutation | – | – | 35 µg/plate | DDT | Probst et al. (1981) |
| | <i>Escherichia coli</i> Q-13 | DNA damage | DNA-cell-binding | NT | – | 150 µM | DDT | Kubinski et al. (1981) |
| | <i>Escherichia coli</i> PQ37 | Mutation | SOS chromotest | – | – | 1000 µg/mL | DDT | Dayan et al. (1987) |

Table 4.7 Genetic and related effects of DDT in non-mammalian systems

| Phylogenetic class | Species, strain, tissue | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------|---|------------|--|------------------------------|---------------------------|----------------------------|------------------|--------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| | <i>Escherichia coli</i> WP2 <i>uvrA</i> | Mutation | Reverse mutation | – | – | 35 µg/plate | DDT | Probst et al. (1981) |
| | <i>Escherichia coli</i> WP2 <i>uvrA</i> | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDT | Glatt & Oesch (1987) |
| | <i>Escherichia coli</i> WP2 <i>hcr</i> | Mutation | Reverse mutation | – | – | 5000 µg/plate | <i>p,p'</i> -DDT | Moriya et al. (1983) |
| | <i>Escherichia coli</i> | Mutation | Reverse mutation Pol-A mutation | – | – | 500 µg/plate | <i>p,p'</i> -DDT | Fluck et al. (1976) |
| | <i>Bacillus subtilis</i> M 45 Rec [–] , H17 Rec ⁺ | Mutation | Rec assay | – | NT | NR | DDT | Shirasu et al. (1976) |
| | <i>Serratia marcescens</i> | Mutation | Host-mediated assay (in vivo) | – | NT | NR | DDT | Buselmaier et al. (1973) |
| Acellular systems | Isolated DNA from ColE1 plasmid | DNA damage | DNA single-strand breaks/alkaline labile sites | – | NT | 100 µg/mL | DDT | Griffin & Hill (1978) |

+, positive; –, negative; ±, equivocal (variable response in several experiments within an adequate study); DDE, dichlorodiphenyl dichloroethylene; DDT, dichlorodiphenyltrichloroethane; HIC, highest ineffective concentration; LEC, lowest effective concentration; NA, not applicable; NR, not reported; NT, not tested

Table 4.8 Genetic and related effects of metabolites of DDT in non-mammalian systems in vitro

| Phylogenetic class | Species, strain | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------------|--|-----------|------------------|------------------------------------|---------------------------------|-------------------------------|------------------|-------------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| Prokaryote (bacteria) | <i>Salmonella typhimurium</i> TA92 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDE | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDE | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 500 µg/plate | <i>p,p'</i> -DDE | Mortelmans et al. (1986) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 5000 µg/plate | <i>p,p'</i> -DDE | De Flora (1981) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 860 µg/plate | <i>p,p'</i> -DDE | Bartsch et al. (1980) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 500 µg/plate | <i>p,p'</i> -DDE | De Flora et al. (1989) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 2500 µg/plate | <i>p,p'</i> -DDE | Brams et al. (1987) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 5000 µg/plate | <i>p,p'</i> -DDE | Moriya et al. (1983) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | – | 370 µg/plate | <i>p,p'</i> -DDE | Van Dijck & Van de Voorde (1976) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDE | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 500 µg/plate | <i>p,p'</i> -DDE | Mortelmans et al. (1986) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 5000 µg/plate | <i>p,p'</i> -DDE | De Flora (1981) |

Table 4.8 Genetic and related effects of metabolites of DDT in non-mammalian systems in vitro

| Phylogenetic class | Species, strain | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------|---|-----------|------------------|------------------------------------|---------------------------------|-------------------------------|------------------|-------------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 860 µg/plate | <i>p,p'</i> -DDE | Bartsch et al. (1980) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 500 µg/plate | <i>p,p'</i> -DDE | De Flora et al. (1989) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 2500 µg/plate | <i>p,p'</i> -DDE | Brams et al. (1987) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 5000 µg/plate | <i>p,p'</i> -DDE | Moriya et al. (1983) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | – | 370 µg/plate | <i>p,p'</i> -DDE | Van Dijck & Van de Voorde (1976) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | NT | – | 2500 µg/plate | <i>p,p'</i> -DDE | McCann et al. (1975) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 1000 µg/plate | DDE | Marshall et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDE | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 500 µg/plate | <i>p,p'</i> -DDE | Mortelmanns et al. (1986) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 5000 µg/plate | <i>p,p'</i> -DDE | De Flora (1981) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 2500 µg/plate | <i>p,p'</i> -DDE | Brams et al. (1987) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 5000 µg/plate | <i>p,p'</i> -DDE | Moriya et al. (1983) |
| | <i>Salmonella typhimurium</i> TA1535 | Mutation | Reverse mutation | – | – | 370 µg/plate | <i>p,p'</i> -DDE | Van Dijck & Van de Voorde (1976) |

