A large body of epidemiological evidence points to vegetables as a dietary source of chemoprotective activity for humans (reviewed by Block et al., 1992; Steinmetz and Potter, 1996). Protective effects due to ingestion of Brassica oleracea vegetables, such as broccoli (B. oleracea, Italica Group), are drawing increased scientific attention and have been the subject of recent reviews (Beecher et al., 1994; Verhoeven et al., 1997). Reevaluation of data from multiple studies by meta-analysis has shown a significant reduction in the odds ratio of colon and rectal cancers associated with the consumption of B. oleracea vegetables (Kohlmeier and Su, 1997). Most recently, Michaud et al. (1999) reported results from a large cohort study in which there was a significant correlation between cruciferous vegetable consumption and reduction in bladder cancer incidence.

Brassica oleracea vegetables, like all cruciferous species, are well known as sources of glucosinolates: β-thiogluco-side N-hydroxysulfates, with an aglycone (or R-group) that is an alkyl, alkenyl, thioalkyl, thioalkenyl, aryl, arylalkyl or indolyl moiety (Fig. 1) (Rosa et al., 1997). Glucosinolates are hydrolyzed to their cognate isothiocyanates by myrosinase which is present in both plant cells and in human gut microflora (Shapiro et al., 1998). More than 20 isothiocyanates have been shown to inhibit the formation of tumors initiated by a variety of chemical carcinogens in several organ systems (Hecht, 1995; Zhang and Talalay, 1994). Of special interest is sulforaphane, the cognate isothiocyanate of glucoraphanin (4-methylsulfinyl-buty1 glucosinolate; GR), that is a potent inducer of mammalian detoxification (Phase 2) enzyme activity. Sulforaphane also protects against tumorigenesis in a rodent mammary tumor model (Fahey et al., 1997; Zhang et al., 1994; Zhang et al., 1992). Broccoli florets and young seedlings or sprouts are the primary dietary sources of sulforaphane and GR (Fahey et al., 1997; Zhang et al., 1994; Zhang et al., 1992).

GR is by far the most abundant thiocyanl glucosinolate present in harvested heads of cultivated broccoli (Fig. 1); however, three other thiocyanl glucosinolates, glucoiberin (GI; 3-methylsulfinylpropyl glucosinolate), glucoraphanin (GR; 4-methylsulfinylbutyl glucosinolate) and 4-methylsulfinylbutyl isothiocyanate (sulforaphane).
glucosinolate), and glucosinolate (GA; 5-methylsulfanylpentyl glucosinolate) have also been identified in broccoli heads. The concentrations of GI range from 0% to 10% that of GR, while those for GE and GA are much less than 5% that of GR (Carlson et al., 1987; Kushad et al., 1999; Shelp et al., 1993). Iberin, erucin, and alyssin (cognate isothiocyanates of GI, GE, and GA, respectively) have only ≈20%, 10%, and 10%, respectively, of the Phase 2 enzyme induction potential of sulforaphane (Prestera et al., 1993; Zhang et al., 1992), and thus account typically for no more than ≈2% of the total glucosinolate-derived Phase 2 induction potency of broccoli. Other glucosinolates present in broccoli heads, such as glucobrassicin (indole-3-ylmethyl glucosinolate), neoglucobrassicin (1-methoxyindole-3-ylmethyl glucosinolate), and 4-hydroxyglucobrassicin (Fig. 1), have no significant Phase 2 inducer potential (Fahey et al., 1997; Fahey et al., 1998).

There is considerable interest in understanding how to control GR levels in broccoli in order to create added value in this vegetable for its development as a functional food, and to enhance its utility in dietary chemoprotection strategies (Giamoustaris and Mithen, 1996; Kushad et al., 1999; Shapiro et al., 1998). Faulkner et al. (1998) have suggested that genetic factors which influence Phase 2 enzyme induction potential of sulforaphane (Prestera et al., 1993; Kushad et al., 1999; Mithen, 1996; Kushad et al., 1999; Shapiro et al., 1998) could be transferred to wild relatives of B. oleracea could be transferred to cultivated broccoli. They have shown that hybrids formed by crossing broccoli inbreds and wild relatives express much higher Phase 2 enzyme induction potential than the broccoli inbreds themselves. Although this approach may ultimately yield a commercially viable cultivar, broccoli lines resulting from the broccoli × wild relative crosses will likely require numerous generations of selection to improve the horticultural phenotype of resulting lines.

