



Published in final edited form as:

*Pediatr Blood Cancer*. 2013 April ; 60(4): 593–598. doi:10.1002/pbc.24366.

## Glutathione S-Transferase P1 Single Nucleotide Polymorphism Predicts Permanent Ototoxicity in Children with Medulloblastoma

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### Abstract

**Background**—Glutathione S-transferase (GST) enzymes are involved in detoxifying chemotherapy and clearing reactive oxygen species formed by radiation. We explored the relationship between the host *GSTP1* 105 A>G polymorphism (rs1695), tumor GSTpi protein expression, and clinical outcomes in pediatric medulloblastoma. We hypothesized that the *GSTP1* 105 G-allele and increased tumor GSTpi expression would be associated with lower progression-free survival and fewer adverse events.

**Procedure**—The study included 106 medulloblastoma/primitive neuroectodermal tumor (PNET) patients seen at Texas Children’s Cancer Center. Genotyping was performed using an Illumina HumanOmni1-Quad BeadChip and GSTpi expression was assessed using immunohistochemistry. We used the Kaplan-Meier method for survival analyses and logistic regression for toxicity comparisons.

**Results**—Patients with a *GSTP1* 105 AG/GG genotype (vs. AA) or who had received high dose craniospinal radiation (≥ 34 Gy vs. <26 Gy) had a greater risk of requiring hearing aids than their counterparts (OR 4.0, 95% CI 1.2-13.6, and OR 3.1, 95% CI 1.1-8.8, respectively, n=69).

Additionally, there was a statistically significant interaction between these variables. Compared with the lowest risk group (*GSTP1* 105 AA-low dose radiation), patients with a *GSTP1* 105 AG/GG genotype who received high dose radiation were 8.4 times more likely to require hearing aids (95% CI 1.4-49.9, p-trend=0.005, n=69). When adjusted for age, cumulative cisplatin dose, and amifostine use, the association remained.

**Conclusions**—The *GSTP1* 105 G-allele is associated with permanent ototoxicity in pediatric medulloblastoma/PNET and strongly interacts with radiation dose. Patients with this allele should be considered for clinical trials employing radiation dose modifications and cytoprotectant strategies.

### Keywords

Glutathione S-Transferase; Polymorphism; Radiation; Ototoxicity; Medulloblastoma

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Conflict of Interest Statement

The authors have no conflicts of interest.

## Introduction

Medulloblastoma is the most common malignant brain tumor in pediatrics, accounting for 20% of central nervous system malignancies in the United States [1,2]. Treatment in children at least three years of age consists of surgical resection, craniospinal radiation (CSI), and chemotherapy [3,4]. This produces long-term overall survival of 60-80% [5]. Despite improving survival, many individuals experience therapy-related morbidities, including neurocognitive impairment, endocrinopathies, and hearing loss [5,6]. Well established prognostic criteria for survival include age and risk grouping - based on the initial disease extent and the degree of surgical resection [1,5]. More recently, the presence of distinctive disease subtypes characterized by molecular classification schemes has been proposed [7,8]. However, for toxicities, it is largely uncertain which patients will develop them and how severe they will be.

One group of enzymes that deserves more attention in medulloblastoma is the glutathione S-transferase (GST) family. GSTs detoxify electrophilic compounds via conjugation of reduced glutathione. These compounds include mutagenic and carcinogenic materials, chemotherapy agents and their metabolites, and free radicals [9]. Genetic variation in these enzymes, including *GSTP1* 105 A>G SNP (rs1695) and whole gene deletions in *GSTM1* and *GSTT1*, have been associated with survival and toxicities in other brain tumors and other cancers treated with platinum agents and radiation [10-15]. Additionally, higher tumor GSTpi expression has been correlated with worse prognosis in adults with high grade glioma and other platinum-responsive malignancies [16-18]. Little is known about the role of *GSTP1* polymorphisms and tumor GSTpi expression in medulloblastoma outcomes.

In this study, we assessed the relationship between the germline *GSTP1* 105 A>G polymorphism, tumor GSTpi expression, and survival and the occurrence of toxicities in patients with medulloblastoma/PNET seen at our institution. We hypothesized that the presence of a G-allele at the *GSTP1* 105 site and increased tumor GSTpi expression would be associated with lower progression-free survival and reduced therapy-related toxicities.

