

NGS molecular profile of paediatric brain tumours research samples: results from 92 consecutive cases treated at Centro Hospitalar Universitário de São João

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Paediatric Cancer and Paediatric CNS tumours

- Paediatric cancer is the leading cause of death, beside accidents, in children aged 1-14y;
- Whilst global survival rates have improved significantly, they continue dismal for specific tumour types and for relapses/refractory disease;
- Moreover, survival rates are attained at the cost of high toxicity due to multimodal therapeutics;
- Paediatric CNS tumours represent ~25% of all paediatric tumours and are a heterogenous and challenging group of tumours;
- Have had tremendous advances in their molecular biology through several large-scale, pan-cancer studies. However, the effective utilization of molecular biomarkers in paediatric oncology clinical practice is still lagging behind what is already routinely performed in many adult cancer patients.

The 2021 WHO Classification of CNS tumours

The 2021 WHO Classification of Tumors of the Central Nervous System: a summary

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brain tumor | central nervous system | classification | diagnosis | World Health Organization

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The 2021 WHO Classification of CNS tumours

6

Louis et al. **The 2021 WHO Classification of CNS Tumors**

Tumor Type	Genes/Molecular Profiles Characteristically Altered ^a
Oligodendroglioma, IDH-mutant, and 1p/19q-codeleted	<i>IDH1</i> , <i>IDH2</i> , 1p/19q, <i>TERT</i> promoter, <i>CIC</i> , <i>FUBP1</i> , <i>NOTCH1</i>
Diffuse astrocytoma, <i>MYB</i> - or <i>MYBL1</i> -altered	<i>MYB</i> , <i>MYBL1</i>
Polymorphous low-grade neuroepithelial tumor of the young	<i>BRAF</i> , <i>FGFR</i> family
Diffuse midline glioma, H3 K27-altered	H3 K27, <i>TP53</i> , <i>ACVR1</i> , <i>PDGFRA</i> , <i>EGFR</i> , <i>EZH1</i>
Diffuse pediatric-type high-grade glioma, H3-wildtype, and IDH-wildtype	IDH-wildtype, H3-wildtype, <i>PDGFRA</i> , <i>MYCN</i> , <i>EGFR</i> (methylome)
Pilocytic astrocytoma	<i>KIAA1549-BRAF</i> , <i>BRAF</i> , <i>NF1</i>
Pleomorphic xanthoastrocytoma	<i>BRAF</i> , <i>CDKN2A/B</i>
Chordoid glioma	<i>PRKCA</i>
Ganglion cell tumors	<i>BRAF</i>
Diffuse glioneuronal tumor with oligodendroglioma-like features and nuclear clusters	Chromosome 14, (methylome)
Rosette-forming glioneuronal tumor	<i>FGFR1</i> , <i>PIK3CA</i> , <i>NF1</i>
Diffuse leptomeningeal glioneuronal tumor	<i>KIAA1549-BRAF</i> fusion, 1p (methylome)
Dysplastic cerebellar gangliocytoma (Lhermitte-Duclos disease)	<i>PTEN</i>
Supratentorial ependymomas	<i>ZFTA</i> , <i>RELA</i> , <i>YAP1</i> , <i>MAML2</i>
Spinal ependymomas	<i>NF2</i> , <i>MYCN</i>
Medulloblastoma, SHH-activated	<i>TP53</i> , <i>PTCH1</i> , <i>SUFU</i> , <i>SMO</i> , <i>MYCN</i> , <i>GLI2</i> (methylome)
Atypical teratoid/rhabdoid tumor	<i>SMARCB1</i> , <i>SMARCA4</i>
CNS neuroblastoma, <i>FOXR2</i> -activated	<i>FOXR2</i>
Desmoplastic myxoid tumor of the pineal region, <i>SMARCB1</i> -mutant	<i>SMARCB1</i>
Solitary fibrous tumor	<i>NAB2-STAT6</i>

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WHO Classification of Pediatric tumours

REVIEW

A Summary of the Inaugural WHO Classification of Pediatric Tumors: Transitioning from the Optical into the Molecular Era

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ABSTRACT

Pediatric tumors are uncommon, yet are the leading cause of cancer-related death in childhood. Tumor types, molecular characteristics, and pathogenesis are unique, often originating from a single genetic driver event. The specific diagnostic challenges of childhood tumors led to the development of the first World Health Organization (WHO) Classification of Pediatric Tumors. The classification is rooted in a multilayered approach, incorporating morphology, IHC, and molecular characteristics. The volume is organized according to organ sites and provides a single, state-of-the-art compendium of pediatric tumor types. A special emphasis was placed on “blastomas,” which variably recapitulate the morphologic maturation of organs from which they originate.

Significance: In this review, we briefly summarize the main features and updates of each chapter of the inaugural WHO Classification of Pediatric Tumors, including its rapid transition from a mostly microscopic into a molecularly driven classification systematically taking recent discoveries in pediatric tumor genomics into account.

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Classification of Pediatric CNS tumours

