

Original Manuscript

Extending (Q)SARs to incorporate proprietary knowledge for regulatory purposes: is aromatic *N*-oxide a structural alert for predicting DNA-reactive mutagenicity?*

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Abstract

(Quantitative) structure–activity relationship or (Q)SAR predictions of DNA-reactive mutagenicity are important to support both the design of new chemicals and the assessment of impurities, degradants, metabolites, extractables and leachables, as well as existing chemicals. Aromatic *N*-oxides represent a class of compounds that are often considered alerting for mutagenicity yet the scientific rationale of this structural alert is not clear and has been questioned. Because aromatic *N*-oxide-containing compounds may be encountered as impurities, degradants and metabolites,

it is important to accurately predict mutagenicity of this chemical class. This article analysed a series of publicly available aromatic *N*-oxide data in search of supporting information. The article also used a previously developed structure–activity relationship (SAR) fingerprint methodology where a series of aromatic *N*-oxide substructures was generated and matched against public and proprietary databases, including pharmaceutical data. An assessment of the number of mutagenic and non-mutagenic compounds matching each substructure across all sources was used to understand whether the general class or any specific subclasses appear to lead to mutagenicity. This analysis resulted in a downgrade of the general aromatic *N*-oxide alert. However, it was determined there were enough public and proprietary data to assign the quindioxin and related chemicals as well as benzo[*c*][1,2,5]oxadiazole 1-oxide subclasses as alerts. The overall results of this analysis were incorporated into Leadscope’s expert-rule-based model to enhance its predictive accuracy.

Introduction

The use of computational or (quantitative) structure–activity relationship [(Q)SAR] methodologies to predict DNA-reactive mutagenicity is increasing due in part to the International Council for Harmonisation (ICH) M7 guideline (Assessment and control of DNA reactive (mutagenic) impurities in pharmaceuticals to limit potential carcinogenic risk) (1). This guideline recommends the use of two complementary (Q)SAR methodologies to predict the results of the reverse bacterial mutagenicity assay [Organisation for Economic Co-operation and Development (OECD) 1997] when no or inadequate experimental mutagenicity data are available for the impurity.

One of these (Q)SAR methodologies is referred to as ‘expert rule-based’ (1,2). This is a computational system that uses a series of structural rules or alerts to predict mutagenicity. When no alerts are present in the test chemical, it is possible to conclude that the compound is non-mutagenic if other factors are satisfied, such as the test chemical being within the model’s applicability domain (3–5) of the alert system, i.e. the region of chemical space where a model makes reliable predictions. Such structural alerts should ideally be based on an understanding of known mechanism(s) of mutagenicity. In addition, it is essential as part of the development of such systems that the alerts are validated based on a sufficient number of relevant chemicals to conclude that the class has an increased likelihood of being mutagenic. As part of this validation process, mutagenic compounds containing the alert may need to be removed from consideration if their mutagenicity can be explained by other structural elements. There should also be a sufficient number of clear examples (defined as at least four clear examples) because anecdotal mutagenic examples may be explained by experimental artefacts or other reasons (6).

The second recommended methodology is referred to as ‘statistical-based’ (1,2). This is based on a mathematical model calculated from a training set containing both mutagenic and non-mutagenic examples. Fragment-based descriptors (e.g. the presence or absence of specific functional groups) are often used to describe the training set examples. Models that use knowledge of DNA-reactive chemical moieties (e.g. structural alerts for mutagenicity from the literature) enhance predictivity and are easier to explain; therefore, it is important to consider such knowledge in the selection of these structural descriptors (6).

One of the first series of structural alerts for mutagenicity, known as the Ashby Tennant alerts, was proposed in the late 1980s (7,8). These expert-rule-based alerts were identified based on an analysis of mutagenic compounds and an understanding of electrophilic theory (9). Ashby and Tennant proposed 18 structural alerts, which were illustrated as part of a large supermolecule in their original publication. Figure 1 shows a portion of this supermolecule to highlight the inclusion of ‘(d) aromatic ring *N*-oxides’ as one of the proposed

alerts. These alerts are based on the potential for entering into an electrophilic reaction with DNA. However, Ashby and Tennant (7) do not discuss any specific mechanistic basis for this alert nor provide any examples of mutagenic aromatic *N*-oxide compounds.

Major structures units ... which led to a chemical being classified as structure-activity positive ... (b) aromatic nitro groups; (c) aromatic azo group ...; (d) **aromatic ring *N*-oxides**; (e) aromatic mono- and dialkylamino groups ... (emphasis [bold] added)

Snodin and McCrossen (10) reviewed available information supporting the use of aromatic *N*-oxides as a DNA-reactive mutagenicity alert and concluded that there is a lack of adequate information to support their use as alerts. However, many current expert-rule-based computational systems use the Ashby–Tennant alerts as a starting point in their development. For example, the aromatic *N*-oxide alert has also been incorporated into the collection of structural alerts described by Benigni *et al.* (11); however, the article indicates that no clear mechanism for the aromatic ring *N*-oxide alert has been identified [Plošnik *et al.* (12) reviews aromatic *N*-oxides and describes the mechanism as redox cycling (p. 176) and assigns the class under the SnAr alert category (p. 177); however, the references they cite have no discussion on aromatic *N*-oxides.]. The article also provides *N*-nitrosornicotine-1-*N*-oxide (CAS 78246-24-9) as an example; however, this is inappropriate because the chemical also contains the well-known DNA reactive alert *N*-nitroso. Separately, the OECD report (13) states that no definitive mechanism can be identified for the aromatic ring *N*-oxide alert. In addition, Galloway *et al.* (14) discusses the link between the aromatic *N*-oxide from the Ashby & Tennant publications (7,8) to the Derek alert ‘Aromatic *N*-oxide or *N*-hydroxy Tautomer’ as part of a comparison of the Ashby alerts to the Derek categories.

An assessment of impurities, metabolites, extractable and leachables showed that many contain aromatic *N*-oxides (15). For example, aromatic *N*-oxide-containing compounds are used as reagents (16).

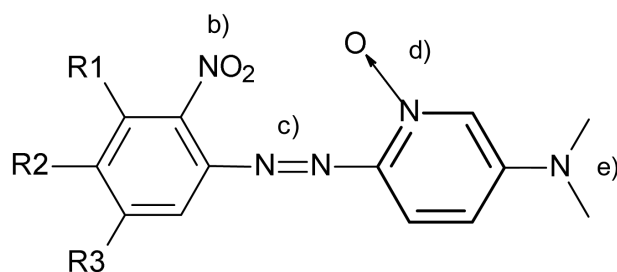


Fig. 1. Portion of the AshbyTennant supermolecular listing mutagenicity alerts.

