

Molecular Targeted Therapies Reveal Glioma Cell Plasticity Linked to Immune Evasion in BRAF-Mutant Brain Tumors

Claudia K. Petritsch, Ph.D.

Today's Speaker



Claudia K. Petritsch, Ph.D. **Stanford School of Medicine**

- Sr. Scientist, Department of Neurology
- Associate Professor in Research, Department of Neurosurgery
- Director, Pediatric Cancer Model Development Center

Agenda

1. *The Human Cancer Model Initiative (HCMI)*
2. *High-Grade Glioma Challenges and Treatment*
3. *Unraveling Molecular Therapy Resistance in Novel BRAF V600E HGG Models*
4. *The HCMI for Pediatric Cancer*

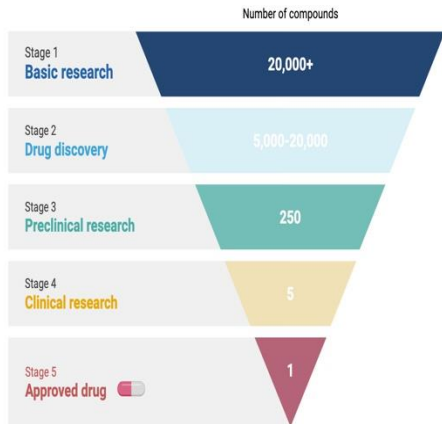
The Human Cancer Model Initiative

Objectives and Achievements in a Nutshell

Barriers for Drug Development in Oncology

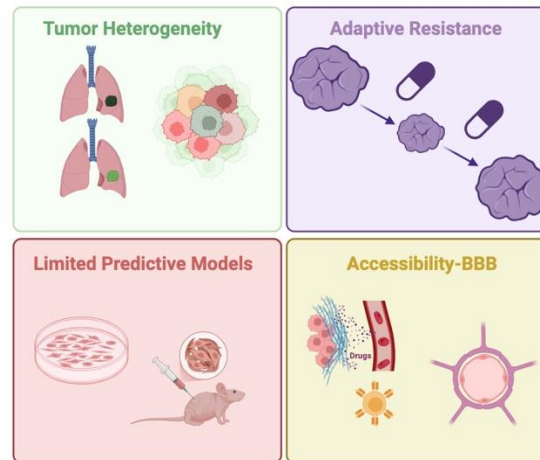
Oncology drug development attrition rate

Causes for lack of efficacy for oncology drugs



Phase Success	2016		2021	
	Phase I to Phase II	N	Phase I to Phase II	N
Ophthalmology	84.8%	66	71.6%	88
Hematology	73.3%	86	69.6%	92
Others	66.7%	96	63.6%	154
Metabolic	61.1%	95	61.8%	136
Infectious disease	69.5%	347	57.8%	403
Allergy	67.6%	37	56.4%	55
Respiratory	65.3%	150	55.9%	179
Autoimmune	65.7%	297	55.2%	413
Psychiatry	53.9%	154	52.7%	150
All indications	63.2%	3582	52.0%	4414
Cardiovascular	58.9%	209	50.0%	214
Oncology	62.8%	1222	48.8%	1628
Neurology	59.1%	462	47.7%	516
Gastroenterology	75.6%	41	46.7%	45
Endocrine	58.9%	299	43.3%	319
Urology	57.1%	21	40.9%	22

Bio, February 2021



~50-60% of drugs fail in the clinic due to lack of efficacy_ up to 85% in oncology)
 FDA Approval can take up to 13 years and cost up to \$2.5b

Conventional vs. Next-Generation Cancer Models

Conventional Models



(monolayer, 2-dimensional (2D) cultures on plastic)

- Cells plated on flat **plastic surfaces**
 - Cells grown in high levels of animal **serum**
 - Poorly recapitulate **tumor heterogeneity and plasticity**
 - Often **overpredict drug efficacy**
 - Often lacking clinical information of **parental tumor origin**
- Simple, scalable, but **low physiological relevance**

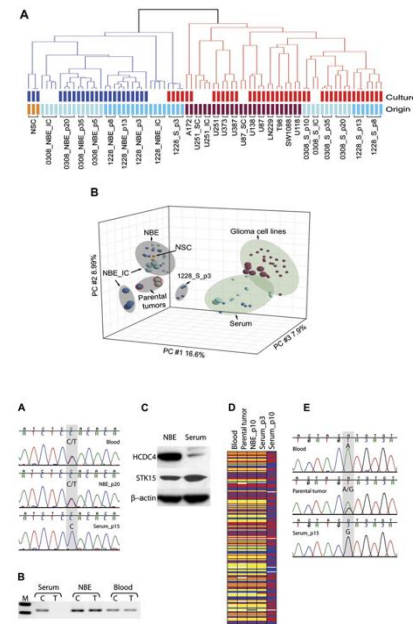
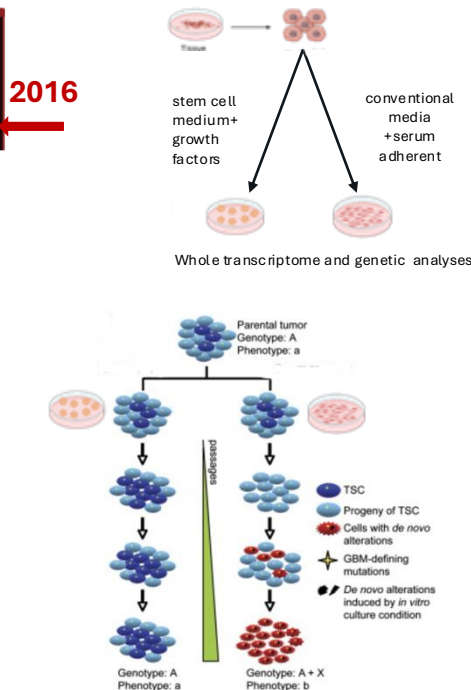
Timeline of Glioblastoma Model Development

1 Timeline of glioblastoma research and models 2 Conventional serum vs 3D cell culture models



Sources: CBTRUS, PMC, Frontiers in Oncology, Cutting Through History (PMC 2024)

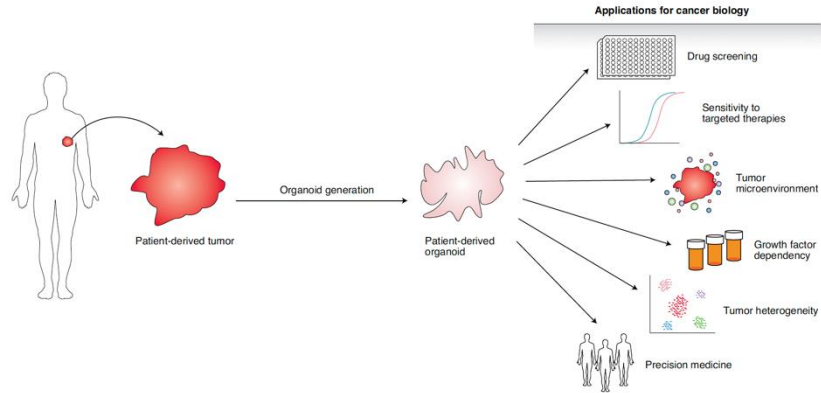
Features	3D/Spheroids w/oSerum	2D/Adherent w/Serum
Proliferation	+++	Limited at early passages
Tumorigenicity	+++	Limited at early passages
Differentiation potential	+++	+
Telomerase activity	+++	Limited at early passages
Infiltrative growth	+++	-
Concordance/transcriptome	+++	+
Concordance/genome	+++	+