Gliomas, glioneuronal, and neuronal tumors

Pediatric-type diffuse low-grade gliomas

Diffuse astrocytoma, MYB or MYBL1-altered **new**

Angiocentric glioma

Polymorphous low-grade neuroepithelial tumor of the young **new**

Diffuse low-grade glioma, MAPK pathway-altered **new**

Pediatric-type diffuse high-grade gliomas defined by H3 status

Diffuse midline glioma, H3 K27-altered

Diffuse hemispheric glioma, H3 G34-mutant **new**

Diffuse pediatric-type high-grade glioma, H3-wild-type and IDH-wild-type **new**

Infant-type hemispheric glioma **new**

Circumscribed astrocytic gliomas

Pilocytic astrocytoma

High-grade astrocytoma with piloid features **new**

Pleomorphic xanthoastrocytoma

Subependymal giant cell astrocytoma

Astroblastoma, MN1-altered

Glioneuronal and neuronal tumors

Ganglioglioma

Desmoplastic infantile ganglioglioma/Desmoplastic infantile astrocytoma

Dysembryoplastic neuroepithelial tumor

Diffuse glioneuronal tumor with oligodendroglioma-like features and nuclear

clusters (DGONC)^a **new**

Diffuse leptomeningeal glioneuronal tumor

Multinodular and vacuolating neuronal tumor **new**

Ependymal tumors

Supratentorial ependymoma

Supratentorial ependymoma, ZFTA fusion-positive

Supratentorial ependymoma, YAP1 fusion-positive **new**

Posterior fossa ependymoma

Posterior fossa ependymoma, Group PFA **new**

Posterior fossa ependymoma, Group PFB **new**

Spinal ependymoma, MYCN-amplified **new**

Myxopapillary ependymoma

Choroid plexus tumors

Choroid plexus papilloma

Atypical choroid plexus papilloma

Choroid plexus carcinoma

CNS embryonal tumors

Medulloblastomas, molecularly defined

Medulloblastoma, WNT-activated

Medulloblastoma, SHH-activated & TP53-wild-type

Medulloblastoma, SHH-activated & TP53-mutant

Medulloblastoma, non-WNT/non-SHH

Medulloblastoma, histologically defined

Medulloblastoma, histologically defined

Other CNS embryonal tumors

Atypical teratoid/rhabdoid tumor

Cribiform neuroepithelial tumor^a **new**

Embryonal tumor with multilayered rosettes

CNS neuroblastoma, FOXR2-activated **new**

CNS tumor with BCOR internal tandem duplication **new**

CNS embryonal tumor NEC/NOS

Pineal region tumors

Pineoblastoma

Melanocytic tumors

Meningeal melanocytosis and melanomatosis

Tumors of the sellar region

Pituitary endocrine tumors

Pituitary adenoma/PitNET

Pituitary blastoma **new**

Craniopharyngiomas

Adamantinomatous craniopharyngioma

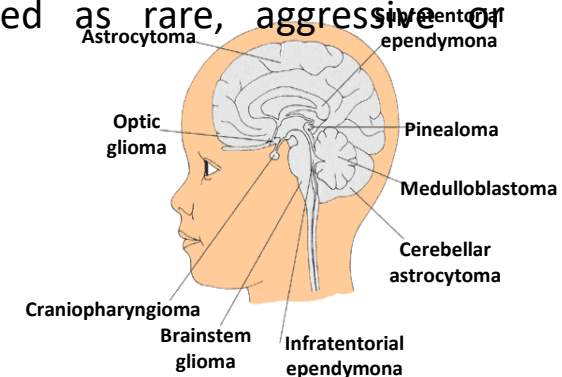
Aims of the Project

Perform a retrospective **comprehensive genetic screening** in all diagnosed **paediatric CNS tumours**, treated at Centro Hospitalar Universitário de São João (Jan 2018-present).

The fundamental goal of this protocol is to evaluate the future value of **a tool to assist in the diagnosis and prognosis**, while supporting the clinician in identifying the best possible therapy.

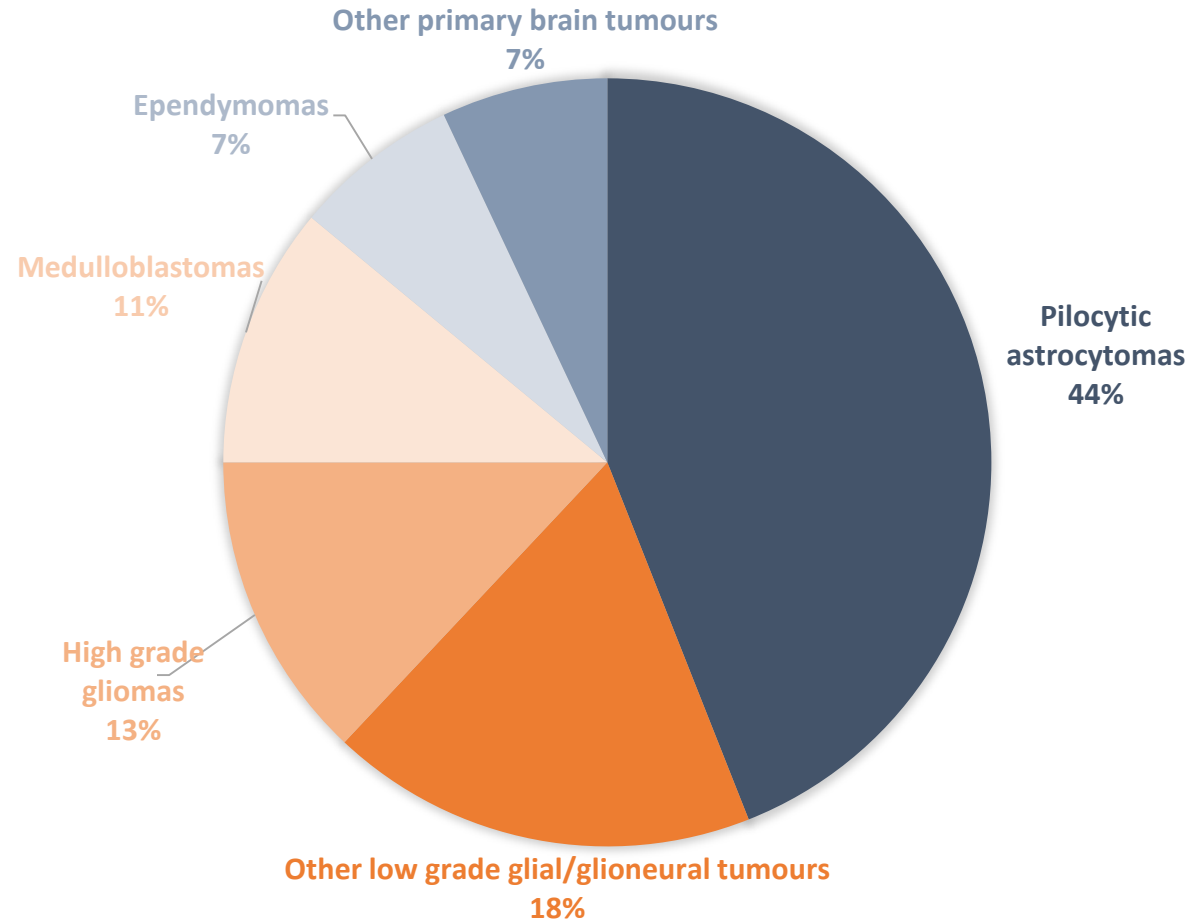
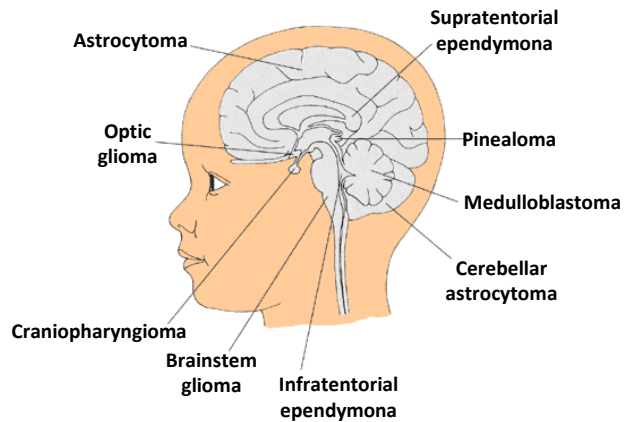
Inclusion criteria are:

- Age 18 years or below;
- Newly diagnosed brain tumour, with histologic confirmation, irrespective of location;
- Previously diagnosed brain tumour, whenever it presented as rare, aggressive or irresectable disease;
- Availability of tumour material;
- Signed informed consent.