Pharmaceutical degradation products and metabolites may also contain aromatic N-oxides (17). In addition, there are examples where aromatic N-oxides are functional groups in active ingredients (18). Furthermore, a retrospective analysis of 1440 ICH M7-compliant (Q)SAR assessments conducted on drug impurities by Food and Drug Administration (FDA)/Center for Drug Evaluation and Research (CDER) found that 24 contained aromatic N-oxides (FDA/CDER, unpublished results). It is therefore important to know whether an aromatic N-oxide or any specific subclass demonstrates a correlation with observed mutagenicity and can serve as an alert for predicting DNA-reactive mutagenicity. As recommended by the ICH M7 guideline, if a compound is predicted to be mutagenic by at least one of the (Q)SAR methodologies, a follow-on

reverse bacterial mutagenicity test may be performed or it may be necessary to control the amount of the impurity in the final product to levels defined by the guideline. This additional work can be both time consuming and costly as alternative synthetic routes may need to be explored. In the case of metabolites, difficult to synthesise, or unstable compounds, it may not be feasible to obtain enough material to test. The ICH M7 guideline also states that, if warranted, an expert review of the (Q)SAR results may be conducted. Such a review may refute any positive, out-of-domain or equivocal results with sufficient supporting evidence (19–21). In addition, such expert review may be supported by publications, such as the current publication, that have analysed the evidence for any DNA-reactive structural alerts.

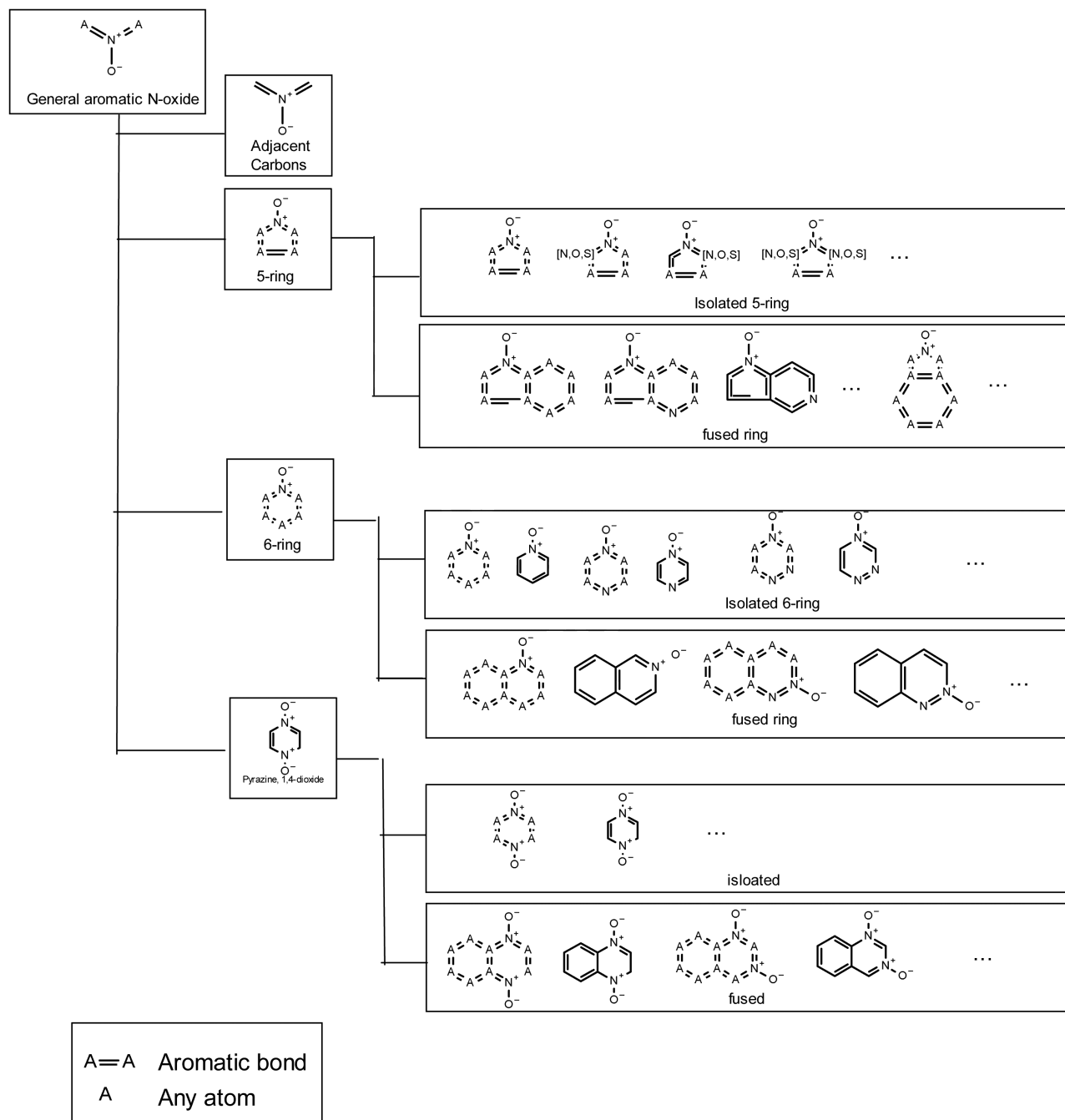


Fig. 2. Hierarchical organisation of selected substructures incorporated into the fingerprint.

The following study analyses available public and proprietary databases to determine whether the aromatic *N*-oxide or any of its subclasses are structural alerts for predicting DNA-reactive mutagenicity.

Methods

Analysis of public data

An initial analysis of public data was performed to understand whether aromatic *N*-oxides or any subclasses are alerts for predicting mutagenicity. This can be broken down into two questions:

- (i) Are there any Ames-positive compounds that can be definitely attributed to the presence of a general aromatic *N*-oxide alert (i.e. there are no other structural alerts most likely responsible for the positive mutagenic experimental call)?
- (ii) Are there experimentally Ames-negative examples of aromatic *N*-oxides?

The reference database from the Leadscope Genetox Expert Alerts version 4 (22), PharmaPendium® (trademark of Reed Elsevier Properties SA, used under license) (23) and PubMed (24) were searched as part of this analysis. The Leadscope reference database includes over 10 600 structured chemicals with detailed bacterial mutagenicity experimental data (6,22). The Leadscope Genetox Expert Alerts version 4 (System: Leadscope Model Applier v2.2.1.1) was used to understand whether any additional alerts for mutagenicity are also present in aromatic *N*-oxide compounds that may explain their mutagenicity. This system is suitable for this task because there are currently 224 alerts for mutagenicity in the Leadscope Genetox Expert Alerts version 4, derived from the literature and from analysis of public and proprietary databases (6,25) and the alerts have been validated using multiple public and proprietary databases covering over 35 000 chemicals. The alerts are annotated with information on the biological mechanisms and, wherever possible, indicate the bacterial tester strains that are sensitive for each alert (26).

Structure–activity relationship fingerprint analysis of public and proprietary data

An analysis of public and proprietary data was performed to see if there was any further evidence, particularly from proprietary data collections, to determine whether aromatic *N*-oxides or any subclasses correlate with observed mutagenicity and hence would serve as alerts for predicting mutagenicity. This analysis used a series of predefined aromatic *N*-oxide substructure searches, as illustrated in Figure 2. This analysis is referred to as structure–activity relationship (SAR) fingerprints (6). These 101 substructure searches are organised hierarchically and are specifically designed to explore the chemical space of aromatic *N*-oxide compounds in order to understand any SARs. A general aromatic *N*-oxide substructure was included to help understand the significance of the general class. A series of isolated 5- and 6-membered rings were included (e.g. pyridinium, 1-oxide-; pyridazinium, 1-oxide-; pyrimidinium, 1-oxide- and pyrazinium, 1-oxide-). These substructures helped to understand the significance of single unfused 5- and 6-membered rings containing an aromatic *N*-oxide, and if the position of any heteroatoms relative to the *N*-oxide was significant. A series of fused aromatic substructures containing an *N*-oxide were included (e.g. quinolinium, 1-oxide-; isoquinolinium, 2-oxide-; quinoxalinium, 1-oxide-; quinazolinium, 3-oxide- and quinazolinium, 1-oxide-). Dioxin-related substructures were also included (e.g. dioxin, quindioxin, quinazolium and quidioxin).

