

Enzymology of bioreductive drug activation

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Keywords: P450 reductase; NQO1; DT-diaphorase; xanthine oxidase; mitomycin C; E09; tirapazamine

Background and current status

The hypoxic fraction of solid tumours is resistant to ionising radiation and some chemotherapeutic agents (Sartorelli et al., 1994). Drugs requiring bioreductive activation have been developed to exploit the oxygen deficiency in these cells on the premise that hypoxic cells should show a greater propensity for reductive metabolism than well-oxygenated cells (Lin et al., 1972). Hypoxia should also limit interactions of the reduced drug with molecular oxygen thereby maintaining the drug in the reduced, cytotoxic form. Much of the recent work in this area has focused on specific enzymes involved in bioreductive drug activation, and this has been more recently described as enzyme-directed bioreductive drug development (Workman and Walton, 1990). Appropriate characteristics of enzyme-directed bioreductive agents have been discussed (Workman, 1994) and these are summarised in Table I. They include knowledge of the role of a particular reductase in bioactivation of a drug and whether there is elevated expression of that reductase in a particular tumour type (Table I). Using this approach a bioreductive drug could be targeted against a particular tumour type. Additional steps for bioreductive drug development such as activity in xenografts and in clinical trials, individual patient profiling for activities of various reductases and the generation of a computerised database are also shown in Table I. A computerised database has proved particularly important in elucidating the role of various reductases in the toxicity of drugs to cell lines in the NCI panel (Paull et al., 1994) and will be discussed below. The enzymology of bioreductive drug activation has posed some problems owing to its complexity, but it has also offered some important opportunities for drug development. In this review, the enzymes which activate the most commonly studied bioreductive drugs and some of the problems associated with bioreductive drug research will be discussed.

Mitomycin C (Figure 1) is the prototype bioreductive alkylating agent (Sartorelli et al., 1994) and a clinically useful agent in the treatment of solid tumours (Crooke and Bradner, 1976). In fact, mitomycin C is the single most active agent for the treatment of non-small-cell lung cancer (Spain, 1993). The reductive activation of mitomycin C was first described in a classic paper by Iyer and Szybalski (1964). Various bioreductive enzymes (Table II) have been implicated in the bioreductive activation of mitomycin C including NAD (P)H:quinone oxidoreductase, also known as NOO1 or DT-diaphorase (Siegel et al., 1990, 1992; Marshall et al., 1989, 1991a; Begleiter et al., 1988, 1989; Dulhanty and Whitmore, NADPH:cytochrome P-450 reductase (Bligh et al., 1990; Keyes et al., 1984; Pan et al., 1984; Komiyama et al., 1982; Bachur et al., 1979), NADH:cytochrome b₅ reductase (Hodnick and Sartorelli, 1993), xanthine oxidase (Pan et al., 1984) and xanthine dehydrogenase (Gustafson and Pritsos, 1992a). The

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roles of another form of NAD(P)H:quinone oxidoreductase, NQO2 (Jaiswal, 1994), and carbonyl reductase (Forrest et al., 1991) in the metabolism of mitomycin C are unclear. Oneelectron reduction of mitomycin C can lead to generation of reactive oxygen species whereas either one- or two-electron reduction can produce DNA monoalkylation products or DNA cross-links (reviewed by Sartorelli et al., 1994). It should be emphasised that bioreductive metabolism is only the first step in expression of cytotoxicity, and many other factors such as the behavior of one-electron and two-electron reduced forms under oxia and hypoxia, the type of cell damage produced (e.g. DNA cross-linking, initiation of apoptosis, etc.) and repair of that damage should be incorporated into any predictive models. This paradigm, although still markedly oversimplified, is shown in Figure 2. Our lack of knowledge of other steps in cellular damage and repair may explain why cells which have markedly increased levels of bioreductive enzymes often show only modest increases in toxicity after treatment with bioreductive agents (Gibson et al., 1992a,b).

