Study Number	ACNS0331		
NCT#	NCT00085735		
Dataset #	D1		
PMID#	34110925		

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Variable Number	Variable Name	Type	Label	Notes
1	USI	Character	Universal Specimen Identifier (USI)	Patient identifier
2	Gender	Character	Gender	
3	Race	Character	Race	
4	AgeOnStudy_years	Numeric	Age at time of study enrollment (in years)	
5	AgeAtDiagnosis_years	Numeric	Age at time of diagnosis (in years)	
6	AgeGroup	Character	Age group at time of study enrollment	This variable equates to stratum in protocol document (3 to 7 years or 8 to 21 years)
7	RTGroup	Character	Radiation group	Involved field radiation (IFRT) or Posterior fossa radiation (PFRT)
8	CSIgroup	Character	Craniospinal irradiation (CSI) group	Low-dose craniospinal irradiation (LDCSI) or Standard-dose craniospinal irradiation (SDCSI)
9	efs	Numeric	Event-free survival (EFS) duration from study enrollment (in days)	
10	efscen	Numeric	Event-free survival censor variable	0=censored, 1=patient had an EFS event
11	os	Numeric	Overall survival (OS) duration from study enrollment (in days)	
12	oscen	Numeric	Overall survival censor variable	0=censored, 1=patient had an OS event/death
13	IsPatientEligible	Character	Is the patient eligible?	
14	ReasonNotEligible	Character	Reason not eligible	A reason not eligible is only provided for ineligible patients (observations where IsPatientEligible=No)

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15	IsPatientEvaluable	Character	Is the patient evaluable for the primary study objectives?	NA denotes ineligible patients
16	ReasonNotEvaluable	Character	Reason not evaluable for primary study objectives	A reason not evaluable is only provided for inevaluable patients (observations where IsPatientEvaluable =No)
17	IsPatientEligandEval	Character	Is the patient both eligible and evaluable for primary study objectives?	For most analyses, only eligible and evaluable patients are included.
18	WasPtMolecularlyProfiled	Character	Was the patient molecularly profiled?	
19	Subgroup	Character	Subgroup based on methylation	NA denotes inconclusive
20	SubgroupForTable1	Character	Subgroup based on methylation as presented in Table 1	In this variable, the following subgroups are recategorized as "Other CNS" to facilitate reporting in Table 1 and Figure 2A: ATRT, MYC; ATRT,SHH; EPN_PF_A; EWS; HGNET,BCOR; PIN_T_PPT.
21	PatientHadWES	Character	Did patient have whole-exome sequencing?	Variable denotes whether a patient had whole- exome sequencing (WES) if profiled
22	TOX_ALT_Indc	Numeric	Did patient have grade 3 or higher alanine aminotransferase increase?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
23	TOX_Anemia_Indc	Numeric	Did patient have grade 3 or higher anemia?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
24	TOX_Anorexia_Indc	Numeric	Did patient have grade 3 or higher anorexia?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
25	TOX_CatheterInfection_Indc	Numeric	Did patient have grade 3 or higher catheter related infection?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
26	TOX_FN_Indc	Numeric	Did patient have grade 3 or higher febrile neutropenia?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.

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27	TOX_Hearing_Indc	Numeric	Did patient have grade 3 or higher hearing impaired?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
28	TOX_Hypokalemia_Indc	Numeric	Did patient have grade 3 or higher hypokalemia?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
29	TOX_Ileus_Indc	Numeric	Did patient have grade 3 or higher ileus?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
30	TOX_Infection_Indc	Numeric	Did patient have grade 3 or higher infection/infestation?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
31	TOX_Lymphocyte_Indc	Numeric	Did patient have grade 3 or higher lymphocyte count decrease?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
32	TOX_MotorNeuropathy_Indc	Numeric	Did patient have grade 3 or higher peripheral motor neuropathy?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
33	TOX_Nausea_Indc	Numeric	Did patient have grade 3 or higher nausea?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
34	TOX_Neutrophil_Indc	Numeric	Did patient have grade 3 or higher neutrophil count decrease?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
35	TOX_Platelet_Indc	Numeric	Did patient have grade 3 or higher platelet count decrease?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
36	TOX_SensoryNeuropathy_Indc	Numeric	Did patient have grade 3 or higher peripheral sensory neuropathy?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.