The SAR fingerprint analysis was used to assess the significance of the aromatic *N*-oxide alert and any subclasses (6). As part of this analysis, all 101 substructures (a subset is illustrated in Figure 2) were matched against a series of public and proprietary databases containing chemical structures with bacterial mutagenicity experimental data. This included the Leadscope Genetox Expert Alerts version 4 reference set, the National Institute of Health Sciences (Tokyo, Japan) databases (27) and a series of proprietary databases maintained by GlaxoSmithKline, Sanofi, Pfizer, Bristol-Myers Squibb and Bayer AG, Pharmaceuticals Division. These tests included both full GLP Ames tests as well as limited strain tests (28). The number of mutagenic (positive) and non-mutagenic (negative) examples from each database matching these substructures was identified using

Substructure	Public negative count	Public positive count	Confidential	Total negative	Total positive	Percentage positive	Binomial score	Assignment
All chemicals	10,047	5,287		27,453	7,984	22.5%	-	No assignment
Substructure A	20	35		158	58	26.9%	0.94	No assignment
Substructure B	15	6		116	10	7.9%	1.3 \times 10 ⁵	Deactivating
Substructure C	0	5		0	16	100%	1	Activating

Fig. 3. Searching the list of substructures over multiple databases (it should be noted that substructure A, substructure B and substructure C are used to illustrate different predefined substructure queries that are incorporated into the SAR fingerprint).

the Leadscope software (29). The number of positive and negative examples across all sources was then summed, and a binomial score (calculated using Microsoft™ Excel) was generated to help understand the statistical significance of the individual chemical classes (6). This analysis uses the percentage of positives across all compounds as a baseline. Subsets containing a significantly greater percentage of positive compounds were designated as activating and those with significantly lower percentage of positive compounds were designated as deactivating. This process is shown conceptually in Figure 3. As part of this exercise, no proprietary information about the chemical structures or the corresponding data were exchanged. Only counts of the number of positive and negative examples per substructure were exchanged.

Results

Analysis of public data

Forty-one relevant mutagenic chemicals were identified from public sources, as shown in Figures 4–12. The chemicals were analysed

using the Leadscope Genetox Expert Alerts version 4 and all contained an alert for mutagenicity. Table 1 lists all the data corresponding to these 41 compounds.

Figure 4 shows a series of isolated and fused mutagenic chemicals containing an aromatic N-oxide that also contain a separate mutagenic structural alert. Compounds 1, 2 and 3 contain an aromatic nitro and compound 4 contains a N-hydroxy aromatic amine.

Figure 5 shows two mutagenic aromatic N-oxide chemicals; however, both contain at least one additional structural alert. Compounds 5 and 6 contain a dibenzofuran alert and compound 5 also contains a strong activating alkyl methyl halide. Similarly, Figure 6 presents three mutagenic aromatic N-oxide chemicals; however, they all contain a polycyclic aromatic system and compounds 7 and 8 also contain an aromatic nitro.

Figure 7 contains a series of mutagenic N-oxides; however, all contain a primary aromatic amine that is alerting in the Leadscope expert alerts. The Leadscope alerts for primary aromatic amines take into consideration many factors that may activate and deactivate aromatic amine mutagenicity, including the position of

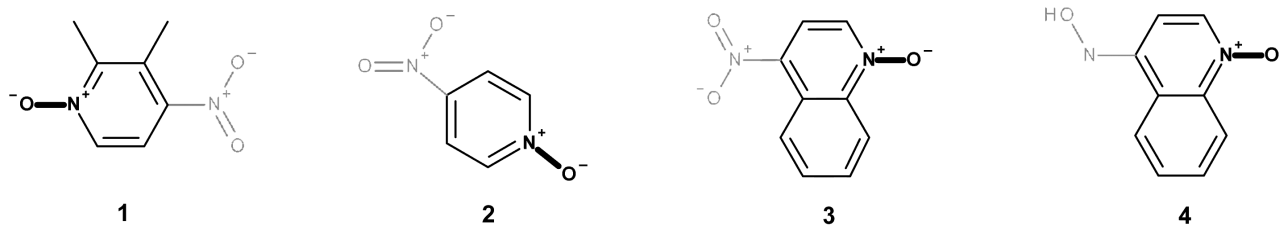


Fig. 4. Mutagenic isolated and fused examples containing both an aromatic N-oxide (highlighted in bold) and one or more structural alerts (highlighted in grey).

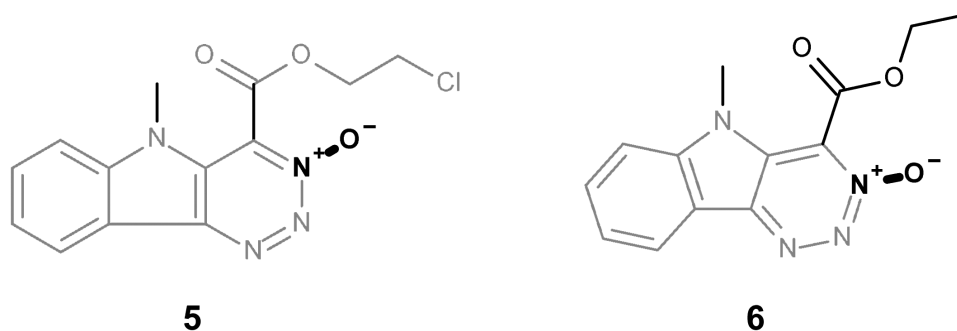


Fig. 5. Mutagenic examples containing both an aromatic N-oxide (highlighted in bold) and one or more structural alerts (highlighted in grey).

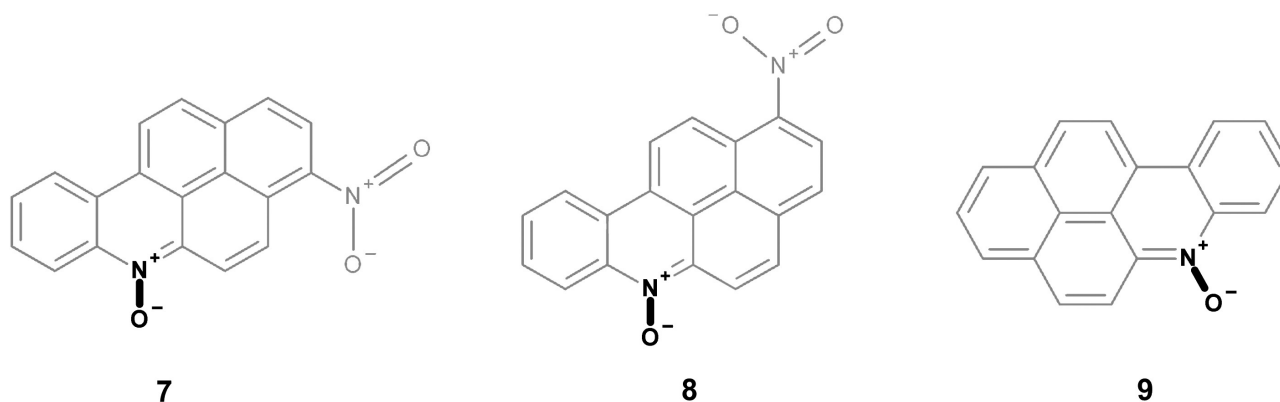


Fig. 6. Mutagenic examples containing both an aromatic N-oxide (highlighted in bold) and a polycyclic aromatic system (highlighted in grey).

heteroatoms and the position of the amine proximate to any fused ring (6).