Lee *et al*, Cancer Cell 2006, PMID:16697959

Opportunities and Challenges for Next-Gen Cancer Model Development

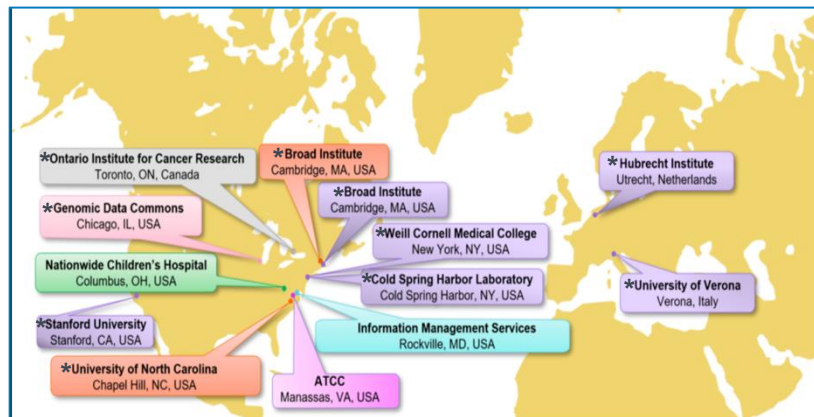
Applications in Oncology
Cancer Modeling, Cancer Biology
Drug Discovery and Tox Studies
Precision Medicine



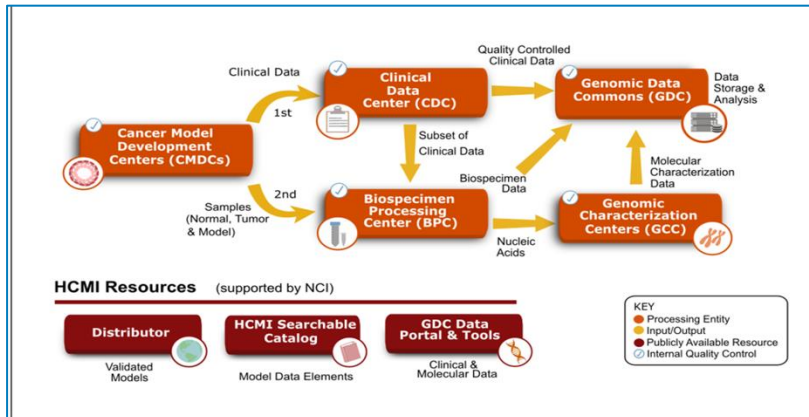
Lo et al, Nature Cancer 2020, PMID:PMC8208643

The Human Cancer Model Initiative (HCMI)

The HCMI is an international consortium founded by the **National Cancer Institute** and dedicated to generating **next-generation, patient-derived cancer models as a community resource** to facilitate cancer research



Source: <https://www.cancer.gov/ccg/research/functional-genomics/hcmi/about/cancer-model-development>



Source: <https://ocg.cancer.gov/programs/hcmi/nci-cancer-model-development>

*Managed by the Frederick National Laboratory for Cancer Research (FNLCR), Leidos Biomedical Research, Inc.

HCMI Models in the ATCC Catalogue - Adult

Human Cancer Models Initiative Searchable Catalog

Model: HCM-BROD-0106-C71 EXPANDED

◀ BACK TO SEARCH ADD MODEL TO MY LIST VIEW LIST

◀ Previous Model 17 of 332 Next ▶

MODEL DETAILS

Model Type: 3-D; Other (e.g. neurosphere, air-liquid interface, etc.)

Split Ratio: 1:2

Time to Split: N/A

Doubling Time: N/A

Tissue Status: Recurrent

MULTIPLE MODELS FROM THIS PATIENT (0)

There are no other models from this patient.

AVAILABLE MOLECULAR CHARACTERIZATIONS (8)

	Model	Tumor	Normal
WGS	✓	✓	✓
WGS	✓	✓	✓
BNA-seq	✓	✓	✗
DNA Methylation	✗	✗	✗

PATIENT DETAILS

Gender: Male

Race: White

Age At Diagnosis (years): 52

Age At Acquisition (years): 56

Disease Status: Progressive disease

Vital Status: Dead

Metastatic Therapy: No

Therapy: • Surgery

Chemotherapeutic Drug List Available: No

Clinical Tumor Diagnosis: Glioblastoma

Histological Subtype: NOS

Primary Site: Brain

Acquisition Site: Brain

Tissue Status: Recurrent

TNM Stage: N/A

Clinical Stage Grouping: N/A

Histological Grade: N/A

MODEL IMAGES (2)

Magnification: 4 x

REPOSITORY STATUS

Date Updated: May 03, 2024

Date Of Availability: September 30, 2019

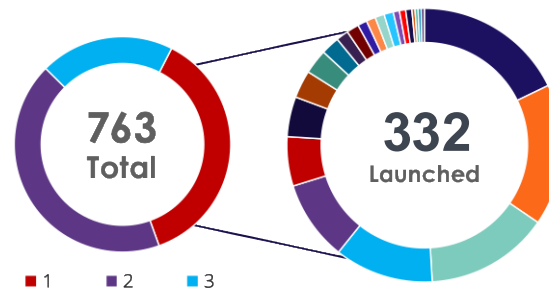
Licensing Required For Commercial Use: Yes

Date Created: September 27, 2019

EXTERNAL RESOURCES

[SEQUENCING FILES](#) [CASE METADATA](#) [MARKED SOMATIC MAP](#)

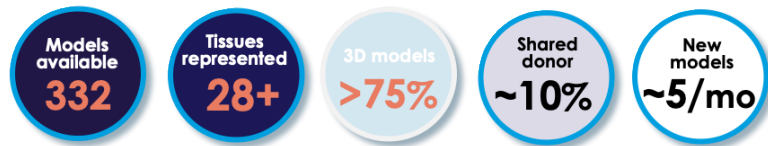
[VISIT PDM.123 TO PURCHASE](#)



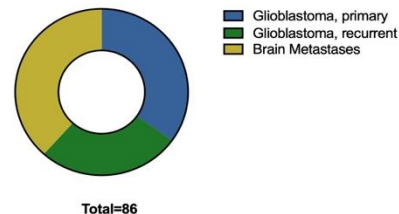
TOP 10 (90% of total)

Colon	55
Pancreas	52
Brain	50
Esophagus	36
Skin	29
Rectum	21
Stomach	18
Lung	7
Breast	7
Connective tissue	5

Collection includes models derived from rare adult and pediatric cancers such as rhabdomyosarcoma, leiomyosarcoma, Ewing sarcoma, and Wilms tumor.