The object of the current breeding study was to better characterize Phase 2 enzyme induction potential of relatively elite and diverse broccoli inbred (doubled-haploid) lines that have horticultural characteristics which approach market quality. If genetic variation exists within an elite broccoli germplasm pool and this variation can be exploited, then cultivar development should proceed rapidly. Kushad et al. (1999) examined a set of 24 F1 hybrid and open-pollinated cultivars and 26 inbred lines of broccoli in a single environment and observed a range of GR at 1.5 to 21.7 µmol·g–1 dry head weight for cultivars and a range of GR at 0.8 to 13.8 µmol·g–1 dry head weight for inbreds. Those authors concluded that significant variation for GR concentration was present in the germplasm examined. However, they did not examine Phase 2 enzyme induction potential of these lines.

In the present study, a larger and more diverse pool of inbred broccoli lines was examined. Specific objectives were to evaluate 1) quinone reductase induction potential (QRIP) diversity among head samples from a population of broccoli lines in a single environment; 2) QRIP of lines selected based on 1996 induction potentials when measured subsequently in 1997; 3) QRIP correlation with head MSAG concentration, head weight, and number of days from transplant to harvest among inbreds; and 4) QRIP in a sample of hybrids made by crossing specific inbreds.

**Materials and Methods**

**Plant material.** Plant materials evaluated in this study included 73 doubled haploid or inbred lines developed at the U.S. Vegetable Laboratory, Charleston, S.C. These inbreds were a subset of lines developed from the commercial hybrid broccoli cultivars Arcadia, Everest, Eureka, Futura, Green Valiant, High Sierra, Marathon, Packman, Southern Comet, Sultan, Symphony, and Viking using standard anther culture techniques (Farnham, 1998) or conventional selfing. These lines were designated as USVL001 through USVL070, USVL073, USVL075, and USVL076. All inbred seed were produced onsite at the U.S. Vegetable Laboratory. The 73 inbreds were selected from a large pool of inbred lines originating in the U.S. Vegetable Laboratory breeding program and they represent a diverse phenotypic and genotypic sample of relatively elite broccoli.

Commercial F1 hybrid cultivars were also grown and evaluated to serve as reference checks in these studies. In addition, several F1 hybrid crosses were generated in a greenhouse during Winter 1996–97 using some of the inbreds described above. Four of the resulting hybrids were evaluated in 1997. All crosses were made following emasculation of females to insure that no selfed seed were produced from the respective crosses.

**Plant culture.** In 1996, 71 inbreds and five hybrid cultivars including Everest, High Sierra, Marathon, Sultan, and Viking were seeded to a commercial potting mix (Metromix 360; Grace Sierra, Milpitas, Calif.) in trays in a greenhouse during the first week of August, and transplanted to the field 7 Sept. All entries were grown in individual plots consisting of a single row of eight to ten plants, and randomly incorporated into a single block of a
field nursery. Additionally, a subset of eight inbred entries and the five hybrid cultivars were transplanted simultaneously to eight to ten plant plots in a second block adjacent to the first. Spacing between rows was 102 cm, and spacing between plants within a row was 15 cm throughout the nursery. The subset of entries with two replicate samples (one from the first and the other from the second block) were treated as a separate experiment (i.e., randomized complete block with two replications) within the 1996 field nursery for statistical analysis. All cultural practices (e.g., cultivation, fertilization, and irrigation) for the 1996 trial were standard for local conditions (Cook and Ezell, 1983). The soil type at the Charleston site is a Yorges loamy sand (fine loamy mixed, thermic Albaqualfs).

In 1997, 21 of the 71 inbreds grown in 1996, two additional inbreds, the five hybrid cultivars Everest, High Sierra, Marathon, Viking, and Futura, and four $F_2$ hybrids produced in Charleston were seeded to the greenhouse 5 Aug. Inbreds grown in 1997 were selected based on their 1996 QRIP values and on their pedigree. These individuals were selected to represent a range of QRIP while maintaining phenotypic and genotypic variation in the second sampling year. The $F_2$ hybrids produced in Charleston were crosses of inbreds that expressed divergent levels of QRIP in 1996. All entries were transplanted to field plots on 4 Sept. The study was designed as a randomized complete block with each of the three blocks containing plots of all entries (6 to 10 plants, depending on available seed supplies). Spacing and cultural practices were the same as in 1996.