Figures 8 and 9 present a series of mutagenic quindioxins [this class is often referred to as quindioxins; however, the article will refer to it as quindioxin to be consistent with its citation in Enoch and Cronin (30)]. It should be noted that quindioxin is an alert within the Leadscope expert alerts v4; however, the examples shown in Figure 8 contain an additional alert (aryl methylene semicarbazones). Figure 9 shows three additional examples with no additional structural alert (based on the Leadscope expert alerts v4).

Figures 10 and 11 contain a series of mutagenic benzo[*c*][1,2,5]oxadiazole 1-oxides, which are alerting in the current Leadscope

expert alerts. In Figure 10, all examples contain an aromatic nitro alert with chemical 32 also containing an aziridine. Although chemicals 34 and 35 are considered Michael acceptors, no chemicals in Figure 11 contain other alerts based on the Leadscope Expert Alerts system v4.

Figure 12 shows two mutagenic examples containing 2,7-naphthyridine 2-oxides, which are alerting in the current Leadscope expert alerts v4 but marked as indeterminate due to the limited number of examples.

A number of diverse non-mutagenic chemicals (Figure 13 and Table 2) that contain an aromatic *N*-oxide were identified. Many of these chemicals were tested in four or more of the standard bacterial

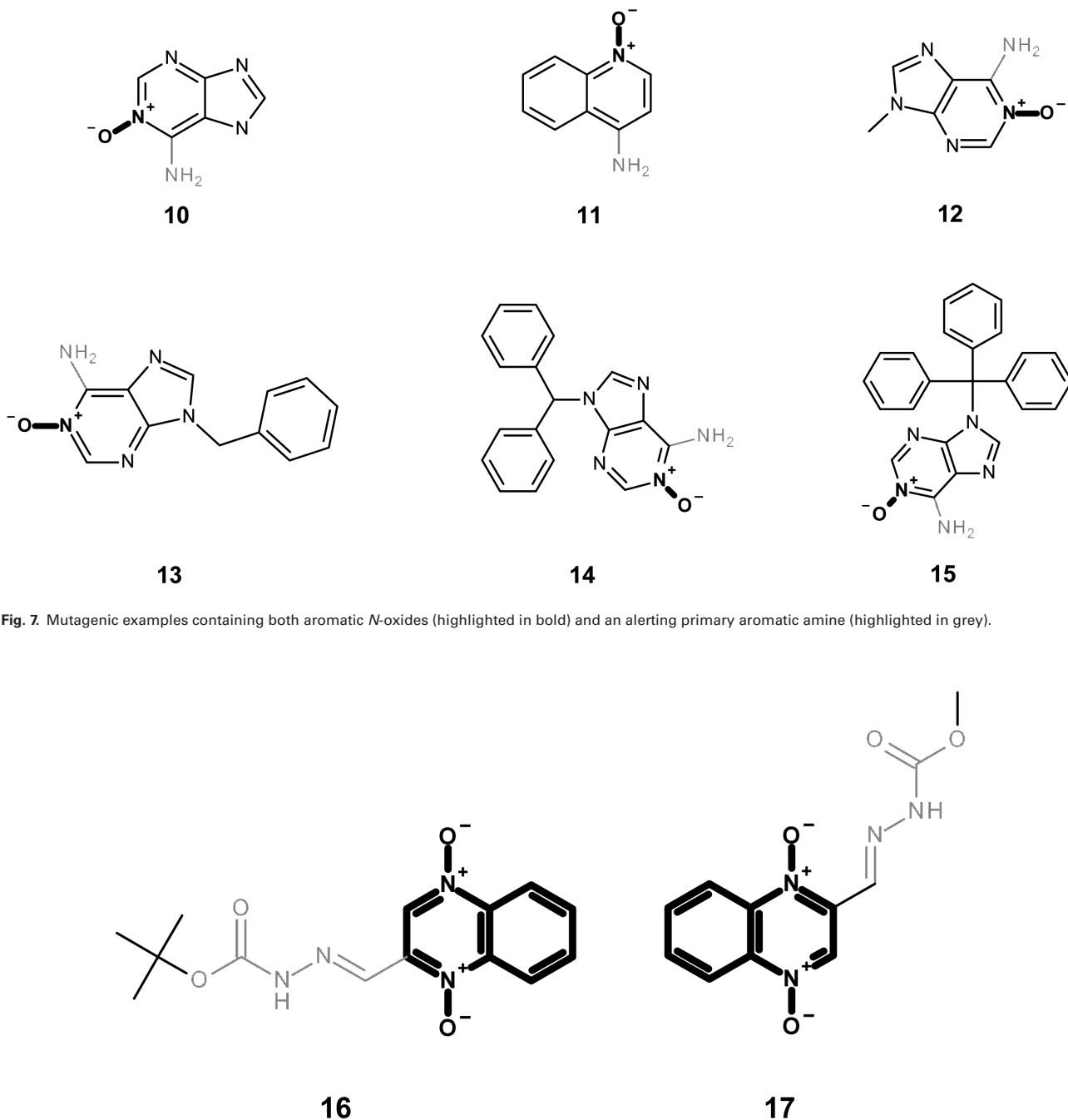


Fig. 8. Mutagenic examples containing a quindioxin (highlighted in bold) and an alerting aryl methylene semicarbazones (highlighted in grey).

tester strains with and without metabolic activation or were part of a submission to a regulatory authority. In addition, all chemicals were predicted to be non-mutagenic using the Leadscape Genetox Expert Alerts version 4 except **50** (although the aromatic nitro alert was deactivated because of the adjacent hydroxyl groups). Minoxidil **42** is a vasodilator and over-the-counter treatment for hair loss in products such as Rogaine® from Johnson and Johnson with negative Ames and rodent carcinogenicity data (48). In addition, *N*-oxides are formed as metabolites of the parent substances. Cotinine *N*-oxide **43** is a metabolite of nicotine in human urine (49). Pyridine *N*-oxide **44** is a significant mammalian metabolite of pyridine, which is detected at around one-third of the urinary radioactivity in human volunteers administered [¹⁴C]pyridine (50). Olbetam **46** is a lipid-lowering agent that is reported as negative for bacterial mutagenicity (51). Both sorafenib and roflumilast form *N*-oxide metabolites (48,49) in patients (52,53). Opicapone **50** is a catechol-*O*-methyltransferase inhibitor for patients with

Parkinson's disease with negative Ames and rodent carcinogenicity data (54), and pyrithione zinc **51** is the active ingredient in shampoos such as Head & Shoulders® and is Ames negative.

Analysis of public and proprietary data using the SAR fingerprint

Table 3 summarises results from the SAR fingerprint exercise analysing public and proprietary databases. Thirty five thousand four hundred and thirty-seven compounds were analysed with 22.5% testing positive in the bacterial reverse mutation assay. For the general class of aromatic *N*-oxides (shown in row 2), there were 242 examples and 27.7% were mutagenic. Specific subclasses are included in this table where they included four or more examples or where they represent a general subclass of aromatic *N*-oxides. An analysis is included in the comments field including cases where positive data are from public sources.