Paediatric CNS tumours study cohort

Paediatric CNS tumours (n=92)



In this cohort, the majority of tumours are **pilocytic astrocytomas** (44%)

Methods: NGS With Oncomine Childhood Cancer Research Assay

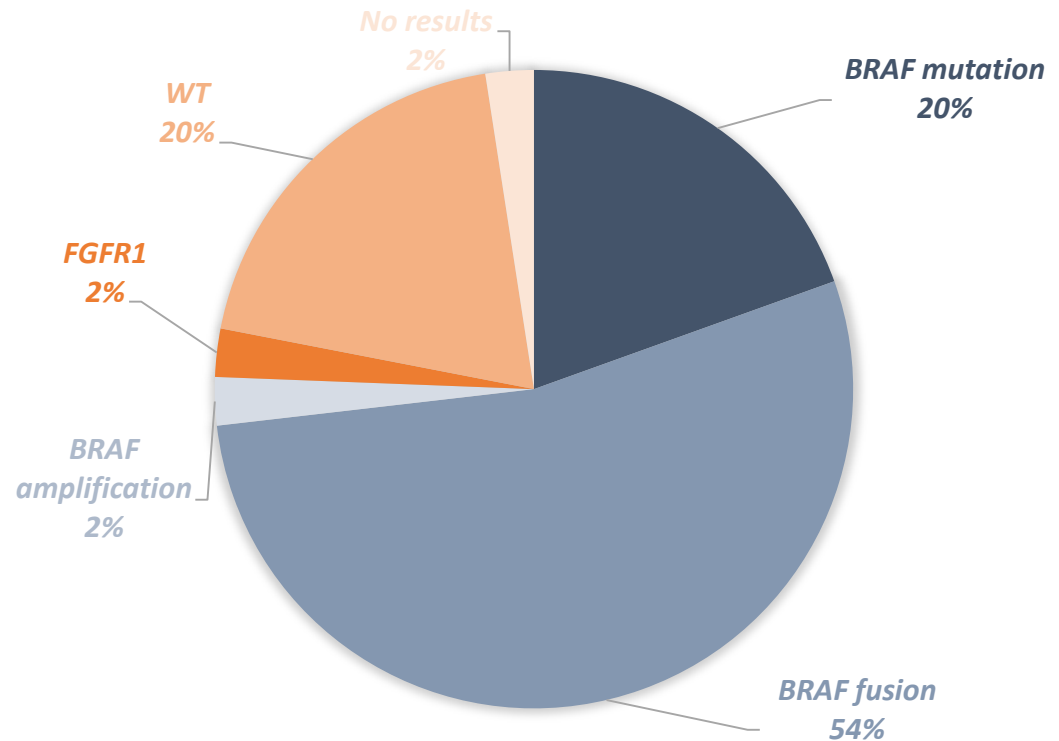
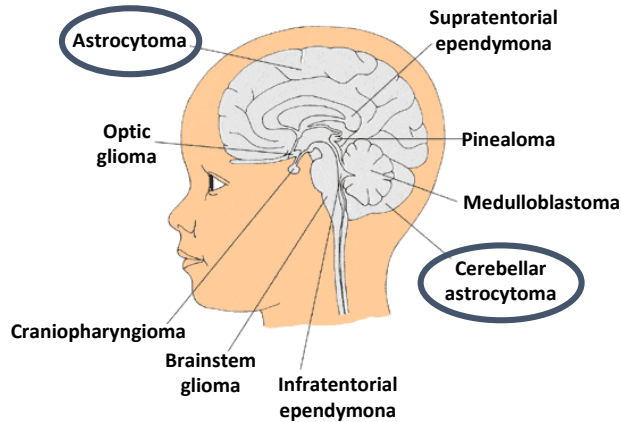
Oncomine Childhood Cancer Research Assay Panel - 203 genes

Comprehensive mutation coverage (86)					CNV (28)		Full exon coverage (44)			Fusion and expression (97)					Gene expression
ABL1	CSF1R	GATA2	MAP2K2	RAF1	ALK	IGF1R	APC	GATA3	RUNX1	ABL1	FGFR2	MEF2B	NUP214	SSBP2	BCL2
ABL2	CSF3R	GNAQ	MET	RET	BRAF	JAK1	ARID1A	GNA13	SMARCA4	ABL2	FGFR2	MET	NUP98	STAG2	BCL6
ALK	CTNNB1	H3F3A	MPL	RHOA	CCND1	JAK2	ARID1B	ID3	SMARCB1	AFF3	FGFR3	MKL1	NUTM1	STAT6	FGFR1
ACVR1	DAXX	HDAC9	MSH6	SETBP1	CDK4	JAK3	ATRX	IKZF1	SOCS2	ALK	FLT3	MLLT10	NUTM2B	TAL1	FGFR4
AKT1	DNMT3A	HIST1H3B	MTOR	SETD2	CDK6	KIT	CDKN2A	KDM6A	SUFU	BCL11B	FOSB	MN1	PAX3	TCF3	IGF1R
ASXL1	EGFR	HRAS	NCOR2	SH2B3	EGFR	KRAS	CDKN2B	KMT2D	SUZ12	BCOR	FUS	MYB	PAX5	TFE3	MET
ASXL2	EP300	IDH1	NOTCH1	SH2D1A	ERBB2	MDM2	CEBPA	MYOD1	TCF3	BCR	GLI1	MYBL1	PAX7	TP63	MYCN
BRAF	ERBB2	IDH2	NPM1	SMO	ERBB3	MDM4	CHD7	NF1	TET2	BRAF	GLIS2	MYH11	PDGFB	TSLP	MYC
CALR	ERBB3	IL7R	NRAS	STAT3	FGFR1	MET	CRLF1	NF2	TP53	CAMTA1	HMGA2	MYH9	PDGFRA	TSPAN4	TOP2A
CBL	ERBB4	JAK1	NT5C2	STAT5B	FGFR2	MYC	DDX3X	PHF6	TSC1	CCND1	JAK2	NCOA2	PDGFRB	UBTF	
CCND1	ESR1	JAK2	PAX5	TERT	FGFR3	MYCN	DICER1	PRPS1	TSC2	CIC	KAT6A	NCOR1	PLAG1	USP6	
CCND3	EZH2	JAK3	PDGFRA	TPMT	FGFR4	PDGFRA	EBF1	PSMB5	WHSC1	CREBBP	KMT2A	NOTCH1	RAF1	WHSC1	
CCR5	FASLG	KDM4C	PDGFRB	USP7	GLI1	PIK3CA	EED	PTCH1	WT1	CRLF2	KMT2B	NOTCH2	RANBP17	YAP1	
CDK4	FBXW7	KDR	PIK3CA	ZMYM3	GLI2		FAS	PTEN	XIAP	CSF1R	KMT2C	NOTCH4	RECK	ZMYND11	
CIC	FGFR2	KIT	PIK3R1				GATA1	RB1		DUSP22	KMT2D	NPM1	RELA	ZNF384	
CREBBP	FGFR3	KRAS	PPM1D							EGFR	LMO2	NR4A3	RET		
CRLF2	FLT3	MAP2K1	PTPN11							ETV6	MAML2	NTRK1	ROS1		
										EWSR1	MAN2B1	NTRK2	RUNX1		
										FGFR1	MECOM	NTRK3	SS18		