Courtesy of Carolina Lucchesi @ ATCC

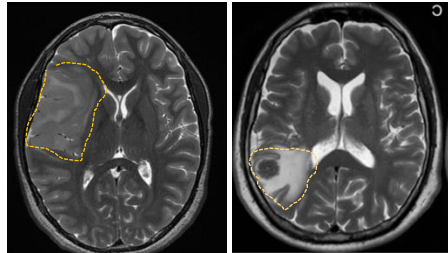


High-Grade Glioma Treatment and Challenges

Therapeutic Challenges and Opportunities

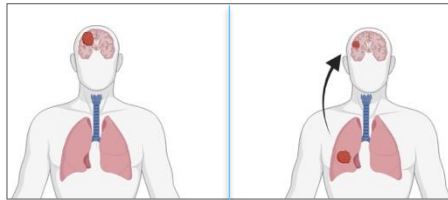
Primary High-Grade Gliomas: Distinction, Frequency, and Mortality

1 Primary vs. Secondary Brain Malignancies 2 Primary Malignant Brain Tumor Incidence, Mortality

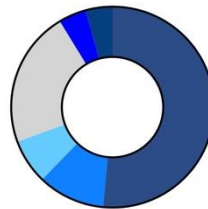
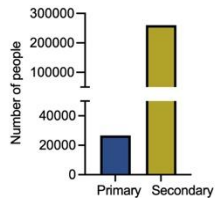


Glioblastoma

Brain Metastasis



Incidence Annual Brain Malignancy Diagnosis in U.S.

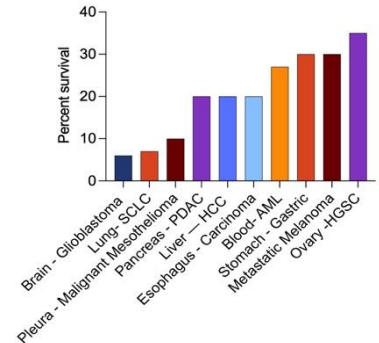


People in the U.S. living with primary, malignant brain tumor: Est. 338,000+

70-80% are glioma

Glioblastoma most common malignant glioma

Tumor Types with Highest Mortality (5-year Survival)

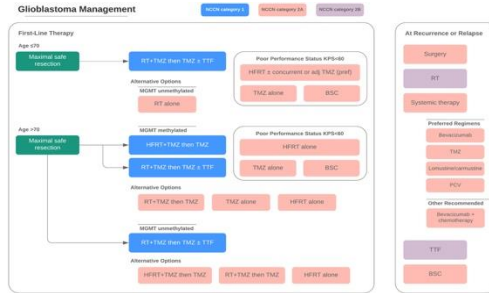


Sources: NCI SEER CSR 1975–2021; SEER Stat Facts; CBTRUS (Ostrom et al., Neuro-Oncology); ACS Cancer Facts & Figures.

High-grade gliomas and in particular glioblastoma <5% 5-year survival rate

Glioblastoma: Standard-of-Care (SoC) and Experimental Therapeutics

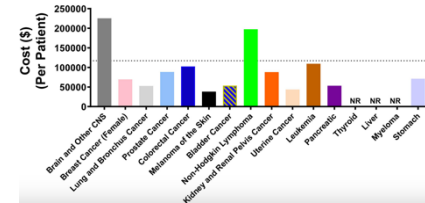
1 Standard-of-Care - Surgery plus Chemoradiotherapy+TTF



CA A Cancer J Clinicians, 2020, DOI: (10.3322/caac.21613)

Mehta, M et al, Critical review in Onc, 2017, PMID:28259296

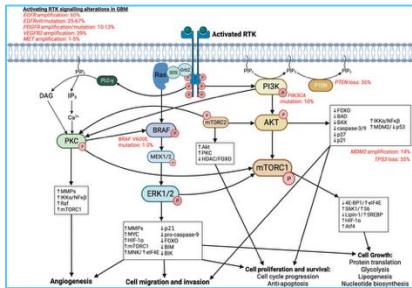
2 Treatment Costs



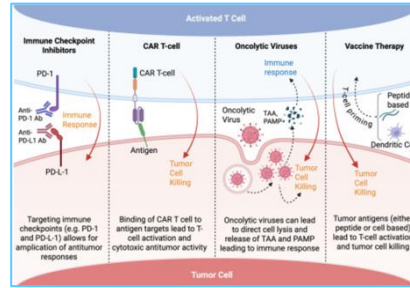
Approx. new U.S. cases/year



2 Targeted and Immunotherapy for glioblastoma in clinical trials



Dewdney et al, Signal Transduct Target Ther 2023
PMCID:PMC10587102



Song et al, Immunity 2025, PMID:40324379

Standard-of-Care is multi-modal

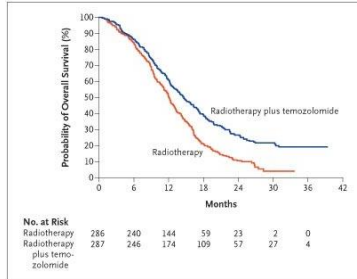
Promising therapies in clinical trials (~2000 clinical trials)

Costs up to \$250K/patient

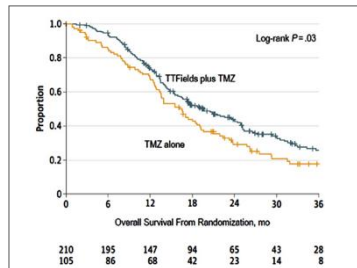
Clinical trials for glioblastoma are expensive (tens of millions of \$)

Survival with SoC and Brain Tumor Drug Approvals

1 Survival rates with SoC

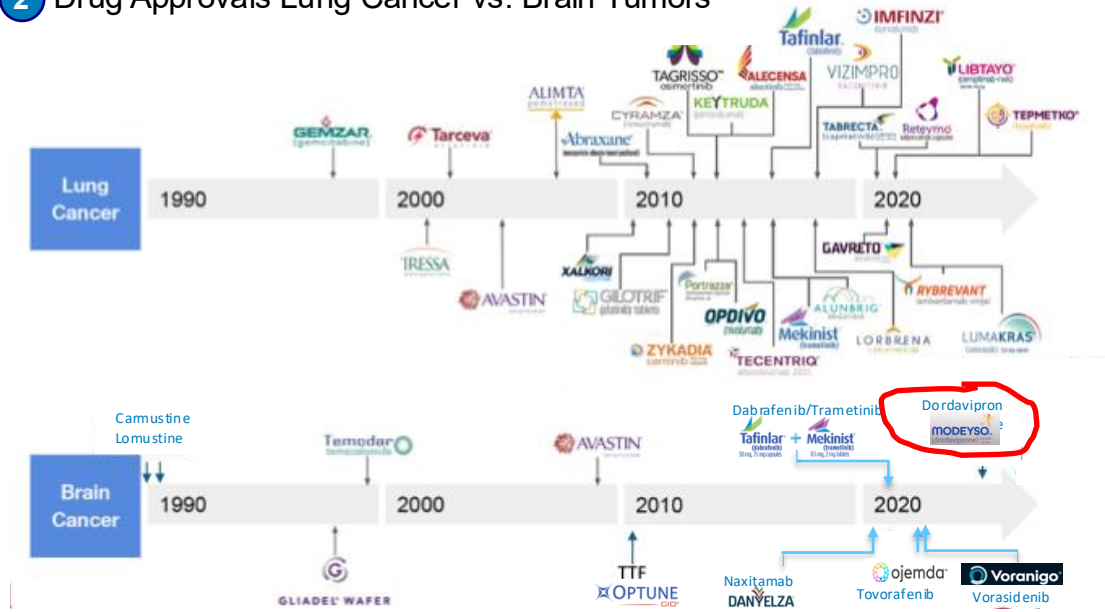


Stupp *et al*, Lancet 2005, PMID1575809



Mehta *et al*, Critical review in Onc 2017, PMID:28259296

2 Drug Approvals Lung Cancer vs. Brain Tumors



Challenges for Development of Malignant Brain Tumor Therapeutics

Tumor Heterogeneity

HGG contains multiple genetically distinct clones. Targeting one subpopulation allows resistant clones to repopulate.