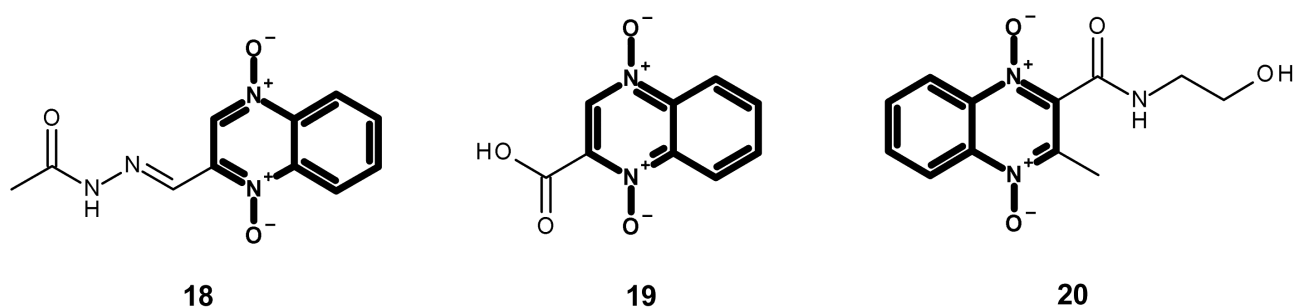


Fig. 9. Mutagenic examples containing a quinoxalin (highlighted in bold) and no additional structural alerts.

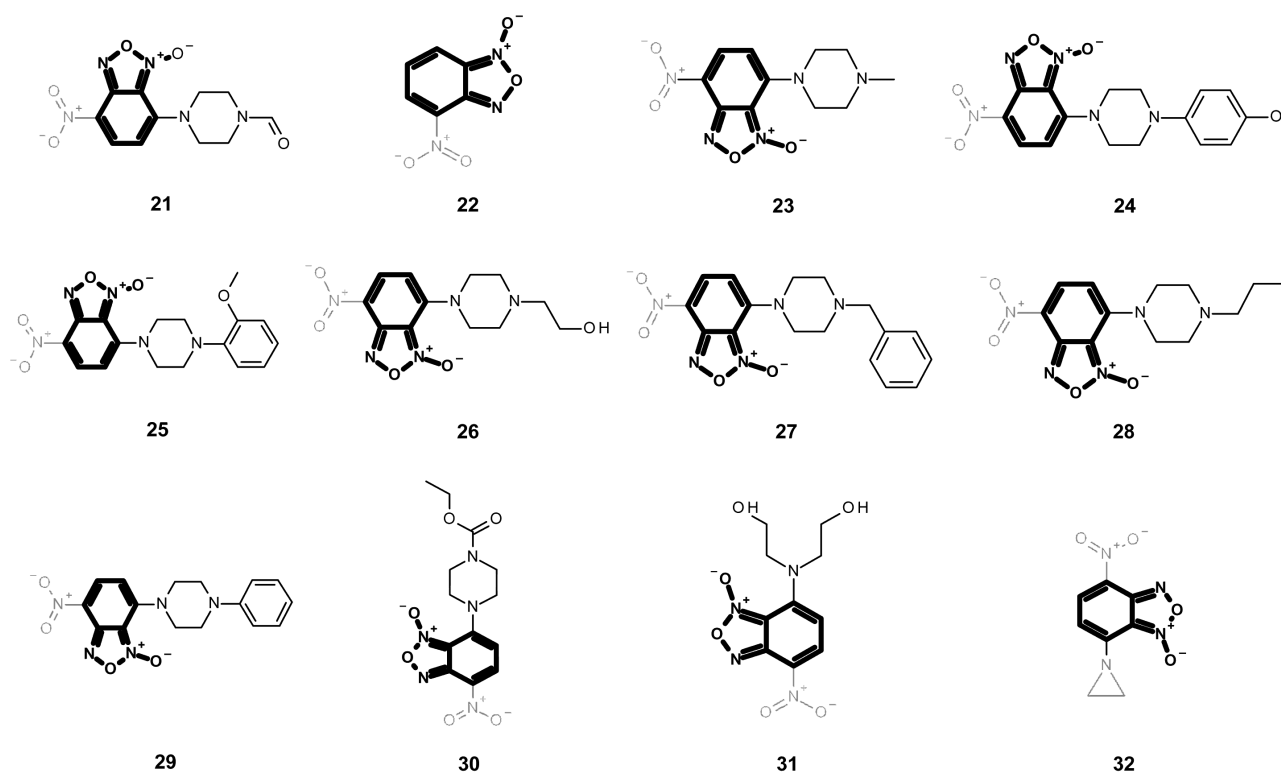


Fig. 10. Mutagenic examples containing a benzo[*c*][1,2,5]oxadiazole 1-oxide (highlighted in bold) that also contain one or more additional structural alerts (highlighted in grey).

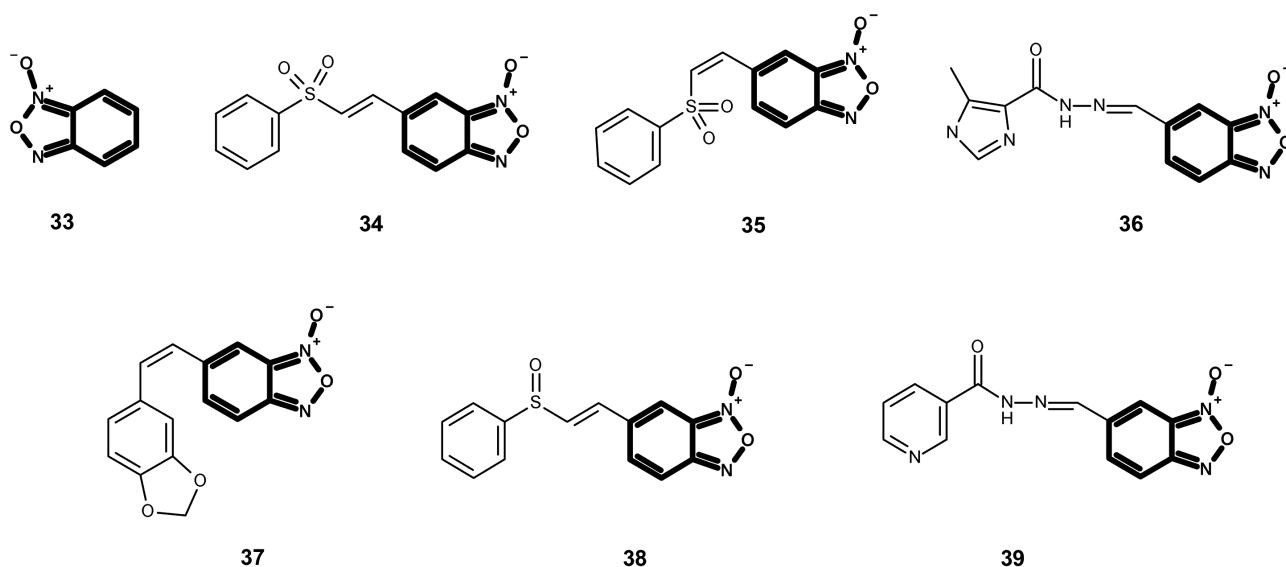


Fig. 11. Mutagenic examples containing a benzo[c][1,2,5]oxadiazole 1-oxide (highlighted in bold) with no additional structural alerts.