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Genomic profiling results

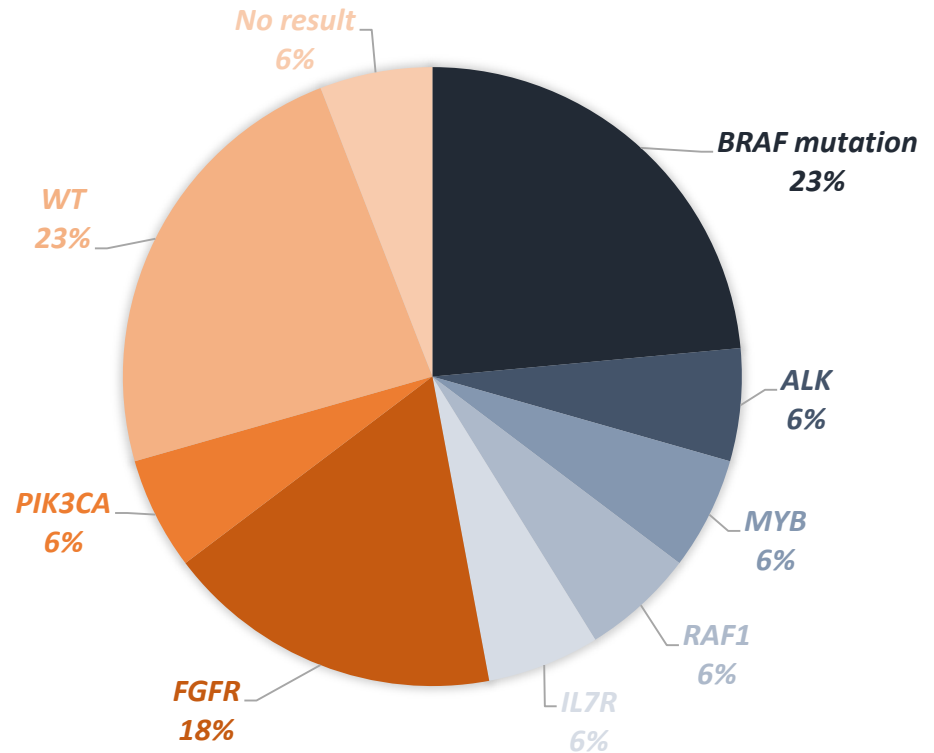
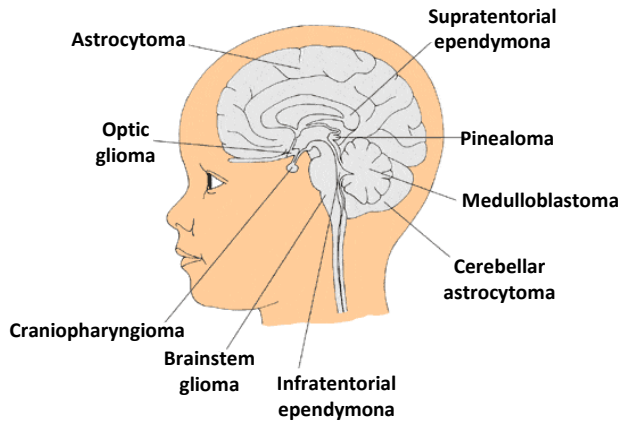
Pilocytic Astrocytomas (n=41)



The majority of variants (54%) were **BRAF** fusions

Genomic profiling results

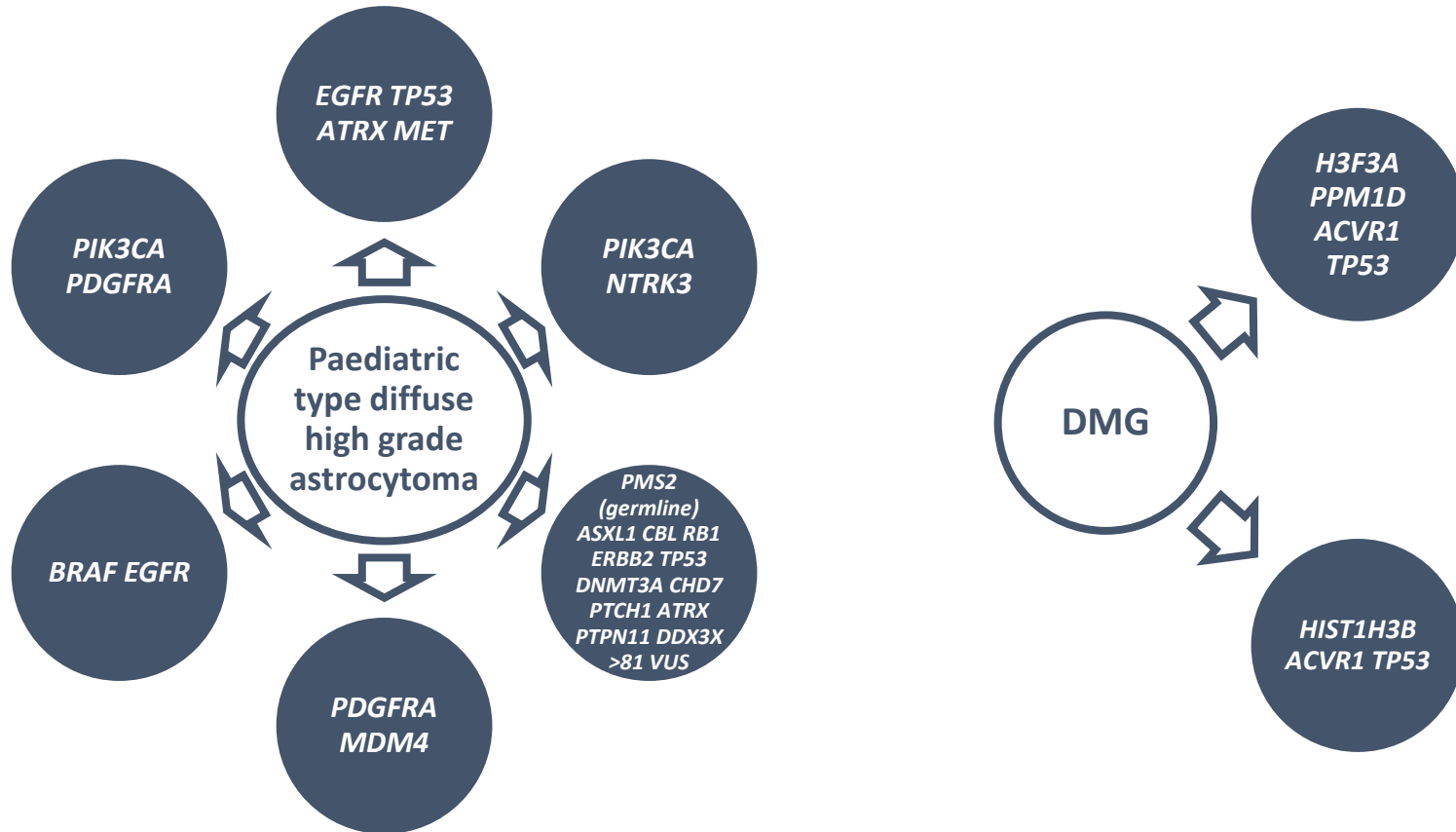
Other Low Grade Glial/ Glioneural Tumours (n=17)