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37	TOX_Vomiting_Inde	Numeric	Did patient have grade 3 or higher vomiting?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
38	TOX_WBC_Indc	Numeric	Did patient have grade 3 or higher white blood cell decrease?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
39	TOX_WeightLoss_Indc	Numeric	Did patient have grade 3 or higher weight loss?	For eligible patients only. 1 indicates that the subject had this toxicity. 0 indicates the subject did not.
40	RTPlanReviewResult	Character	Review of radiation (RT) plan	
41	ProtonIndc	Character	Proton therapy?	
42	InHearingLossAnalysis	Character	Was patient included in the analysis of hearing loss?	
43	CDDP_mgm2_cum	Numeric	Cumulative dose of cisplatin in milligrams per meter squared (mg/m2)	For patients included in the hearing loss analyses only
44	EndofTxHearingLossCategory	Character	Hearing loss category at end of treatment	For patients included in the hearing loss analyses only
45	PretherapyHearingLossIndc	Character	Did patient have hearing loss at the time of baseline or pretherapy?	This variable denotes whether a patient had hearing loss at the time of baseline/pretherapy (for patients included in the hearing loss analyses only). (Tables S5, S6, S7)
46	inQOLanalyses	Character	Was patient included in the analysis of quality of life/neurocognitive outcomes?	
47	QOL_PaymentMethod	Character	Payment method	For patients included in quality of life analyses

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48	QOL_T1_inDenominator	Character	Was patient evaluable for quality of life at time 1?	Yes denotes that the patient was in the time 1 denominator for quality of life/neurocognitive analyses.
49	QOL_T1_timing_mos	Numeric	Timing of time 1 quality of life assessment (in months)	
50	QOL_T1_IQ	Numeric	Intelligence quotient (IQ) at time 1	
51	QOL_T1_PSI	Numeric	Processing speed index (PSI) at time 1	
52	QOL_T2_inDenominator	Character	Was patient evaluable for quality of life at time 2?	Yes denotes that the patient was in the time 2 denominator for quality of life/neurocognitive analyses.
53	QOL_T2_timing_mos	Numeric	Timing of time 2 quality of life assessment (in months)	
54	QOL_T2_IQ	Numeric	Intelligence quotient (IQ) at time 2	
55	QOL_T2_PSI	Numeric	Processing speed index (PSI) at time 2	
56	QOL_T3_inDenominator	Character	Was patient evaluable for quality of life at time 3?	Yes denotes that the patient was in the time 3 denominator for quality of life/neurocognitive analyses.
57	QOL_T3_timing_mos	Numeric	Timing of time 3 quality of life assessment (in months)	
58	QOL_T3_IQ	Numeric	Intelligence quotient (IQ) at time 3	
59	QOL_T3_PSI	Numeric	Processing speed index (PSI) at time 3	

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60	FirstEventType	Character	First event type	
61	SecondMalignancyIndc	Character	Second malignancy indicator	
62	YearstoSM1	Numeric	Years from enrollment to date of first subsequent malignancy	
63	YearstoSM2	Numeric		For patients with >1 subsequent malignancy (n=1)
64	YearstoSM3	Numeric		For patients with >2 subsequent malignancies (n=1)
65	SM1_Histology	Character	Histology of first subsequent malignancy	
66	SM2_Histology	Character	Histology of second subsequent malignancy	For patients with >1 subsequent malignancy (n=1)
67	SM3_Histology	Character	Histology of third subsequent malignancy	For patients with >2 subsequent malignancies (n=1)
68	SurvivalfromFirstSM_months	Numeric	Time from first subsequent malignancy to death or last follow-up (in months)	This variable is used in Supplemental Table 3
69	SM_FocalTP53alteration	Character	Focal TP53 alteration for secondary malignancy	This variable is used in Supplemental Table 3
70	InPatternsofFailureAnalyses	Character		All eligible and evaluable patients contributed to patterns of failure analyses unless the pattern of failure was unknown/not available.
71	RLPDType	Character		Data are provided only for subjects with InPatternsofFailureAnalyses='Yes'
72	LocalFailure_time	Numeric	Time from enrollment for local failure analyses (in days)	Time variable for local failure analyses in days. Data are provided only for subjects with InPatternsofFailureAnalyses='Yes'.