The indoloquinone EO9 (Figure 1) is a promising bioreductive drug which is currently in phase I/II evaluation with the EORTC (Plumb et al., 1994). EO9 is a much better substrate for NQO1 than mitomycin C, and the reduced form of EO9 redox cycles more efficiently than the reduced form of mitomycin C (Beall et al., 1995). DNA single-strand breaks have been reported following reduction of EO9 by NQO1 (Walton et al., 1991), and more recently, it was shown that reductive activation of EO9 by NQO1 and xanthine oxidase produced DNA cross-links under hypoxic conditions (Maliepaard et al., 1995). A series of aziridinyl benzoquinones has been studied as potential bioreductive drugs (Gibson et al., 1992a) including diaziquone which has been investigated clinically (Alley et al., 1994). MeDZQ (Figure 1), an analogue of diaziquone, is a better substrate for NQO1 than diaziquone (Beall et al., 1994), and it shows an absolute requirement for reduction to cross-link DNA (Lee et al., 1992). Streptonigrin (Figure 1) is a hydroxyl radical-generating quinone (Cone et al., 1976) that is efficiently activated by NQO1 (Beall et al., 1995).

Other compounds such as the aromatic heterocyclic di-Noxides are bioactivated by one-electron reduction, and two-electron reduction may represent a detoxification reaction (Walton and Workman, 1993). Bioactivation of tirapazamine

Table I Appropriate characteristics of enzyme-directed bioreductives and steps in drug development

Can the enzyme activate the drug?
Are there increased enzyme activities in tumours?
Is there a relationship between enzyme activity and a biochemical end point of cell damage or cytotoxicity?
Is there a similar relationship in xenografts?
Clinical trails?
Individual patient profiling?
Computerised database?



Figure 1 Structures of bioreductive drugs.

(Figure 1), the prototype of this series, can be accomplished with cytochrome P-450 reductase, cytochrome P450 (various isozymes), and xanthine oxidase while NQO1 can detoxify tirapazamine (Patterson et al., 1994). Single- and doublestrand breaks in DNA have been reported following activation of tirapazamine by xanthine oxidase under hypoxia (Laderoute et al., 1988). The dinitrophenylaziridines represent another class of bioreductive drugs. CB1954 (Figure 1) is the lead compound in this series in which the 4-nitro substituent is reduced to a 4-hydroxylamino group in a fourelectron reduction, although this compound is a much less efficient substrate for NQO1 than mitomycin C (Beall et al., 1994). NQO1 has been implicated in this reaction both in air and under hypoxia whereas cytochrome P-450 reductase and xanthine oxidase, which are one-electron reductases, generate the nitro radical anion which redox cycles under aerobic conditions (Knox et al., 1988). Acetylation of the hydroxylamino group generates a bifunctional alkylating agent, the proposed ultimate DNA cross-linking species (Knox et al.,

NQO1 has generated the most interest of the potential bioactivating enzymes because of its marked overexpression in many tumours and tumour cell lines (Ross et al., 1994; Workman, 1994). Elevations in NQO1 activity or mRNA have been observed in preneoplastic tissues, established tumours or tumour cell lines from human lung, liver, colon, breast and brain (Schlager and Powis, 1990; Malkinson et al., 1992; Cresteil and Jaiswal, 1991; Workman et al., 1991; Berger et al., 1985). Other tumour tissues, including human kidney and stomach, have been found to have decreased

NQO1 activity relative to normal tissue (Schlager and Powis, 1990). The NQO1 activity in human non-small-cell lung tumours and paired normal tissue is shown in Figure 3, illustrating the differences in NQO1 levels between normal and tumour tissue. It should be stressed, however, that although NOO1 is elevated in some tumours such as nonsmall-cell lung cancer (Malkinson et al., 1992), there is a marked variation in NQO1 activity within a specific tumour type (Figure 4). The variation in NQO1 activity in human tumours is also readily apparent from the data shown in Figure 3.

NQO1 is one of the four diaphorases (diaphorase-4) identified by Edwards et al. (1980) corresponding to four gene loci. A diaphorase, or 'coenzyme factor', catalyses the transfer of electrons from reduced pyridine nucleotides to oxidation-reduction indicators (Dewan and Green, 1938; Corran et al., 1939). NQO1 is a two- (Iyanagi and Yamazaki, 1970) or four-electron reductase (Huang et al., 1979) characterised by its unique ability to use both NADH and NADPH as reducing cofactors and its inhibition by dicumarol (Ernster, 1967). A flavoprotein, NQO1 exists as a homodimer with 1 mol of FAD per mol of NQO1 (Lind et al., 1990). Human NQO1 has been cloned and sequenced (Jaiswal et al., 1988), and the crystal structure of the rat liver homologue has recently been reported (Li et al., 1995). Human NQO1 is similar to rat NQO1 with 83% homology for the cDNA and 85% homology for the protein (Jaiswal et al., 1988). Another form of NAD(P)H:quinone oxidoreductase has been isolated, cloned and sequenced and has been designated NQO2 (Jaiswal et al., 1990).