The majority of variants (23%) were **BRAF** mutations

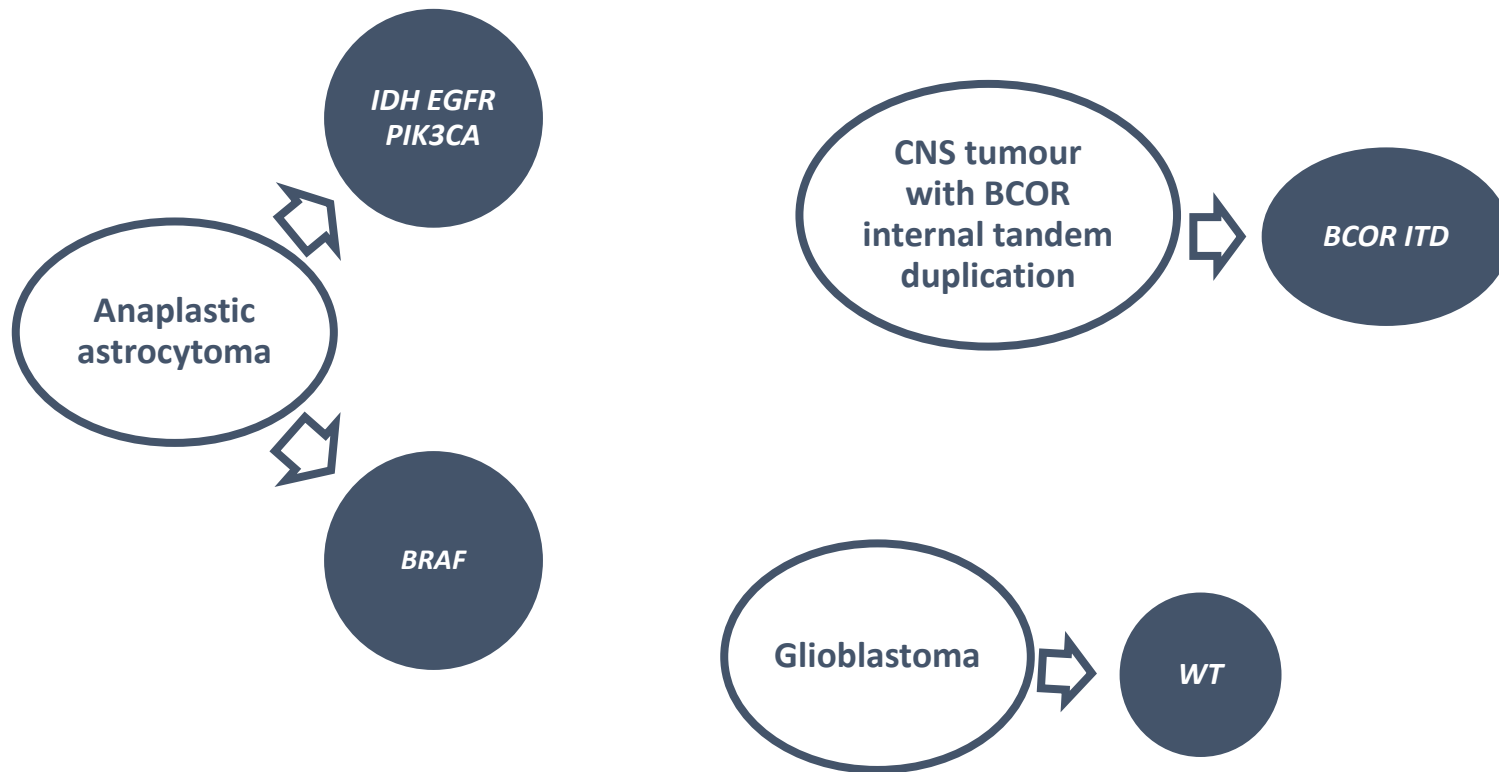
Genomic profiling results

High Grade Gliomas (n=12)