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73	LocalFailure_censorvar	Numeric	Censor variable for local failure analyses	Data are provided only for subjects with InPatternsofFailureAnalyses='Yes'. Censor variable for local failure analyses (0=censored, 1=patient had local failure or local+distant failure [event of interest], 2=patient had distant failure only, 3=patient had second malignancy prior to progression, 4=patient died prior to progression).
74	IsolatedLocalFailure_time	Numeric	Time from enrollment for isolated local failure analyses (in days)	Time variable for isolated local failure analyses in days. Data are provided only for subjects with InPatternsofFailureAnalyses='Yes'.
75	IsolatedLocalFailure_censorvar	Numeric	Censor variable for isolated local failure analyses	Data are provided only for subjects with InPatternsofFailureAnalyses='Yes'. Censor variable for isolated local failure analyses (0=censored, 1=patient had local failure only [event of interest], 2=patient had distant failure or local+distant failure, 3=patient had second malignancy prior to progression, 4=patient died prior to progression).
76	DistantFailure_time	Numeric	Time from enrollment for distant failure analyses (in days)	Time variable for distant failure analyses in days. Data are provided only for subjects with InPatternsofFailureAnalyses='Yes'.
77	DistantFailure_censorvar	Numeric	Censor variable for distant failure analyses	Data are provided only for subjects with InPatternsofFailureAnalyses='Yes'. Censor variable for local failure (0=censored, 1=patient had distant or local+distant failure [event of interest], 2=patient had local failure only, 3=patient had second malignancy prior to progression, 4=patient died prior to progression).

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78	IsolatedDistantFailure_time	Numeric	Time from enrollment for isolated distant failure analyses (in days)	Time variable for isolated distant failure analyses in days. Data are provided only for subjects with InPatternsofFailureAnalyses='Yes'.
79	IsolatedDistantFailure_censorvar	Numeric	Censor variable for isolated distant failure analyses	Data are provided only for subjects with InPatternsofFailureAnalyses='Yes'. Censor variable for isolated distant failure (0=censored, 1=patient had distant failure only [event of interest], 2=patient had local failure or local+distant failure, 3=patient had second malignancy prior to progression, 4=patient died prior to progression).
80	FocalAlt_PTCH1	Character	PTCH1 focal alteration	These data are provided for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroups MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA indicates data for this variable were not available or inconclusive.
81	FocalAlt_GLI2	Character	GLI2 focal alteration	These data are provided for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroups MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA indicates data for this variable were not available or inconclusive.
82	FocalAlt_MYCL	Character	MYCL focal alteration	These data are provided for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroups MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA indicates data for this variable were not available or inconclusive.

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83	FocalAlt_MYCN	Character	MYCN focal alteration	These data are provided for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroups MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA indicates data for this variable were not available or inconclusive.
84	FocalAlt_CCND2	Character	CCND2 focal alteration	These data are provided for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroups MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA indicates data for this variable were not available or inconclusive.
85	FocalAlt_PPM1D	Character	PPM1D focal alteration	These data are provided for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroups MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA indicates data for this variable were not available or inconclusive.
86	FocalAlt_MDM2	Character	MDM2 focal alteration	These data are provided for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroups MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA indicates data for this variable were not available or inconclusive.
87	MYCamplification	Character	MYC focal amplification	These data are provided for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroups MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA indicates data for this variable were not available or inconclusive.