Table II Bioreductive enzymes

NAD(P)H:quinone oxidoreductase1, NQO1 NAD(P)H:quinone oxidoreductase2, NOO2 NADPH:cytochrome P-450 reductase NADH:cytochrome b₅ reductase Xanthine oxidase Xanthine dehydrogenase Carbonyl reductase

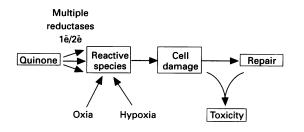
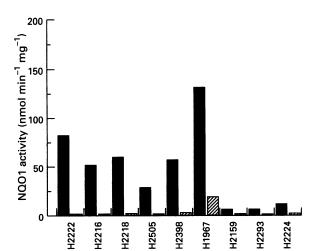


Figure 2 Paradigm for bioreductive activation of quinones.



Cell line

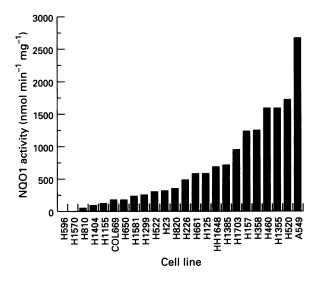


Figure 4 NQO1 activity in non-small-cell lung cancer cell lines (Malkinson et al., 1992).

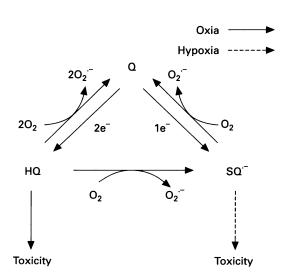


Figure 5 Bioactivation of quinones under aerobic and hypoxic conditions. Q, quinone; HQ, hydroquinone; SQ, semiquinone.



The role of NQO1 in the bioactivation and toxicity of mitomycin C has been controversial. Although metabolism of mitomycin C by NQO1 is pH-dependent owing to pHdependent inactivation of the enzyme (Siegel et al., 1993), mitomycin C can be bioactivated by NQO1 in the physiological pH range of 7.0-7.4 (Siegel et al., 1990, 1992). These results indicate that NQO1 can bioactivate mitomycin C at physiological pH, but do not imply that NQO1 is the sole reductase catalysing bioactivation of mitomycin C in cellular systems. The role of NQO1 in the toxicity of mitomycin C in tumour cells is controversial. Evidence for an association between NQO1 activity and mitomycin C sensitivity has been reported in both cell lines (Siegel et al., 1990; Begleiter et al., 1989; Dulhanty and Whitmore, 1991; Marshall et al., 1991a), and xenografts (Malkinson et al., 1992), but the role of NQO1 has been questioned because of a lack of correlation between mitomycin C sensitivity and NQO1 activity in a tumour cell line panel (Robertson et al., 1992). More recently, however, we demonstrated a significant correlation between NQO1 activity and mitomycin C sensitivity in human lung and breast cancer cell lines (Beall et al., 1995), and a highly significant correlation (r = 0.424, P < 0.0005) was reported in the NCI panel of 69 human tumour cell lines (Fitzsimmons et al., 1996). Of all of these observations the latter provides the strongest support for a potential role of NQO1 in bioactivation of mitomycin C but this evidence is, by nature, indirect. It could also be reasonably argued that, from a toxicological perspective, one might not expect a correlation of NQO1 activity with toxicity, but the situation may be better represented by a threshold model. Correlational studies, however, do not imply causality. The only mechanistic evidence supporting a role for NQO1 in the toxicity of mitomycin C in cells is the data of Hodnick et al. (1995) who showed that both mitomycin C and porfiromycin toxicity could be potentiated in CHO cells transfected with NQO1 cDNA. This observation is the first direct demonstration of the role of NQO1 in the bioactivation of mitomycin C in cells. However, in view of the lower catalytic efficiency of human NQO1 as compared with rat NQO1 (see below), these experiments need to be performed using human NQO1.