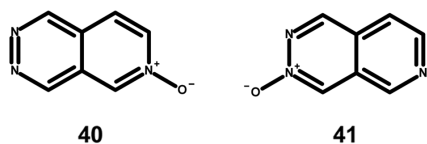


Fig. 12. Mutagenic examples containing a 2,7-naphthyridine 2-oxide (highlighted in bold).

Discussion

Are aromatic *N*-oxides, in general, associated with DNA-reactive mutagenicity?

The initial analysis of available public data found no evidence that the general class of aromatic *N*-oxides is a structural alert for DNA-reactive mutagenicity. All mutagenic examples contained another alert most likely responsible for the mutagenicity (e.g. presence of an aromatic nitro alert in compound **1** in Figure 2). There are also structurally diverse examples of non-mutagenic compounds containing aromatic *N*-oxides, and for many the data have been accepted by regulatory authorities. Moreover, a review of the published literature did not identify a plausible mechanism of mutagenicity for simple aromatic *N*-oxides.

An additional retrospective analysis of 24 ICH-M7-compliant (Q)SAR evaluations of impurities containing aromatic *N*-oxides conducted by FDA/CDER with the application of expert knowledge found that 6 were predicted to be mutagenic and 18 predicted to be non-mutagenic (not shown in the article). Those predicted to be positive were found to contain other confounding structural alerts that could explain their mutagenic activity. The negatives consisted of simple, monocyclic *N*-oxides, which appeared to be accurately predicted by the models. Although not experimentally verified, these observations are supportive of the hypothesis that aromatic *N*-oxides are not generally an alerting class and consistent with the data from Table 3.

The combined analysis of public and proprietary data confirmed this finding. In fact, if the 4 negative and 33 positive examples are removed from consideration based on the two classes shown in Figure 14, the number of non-mutagenic aromatic *N*-oxide examples (covering

monocyclic and fused aromatic *N*-oxides) is 171 and the number of mutagenic aromatic *N*-oxide examples is 34. The proportion of mutagenic examples is 16.6%, which is statistically significantly lower (P -value = 0.023) than the baseline computed across all compounds of 22.5%. This is comparable with other functional groups that are generally considered to be non-mutagenic. For example, the non-mutagenic carboxylate functional group was included in this SAR fingerprinting exercise and the proportion of mutagenic chemicals for these 5000 examples was 16.7%. The remaining positive rate can be explained by other factors responsible for the positive test results, such as additional mutagenic functional groups, impurities of the tested compound and unstable compounds resulting in mutagenic cleavage products.

Using the information in Table 3, the SAR fingerprint analysis provides additional affirmative support, because an isolated 6-membered ring with an *N*-oxide is assigned as deactivating based on the 7.7% positive rate from the 130 examples (P -value < 0.001). There was also no evidence that five-membered isolated aromatic rings (e.g. furan, pyrrole) containing an *N*-oxide are mutagenic.

As discussed earlier, a structural alert must be based on a sufficient number of examples to conclude that chemicals in a class have an increased likelihood of being active (e.g. mutagenic). Strege *et al.* (17) identified two chemicals containing aromatic *N*-oxides as being mutagenic. However, when combined with the other data reported in this article, it is not sufficient to support assigning aromatic *N*-oxides as alerts.

Are subclasses of aromatic *N*-oxide alerts DNA-reactive mutagens?

On the basis of the analysis of public and proprietary data, there is sufficient evidence to conclude and define a DNA-reactive mutagenic alert for quindioxin and related compounds (see Figure 14 and Table 4). Out of the 21 identified examples, 20 were positive and the association between chemicals containing this class and positive mutagenicity data was determined to be statistically significant (P -value < 0.001 based on the binomial score as discussed previously). There are 17 positive examples (public and proprietary) with no additional alerts (shown in Figure 14). The public compounds had similar positive tester strains suggesting a common mechanism.

Table 1. Summary of data and matching alerts for the 35 public mutagenic aromatic N-oxide chemicals