## Methods

### Study population

There were 106 patients with medulloblastoma or supratentorial primitive neuroectodermal tumor (PNET) seen at Texas Children's Cancer Center included in this study. Peripheral blood samples were obtained for genotyping from 86 patients. Tumor samples were obtained for immunohistochemical (IHC) staining from 51 patients. Paired peripheral blood and tumor samples were available on 32 patients. The parents/legal guardians of all subjects provided informed consent for an IRB-approved institutional protocol for SNP-cancer outcome associations. Patients were diagnosed between March 1987 and July 2009 and were younger than 19 years at diagnosis. Seventy patients (66%) were treated on the SJMB96/03 or CCG-A9961 protocols with the remainder mostly treated with vincristine, cisplatin, and cyclophosphamide based regimens [3,4]. Medical record abstractions were performed for patient and therapy characteristics and outcomes including survival events and the occurrence of acute toxicities/late effects (i.e., present or absent). Acute toxicities included were myelosuppression, renal insufficiency, and neurotoxicity. They were deemed present if they led to at least one instance of chemotherapy dose reduction, delay, or omission. Late effects included were hypothyroidism, growth hormone deficiency, chronic renal insufficiency, and ototoxicity. These were defined as present if they were noted in the medical record greater than one year after the end of the initial therapy regimen and led to an intervention. These interventions were defined for hypothyroidism and growth hormone

deficiency as hormone supplementation, for chronic renal insufficiency as ongoing nephrology care, and for ototoxicity as the need for hearing aids.

### Laboratory methods

Peripheral blood samples were collected in EDTA vacutainers and genomic DNA was extracted from mononuclear cells using the QIAamp DNA blood mini kit (Qiagen, Valencia, CA) according to the manufacturer's protocol. DNA concentrations were determined using the PICOgreen method and DNA quality was determined by gel electrophoresis of extracted products. *GSTP1* 105 A>G genotyping was performed as part of another study involving these subjects. For that study, the Illumina HumanOmni1-Quad BeadChip (Illumina, San Diego, CA) was used according to the manufacturer's protocol.

Immunohistochemical staining of formalin-fixed, paraffin-embedded tumor samples was performed to assess GSTpi expression. Sections were rehydrated in PBS and then incubated with a polyclonal rabbit anti-human GSTpi antibody at a 1:500 dilution. The slides were then rinsed with PBS and incubated with an avidin-conjugated mouse anti-rabbit antibody. After further PBS rinsing, the slides were treated with a biotinylated peroxidase solution (Vector Laboratories, Burlingame, CA) and developed with 0.05% diaminobenzidine and 0.01% H<sub>2</sub>O<sub>2</sub> in 50 mM Tris-HCL buffer (pH 7.5). GSTpi expression was quantified for staining intensity and proportion of positive cells. Intensity was assessed in 600 cells (200-fold magnification). Based on the cytoplasmic staining intensity of at least 70% of cells, the intensity was categorized in quartiles. In the same microscopic fields, the proportion of cells expressing GSTpi was evaluated and also categorized into quartiles. An IHC score was defined as the product of the category values for the staining intensity and proportion.

### Statistical analyses

Descriptive analyses were performed for patient, disease, and therapy characteristics, the *GSTP1* 105 A>G genotype, and the GSTpi IHC staining variables. The consistency of the genotypes with Hardy-Weinberg equilibrium and published results was assessed. All SNP analyses were performed using co-dominant and dominant modeling. Spearman rank correlation methods were used to assess the correlation,  $r_s$ , between the genotyping and IHC staining. Since radiation therapy is typically delayed, reduced, or omitted for patients younger than three years, separate survival analyses were performed for patients diagnosed before three years of age. Survival analyses were performed using the Kaplan-Meier method with log-rank comparisons. Progression-free survival (PFS) and overall survival (OS) were calculated from the date of diagnosis to progression/recurrence or death, respectively, or for survivors, the date of the last contact.

Patients of all ages were included in the toxicity analyses with some exceptions: 1) those having the outcome of interest as a pre-existing condition (e.g., pre-therapy hearing loss), 2) those not receiving toxicity-inducing therapies (e.g., CSI unexposed subjects were excluded from endocrinopathy analyses), 3) and those lacking adequate medical documentation for evaluation of the toxicity. Toxicity analyses were performed with Chi-Square and Fisher's Exact tests, as appropriate. Odds ratios with 95% confidence intervals were calculated using univariate logistic regression; multivariable logistic regression models were constructed in a forward selection manner. For the ototoxicity outcome, exploratory interaction analyses were conducted by creating combined genotype-CSI dose groups and comparing them with the Chi-Square test of homogeneity and score test for trend. These analyses were performed unadjusted and adjusted for age group (divided at median), cumulative cisplatin dose group (divided at median=300 mg/m<sup>2</sup>), and amifostine use (yes or no). Cumulative cisplatin doses were standardized using a 1 mg/kg=30 mg/m<sup>2</sup> conversion. Statistical significance was defined as a p-value less than 0.05 and all tests of significance performed were two-sided.

No adjustments for multiple testing were implemented since only a single polymorphism was studied. For all analyses, the Stata 11 software package was used (StataCorp LP, College Station, TX).