The importance of NQO1 in EO9 metabolism is somewhat clearer. EO9 can be bioactivated by NQO1 (Walton $et\ al.$, 1991), and correlations between sensitivity to EO9 and NQO1 levels have been reported in tumour cell line panels (Robertson $et\ al.$, 1992, 1994; Smitskamp-Wilms $et\ al.$, 1994). In the NCI panel, EO9 also had a highly significant correlation with NQO1 activity (r=0.446, P<0.0013; Fitzsimmons $et\ al.$, 1996). A role for NQO1 in the mechanism of toxicity of both EO9 and mitomycin C has been suggested in a study using NSCLC lines made resistant to mitomycin C which were also cross-resistant to EO9 (Shibata $et\ al.$, 1995).

The situation becomes much more complex when chemosensitivity studies are conducted under aerobic and hypoxic conditions. Work by Sartorelli's group in the 1980s suggested that NQO1 may bioactivate mitomycin C in welloxygenated cells, but may protect against mitomycin Cinduced cytotoxicity in hypoxic cells (Keyes et al., 1984, 1985, 1989). These studies used the NQO1 inhibitor dicumarol which may have other effects on cellular metabolism as well (reviewed by Ross et al., 1993). In fact, dicumarol was found to increase the hypoxic toxicity of mitomycin C to L1210 murine leukaemia cells which have no measurable NQO1 activity (Keyes et al., 1989). Perhaps most notably, dicumarol has been shown to exert marked stimulatory effects on xanthine dehydrogenase-mediated metabolism of mitomycin C (Gustafson and Pritsos, 1992b) and inhibitory effects on cytochrome b₅ reductase (Hodnick and Sartorelli, 1993). More recent work has used two human colon carcinoma cell lines; HT-29 with high NQO1 activity and BE with undetectable NQO1 activity due to a polymorphism in the NQO1 gene (Traver et al., 1992). Under aerobic conditions, it was found that mitomycin C was much more toxic to the



high NQO1 HT-29 cells than the NQO1-deficient BE cells. Under hypoxic conditions, sensitivity was essentially unchanged in the HT-29 cells, but was markedly enhanced in the BE cells suggesting the involvement of a one-electron reductase in the enhancement of mitomycin C toxicity in BE cells (Beall et al., 1994). Plumb and Workman (1994) reported similar results for mitomycin C, but the differences in sensitivity were more dramatic with EO9. While sensitivity to EO9 in the HT-29 cells was moderately increased under hypoxia, EO9 toxicity in the BE cells was enhanced by over 1000-fold in hypoxic vs aerobic conditions (Plumb and Workman, 1994). Other reductases in these two cell lines are present at low levels with similar activities (Beall et al., 1994; Plumb and Workman, 1994).

From the data above, it seems clear that the enhanced hypoxic toxicity of quinones is not a function of NQO1mediated metabolism, but depends on the activity of oneelectron reductases. It is not clear, however, whether the oneelectron or two-electron reduced forms of quinones or both are responsible for cytotoxicity, and mechanisms may be proposed for either the semiquinone or hydroquinone as the toxic species. The mechanism is complicated not only by interactions of the two reduced species with molecular oxygen, but also by disproportionation and comproportionation reactions between the quinones and their two reduced forms. A possible scheme for bioactivation of quinones is presented in Figure 5. Disproportionation and comproportionation reactions have not been included for simplicity.