**Head harvest, sampling, and extraction.** As plots approached maturity in both years, trials were checked every 2 to 3 d to identify plants ready for harvest, until studies were completed. In all trials and with all entries, heads were harvested when head diameter reached 10 to 12 cm. Two heads per plot were sampled at random, and subcutting stalks were cut to a 15-cm length. Sample date was recorded for calculation of the mean number of days from transplant to harvest (DTH). Heads were placed immediately on ice in a cooler and within 30 min of field harvest, fresh weights were recorded, ca. half of the florets were cut from the stem, placed in an individual sealable freezer bag, and frozen at –80°C. At the end of each growing season, samples were transported to Baltimore, Md., on dry ice for storage at –80°C before analysis. In processing samples for analysis, frozen florets ($\geq 30$ to 40 g) were removed from storage, weighed, and extracted as described by Fahey et al. (1997). Extracts were then stored at –20°C until needed for bioassay of QRIP and for direct quantitation of glucosinolates by paired ion chromatography.

**Bioassay of quinone reductase induction potential (QRIP).** Bioassay of QRIP was performed using Hepa 1c1c7 cells as described originally by Prochaska et al. (1992) and modified by Fahey et al. (1997). Excess myrosinase, [0.0003 units/mL of cell culture medium; purified according to the methods of Shikita et al. (1999)], was added at the time of dosing along with 500 μM ascorbate, in order to achieve complete hydrolysis of glucosinolates to their cognate isothiocyanates. Conversion of glucosinolates in broccoli extracts, to their cognate isothiocyanates, is essentially quantitative by this procedure (Fahey, unpublished results). One unit of inducer activity is the amount that doubles the QR activity in a microtiter well containing 150 μL of medium. Hence, a compound with a CD (the concentration of a compound required to double the QR specific activity in Hepa 1c1c7 murine hepatoma cells) of 1.0 μM has 6,667 units of inducer activity/μmol. We expressed the inducer potency of plant extracts as units per gram fresh weight (FW).

**Paired-ion chromatography of glucosinolates.** Plant extracts were chromatographed isocratically in 1 acetonitrile : 1 water (v/v) containing 5 mM tetracylammonium bromide (TDAB) at a flow rate of 3 mL·min⁻¹, on a reverse-phase column (Whatman Partisil 10 ODS-2, 250 × 4 mm; Whatman, Inc., Clifton, N.J.) using a Waters (Waters Corp., Milford, Mass.) high-performance liquid chromatography (HPLC) system equipped with a photodiode array detector (Prestera et al., 1996). Sinigrin (allyl glucosinolate) was used as a standard. When compared to equimolar concentrations of sinigrin, the relative integrated absorbance areas for alkyl glucosinolates (GR, GI, and GE), glucobrassicin, and neoglucobrassicin at 235 nm were 1.00-, 1.22-, and 2.70-fold greater. The use of TDAB, an otherwise ideal paired-ion solvent system, does not permit resolution of GR from GI, which has been identified as a minor component in most broccoli samples. In this study, a subset of representative germplasm from both trial years was subjected to chromatography using 5 mM tetracylammonium bromide (TMAB) in water at a flow rate of 2 mL·min⁻¹ on the same column and HPLC system described in the preceding paragraph. Glucosinolate identities were confirmed using a complimentary, normal-phase HPLC method (J. Troyer, personal communication). GI was not detected in over half of these samples and on average, accounted for <4% of the GR/GI peak obtained using TDAB as a paired ion. Based on these observations, although we report these data as MSAG (GR+GI) concentrations, GI is considered a minor component in all lines examined herein, while GR comprises well over 90% of the MSAG content.

**Data analysis.** Analyses of variance (ANOVA) of the subset of replicated samples from 1996 and the entire 1997 trial were performed using PROC GLM of SAS (release 6.12, SAS Inst., Inc., Cary, N.C.). Entry means were compared using Fisher’s protected LSD. Pearson correlation coefficients were calculated for all pairs of QRIP, MSAG concentration, head weight, and DTH. Reported values are followed by the se where appropriate.

**Results**

The QRIP of all broccoli heads harvested in 1996 ranged from essentially zero to 150,000 units/g FW and the overall mean was 34,000 ± 3,000 units/g FW (Fig. 2A). The 1996 range for MSAG concentration was 0.04 to 2.94 µmol·g⁻¹ FW (mean of 0.88 ± 0.07 µmol·g⁻¹ FW) for this same set of broccoli entries (Fig. 2B). A subset of entries for which replicate measurements of QRIP were obtained had a range of QRIP from 5,500 to 69,100 units/g FW (Fig. 3), with significant differences among entries (LSD₀.₀₅ of 0.29 µmol·g⁻¹ FW). As with QRIP, there were significant differences in MSAG concentration for this subset (range 0.15 to 2.05 µmol·g⁻¹ FW, LSD₀.₀₅ of 0.29 µmol·g⁻¹ FW).