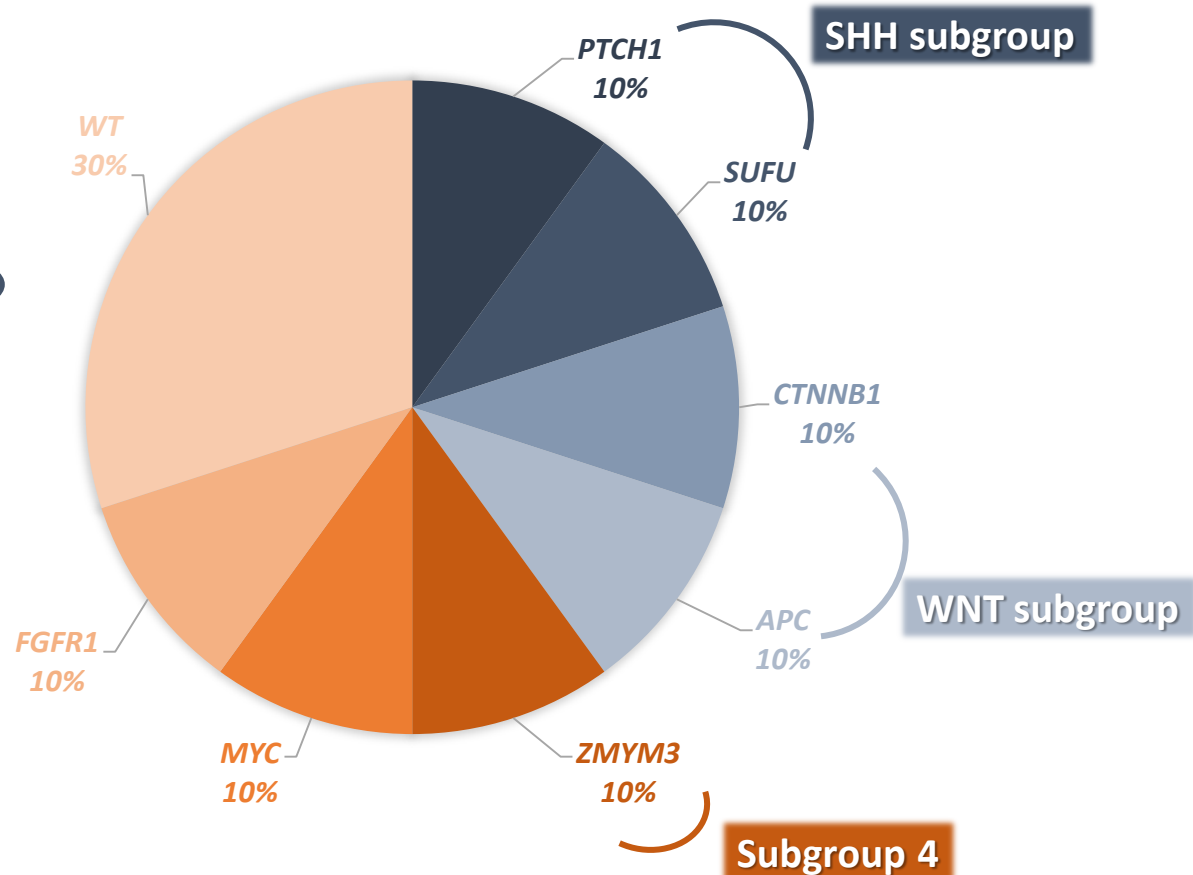
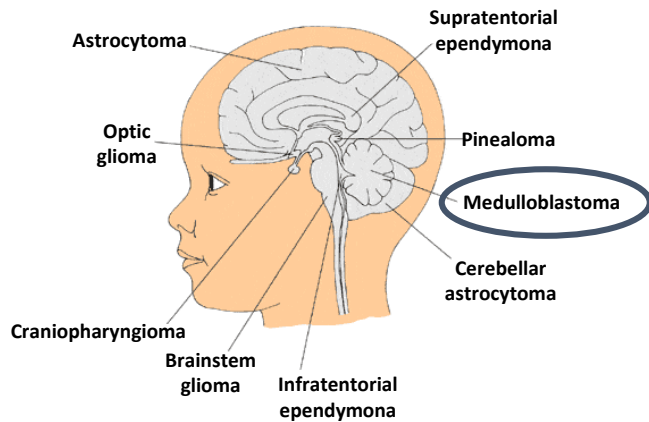
Genomic profiling results

High Grade Gliomas (n=12)



Genomic profiling results

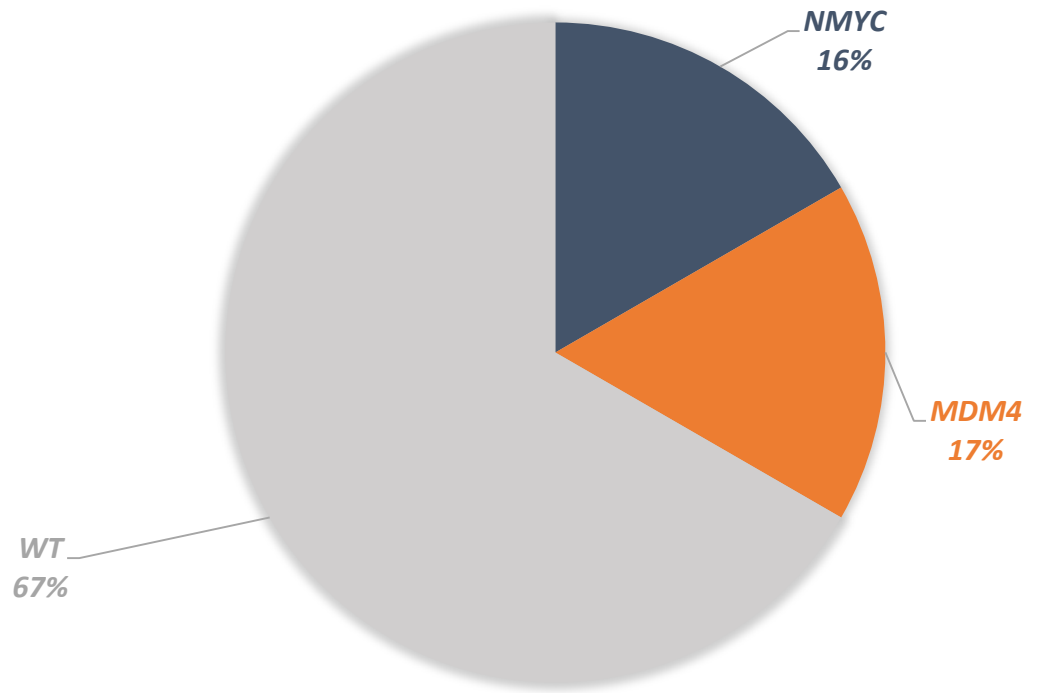
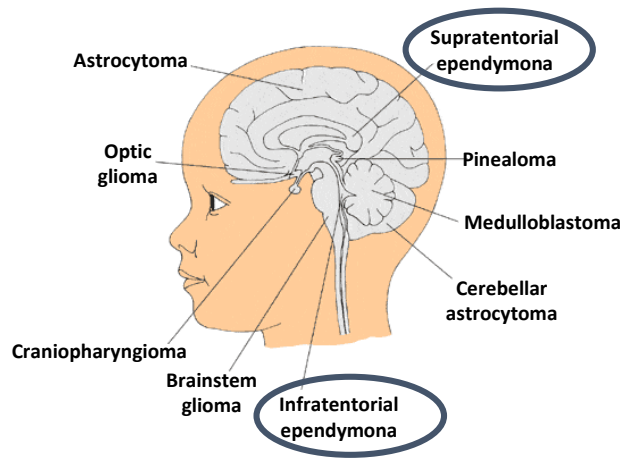
Medulloblastomas (n=10)