A recent study (Plumb et al., 1994) supports the hypothesis that it is the semiquinone that is the proximate toxin. In a panel of cell lines, they demonstrated that the expected correlation between NQO1 activity and EO9 toxicity did exist, but they also showed that there was an inverse correlation under hypoxia, that is, sensitivity decreased with increasing NQO1 activity (Plumb et al., 1994). This suggests that NQO1 protects the cell from EO9 toxicity under hypoxia because autooxidation of the hydroquinone to the more toxic semiquinone is inhibited by the decreased oxygen concentration. The reduced form of EO9 is known to be unstable to oxygen (Beall et al., 1995; J Butler, personal communication). Moreover, in the presence of dicumarol, the inverse correlation observed under hypoxia was lost providing additional evidence for the role of NQO1 in protecting the cells in hypoxic conditions. They also observed that there was a hypoxic/oxic differential for EO9 toxicity in all cell lines with the exception of the A549 lung tumour cell line (highest NQO1 activity) and that the highest differentials were found in those cell lines with the lowest NQO1 activities (Plumb et al., 1994). This could support the premise that the semiquinone is the most toxic species, as diminished oxygen levels would stabilise the semiquinone leading to greater toxicity under hypoxia especially in those cell lines where NOO1 levels are low and the putative protective effects of this enzyme are absent.

The aerobic/hypoxic controversy can also be extended to the one-electron reductase, cytochrome P-450 reductase. Surprisingly, cytochrome P-450 reductase has been implicated in the activation of mitomycin C under aerobic, but not hypoxic conditions (Hoban et al., 1990; Bligh et al., 1990).

Table III Rates of reduction of bioreductive drugs by rat and human recombinant NQO1 (Beall et al., 1995) and percentage efficiency of the human enzyme vs the rat enzyme

	Velo (μmol m	Velocity ^a (μmol min ⁻¹ mg ⁻¹)	
Compound	Rat NQO1	Human RQ01	(%) Human/rat
Streptonigrin	243 ± 42	51 ± 4	21
MeDZQ	104 ± 31	25 ± 4	24
EO9	181 ± 19	7.7 ± 2.0	4
Mitomycin C	0.68 ± 0.02	0.15 ± 0.01	22
CB1954 ^b	0.036 ± 0.003	0.010 ± 0.005	28

^aMean \pm s.d. for at least n=3 reactions. ^b100 μ M CB1954; all others $50 \mu M$.

However, Belcourt et al. (1995) found that CHO cells which were transfected with cytochrome P-450 reductase were more sensitive to mitomycin C and porfiromycin under both aerobic and hypoxic conditions. Sensitivity of the clones to mitomycin C and porfiromycin was greater under hypoxic conditions than in air, and they concluded that cytochrome P-450 reductase contributed to the hypoxic/oxic differential sensitivity of the cells to mitomycin C and porfiromycin (Belcourt et al., 1995).

Search for efficient substrates for bioreductive enzymes

In the search for efficient substrates for the bioreductive enzymes, one approach has been to attempt to correlate reduction potentials to rates of reduction (reviewed by Cadenas, 1995). For the one-electron reductase, xanthine oxidase, Clarke et al. (1980, 1982) determined that there was a good correlation between one-electron reduction potential and rates of reduction of nitroimidazoles. A relationship between one-electron reduction potential and rates of reduction by cytochrome P-450 reductase (Butler and Hoey, 1993; Powis and Appel, 1980) and cytochrome b₅ reductase (Powis and Appel, 1980) have also been reported. However, when a similar approach was attempted for NQO1, no correlation was found between reduction potential and rates of reduction by NQO1 for a series of naphthoquinones (Buffington et al., 1989) or a series of aziridinylbenzoquinones (Gibson et al., 1992a). In the naphthoguinone study (Buffington et al., 1989), rates of reduction by NQO1 also did not correlate with octanol-water partition coefficients which further precludes the ability to predict substrate efficiency for NQO1. In view of these data, there are basically two approaches to identifying efficient substrates for NQO1. The first is an experimental approach using recombinant rat and human NQO1 (Ross et al., 1994), whereas the second is to use the crystal structure of NQO1. The crystal structure of rat liver NQO1 has been determined to 2.1Å resolution and has been published in preliminary form (Li et al., 1995). Using the experimental approach several excellent substrates have been identified and assayed using purified rat and human recombinant NQO1 (Table III). Streptonigrin, MeDZQ and EO9 are rapidly reduced by both the rat and human forms of NQO1, and with the exception of EO9, the rates of reduction of the bioreductive agents by human recombinant NQO1 are between 20% and 30% of the rates of reduction by rat recombinant NQO1 (Beall et al., 1995).