## Results

### Patient, disease, and therapy characteristics

The patient, disease, and therapy characteristics are summarized in Table I. The median follow-up in survivors was 7.3 years (range 1.7-23.5 years, n=78). The mean and median ages at diagnosis were 7.0 and 6.0 years, respectively (range 0.5-18 years, n=106). Eighty-six patients (81%) were at least three years of age at diagnosis. Seventy-four patients (70%) were male and 47 (45%) were non-Hispanic white. In addition to 97 patients with medulloblastoma, 9 patients with supratentorial PNET were included. Fifty-two patients (61%) were classified as average-risk and 33 patients (38%) were high-risk, including the 5 patients with supratentorial PNET who were at least three years of age. All high-risk patients received high dose CSI (median 36 Gy, range 34.2-39.6 Gy), while 50/52 (96%) of the average-risk patients received low dose CSI (median 23.4 Gy, range 18-25.2 Gy).

### Genotyping and immunohistochemical staining

The frequencies of AA, AG, and GG were 0.37 (32/86), 0.41 (35/86), and 0.22 (19/86), respectively, yielding a minor allele frequency (MAF) for the G-allele of 0.42. These data are consistent with Hardy-Weinberg equilibrium and with allelic frequencies reported in public databases ([http://www.ncbi.nlm.nih.gov/projects/SNP/snp\\_ref.cgi?rs=1695](http://www.ncbi.nlm.nih.gov/projects/SNP/snp_ref.cgi?rs=1695)). For the IHC staining, samples were more likely to have absent/mild staining intensity (59%) and over 50% of the cells staining positive (55%) (n=51). The *GSTP1* 105 A>G genotype did not correlate with the GSTpi expression variables ( $-0.1 < r_s < 0.05$ ;  $p = 0.48$ ).

### Survival

Survival was assessed by age and risk grouping (Table II). Eighty-six patients were at least three years of age at diagnosis and 20 were less than three years. Those younger than three years had significantly worse 5-year PFS, but not 5-year OS ( $p=0.001$  and  $0.12$ , respectively). For risk grouping, the 5-year PFS and OS did not significantly differ between high-risk (n=33) and average-risk (n=52) categories ( $p=0.22$  and  $0.74$ , respectively).

In patients at least 3 years of age, there were no significant differences in PFS based on the *GSTP1* 105 A>G genotype (n=68) or GSTpi expression (n=46) in univariate and multivariable analyses. Among the 41 patients in the *GSTP1* 105 AG/GG group, there were 7 progressions (5-year PFS 80.7, 95%CI 63.3-90.4) and in the 27 patients with the AA genotype, there were 2 progressions (5-year PFS 92.2%, 95%CI 72.2-98.0), ( $p=0.38$ ).

### Toxicities

In univariate analyses, only the *GSTP1* 105 A>G genotype and craniospinal radiation dose were significantly associated with hearing loss requiring hearing aids (n=69). When compared to patients with the AA genotype (n=24), patients with the AG (n=30) and GG (n=15) genotypes had greater risks of severe ototoxicity ( $OR_{AG/AA} 3.8$ , 95%CI 1.0-14.9;  $OR_{GG/AA} 4.4$ , 95%CI 0.9-21.3,  $p=0.04$ ) (Figure 1). With dominant modeling, patients in the AG/GG group had a greater risk of severe hearing impairment when compared to patients with the AA genotype ( $OR 4.0$ , 95%CI 1.2-13.6,  $p=0.03$ ) (Figure 2). Compared to those receiving low dose CSI, patients who received high dose, were more likely to experience hearing loss requiring hearing aids ( $OR 3.1$ , 95%CI 1.1-8.8,  $p=0.04$ ) (Figure 3). Ototoxicity did not differ by cumulative cisplatin dose ( $OR_{high/low} 0.3$ , 95%CI 0.1-1.2,  $p=0.08$ ; n=67) or amifostine use ( $OR_{yes/no} 0.5$ , 95%CI 0.2-1.5,  $p=0.2$ , n=69). Multivariable analyses showed

elevated ORs for *GSTP1* 105 A>G genotype and CSI dose without statistical significance (OR<sub>SNP</sub> 3.5, 95%CI 0.99-12.1, p=0.05; OR<sub>CSI</sub> 2.6, 95%CI 0.89-7.8, p=0.08; n=69).

To explore a potential interaction between the *GSTP1* 105 A>G genotype and CSI dose, patients were divided into four groups using all 2x2 combinations, ordered by a proposed risk for severe ototoxicity (n=69) (Table III). In the lowest risk group were individuals with the AA genotype who received low dose CSI and in the highest risk group were those in the AG/GG group who received high dose CSI. Using the lowest risk group as a reference, a significant dose-response pattern was observed (OR<sub>AA-High</sub>=1.3, 95%CI 0.1-17.4; OR<sub>AG/GG-Low</sub>=2.7, 95%CI 0.6-12.1; OR<sub>AG/GG-High</sub>=8.4, 95%CI 1.4-49.9, p-homogeneity=0.03, p-trend=0.005). The trend remained significant when adjusting for age group, cumulative cisplatin dose group, and amifostine use (p=0.02, n=67).