Compound ID	Leadscope ID	Name	Positive tester strains	Reference
1	LS-131487	2,3-Dimethyl-4-nitropyridine 1-oxide	Positive tester strains not reported (conducted according to the OECD TG471 guideline)	(31)
2	LS-131865	4-Nitropyridine N-oxide	TA102 ± S9; TA100 ± S9; TA1538 + S9	(32–35)
3	LS-1376	4-Nitroquinoline N-oxide	WP2uvra/pKM101 ± S9; TA102 ± S9; TA100 ± S9; TA98 ± S9; TA1535 – s9; TA1537 – s9	(36,37)
4	LS-1278	4-(Hydroxyamino)quinoline 1-oxide	WP2uvra ± S9; TA100 ± S9; TA1537 ± S9	(34,35, 37–39)
5	LS-189132	2-Chloroethyl 1,2,3-triazino[5,4- <i>b</i>]indole-4-carboxylate N(3)-oxide	TA97 ± S9; TA98 ± S9	(40)
6	LS-189651	Ethyl 1,2,3-triazino[5,4- <i>b</i>]indole-4-carboxylate N(3)-oxide	TA102 – S9; TA100 + S9; TA98 ± S9	(40)
7	LS-94153	3-Nitro-6-azabenz[<i>a</i>]pyrene N-oxide	TA98 – S9	(41)
8	LS-94152	1-Nitro-6-azabenz[<i>a</i>]pyrene N-oxide	TA98 – S9	(41)
9	LS-212651	6-Azabenz[<i>a</i>]pyrene N-oxide	TA100 – S9; TA98 – S9	(40)
10	LS-15080	Adenine 1-N-oxide	TA98 + S9	(42)
11	LS-141320	4-Aminoquinoline 1-oxide	Positive tester strains not reported	(43)
12	LS-188416	9-Methyladenine 1-N-oxide	TA98 + S9	(42)
13	LS-188320	9-Benzyladenine 1-N-oxide	TA98 + S9	(42)
14	LS-189066	9-Benzhydryl adenine 1-N-oxide	TA98 + S9	(42)
15	LS-189065	9-Trityl adenine 1-N-oxide	TA98 + S9	(42)
16	LS-76683	<i>tert</i> -Butyl-3-(2-quinoxalinylmethylene)carbazate 1,4-dioxide	WP2uvra/pKM101 ± S9; TA100 ± S9; TA98 ± S9	(44)
17	LS-51665	Carbadox	WP2uvra/pKM101 ± S9; TA98 ± S9	(44)
18	LS-12821	1-Acetyl-2-(2-quinoxalinylmethylene)hydrazine 1,4-dioxide	WP2uvra/pKM101 ± S9; TA100 ± S9; TA98 ± S9	(44)
19	LS-143005	2-Quinoxalinecarboxylic acid 1,4-dioxide	WP2uvra/pKM101 ± S9; TA100 ± S9; TA98 ± S9	(44)
20	LS-142996	Bayonox; N-(2-hydroxyethyl)-3-methyl-4-oxido-1-oxoquinoxalin-1-ium-2-carboxamide	WP2uvra/pKM101 ± S9; TA100 ± S9; TA98 ± S9	(44)
21	LS-110641	4-(4-Formylpiperazinyl)-7-nitrobenzofuroxan	Positive tester strains not reported	(37)
22	LS-35380	Benzofurazan, 4-nitro-, 1-oxide	Positive tester strains not reported	(37)
23	LS-35377	Benzofurazan, 7-(4-methyl-1-piperazinyl)-4-nitro-, 1-oxide	Positive tester strains not reported	(37)
24	LS-35368	4-(4-(4-Methoxyphenyl)-1-piperazinyl)-7-nitrobenzofurazan 3-oxide	Positive tester strains not reported	(37)
25	LS-35367	4-(4-(2-Methoxyphenyl)-1-piperazinyl)-7-nitrobenzofurazan 3-oxide	Positive tester strains not reported	(37)
26	LS-35364	7-(4-(2-Hydroxyethyl)-1-piperazinyl)-4-nitrobenzofurazan 1-oxide	Positive tester strains not reported	(37)
27	LS-35355	Benzofuran, 4-nitro-7-(4-(phenylmethyl)-1-piperazinyl)-, 1-oxide	Positive tester strains not reported	(37)
28	LS-35386	Benzofuran, 4-nitro-7-(4-propyl-1-piperazinyl)-, 1-oxide	Positive tester strains not reported	(37)
29	LS-35383	4-Nitro-7-(4-phenyl-1-piperazinyl)benzofurazan 1-oxide	Positive tester strains not reported	(37)
30	LS-201311	7-(4-(Ethoxycarbonyl)piperazin-1-yl)-4-nitrobenzo[<i>c</i>][1,2,5]oxadiazole 1-oxide	Positive tester strains not reported	(43)
31	LS-201438	7-(Bis(2-hydroxyethyl)amino)-4-nitrobenzo[<i>c</i>][1,2,5]oxadiazole 1-oxide	Positive tester strains not reported	(43)
32	LS-35354	4-Ethylenimino-7-nitrobenzofuroxan	Positive tester strains not reported	(37)
33	LS-201105	Benzo[<i>c</i>][1,2,5]oxadiazole 1-oxide	Positive tester strains not reported	(45)
34	LS-460268	(<i>E</i>)-6-(2-(Phenylsulfonyl)vinyl)benzo[<i>c</i>][1,2,5]oxadiazole 1-oxide	TA98 ± S9	(46)
35	LS-460269	(<i>Z</i>)-6-(2-(Phenylsulfonyl)vinyl)benzo[<i>c</i>][1,2,5]oxadiazole 1-oxide	TA98 ± S9	(46)
36	LS-460270	6-((2-(5-Methyl-1 <i>H</i> -imidazole-4-carbonyl)hydrazineylidene)methyl)benzo[<i>c</i>][1,2,5]oxadiazole 1-oxide	TA98 ± S9	(46)
37	LS-417996	(<i>Z</i>)-6-(2-(Benzo[<i>d</i>][1,3]dioxol-5-yl)vinyl)benzo[<i>c</i>][1,2,5]oxadiazole 1-oxide	TA98 ± S9	(46)
38	LS-460271	(<i>E</i>)-6-(2-(Phenylsulfonyl)vinyl)benzo[<i>c</i>][1,2,5]oxadiazole 1-oxide	TA98 ± S9	(46)
39	LS-460272	6-((2-(Nicotinoyl)hydrazineylidene)methyl)benzo[<i>c</i>][1,2,5]oxadiazole 1-oxide	TA98 ± S9	(46)
40	LS-189357	Pyrido[3,4- <i>d</i>]pyridazine 3-oxide	TA98 ± S9	(47)
41	LS-189356	Pyrido[3,4- <i>d</i>]pyridazine 6-oxide	TA100 + S9; TA98 ± S9	(47)

Table 3. Summary of results from the SAR fingerprints


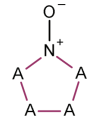
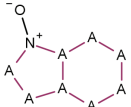
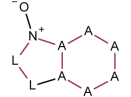
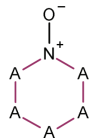
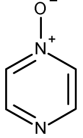
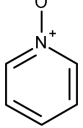
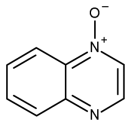
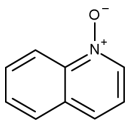
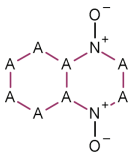
Row id	Substructure name	Substructure query	Number of negative	Number of positive	Total	Percentage of positives (%)	Comments
1	All compounds		27 453	7 984	35 437	22.5	
2	Aromatic N-oxide		175	67	242	27.7	35 of the positives are attributable to subclasses shown in row ids 5, 11, 13
3	Aromatic N-oxide- (5-ring isolated; any atom in ring)		4	0	4	0.0	No isolated 5-membered rings containing an aromatic N-oxide are mutagenic. It should be noted that these examples are from a single proprietary source
4	5,6 fused rings N-Oxide (any atom in ring)		17	16	33	48.5	13 of the 16 positive examples are from row id 5
5	1H-[2,3- [azaloxolthia]] indole 1-oxide		3	13	16	81.3	Many positive examples are from the public domain and contain other alerts. The negative proprietary examples are not members of the class benzo[c][1,2,5]oxadiazole 1-oxide
6	Aromatic N-oxide- (6-ring isolated; any atom in ring)		120	10	130	7.7	Strong evidence that isolated 6-membered aromatic rings with an aromatic N-oxide attached are not mutagenic (P-value < 0.001)
7	Pyrazinium, 1-oxide-		4	0	4	0.0	No evidence that pyrazinium, 1-oxide- subclass is a mutagenic alert
8	Pyridinium, 1-oxide-		114	10	124	8.1%	No evidence that pyridinium, 1-oxide-subclass is a mutagenic alert
9	Quinoxalinium, 1-oxide-		4	1	5	20.0	No evidence that quinoxalinium, 1-oxide- is a mutagenic alert
10	Quinolinium, 1-oxide-		9	6	15	40.0	The three positive public examples (structure ids: 3, 4, 11) contain other alerts; two proprietary positive examples also contained other alerting structures
11	Quindioxin and related		1	20	21	95.2	Strong evidence that quindioxin and related chemicals are mutagenic (P-value < 0.001)

Table 3. (Continued)

12	Isoquinolinium, 2-oxide-		7	3	10	30.0	Two positive examples contain the substructure from row id 13
13	2,7-[1,3,4,4a,5,6,8,8a- any]naphthyridine 2-oxide		0	2	2	100.0	There are only two public mutagenic examples(40,41)
14	6,5 Fused ring N-oxide		3	0	3	0.0	No evidence for assignment as a mutagenic alert
15	6,5 Fused ring N-oxide		1	7	8	87.50	All 7 positive examples (structure ids: 5, 6, 10, 12, 13, 14, 15) contain other alerts

A is any atom; L is N or S or O; A-A, A-L, L-L, A-N and A-N⁺ are aromatic bonds; all P-values are calculated based on the binomial score.

Alerting class	Public examples with no other alerts	Proprietary examples
 quindioxin and related	 18 19 20	14 positive proprietary examples
 benzo[c][1,2,5]oxadiazole 1-oxide	 33 34 35 36 37 38 39	No positive proprietary examples

Fig. 14. Summary of two validated alerts with mutagenic examples.