There was a highly significant correlation ($r = 0.93$, $P < 0.0001$) between QRIP and MSAG of entries assayed in 1996. In addition, there was a significant positive correlation between DTH and QRIP ($r = 0.78$, $P < 0.0001$) and between DTH and MSAG ($r = 0.74$, $P < 0.0001$) of entries.

The set of entries from 1996 that were chosen for retest in 1997, had a 1996 range for QRIP of 1,000 to 142,000 units/g FW and a mean of 42,800 ± 6,100 units/g FW. In addition, this same set of entries had a 1996 MSAG range of 0.15 to 2.72 µmol·g⁻¹ FW and mean of 1.10 ± 0.13 µmol·g⁻¹ FW. These 1996 ranges and means of this selected subset of entries are similar to the ranges and means for the entire sample of entries grown in 1996.

The QRIP of entries grown in 1997 correlated with QRIP of...
those same entries grown in 1996 ($r = 0.73, P < 0.0001$; Fig. 4A). The range of QRIP values was 15,800 to 150,200 units/g FW in 1997 and the mean QRIP for all entries was 63,200 units/g FW. ANOVA indicated highly significant differences among the entries in this second year.

As with QRIP, MSAG means for entries grown in 1997 also correlated with values for the same entries grown in 1996 ($r = 0.79, P < 0.0001$; Fig. 4B). The range (0.24 to 2.99 µmol·g$^{-1}$ FW) for MSAG concentration in 1997 was nearly identical to that for entries grown the previous year. The overall MSAG mean for 1997 was 1.33 µmol·g$^{-1}$ FW, and ANOVA indicated significant differences among entries.

As observed in the 1996 trial, there was a significant and positive correlation in 1997 between entry means for QRIP and MSAG concentration ($r = 0.90, P < 0.0001$). In addition, DTH among entries was significantly and positively correlated with QRIP ($r = 0.79, P < 0.0001$) and MSAG concentration ($r = 0.78, P < 0.0001$). On the contrary, no significant correlations were observed between head weight and the other measured characters.

Of four hybrid crosses evaluated in 1997, only two (1 and 2) were actually made using parents with significantly different QRIP means (Table 1). The F$_1$ mean for Cross 1 was intermediate between that of the high and low parents, and the F$_1$ mean for Cross 2 was lower than that of the low parent. For cross 3, the F$_1$ mean was again intermediate between the high and low parents. Cross 4 had parental and F$_1$ QRIP means that were all similar.

All four crosses evaluated involved combinations of parents with significantly different MSAG means (Table 1). In all cases, the F$_1$ mean was intermediate between means of the high and low parents. In two crosses (1 and 4), the F$_1$ was closer to the high parent while for the other two (2 and 3), the F$_1$ was closer to the low parent.

Similar to observations for MSAG concentration, all crosses involved combinations of inbred parents with significantly different mean DTH (Table 1). For two crosses (1 and 3), the DTH mean of the F$_1$ fell in between that of high and low parents but closer to the low parent, while in the other two crosses (2 and 4), the F$_1$ mean was lower than the mean of the low parent.

**Discussion**

This is the first report in which Phase 2 detoxication enzyme (quinone reductase) induction potential of broccoli heads has been examined using a relatively large and diverse pool of inbred (primarily doubled-haploid) lines. This work is consistent with
the hypothesis that the primary broccoli inducer of detoxication enzymes is glucoraphanin, concentrations of which were elevated concomitantly with QRIP in inbreds and hybrids created from them. Faulkner et al. (1998) also found that MSAG (primarily glucoraphanin) and QRIP were interrelated in broccoli; however, results described herein, are based on a much larger pool of broccoli entries, a more elite pool of germplasm, and samples harvested from field-grown plants. Whereas other recent studies on the phytochemical attributes of broccoli and other *B. oleracea* crops focus primarily on glucosinolate content (i.e., Kushad et al., 1999), our primary focus is the potential for these vegetables to induce Phase 2 enzyme activity in a versatile in vitro mammalian cell bioassay system (Talalay et al., 1995). Whereas Faulkner et al. (1998) have suggested that increased Phase 2 induction potential can be transferred from related wild species to broccoli, results herein suggest that diversity already extant within elite broccoli germplasm might be exploited in a program of hybridization and rapid development of enhanced doubled-haploid lines.