Of the 86 patients genotyped, 17 patients were excluded from the ototoxicity analyses: 4 with pre-existing hearing loss, 9 who were lost to follow-up prior to one year post-primary therapy, and 4 who received focal radiation alone. Among the 69 patients in the ototoxicity analyses, 68 (99%) received cisplatin and 69 (100%) received CSI with 24 of 69 patients (35%) requiring hearing aids. In analyses involving the cumulative cisplatin dose, 2 children, for whom dose information was unavailable, were excluded. The median cumulative cisplatin dose was 300 mg/m<sup>2</sup> (range 0-720 mg/m<sup>2</sup>). The mean cumulative cisplatin doses for those requiring hearing aids (302.6 mg/m<sup>2</sup>, 95%CI 263.1-342.1) and those not (328.5 mg/m<sup>2</sup>, 95%CI 294.4-362.1) were not statistically different (p=0.31).

There were no differences in other toxicities based on the *GSTP1* 105 A>G genotype. Additionally, no relationship between the toxicities and the GSTpi IHC score was demonstrated.

## Discussion

We observed that patients in the *GSTP1* 105 AG/GG group were 4 times more likely to require hearing aids than those with the AA genotype. Moreover, exploratory analyses, based on combinations of the *GSTP1* 105 A>G genotype and CSI dose group, suggests a novel interaction, characterized by the risk allele conferring greater risk in individuals treated with high radiation doses than those treated with low doses.

The primary mechanism of platinum-induced ototoxicity occurs through the generation of free radicals, leading to apoptosis of the cochlear outer hair cells [19]. This produces sensorineural hearing loss (SNHL) with the high frequencies most likely affected. The hearing loss is typically permanent and may occur during therapy or in the months afterwards. Months to years later, radiation therapy may also cause SNHL, usually high-frequency, via cochlear injury mediated by reactive oxygen species [20,21]. Genetic variations in GSTs, a major enzyme family involved in clearing free radicals, are an ideal mechanism to explain the differences in ototoxicity susceptibility after medulloblastoma therapy. Supporting the role of GSTs in medulloblastoma toxicities is a prior study from our group. We demonstrated that children with medulloblastoma (n=42) having *GSTM1* and *GSTT1* deletions were at significantly greater risk for any acute toxicity (grade 3) requiring therapy modification (1 null genotype: OR 6.4, 95% CI 1.2-34) and cognitive impairment (27.2 point post-radiation full-scale IQ drop, p=0.002) [22]. No association between ototoxicity and *GSTM1* and *GSTT1* deletions was detected.

While the influence of the *GSTP1* 105 A>G genotype on ototoxicity in medulloblastoma has not been previously explored, its role in ototoxicity in other malignancies has been studied. In 173 adults with testicular cancer treated with cisplatin-based regimens, hearing was evaluated with the threshold for impairment set at the 75<sup>th</sup> percentile, based on Norwegian

men in their fifties in the general population [23]. The investigators found a greater risk of hearing loss in the AG/AA group when compared to those with the GG genotype (OR 4.3, 95% CI 1.3-14.4). The relationship between the *GSTP1* 105 A>G genotype and ototoxicity was evaluated in two other studies with no association discovered [24,25]. In both studies, children and young adults with cancer (e.g., osteosarcoma, neuroblastoma, brain tumors) treated with cisplatin-based regimens were assessed.

None of the above studies have compared the role of *GSTP1* genotype between those who received radiation and those who did not. Moreover, few studies have assessed the role of the *GSTP1* 105 A>G genotype on radiation-associated toxicities. Two such studies were conducted in women with breast cancer who received adjuvant radiation [26,27]. A greater risk of acute skin toxicities was reported in women with the *GSTP1* 105 GG genotype (HR 2.3, 95% CI 1.04-5.0) and a slightly greater proportion of late pleural thickening was observed in women with the GG genotype (OR 1.1, p=0.004). More recently, in patients with esophageal adenocarcinoma treated with cisplatin-based radiochemotherapy, those in the *GSTP1* 105 AG/GG group were more likely to experience grade 3-4 radiation-attributable toxicities (e.g., dysphagia) than those with the AA genotype (31% vs. 0%, p=0.005) [28].

The *GSTP1* 105 A>G SNP is characterized by an adenine or guanine at position 313 resulting, respectively, in an isoleucine or valine at codon 105. Its functional significance varies by the exposure with the G-allele conferring increased GSTpi activity to platinum agents *in vitro* [29]. The variable metabolism of cisplatin by *GSTP1* 105 A>G genotype may not be mediated by the GSTpi free radical detoxification activity, but rather may depend on interactions with other pathways [30]. For example, JNK/SAPK pathway activity, which may be modulated by GSTpi expression, has been shown to influence cisplatin-mediated cell death. There are no data on the functional implications of the *GSTP1* 105 A>G genotype on radiation-associated reactive oxygen species, although it may be inferred from clinical studies that the G-allele leads to decreased GSTpi activity [26-28].