Table 4.8 Genetic and related effects of metabolites of DDT in non-mammalian systems in vitro

| Phylogenetic class | Species, strain | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------|--|-----------|------------------|------------------------------------|---------------------------------|-------------------------------|----------|-------------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| | <i>Salmonella typhimurium</i> TA1536 | Mutation | Reverse mutation | – | – | 1000 µg/plate | DDE | Marshall et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | – | – | 1000 µg/plate | DDE | Marshall et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDE | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | – | – | 500 µg/plate | p,p'-DDE | Mortelmans et al. (1986) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | – | – | 5000 µg/plate | p,p'-DDE | De Flora (1981) |
| | <i>Salmonella typhimurium</i> TA 1537 | Mutation | Reverse mutation | – | – | 2500 µg/plate | p,p'-DDE | Brams et al. (1987) |
| | <i>Salmonella typhimurium</i> TA 1537 | Mutation | Reverse mutation | – | – | 5000 µg/plate | p,p'-DDE | Moriya et al. (1983) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | – | – | 370 µg/plate | p,p'-DDE | Van Dijck & Van de Voorde (1976) |
| | <i>Salmonella typhimurium</i> TA1537 | Mutation | Reverse mutation | NT | – | 2500 µg/plate | p,p'-DDE | McCann et al. (1975) |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | – | – | 1000 µg/plate | DDE | Marshall et al. (1976) |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDE | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | – | – | 5000 µg/plate | p,p'-DDE | De Flora (1981) |
| | <i>Salmonella typhimurium</i> TA 1538 | Mutation | Reverse mutation | – | – | 5000 µg/plate | p,p'-DDE | Moriya et al. (1983) |

Table 4.8 Genetic and related effects of metabolites of DDT in non-mammalian systems in vitro

| Phylogenetic class | Species, strain | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------|--|-----------|------------------|------------------------------------|---------------------------------|-------------------------------|--|----------------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | – | – | 370 µg/plate | <i>p,p'</i> -DDE | Van Dijck & Van de Voorde (1976) |
| | <i>Salmonella typhimurium</i> TA1538 | Mutation | Reverse mutation | NT | – | 2500 µg/plate | <i>p,p'</i> -DDE | McCann et al. (1975) |
| | <i>Salmonella typhimurium</i> TA1950 | Mutation | Reverse mutation | – | – | 370 µg/plate | <i>p,p'</i> -DDE | Van Dijck & Van de Voorde (1976) |
| | <i>Salmonella typhimurium</i> TA1978 | Mutation | Reverse mutation | – | – | 370 µg/plate | <i>p,p'</i> -DDE | Van Dijck & Van de Voorde (1976) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | – | NT | 0.10 µg/plate | Metabolite 2,2-bis(<i>p</i> -chlorophenyl)-2-chloroacetaldehyde (α C1-DDCHO) | Gold et al. (1981) |
| | <i>Salmonella typhimurium</i> TA98 | Mutation | Reverse mutation | – | + | 3.15 µg/plate | Metabolite 1,1-bis(<i>p</i> -chlorophenyl)-2,2-dichloroethane was mutagenic when norharman was added to the metabolizing system | Glatt & Oesch (1987) |
| | <i>Salmonella typhimurium</i> TA100 | Mutation | Reverse mutation | + | NT | 0.025 µg/plate | Metabolite 1-Chloro-2,2-bis(<i>p</i> -chlorophenyl)ethene (DDMU)-epoxide | Gold et al. (1981) |
| | <i>Escherichia coli</i> WP2 <i>uvrA</i> | Mutation | Reverse mutation | NT | – | 1000 µg/plate | DDE | Glatt & Oesch (1987) |
| | <i>Escherichia coli</i> WP2 <i>uvrA</i> | Mutation | Reverse mutation | NT | – | 1000 µg/plate | <i>p,p'</i> -DDE | Mamber et al. (1984) |