In an initial screen of broccoli inbreds in 1996, a wide range of QRIP was observed (0 to 150,000 units/g FW), based on single plant samples. A mean of 34,000 units/g FW for this 1996 sampling of inbreds was virtually identical to that reported by Fahey et al. (1997) for a random sample of broccoli heads obtained from supermarkets. The magnitude of the SD for the sampled population and of the SE for QRIP indicates that differences among tested entries in 1996 were likely due in part to genotypic effects. The importance of entry (and presumably genotype) in 1996 was supported by detection of significant differences for QRIP among a subset of replicated entries.

Line selection for QRIP based on individual samples in an initial test effectively identified lines with differing phenotypic expression of QRIP in a subsequent environment. This is best illustrated by comparing means of the five lines selected for highest QRIP (88,400 units/g FW in 1996 and 103,100 units/g FW when regrown in 1997) to means of the five lines selected for lowest potential (9,900 units/g FW in 1996, and 33,700 units/g FW when regrown in 1997). The 1996 QRIP mean of all entries was 34,000 units/g FW and the 1997 mean was 63,200 units/g FW. Higher QRIP values for entries in 1997 compared to 1996, are probably due in part to environmental factors. These differences between years are also partially explained by the fact that individuals selected to be grown in 1997 had QRIP levels in 1996 that were evenly distributed along the range of values for the trait; whereas, the 1996 sample was weighted with entries that had moderate to low levels of induction potential. However, it is clear that differential expression between high and low QRIP inbreds was consistent in both years. MSAG concentration differentials between the above-described five high versus five low lines were also concordant: 2.13, 0.35, 2.13, and 0.58 µmol·g⁻¹ FW for the high and low lines of 1996 and 1997, respectively. The QRIP bioassay thus identifies lines divergent for MSAG concentration.

Few studies have measured the genetic versus environmental contribution to MSAG content of broccoli. Carlson et al. (1987) and Kushad et al. (1999) found very low levels of GI compared to GR, and significant differences among broccoli entries for GR concentration, but looked only at a single environment. Shelp et al. (1993) grew two cultivars of broccoli in several environments, reported only GR levels, and found a significant effect of genotype and environment on the concentration of this glucosinolate. However, those authors utilized two relatively early maturing cultivars, Commander and Baccus, and the magnitude of differences between the two cultivars was small compared to the ranges observed by Carlson et al. (1987), Kushad et al. (1999), and those described herein. Conclusions from the present study are based on only two autumn harvest environments, but results indicate a significant role of genotype in expression of GR levels, in broccoli heads.

It is noteworthy, that Kushad et al. (1999) evaluated inbred lines developed by one of the present authors (M.W. Farnham), citing ‘VI158’ as a line with about six times more GR than ‘EV2-1’. In our 1996 trial, line ‘VI158’ (USVL068) had ≈7-fold higher GR than USVL003, which is a sister line of ‘EV2-1’. Inbreds USVL066 and USVL069, which exhibited the highest QRIP and MSAG levels in this study are closely related to ‘VI158’ and are derived from ‘Viking’. In addition, several inbreds, such as USVL003 and USVL013, with low QRIP and MSAG in the trials reported herein, as well as ‘EV2-1’, are all derived from ‘Everest’. These observations provide circumstantial evidence that geno-

![Fig. 4. (A) Mean quinone reductase induction potential (QRIP) and (B) methylsulphinylalkyl glucosinolate (MSAG) concentration of entries grown in 1997 (y-axis) as a function of entry phenotype in 1996 (x-axis). Vertical bar in each graph represents the LSD₀.₀₅ for differentiating entry means in 1997.](image-url)

Fig. 4. (A) Mean quinone reductase induction potential (QRIP) and (B) methylsulphinylalkyl glucosinolate (MSAG) concentration of entries grown in 1997 (y-axis) as a function of entry phenotype in 1996 (x-axis). Vertical bar in each graph represents the LSD₀.₀₅ for differentiating entry means in 1997.
Table 1. Entry means for quinone reductase induction potential (QRIP), methylsulphynyl-alkyl glucosinolate (MSAG) concentration, and number of days from transplant to harvest (DTH) for the high parent (HP) and low parent (LP), and the F1 formed by crossing the two parents for four different hybrid combinations.