Our finding of the *GSTP1* 105 G-allele as a risk factor for severe ototoxicity in children with medulloblastoma/PNET may appear to contradict results observed by others [23,29]. However, previous clinical investigations included subjects on cisplatin-based regimens without radiation, a central component of medulloblastoma therapy [23-25]. Additionally, prior *in vitro* functional studies were performed with cisplatin alone, without radiation, which may interact with GSTpi [29,31]. We found that severe ototoxicity is associated with the combined exposure of the *GSTP1* 105 G-allele with high dose CSI. This is consistent with the clinical evidence showing that the G-allele is associated with greater radiation-related toxicities [26-28]. We did not find an association between severe hearing loss and the cumulative cisplatin dose. Several groups have suggested that the correlation between ototoxicity and the cumulative cisplatin dose is more pronounced above 400 mg/m<sup>2</sup> [32,33]. The importance of CSI as opposed cisplatin in our study may be partly explained by the relatively lower cumulative cisplatin doses to which our population was exposed (median 300 mg/m<sup>2</sup>). Moreover, cisplatin dose modifications, which were widespread in our study, confounded the association between cumulative cisplatin dose and ototoxicity.

We also evaluated the relationship between tumor GSTpi expression and outcomes in medulloblastoma/PNET and found no associations. IHC staining may not be the best method of assessing GSTpi activity, particularly when using tumor tissue in which the activity may be dysregulated. Instead, substrate-specific activity assays in normal tissue may be preferable.

Hearing loss remains a significant problem for patients with medulloblastoma. As many as half experience impaired hearing with roughly one-quarter of all patients needing hearing aids [34]. Hearing loss can substantially diminish quality of life. Cisplatin dose modifications for early ototoxicity, conformal radiation techniques, and amifostine prophylaxis have decreased this impairment, but have not eliminated it [35,36]. If our finding of a *GSTP1* 105 A>G SNP-ototoxicity link and its association with radiation dose can be validated, we can identify a group of patients at elevated risk for severe hearing loss. With this group, more sensitive screening methods for early ototoxicity may be applied, radiation dose reductions may be considered, or more targeted cytoprotectant strategies than are currently being employed with amifostine may be implemented.

This study is the first evaluating the relationship between the germline *GSTP1* 105 A>G genotype, tumor GSTpi expression, and ototoxicity in patients with medulloblastoma/PNET. Our population was largely uniformly treated with approximately 66% treated on the SJMB96/03 or CCG-A9961 protocols with the remainder mostly treated with regimens including the same chemotherapy. Additionally, with a median follow up in survivors of over 7 years, adequate time elapsed to assess the outcomes of interest. The analyses accounted for the potential confounders of age, cumulative cisplatin dose, and amifostine use. Although genetic admixture was not included in the analyses, the patients were almost exclusively non-Hispanic Caucasian, Hispanic, or African-American and the reported allelic frequencies for these three groups is very similar (MAF 39-42%) and was consistent with those observed in our study.

Our study had several limitations. This was a retrospective study spanning over two decades, during which time radiation techniques have progressed and toxicity monitoring and interventions have evolved. Also, radiation oncology reports specifying the cochlear doses were not available for a substantial proportion of the subjects. However, since most patients received conformal radiation, it was presumed that the radiation dose to the cochlea could be reasonably approximated by the minimum dose to which all tissues in the craniospinal axis were exposed (i.e., the CSI dose). The need for using the CSI dose as a surrogate for the cochlear dose will be avoided in our future prospective studies. Additionally, while patients with known pre-existing hearing loss were excluded from the analyses, it is possible that some patients with prior mild hearing loss were included due to lacking documentation. Lastly, the study sample size was small and limited the precision of our statistical estimates.

In the future, we plan to more extensively investigate the relationship between clinical outcomes in medulloblastoma and glutathione S-transferases and other detoxification/free radical clearance enzymes including glutathione peroxidases, glutathione reductase, superoxide dismutases, catalase, and metallothioneins. These studies will be performed with a larger, independent sample of patients and may provide a more comprehensive understanding of the role that detoxification and free radical clearance enzymes play in medulloblastoma outcomes and may ultimately influence future therapeutic strategies.

## Acknowledgments

This work was supported by research grants from the National Institutes of Health (P50CA127001-04), the Gillson Longenbaugh Foundation, the John S. Dunn Research Foundation, the Robert J. Kleberg, Jr. and Helen C. Kleberg Foundation, and the American Brain Tumor Association.