Therapy Resistance, Plasticity, Glioma Stem Cells

A subpopulation of cells is resistant to radiation and chemotherapy, showing plasticity, driving tumor re-initiation after treatment.

Diffuse Infiltration

Tumor cells invade surrounding brain tissue far beyond the visible mass, making complete surgical resection impossible.

Near-Universal Recurrence

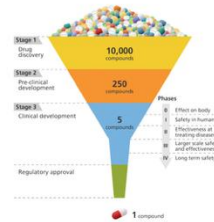
Nearly all HGGs recur, often with acquired resistance to TMZ and altered molecular profiles after treatment.

Blood-Brain Barrier

Tight junctions prevent most systemic drugs from reaching the tumor, severely limiting chemotherapy options.

Immunosuppressive TME

The tumor microenvironment is deeply immunosuppressive, blunting CAR-T and checkpoint inhibitor efficacy.



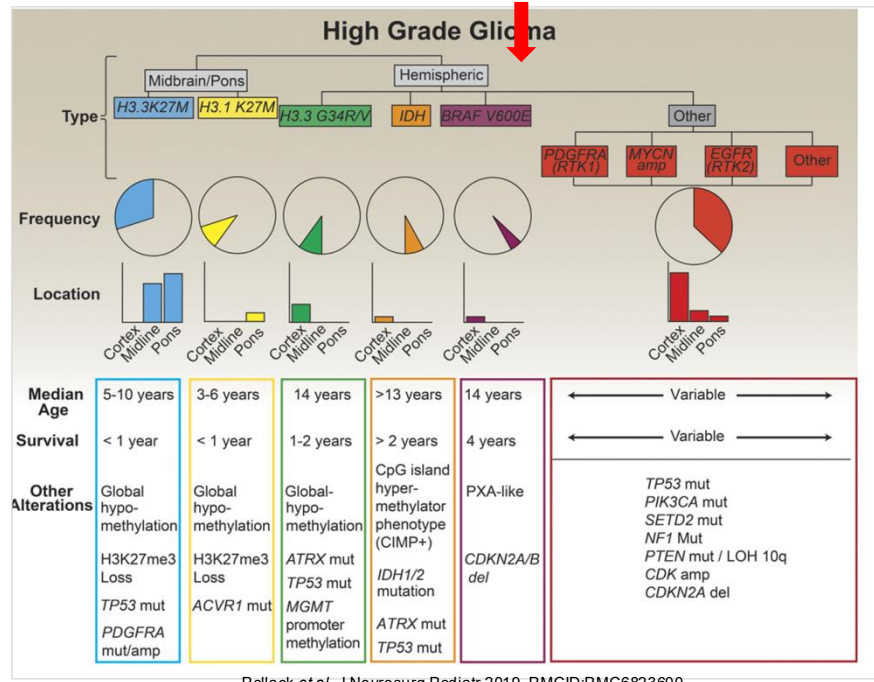
Several key features of malignant brain tumors contribute to difficulties in developing new treatments

Next-generation models recapitulating these key features will facilitate impactful research & pave the way for developing new therapies with high clinical efficacy

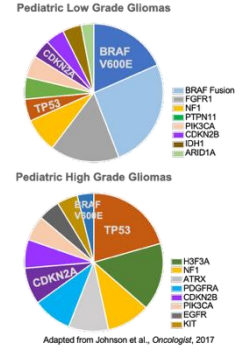
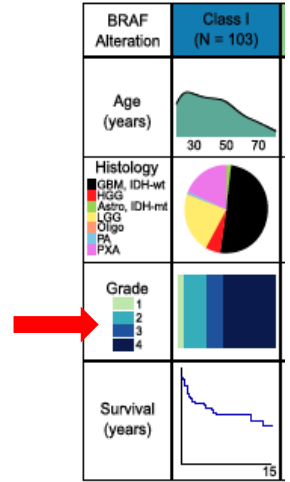
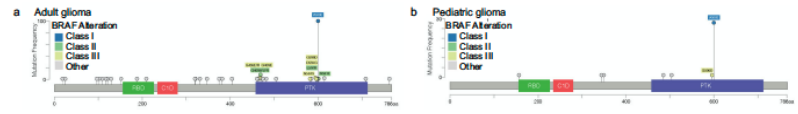
Malignant brain tumors are very diverse and have many subtypes

BRAF V600E-Mutant Glioblastoma as a Paradigm for Precision Medicine in Brain Tumors

1 BRAF V600E-altered glioma occur in pediatric + adult patients but in different types



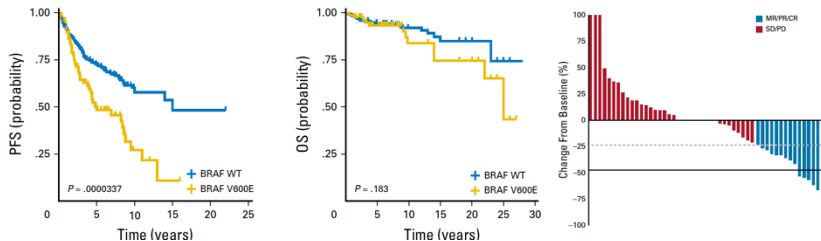
Pollack et al, J Neurosurg Pediatr 2019, PMID:PMC6823600



Schreck et al, NPJ Precis Oncol 2023, PMID:PMC9975216

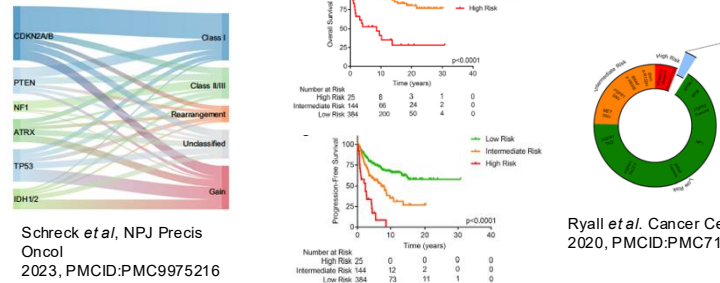
The BRAF V600E-mutant glioma frequency, types, and prognosis