Table 4.8 Genetic and related effects of metabolites of DDT in non-mammalian systems in vitro

| Phylogenetic class | Species, strain | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|---|---|--------------------|--------------------------------------|------------------------------------|---------------------------------|-------------------------------|------------------|---|
| | | | | Without metabolic activation | With metabolic activation | | | |
| | <i>Escherichia coli</i> WP2 <i>her</i> | Mutation | Reverse mutation | – | – | 5000 µg/plate | <i>p,p'</i> -DDE | Moriya et al. (1983) |
| | <i>Escherichia coli</i> Pol-A | Mutation | Reverse mutation | – | – | 500 µg/plate | <i>p,p'</i> -DDE | Fluck et al. (1976) |
| | <i>Escherichia coli</i> Pol-A | Mutation | Reverse mutation | – | – | 500 µg/plate | <i>p,p'</i> -DDD | Fluck et al. (1976) |
| | <i>Serratia marcescens</i> a21 <i>Leu</i> ^r | Mutation | Reverse mutation | + | NT | NR | DDD | Buselmaier et al. (1973) |
| | <i>Serratia marcescens</i> a31 <i>His</i> ^r | Mutation | Reverse mutation | + | NT | NR | DDD | Buselmaier et al. (1973) |
| | <i>Serratia marcescens</i> | Mutation | Host-mediated assay (in vivo) | – | NT | NR | DDE | Buselmaier et al. (1973) |
| | <i>Serratia marcescens</i> | Mutation | Host-mediated assay (in vivo) | – | NT | NR | DDA | Buselmaier et al. (1973) |
| Lower eukaryote (yeast, mould, fungi) | <i>Saccharomyces cerevisiae</i> | Mutation | Mitotic gene conversion | – | NT | NR | DDE | Fahrig (1974) |
| | <i>Saccharomyces cerevisiae</i> | Mutation | Mitotic gene conversion | – | NT | NR | DDD | Fahrig (1974) |
| | <i>Saccharomyces cerevisiae</i> | Mutation | Mitotic gene conversion | – | NT | NR | DDA | Fahrig (1974) |
| | <i>Saccharomyces cerevisiae</i> RS112 | Chromosomal damage | Intrachromosomal recombination assay | + | – | 100 µg/mL | <i>p,p'</i> -DDE | Schiestl (1989); Schiestl et al. (1989) |
| Invertebrates | Zebra mussel <i>Dreissena polymorpha</i> | DNA damage | Comet assay | + | NA | 0.1 µg/L for 168 h | <i>p,p'</i> -DDE | Binelli et al. (2008a) |

Table 4.8 Genetic and related effects of metabolites of DDT in non-mammalian systems in vitro

| Phylogenetic class | Species, strain | End-point | Test | Results | | Concentration (LEC or HIC) | Comments | Reference |
|--------------------|--|--------------------|-------------------------------|------------------------------------|---------------------------------|--------------------------------|------------------|---------------------------|
| | | | | Without metabolic activation | With metabolic activation | | | |
| Insects | Zebra mussel <i>Dreissena polymorpha</i> | Chromosomal damage | Micronucleus formation | + | NA | 0.1 µg/L for 168 h | <i>p,p'</i> -DDE | Binelli et al. (2008a, b) |
| | Zebra mussel <i>Dreissena polymorpha</i> | Chromosomal damage | Micronucleus formation | + | NA | 0.1 µg/L for 168 hours | <i>p,p'</i> -DDD | Binelli et al. (2008b) |
| | <i>Drosophila melanogaster</i> Canton-S, male, germ-line cells | Mutation | Sex-linked recessive lethal | + | NA | 10 000 µg/mL, in feeding media | <i>p,p'</i> -DDE | Valencia et al. (1985) |
| | <i>Drosophila melanogaster</i> , male, germ-line cells | Mutation | Sex-linked recessive lethal | + | NA | 3.4 mM, in feeding media | DDA | Vogel (1972) |
| | <i>Drosophila melanogaster</i> , male, germ-line cells | Mutation | Sex-linked recessive lethal | - | NA | 3.1 mM, in feeding media | DDE | Vogel (1972) |
| | <i>Drosophila melanogaster</i> , male, germ-line cells | Mutation | Sex-linked recessive lethal | - | NA | 0.3 mM, in feeding media | DDD | Vogel (1972) |
| | <i>Drosophila melanogaster</i> , male, germ-line cells | Mutation | Sex-linked recessive lethal | - | NA | 3.7 mM, in feeding media | DDOM | Vogel (1972) |
| | <i>Drosophila melanogaster</i> Canton-S, male, germ-line cells | Chromosomal damage | Heritable translocation assay | - | NA | 10 000 µg/mL, in feeding media | <i>p,p'</i> -DDE | Valencia et al. (1985) |

+, positive; -, negative; DDA, 2,2,-bis(4-chlorophenyl)-acetic acid; DDD, trichlorodiphenyldichloroethane; DDE, dichlorodiphenyldichloroethylene; DDOM, 2,2-bis(*p*-chlorophenyl-ethane); DDT, dichlorodiphenyltrichloroethane; HIC, highest ineffective concentration; LEC, lowest effective concentration; NA, not applicable; NR, not reported; NT, not tested