For medulloblastoma, **3 subgroups (SHH, WNT and ZMYM3)** were identified

Genomic profiling results

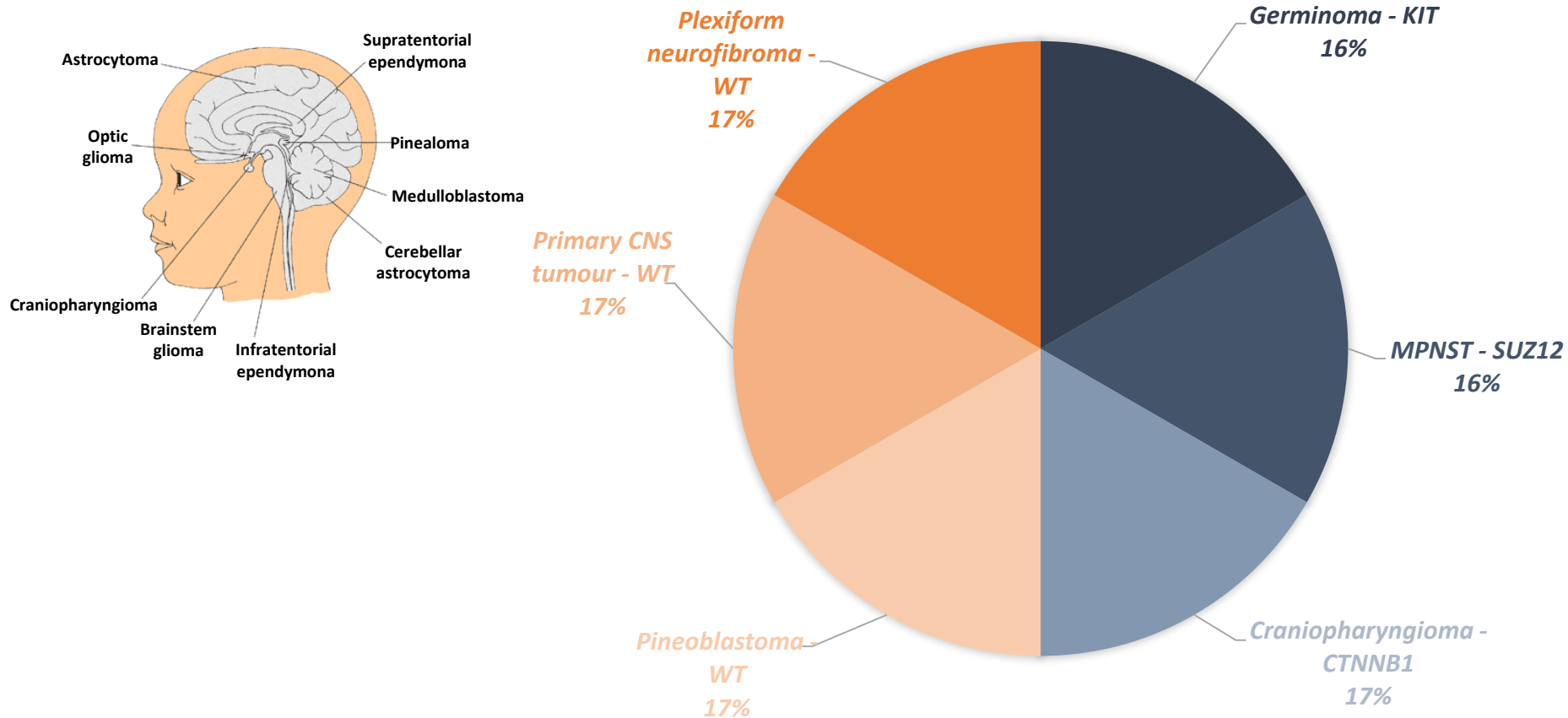
Ependymomas (n=6)



In **33%** of ependymoma cases a relevant genetic variant was identified

Genomic profiling results

Other Brain Tumours (n=6)