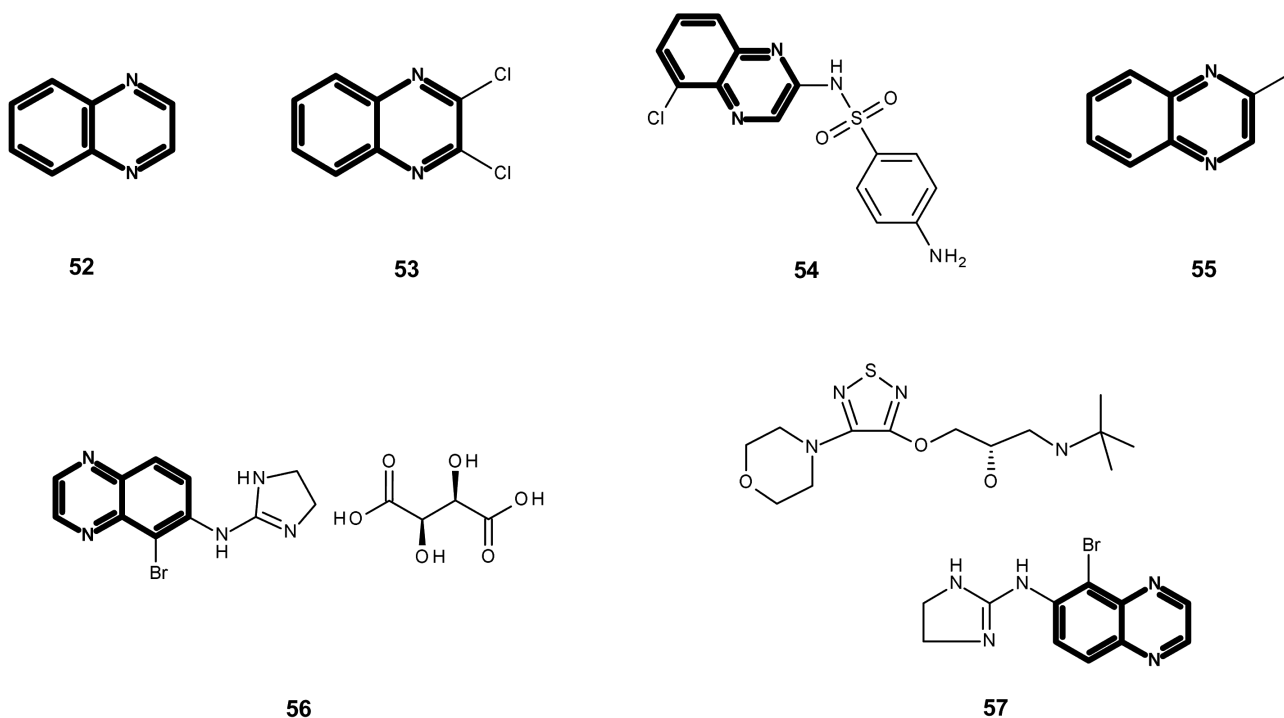
(including Fe, Cu, Cr, As and V) and peroxisome proliferators where this type of mechanism has been proposed. To a broad generalisation, the examples suggest that carcinogenic effects are probably thresholded because high/prolonged exposure is required in order to deplete/overcome endogenous defence mechanisms (68). Hydrogen peroxide, which is a direct-acting mutagen in multiple bacterial strains and a weak local carcinogen in the gastrointestinal tract in the mouse (69), is a well-established source of hydroxyl radicals, yet its cosmetic use as a tooth whitener is considered to be without significant risk and it is not classifiable as to its carcinogenicity in humans (70).

Analysis of the free form of quindioxins

To help verify that the mutagenicity is attributable to the quindioxin or the free form, a separate analysis of compounds without the oxide was performed, shown in Figure 15 and Table 4. All examples were negative, with the exception of quinoxaline (49) where there are conflicting study results for strain TA98 (but concordant negative in TA100 as well as negative results in TA102, TA1535, TA1537 and TA1538). In this case, quinoxaline may be positive in TA98 with metabolic activation; however, there are also conflicting negative data present. Therefore, this analysis did not identify any association with the free form of quindioxin and mutagenicity.

Table 4. Study summary for compounds containing the free form of quindioxins

Compound id	Leadscope id	Name	Study result	Strains tested	Positive tester strain	Reference
52	LS-142960	Quinoxaline	Positive	TA98(2); TA100(2); TA102(2)	TA98 + S9	(71)
52	LS-142960	Quinoxaline	Negative	TA98(3); TA100(3); TA1535(3); TA1537(3); TA1538(3)		(72)
53	LS-1019	2,3-Dichloroquinoxaline	Negative	TA100(6); TA1535(6); TA1537(6); TA98(6)		(73)
53	LS-1019	2,3-Dichloroquinoxaline	Negative	TA98(3); TA100(3); TA1535(3); TA1537(3); TA1538(3)		(74)
53	LS-1019	2,3-Dichloroquinoxaline	Negative	TA100(3); TA1535(3); TA1537(3); TA98(3)		(75)
54	LS-31236	Chlorsulfaquinoxaline	Negative	TA98(3); TA100(3); TA1535(3); TA1537(3); TA1538(3)		(76)
55	LS-143043	2-Methylquinoxaline	Negative	TA98(2); TA100(2); TA102(2)		(71)
56	LS-177484	Brimonidine tartrate	Negative	TA1535(2); TA1537(2); TA1538(2); TA98(2); TA100(2)		(77)
56	LS-177484	Brimonidine tartrate	Negative	TA98(2); TA1537(2); TA1538(2); TA1535(2); TA100(2)		(78)
57	LS-445000	Brimonidine tartrate and timolol maleate	Negative	TA98(4); TA100(4); TA102(4); TA1535(4); TA1537(4)		(79)

**Fig. 15.** Compounds containing the free form of quindioxin (shown in bold).

SAR fingerprint analysis

This article describes how information from public and proprietary sources can be combined to refine alerts for predicting mutagenicity. Through refinement of the set of mutagenicity alerts, it is possible to

streamline the assessment of impurities, degradants and other types of chemicals to avoid unnecessary testing and implementation of control strategies. This project made use of proprietary data using a much larger set of compounds than previously available for analysis

of aromatic *N*-oxides to better define the structure–activity (mutagenicity) relationships for this class. It should be noted that it was not necessary to release any proprietary information on individual compounds or data, because only derived aggregated information on chemical classes was used in this analysis. It was, however, possible to ask questions to the owners of the proprietary data to support this analysis. For example, questions were asked for a number of entries to determine whether, for specific classes, they contained other functional groups most likely responsible for DNA-reactive mutagenicity (e.g. Table 3, row 10).

Conclusion

On the basis of the analysis of aromatic *N*-oxide structure–activity relationships for bacterial mutagenicity presented in this article, we conclude that the general class of aromatic *N*-oxide is not predictive of DNA-reactive mutagenicity. Two subclasses were determined to be predictive for DNA-reactive mutagenicity: quindioxin and benzo[*c*] [1,2,5]oxadiazole-1-oxide. Currently available evidence suggests Ames-positive aromatic *N*-oxides may be indirectly DNA-reactive by a mode of action involving hydroxyl radicals. The conclusions from this article may be used to support the development of (Q)SAR methods as well as support an expert review of the (Q)SAR results, mentioned in the ICH M7 guideline, where an aromatic *N*-oxide is highlighted as responsible for a DNA-reactive mutagenic prediction.

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