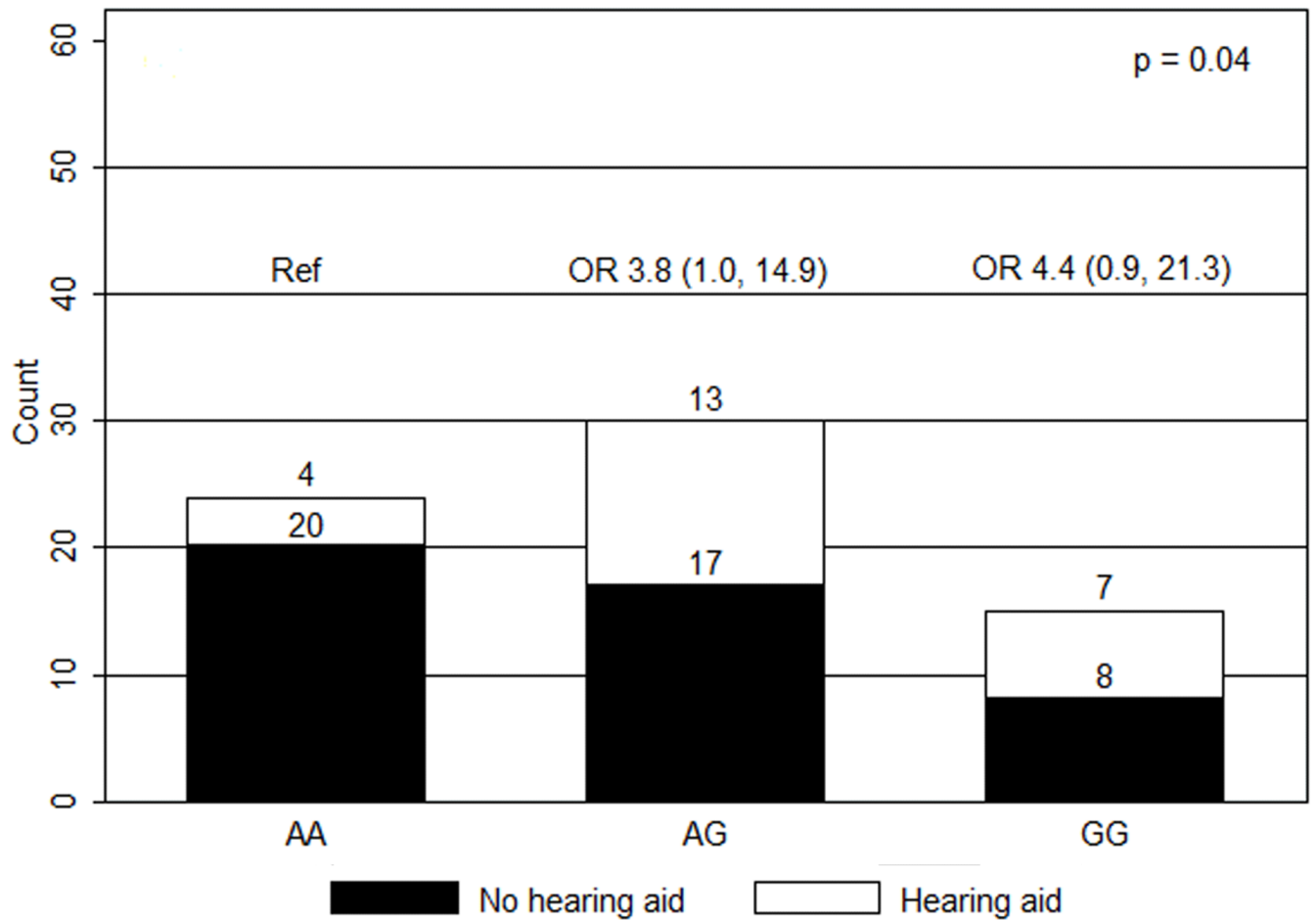
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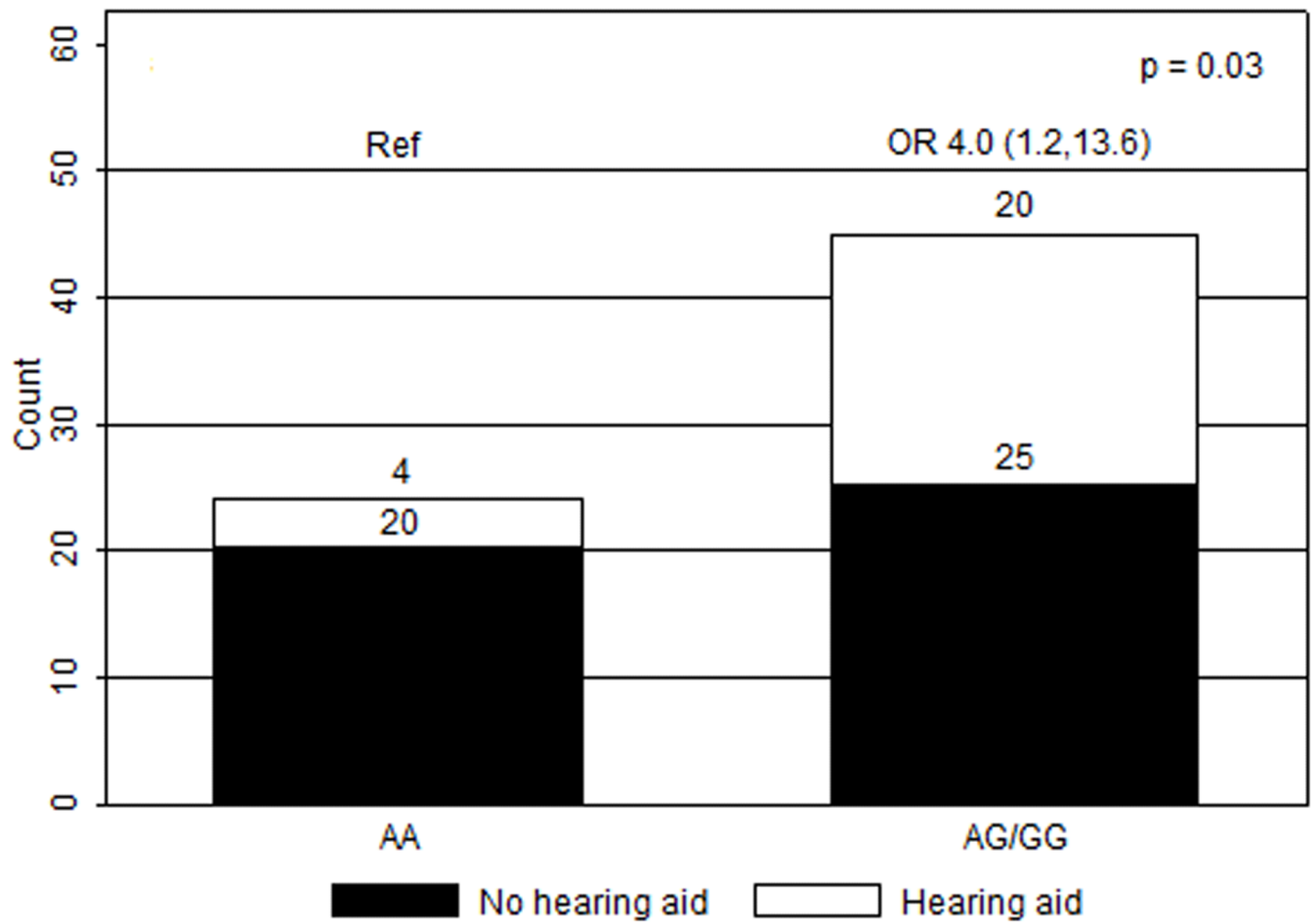
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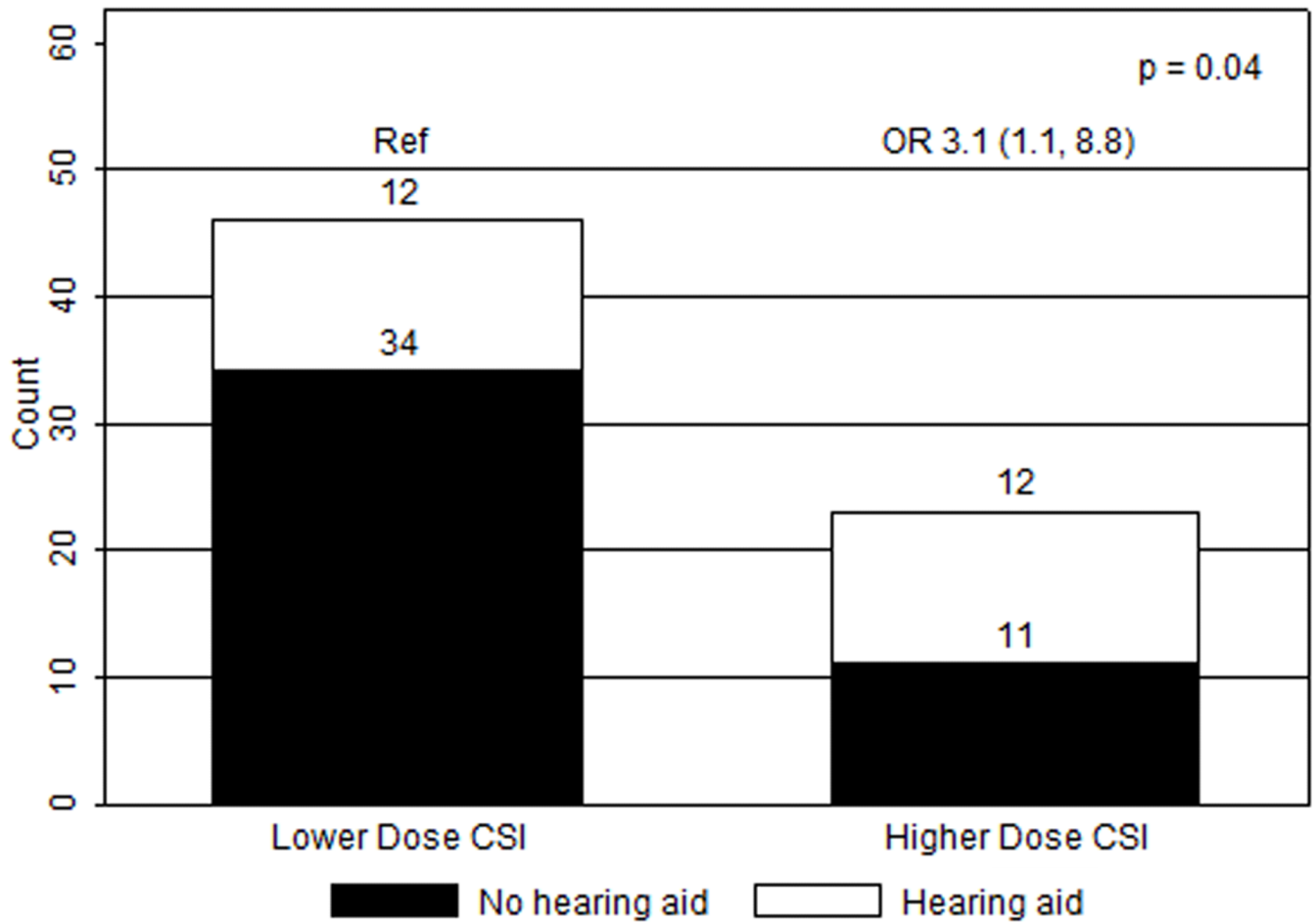
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**Figure 1.** Ototoxicity by GSTP1 105 A>G genotype (co-dominant model). OR, odds ratios with 95% confidence intervals in parentheses; Ref, reference group.