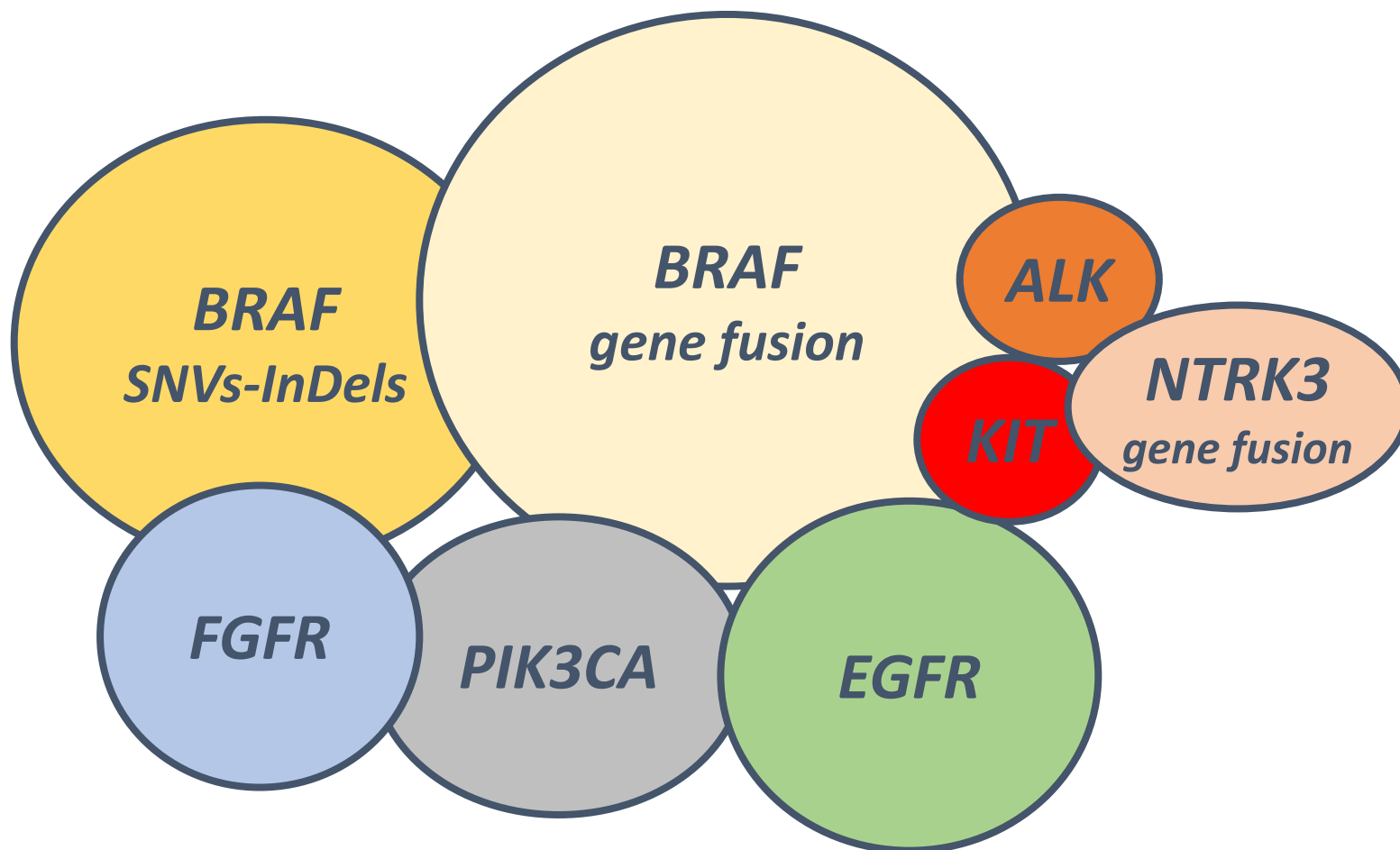


In other brain tumors there are **heterogeneity** in terms of **genomic findings**

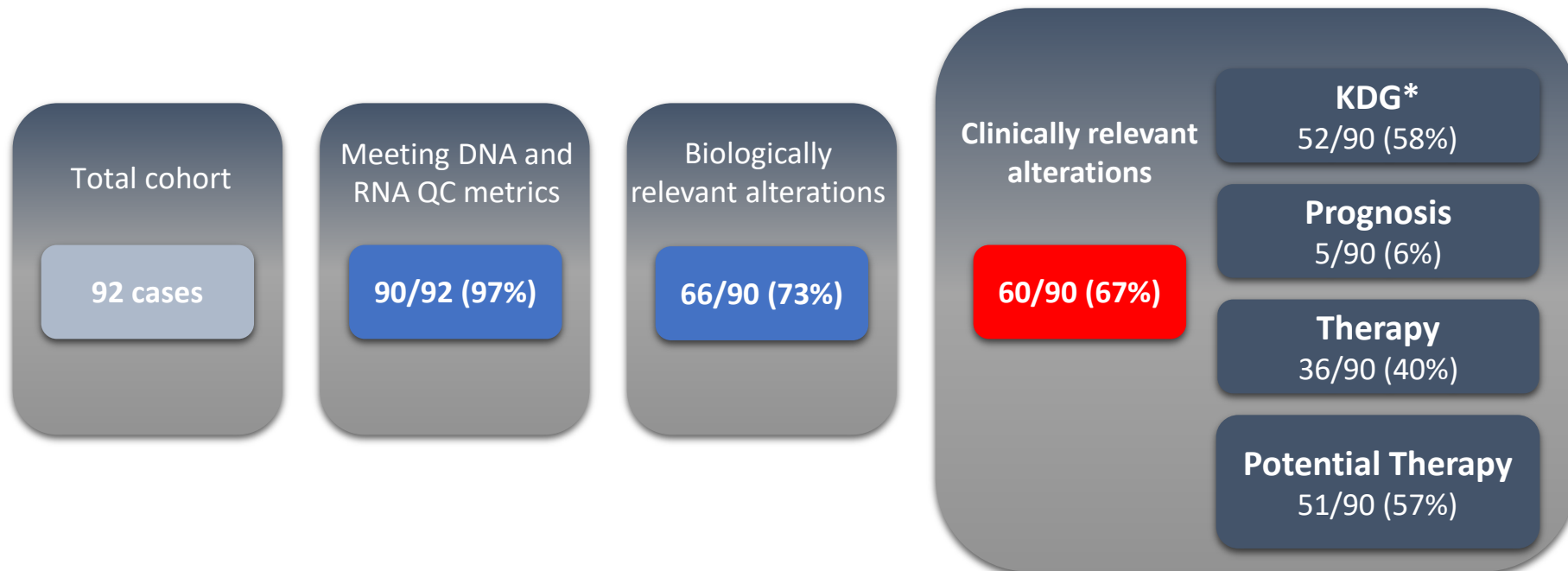
Notable Cases

Diagnosis	Alteration #1	Alteration #2	Alteration #3
PA	KIAA1549-BRAF	PTPRZ1-MET	EGFRVIII
PA	KIAA1549-BRAF	PAX8-PPARg	-
PA	KIAA1549-BRAF	CBL c.1096-1G>T	EGFRVIII
PA	KIAA1549-BRAF	ZCCHC8-WHSC1	C2orf44-FER
Ependymoma	<i>MYCN</i>	-	-
PTDHGA	<i>PMS2</i> (homoz)	hypermutable	

Alterations with potential actionable value



Study Overall Performance



*KDG: key diagnostic gene (WHO 2021)

Research cases with *BRAF*-V600 Mutant tumours

Pt	Age Dx	Age Tr	Sex	Topog.	Histology	option	Tr time	Response	Tox
1	12y	12y	F	IV vent / invasion BST	PA	first line	47 m	improved	-
2	8m	4y	F	Sup. selar	PA	rescue	42 m	mixed	Rash g1
3	4y	5y	F	Hemisph.	Met. GBM	rescue	2 m	progr. died	-
4	2y	12y	F	Sup. selar	PA	rescue	41 m	improved	Rash g1
5	4y	4y	M	Sup. selar	GG	first line	43 m	improved	Rash g1

Pt: Patient; Dx: Diagnosis; Tr: Treatment; Tox: Toxicity; PA: Pilocytic Astrocytoma; GBM: Glioblastoma Multiforme; GG: Glioma

Conclusions

- **NGS proved to be a valuable tool in the setting of PBT**, resulting from its significant capacity in revealing clinically meaningful molecular alterations, that can be potentially used for the diagnosis, prognosis and therapy selection
- **Low and high grade gliomas were the tumour types that benefitted more from NGS**, while ependymomas were the tumours less likely to harbour molecular alterations
- In terms of potential therapeutic targets, **BRAF** stands out, but **FGFR** genes might also be a potentially relevant target in LGG. In high grade gliomas, there were several potential targets (**EGFR**, **PIK3CA**, **PDGFRA**, **MET**) although there is still little data about the efficacy of therapeutic agents directed at these targets

Thank you



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