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88	TP53Alteration	Character	TP53 focal alteration	These data are available for eligible patients who were molecularly classified and had whole exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
89	GeneAlt_CTNNB1	Character	CTNNB1 gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
90	GeneAlt_DDX3X	Character	DDX3X gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
91	GeneAlt_PTCH1	Character	PTCH1 gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
92	GeneAlt_TP53	Character	TP53 gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.

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93	GeneAlt_SUFU	Character	SUFU gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
94	GeneAlt_SMO	Character	SMO gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
95	GeneAlt_SMARCA4	Character	SMARCA4 gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
96	GeneAlt_KBTBD4	Character	KBTBD4 gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
97	GeneAlt_MYC	Character	MYC gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.

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98	GeneAlt_KMT2C	Character	KMT2C gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
99	GeneAlt_KDM6A	Character	KDM6A gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
100	GeneAlt_CREBBP	Character	CREBBP gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
101	GeneAlt_ZMYM3	Character	ZMYM3 gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
102	GeneAlt_GSE1	Character	GSE1 gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.

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103	GeneAlt_MYCN	Character	MYCN gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
104	GeneAlt_TBR1	Character	TBR1 gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
105	GeneAlt_CTDNEP1	Character	CTDNEP1 gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
106	GeneAlt_KMT2D	Character	KMT2D gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
107	GeneAlt_PPM1D	Character	PPM1D gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.

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108	GeneAlt_ARID1A	Character	ARID1A gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
109	GeneAlt_MYCL	Character	MYCL gene alteration	These data are available for eligible and evaluable patients molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT) who had whole-exome sequencing (WES). NA indicates data for this variable were not available or inconclusive.
110	ChrAlt_1q	Character	1q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
111	ChrAlt_3q	Character	3q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
112	ChrAlt_6p	Character	6p chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.

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Data can be used to approximate published study findings, but exact reproduction of previous manuscripts may not be possible in some cases (e.g., when data must be modified for deidentification purposes or have undergone further data cleaning).

Variables in this dataset that contain limited high-dimensional "omic" data are provided in addition to the clinical data in an "as-is" state without all the same quality-control procedures that are performed on the clinical data by the NCTN/NCORP Data Archive. It may not be possible to fully reproduce the results of the genomic analyses included in the publication. For example, the "omic" data provided may be a subset of the data used in the publication or may not reflect certain aspects of the data

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113	ChrAlt_6q	Character	6q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
114	ChrAlt_7p	Character	7p chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
115	ChrAlt_7q	Character	7q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
116	ChrAlt_8p	Character	8p chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
117	ChrAlt_8q	Character	8q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.

Study Number	ACNS0331
NCT#	NCT00085735
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118	ChrAlt_9p	Character	9p chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available
				or inconclusive.
119	ChrAlt_9q	Character	9q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA
				denotes data for this variable was not available or inconclusive.
120	ChrAlt_10q	Character	10q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
121	ChrAlt_11p	Character	11p chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
122	ChrAlt_11q	Character	11q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.

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123	ChrAlt_14q	Character	14q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
124	ChrAlt_16q	Character	16q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
125	ChrAlt_17p	Character	17p chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
126	ChrAlt_17q	Character	17q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
127	ChrAlt_18p	Character	18p chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.

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128	ChrAlt_18q	Character	18q chromosomal alteration	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
129	Isochromosome17	Character	Isochromosome 17	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
130	Chr10qLossOnly	Character	Chromosome 10q loss only	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
131	Chr11Loss	Character	Chromosome 11 loss	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
132	Chr17Gain	Character	Chromosome 17 gain	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.

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133	Chr11LossORChr17Gain	Character	chromosome 17 gain	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.
134	MYCamplificationORIso17	Character	isochromosome 17	These data are available for eligible and evaluable patients who were molecularly profiled as medulloblastoma (subgroup MB,G3 or MB,G4 or MB,SHH or MB,WNT). NA denotes data for this variable was not available or inconclusive.