Chemosensitivity studies have also been carried out using streptonigrin, MeDZQ, EO9 and other bioreductive agents (Beall et al., 1995). Two non-small-cell lung cancer cell lines were used: H460 which has high NQO1 activity (1360±100 nmol min⁻¹ mg⁻¹) and H596 which has undetectable NQO1 activity. Selectivity ratios (SRs) were calculated as the concentration at which survival equals 50% of control (IC₅₀) for H596 cells divided by the IC₅₀ for H460 cells. A higher SR implies a greater selectivity for cells with elevated NQO1 activity. In this system, streptonigrin was the most selective of the bioreductive drugs that were tested (SR = 86) followed by EO9 (SR = 62) and MeDZQ (SR = 17). This is in agreement with the ability of streptonigrin to serve as the most efficient substrate for NQO1 that we have tested to date (Beall et al., 1995). In addition, streptonigrin showed the best correlation between NQO1 activity and cytotoxicity of over 31 000 compounds tested in the NCI human tumour cell line panel (Paull et al., 1994).

It seems clear that compounds which are efficient substrates for bioreductive enzymes will be identified either on the basis of reduction potentials, experimental studies using purified enzymes or using crystal structures. Whether compounds should be directed against a single reductase such as NQO1 or whether they should be targeted to multiple reductases is an important question. In addition, a critical step in identifying appropriate compounds which should progress to the clinic is the selection of appropriate and predictive preclinical models.

In this regard, xenografts (Langdon et al., 1994) and perhaps tumour spheroids may be more predictive models than simply using cell lines (Figure 6).

Potential problems with the bioreductive approach

Although NQO1 appears to be a useful target for bioreductive drug research, there are many potential problems with the bioreductive approach (Table IV). Tumour heterogeneity could be considered a fundamental flaw in the bioreductive approach since there may be marked differences in bioreductive enzymes in different cells within the tumour. Differences in oxygen tension, blood flow and pH may also modulate bioreductive enzyme activity within a tumour. For example, hypoxia has been shown to modulate enzyme activity by transcriptional induction and increased stability of mRNA (O'Dwyer et al., 1994). In addition, a study using multicellular spheroids demonstrated that NQO1 activity and gene expression increased with increasing depth within the spheroid. This trend was reversed, however, in layers near the necrotic centre of the spheroid (Phillips et al., 1994). Differences in cellular pH within a tumour can affect the activation of bioreductive drugs such as mitomycin C (Kennedy et al., 1985; Pan and Hipsher, 1993; Siegel et al., 1993). Environmental factors must also be considered since xenobiotics and dietary constituents can influence bioreductive enzyme activity. NQO1 and other detoxification enzymes are induced by a wide variety of chemical agents including bifunctional enzyme inducers (dioxin, polycyclic aromatic hydrocarbons, azo dyes, flavonoids) and monofunctional enzyme inducers (diphenols, isothiocyanates, dithiolthiones and thiocarbamates) (Prochaska et al., 1985; Prochaska and Talalay, 1988). An important consideration is the effect that cigarette-smoking history has on bioreductive enzyme levels. Large differences in NQO1 activity have been reported between non-smokers, past smokers and present smokers in normal and tumour tissues in the colon and lung with the most dramatic differences seen in lung tumour tissues (Schlager and Powis, 1990). Elevated enzyme activities in normal tissues such as kidney (Schlager and Powis, 1990) may also be problematic and could potentially lead to unwanted bioreductive drug toxicity.

Many of the early studies in bioreductive drug research were carried out using reductases and cell lines of rodent origin. Extrapolation of data obtained using rodent models to human systems could be misleading. For example, rodent enzymes may catalyse bioreduction at very different rates than enzymes of

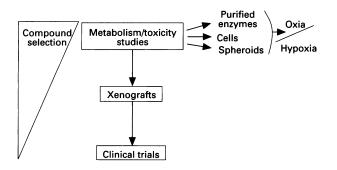


Figure 6 Possible test systems for bioreductive drug development.

Table IV Potential problems with the bioreductive approach

Tumour heterogeneity. Elevated enzyme activities in normal tissues. Influence of environmental factors on bioreductive enzymes. Rodent/human differences. Polymorphisms in bioreductive enzymes.

human origin (Table III). Work with CB1954 in rodent cell lines revealed another potential problem with rodent/human extrapolations. CB1954 was found to be between 500 and 5000 times less toxic to human cell lines than to rat cell lines with similar NQO1 activities (Boland et al., 1991). The availability of purified recombinant human proteins, human tumour cell lines and xenografts should facilitate the design of future studies using human systems.