1 BRAF V600E-altered glioma are chemotherapy-resistant

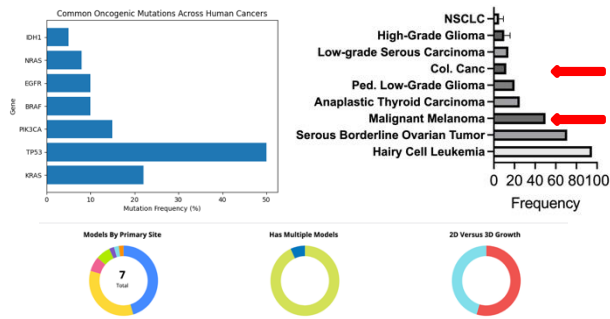


Lassaletta *et al.*, JCO 2017, PMID:PMC5791837

2 BRAF V600E co-mutations affect risk for progression

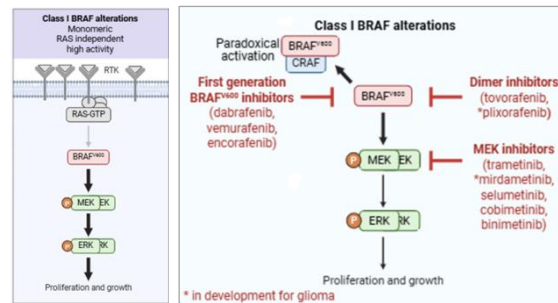


3 BRAF V600E mutations found in various cancer



BRAF V600E models in HCM1 catalogue

4 BRAF/MEK inhibitors in the clinical arena

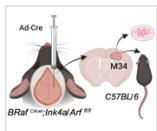


Courtesy of Karisa Schreck

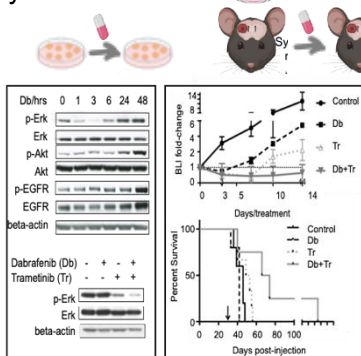
Research Objectives Based on Low-Response Rates

- 1 MAPK pathway reactivation after BRAF inhibition
- 2 Combined BRAF and MAPK inhibition response rates

Syngeneic Mouse Models For BRAF V600E-mutant High-Grade Glioma:



Grossauer *et al.* Oncotarget, 2016, PMID:PMC5342782



Dabrafenib (Db)=BRAF inhibitor
Trametinib (Tr)=MEK inhibitor

Understand how BRAF V600E mutated high-grade gliomas respond to clinically relevant, molecular-targeted inhibition (BRAF V600E and MEK inhibition)

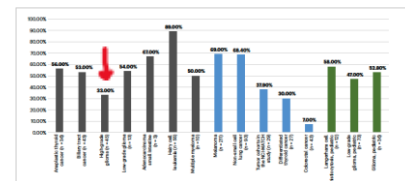
Identified potential mechanisms of therapy escape

Find novel therapeutic opportunities to combine with BRAF V600E- targeted therapy to overcome resistance

- 2 Combined BRAF and MAPK inhibition response rates

	Grade III (n=13)	Glioblastoma (n=31)	Age 18-39 years (n=22)	Age ≥40 years (n=23)
Objective response rate by investigator, % (95% CI)	38 (13.9-68.4)	32 (16.7-51.4)	50 (28.2-71.8)	17 (5.0-38.8)
Patients responding at 12 months by investigator assessment, % (95% CI)	100	67 (28-87.8)	89 (43-98.4)	50 (5-84.5)
Median progression-free survival by investigator, months (95% CI)	3.8 (1.7-NR)	2.8 (1.8-13.7)	19.5 (5.5-41.4)	1.7 (0.9-2.5)
Median overall survival, months (95% CI)	45.2 (6.3-NR)	13.7 (8.4-25.6)	45.2 (17.9-NR)	8.7 (3.7-11.7)

Wen *et al.*, The Lancet Oncology 2022, PMID:34838156



Subbiah *et al.*, Nat Med 2023, PMID:PMC10202803

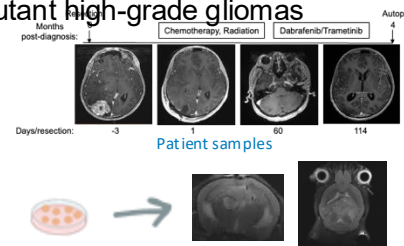
- 3 Model Development for BRAF V600E-mutant high-grade gliomas



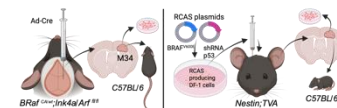
Patient samples

- 1 conventional cell line
- 1 xenograft model
- 1 orthotopic mouse model

Lerner *et al.* Cancer Res 2015, PMID:PMC4698003



3 patient-derived spheroid lines
+ matched patient-derived xenografts



2 syngeneic mouse models- fully immunocompetent

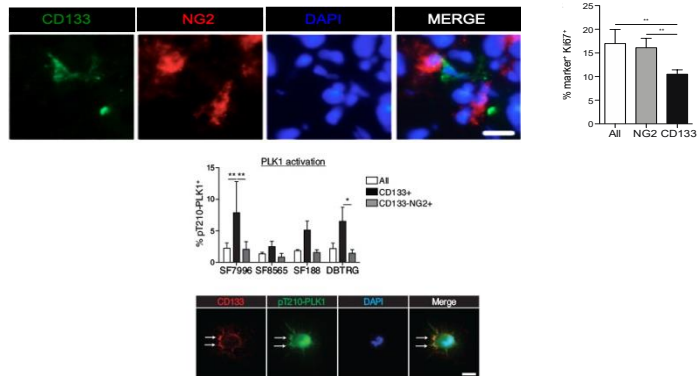
Xing, Panovska *et al.* Cell Reports Med 2025, PMID:PMC12208339

Unraveling Molecular Therapy Resistance in Novel BRAF V600E HGG Models

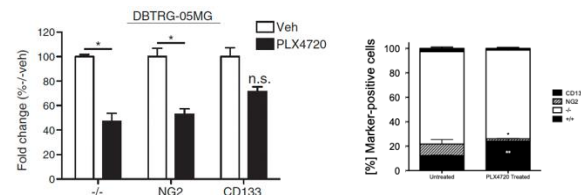
*Identifying novel therapeutic combinations to
overcome resistance*

Targetable Therapy Escape Mechanisms in BRAF V600E-mutant Glioblastoma

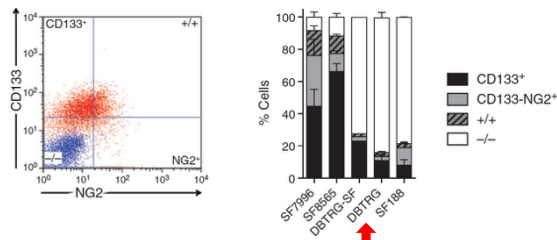
1 Intratumoral heterogeneity of BRAF V600E-mut glioblastoma