**Figure 2.** Ototoxicity by GSTP1 105 A>G genotype (dominant model). OR, odds ratios with 95% confidence intervals in parentheses; Ref, reference group.



**Figure 3.** Ototoxicity by craniospinal radiation (CSI) dose group. Low dose CSI, 18 - 25.2 Gy (median 23.4 Gy); high dose CSI, 34.2 - 39.6 Gy (median 36 Gy); OR, odds ratios with 95% confidence intervals in parentheses; Ref, reference group.

**Table I**  
**Clinical Data for Study Patients**

Variable	All Patients (n = 106)		Patients 3 years (n = 86)	
	N	%	N	%
Study subset				
Genotyping	86	81	68	79
IHC Staining	51	48	46	53
Overlap	32	30	29	34
Gender				
Male	74	70	60	70
Female	32	30	26	30
Race/Ethnicity				
Non-Hispanic white	47	44	38	44
Hispanic	37	35	28	33
Other	22	21	20	23
Disease				
Medulloblastoma	97	92	81	94
PNET	9	8	5	6
Risk				
High	NA	NA	33	38
Average	NA	NA	52	61
Unknown	NA	NA	1	1
Radiation Dose				
High	42	39	35	41
Low	57	54	50	58
Focal only	3	3	0	0
None	3	3	0	0
Unknown	1	1	1	1
Chemotherapy				
SJMB/CCG-A9961	70	66	68	79
Other regimen	34	32	16	19
None	2	2	2	2

IHC, immunohistochemistry; PNET, primitive neuroectodermal tumor; Craniospinal radiation dose: high, 34.2-39.6 Gy (median 36 Gy), low, 18-25.2 Gy (median 23.4 Gy); Chemotherapy (other regimen), non-SJMB96/03 and non-CCG-A9961 treatment regimen.

**Table II**  
**5-year Progression-Free Survival and 5-year Overall Survival for Known Prognostic Factors in Medulloblastoma**

Group	N	Events	PFS	95%CI	p	Events	OS	95%CI	p
3 years	86	18	77.5	66.5 - 85.3	0.001	14	80.1	68.3 - 87.9	0.12
< 3 years	20	10	48.1	24.7 - 68.3		6	66.4	38.9 - 83.7	
High-risk	33	9	70.9	51.1 - 83.8	0.22	6	79.2	59.1 - 90.1	0.74
Average-risk	52	9	81.3	67.0 - 89.8		8	79.5	61.8 - 89.6	

PFS, 5-year progression-free survival; OS, 5-year overall survival.

**Table III**  
**Risk of Severe Ototoxicity by Combination of GSTP1 105 A>G Genotype and Craniospinal Radiation Dose Group**

Genotype - CSI dose group	N (hearing aid)	N (No hearing aid)	OR	95%CI
AA-Low	3	16	1 (Ref)	NA
AA-High	1	4	1.3	0.1 - 17.4
AG/GG-Low	9	18	2.7	0.6 - 12.1
AG/GG-High	11	7	8.4	1.4 - 49.9

CSI, craniospinal radiation dose: high 34.2-39.6 Gy (median 36 Gy), low, 18-25.2 Gy (median 23.4 Gy); Ref, reference group. In crude analyses (above), p-homogeneity=0.03 and p-trend=0.005. After adjusting for age group, cumulative cisplatin dose group, and amifostine use, p-trend=0.02.