| Cross | Entry type | QRIP (units/g FW) | MSAG (µmol·g−1 FW) | DTH (d) |
|-------|------------|------------------|--------------------|---------|
| 1     | HP         | 106,000          | 1.65               | 103     |
|       | LP         | 27,700           | 0.42               | 49      |
|       | F1         | 45,800           | 1.25               | 62      |
| 2     | LP         | 63,900           | 1.07               | 86      |
|       | F1         | 42,100           | 1.17               | 76      |
| 3     | LP         | 45,200           | 1.01               | 85      |
|       | F1         | 36,400           | 0.69               | 59      |
| 4     | LP         | 51,700           | 1.32               | 87      |
|       | F1         | 46,900           | 0.66               | 69      |
|       | F1         | 57,600           | 1.11               | 65      |

LSD 0.05 24,700 0.60 6

\(^{3}\text{FW} = \text{fresh weight.}\)

Table 1 entry means for QRIP, MSAG, and DTH.

The bioassay for QRIP was used initially to identify chemoprotective compounds, both natural and synthetic (Prochaska et al., 1992), and has been used successfully to identify Phase 2 enzyme inducing constituents in broccoli (sulforaphane), red wine (resveratrol), broccoli sprouts (GR), and tomatillos (withanthiolides). (Fahey et al., 1997; Jang et al., 1997; Kennelly et al., 1997; Zhang et al., 1992). It has also been used to evaluate the potency of various purified phytochemicals and synthetic compounds (Dinkova-Kostova and Talalay, 1999; Khachick et al., 1999; Posner et al., 1994).

Type plays a significant role in expression of GR content, and thus, QRIP in broccoli.

Giamoustaris and Mithen (1996) provided evidence for a model in *Brassica* L., whereby two primary loci control the balance of alkyl glucosinolate forms in this genus. Their model suggests that the Gsl-elong locus controls elongation of the glucosinolate amino acid-derived side chain, and that the Gsl-alk locus controls formation of alkyl glucosinolates from alkyl glucosinolates. They hypothesize a functional Gsl-elong locus and a null allele at the Gsl-alk locus in broccoli that would result in accumulation of GR. The evidence for this model is very strong, but it does not explain entirely the wide variation in broccoli head GR levels that have been observed by others and are described herein. It is likely that genes other than those hypothesized, also influence MSAG levels in broccoli.

Nearly all broccoli consumed in the United States is harvested from hybrid production fields. Although it is important to evaluate Phase 2 induction potential and GR content of inbreds, attention must be focused on expression of these traits in hybrid combinations. Thus, the current study, a limited sample of hybrids formed from USVL inbreds were examined in order to identify possible trends. In most cases, F1 hybrid means for QRIP and MSAG concentration were intermediate between the means of the two parents but closer to the mean of the low parent. A similar trend was observed for DTH, although with two crosses, F2 hybrid means were actually lower than means of the low parents. These observations for DTH reported herein are consistent with results of others (Borchers, 1968; Hulbert and Orton, 1984) who have concluded that days to maturity in broccoli is primarily a consequence of additive gene action with some contribution of dominance to variance that tends in the direction of earliness. Similar, although less consistent observations for the correlation of QRIP and MSAG with DTH may also point to a close relationship between maturity and the level of GR in broccoli heads, which in turn controls QRIP of this vegetable. Future studies must examine a larger sample of F1 hybrids made by crossing inbreds of known QRIP and MSAG level to determine if trends for these characters observed herein are consistent in a larger population of hybrid crosses and whether gene action similar to that for DTH might control QRIP and MSAG as well.

The bioassay for QRIP was used initially to identify chemoprotective compounds, both natural and synthetic (Prochaska et al., 1992), and has been used successfully to identify Phase 2 enzyme inducing constituents in broccoli (sulforaphane), red wine (resveratrol), broccoli sprouts (GR), and tomatillos (withanthiolides). (Fahey et al., 1997; Jang et al., 1997; Kennelly et al., 1997; Zhang et al., 1992). It has also been used to evaluate the potency of various purified phytochemicals and synthetic compounds (Dinkova-Kostova and Talalay, 1999; Khachick et al., 1999; Posner et al., 1994).

Results of the present study demonstrate a high correlation between QRIP and direct paired-ion HPLC of plant extracts for quantitation of GR. Although the use of analytical chemical assays (e.g., paired-ion HPLC) to quantify GR in broccoli has proven a straightforward means of analysis, our results indicate that assessment of the Phase 2 (QR) induction potential of broccoli might be an additional and useful character on which to focus in future efforts to enhance chemoprotective attributes of broccoli cultivars.

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