2 BRAF inhibitor treatment does not eliminate CD133+ stem-like cells

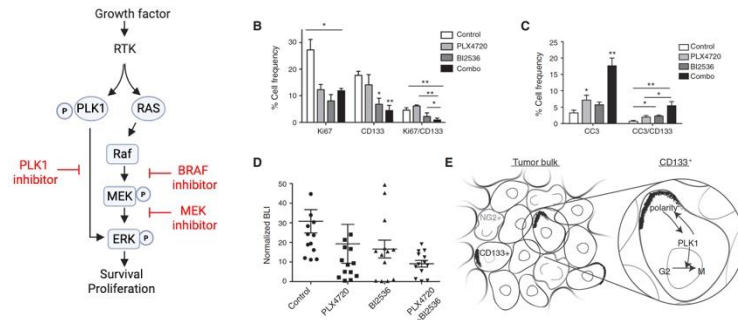


4 BRAF V600E-mutant glioblastoma cell line (serum) is homogeneous



- **Low plasticity and heterogeneity (few CD133+ and NG2+ cells)**

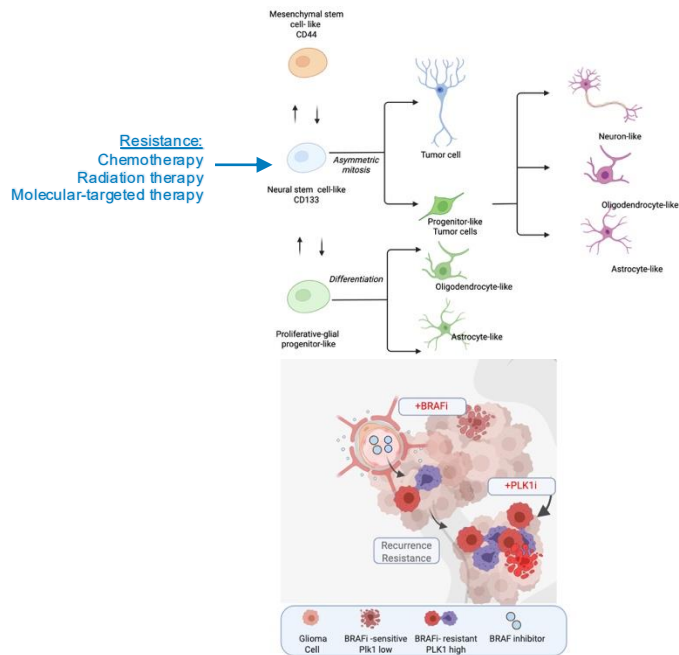
3 Combination therapy eliminates CD133+ cells = cancer stem cells and increases survival in xenografts



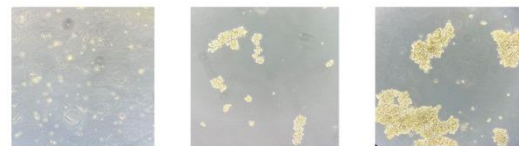
Lerner *et al.* Cancer Res 2015, PMID:PMC4698003

Next-gen BRAF V600E-mutant Glioblastoma Models are Sensitive to PLK1 inhibition

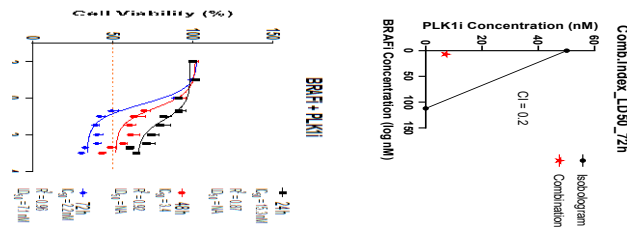
1 High-grade glioma exhibit plasticity, intra-tumoral heterogeneity, differentiation



2 BRAF V600E-mutant 3D/spheroid glioblastoma models sensitive to PLK1i



Morphology of an epithelioid glioblastoma tumor 3D spheroid cell line HCM-STAN-1297-C71. Morphology. left. 2 h after thawing; center. at low density; right. at high density.



Cell line	Type	IC50 BRAF1 (nM)	IC50 PLK1I (nM)	IC50 Combo
Patient-derived cell line 2- BRAF V600E	Human adult glioblastoma	71.13	172.8	43.83
HCM-STAN-1297-C71_A- BRAF V600E	Human pediatric glioblastoma	1.73	~2469	3.55
2341-luc-BRAF V600E	Murine high-grade glioma	0.63	NR	0.44

Petrtsch Lab, Unpublished Data, Do not post

PLK1 inhibition in the clinical arena

1 PLK1 inhibitors FDA-approved for non-brain cancers

TABLE 1 | Clinical trials based on PLK1 inhibitors.

NCT number	Phase	Disease	PLK1i	Ref
NCT02211859	I	Advanced solid tumors	BI2536	(127)
NCT02211872	I	Advanced solid tumors	BI2536	(128)
NCT00412880	II	Small cell lung cancer	BI2536	(129)
NCT01662505	I	Acute myeloid leukemia	BI6727 (Volasertib)	(133)
NCT00904856	IIIa	Acute myeloid leukemia	BI6727 (Volasertib)	(131)
NCT01023958	II	Urothelial cancer	BI6727 (Volasertib)	(132)
NCT00824408	II	NSCLC	BI6727 (Volasertib)	(133)
NCT01014429	I	Advanced solid tumors	NMS-128937 (Onvasertib)	(134)
NCT03598035	I	Advanced solid tumors	GS441364	(135)
NCT01173099	I	Advanced solid tumors	TAK980	Not yet published
NCT01538537	I	Advanced solid tumors	ON01910 (Rigosertib)	(136, 137)
NCT00654646	III	Myelodysplastic syndrome	ON01910 (Rigosertib)	(138)
NCT01168011	I	Acute myeloid leukemia; Advanced solid tumors	ON01910 (Rigosertib)	(138)

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7

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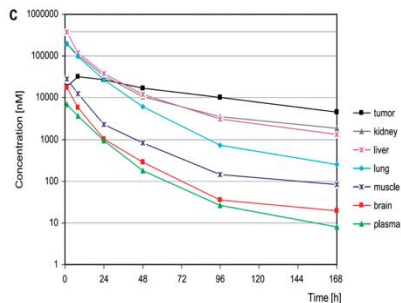
Plk1 inhibitors are in clinical trials against several cancer types

Volasertib: FDA rare disease breakthrough and orphan drug designation
-stable in brain
-high systemic toxicity

New inhibitors in early stages of development for brain tumors (e.g., onvasertib) showing promising results in pediatric patients

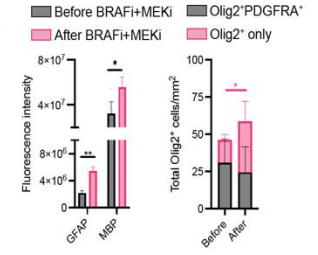
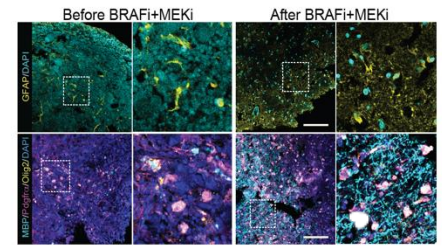
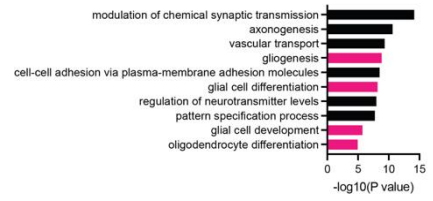
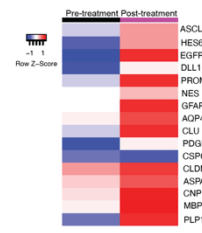
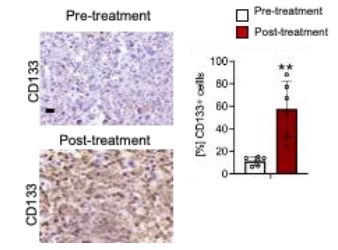
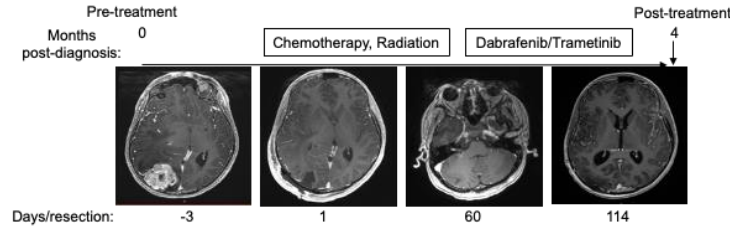
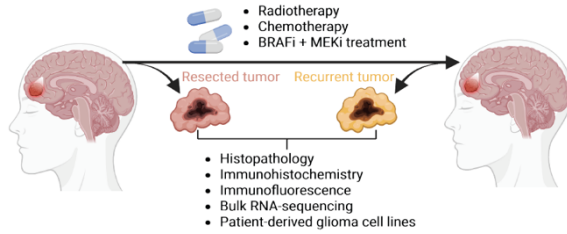
Combination with BRAF/MEK inhibitors to be determined