Finally, the presence of a genetic polymorphism in the NQO1 gene, and possibly other reductase genes, may necessitate implementation of pretreatment biopsies to determine if a functional bioreductive enzyme is present in the target tumour (Ross et al., 1994). The absence of functional NQO1 activity in the BE colon carcinoma cell line led to the discovery of a C to T point mutation at position 609, exon 6 of the NQO1 cDNA which codes for a proline to serine change at position 187 of the NQO1 protein (Traver et al., 1992). The same mutation has subsequently been identified in the H596 non-small-cell lung cancer cell line which also lacks detectable NQO1 activity (Traver et al., 1995). These two cell lines have abundant NQO1 mRNA, but they are not immunoreactive to a monoclonal antibody to human NQO1. Since the same monoclonal antibody is immunoreactive to recombinant mutant NQO1 (Siegel et al., 1995), this suggests that the NQO1 mutant protein is either not made, or it is made and rapidly degraded. The use of PCR to determine enzyme expression before bioreductive drug treatment has been suggested as a potential approach to individual patient enzyme profiling. The NQO1 polymorphism would present a problem in this regard since elevated mRNA expression, detected using PCR, would not determine the presence of the mutation. The incidence of the polymorphism in the NQO1 gene is as yet unknown, but one early study showed that NQO1 was absent in 4% of samples from a British population (Edwards et al., 1980). Since the point mutation was discovered, Rosvold et al. (1993) determined that the mutation had a frequency of 0.11 in a reference population and noted that it was significantly overrepresented in lung cancer. An absence of NQO1 was also identified in approximately 10% of renal cancer and associated normal tissue (Eickelmann et al., 1994). As previously discussed, NQO1 is classified as a detoxification and cancer-protective enzyme and whether a lack of NQO1 activity predisposes to cancer is an important question (Marshall et al., 1991b). Work is currently underway to document the frequency and ethnic distribution of this potentially important polymorphism and its role, if any, in predisposition to cancer.

Conclusions

- Present bioreductives are activated by multiple reductases, via both one- and two-electron reduction.
- Some reductases, such as NQO1, are elevated in certain tumours, but there is a wide variation even within tumour groups.
- Relationships between NQO1 activity and aerobic drug sensitivity have been reported for EO9 and mitomycin C.
- Metabolism by NQO1 is oxygen insensitive. Increased toxicity of bioreductive quinones under hypoxia is mediated by other enzymes, presumably one-electron reductases.
- NQO1 has been reported to protect cells from the hypoxic toxicity of EO9.
- Potential problems with the enzyme-directed approach include tumour heterogeneity, elevated enzyme activities in normal tissue, influence of environmental factors on bioreductive enzymes, rodent/human differences and polymorphisms in bioreductive enzymes.

Future directions

- 1. Characterise reactive species generated, how they are produced and how they react under aerobic and hypoxic conditions.
- 2. Determine whether the protective effect of NQO1 against the hypoxic toxicity of EO9 extends to other bioreductive drugs and in vivo situations.
- 3. Transfection studies in human cell systems should be performed with human cDNAs.
- Exploit the NCI database to identify new targets. Currently, the NCI panel only performs toxicity assays under aerobic conditions and selected assays should also be performed under hypoxia.
- 5. Incorporate parameters other than bioactivation such as DNA repair into predictive models.
- 6. Determine the incidence of polymorphisms in key enzymes such as NQO1 in different populations.
- Use more predictive preclinical assays. Different test systems such as xenografts should be incorporated into the testing paradigm.

Key questions

- Should we be developing 'single-enzyme' agents or continue with our present 'multiple enzyme' compounds?
- In view of the potential problems, should we even be attempting the bioreductive approach without patient profiling?
- How are bioreductive enzymes regulated in normal and tumour tissue?
- Do these compounds work?

Acknowledgements

The authors are extremely grateful to Drs NW Gibson, J Butler and RM Phillips for their assistance and critical input. This work was supported by HHS R01 CA51210.

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