2 PLK1 inhibitor Volasertib is stable in the brain

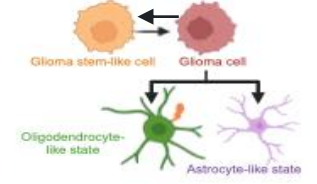


Rudolph et al, Clin Can Res 2009, PMC19383823

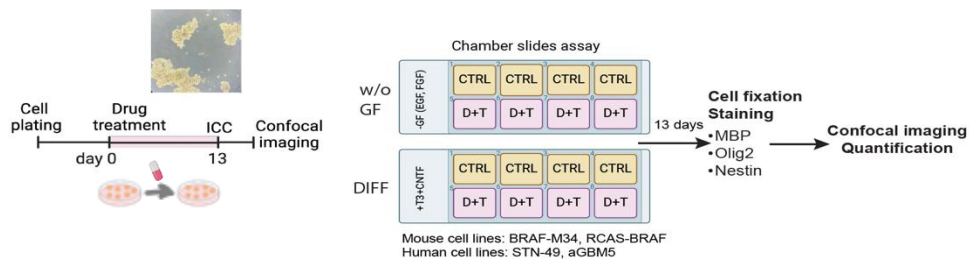
Analyses of Pre- and Post-treatment BRAF V600E-mutant Glioblastoma Samples



BRAF/MEK inhibition in patients upregulated not only stem-cell markers but also glial differentiation, indicative of therapy-induced increases in cell plasticity and differentiation



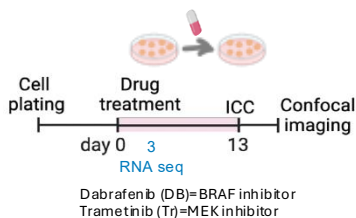
BRAF/MEK Inhibition Direct Effects on Cell States and Differentiation in Patient-Derived 3D Tumor Models



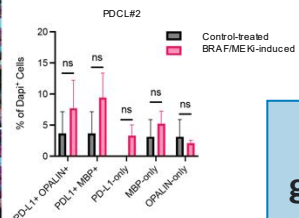
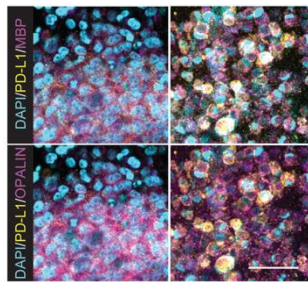
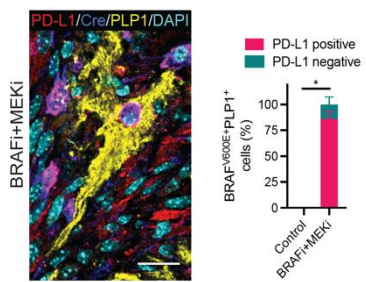
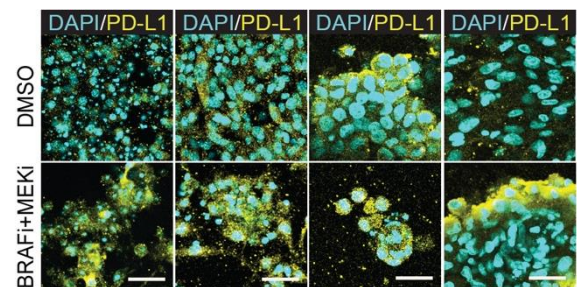
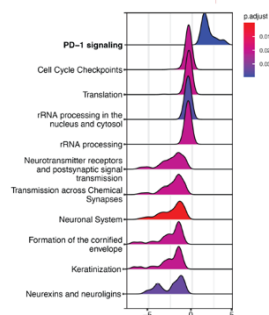
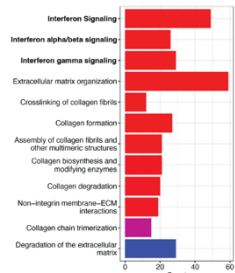
BRAF/MEK inhibition upregulated not only stem-cell markers but also glial differentiation, in cell culture-based assays, indicative of direct effects on tumor cell plasticity and differentiation

Xing, Panovska et al, Cell Reports Med 2025, PMID:PMC12208339

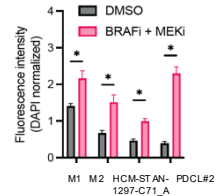
Immune-modulatory Effects of BRAF/MEK inhibitors?



Xing, Panovska et al, Cell Reports Med 2025, PMID:PMC12208339

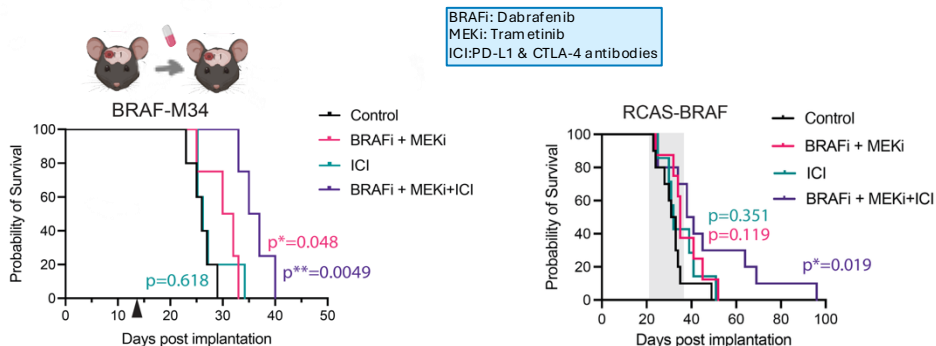


Petritsch Lab, Unpublished, Do not post



BRAF+MEK inhibitors induced glioma cell differentiation along with upregulated antigen presentation & up-expression of immune checkpoint inhibitors

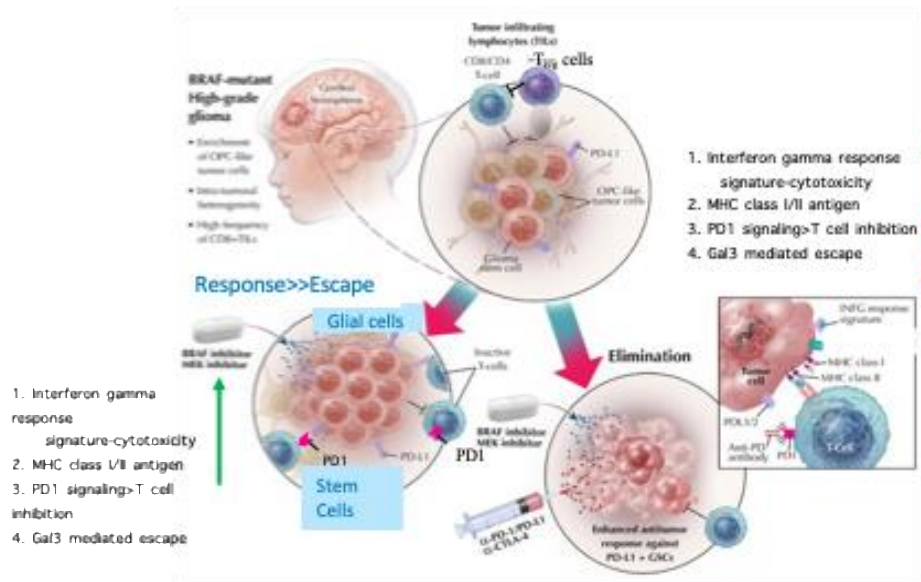
Immune Checkpoint Inhibition Enhancing BRAF/MEK inhibition Anti-Tumor Effects?



BRAF/MEK inhibitor treatment **inactivates T cells** mediating immune escape

BRAF/MEK inhibition Synergizes with **Immune Checkpoint Inhibition** to Reactivate **T cells** and Overcome Therapy Resistance.

Summary



MAIN FINDINGS

- BRAF/MEK inhibition induces **glioma cell state transitions** that triggers increases in stem cells, and immune evasion
- BRAFi+MEKi activates the interferon response and anti-tumor immunity, while simultaneously **suppressing T cells via PD-L1 upregulation in glial cells**
- Glial differentiation and immune evasion could be mediated by therapy-induced **secretion of immune modulatory factors**
- High PD-L1 expression in BRAF-mutant GBM provides a criterion for anti-PD-1 therapy
- **Concurrent BRAF/MEK and checkpoint inhibition** enhances anti-tumor immunity and survival

CLINICAL IMPLICATION

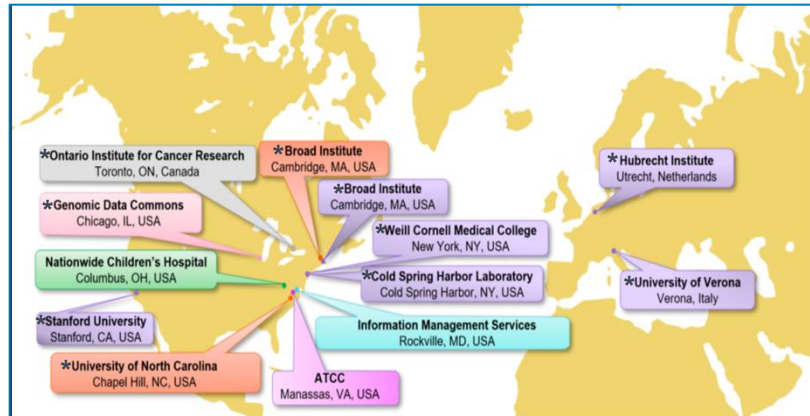
Our preclinical findings highlight the potential of integrating BRAFi+MEKi treatment with PLK1 inhibition and with ICI, with emphasis on concurrent treatment

The HCMI Pediatric Cancer Models

Introducing the developing collection of models

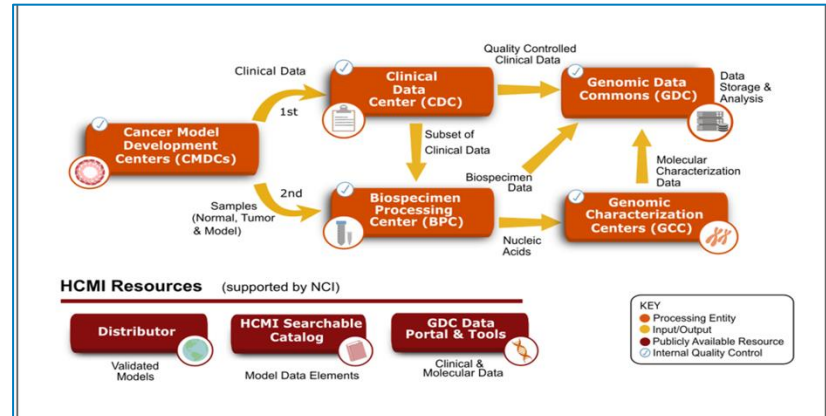
The Human Cancer Model Initiative (HCMI)

The HCMI is an international consortium founded by the **National Cancer Institute** and dedicated to generating next-generation, patient-derived cancer models as a community resource to facilitate cancer research



(Source: <https://www.cancer.gov/ccg/research/functional-genomics/hcmi/about/cancer-model-development>)

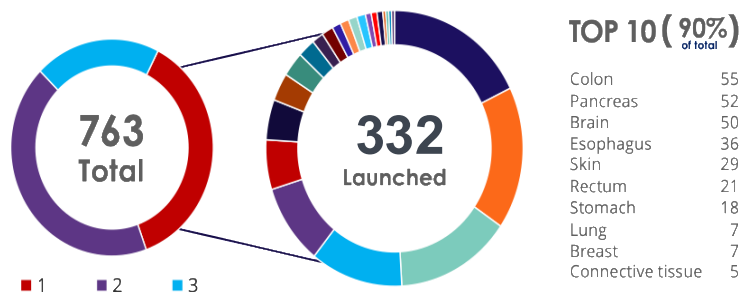
(* Managed by the Frederick National Laboratory for Cancer Research (FNLCR), Leidos Biomedical Research, Inc.)



(Source: <https://ocg.cancer.gov/programs/hcmi/nci-cancer-model-development>)

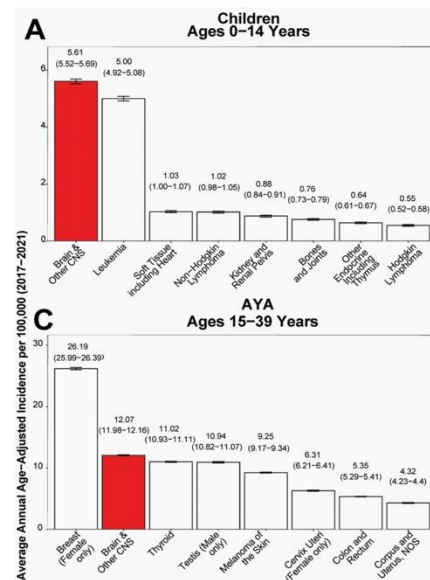
HCFI Has Focused on Adult Cancer

- Current HCFI models are mainly from adult tumors



Collection includes models derived from rare adult and pediatric cancers such as rhabdomyosarcoma, leiomyosarcoma, Ewing sarcoma, and Wilms tumor.

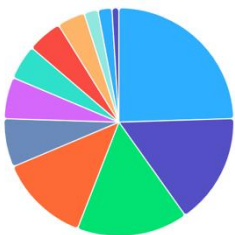
12 models are from pediatric patients



Neuro Oncol, Volume 26, Issue Supplement_6, October 2024, Pages vi1-vi85, <https://doi.org/10.1093/neuonc/noae145>.

Rationale for Developing Next-Generation Pediatric Solid Tumor Models

Childhood Cancer Incidence, Age 0-19, 2017-2021



- Leukemia: 25%
- Lymphoma: 16%
- Brain and Central Nervous System: 16%
- Epithelial Neoplasms and Melanomas: 13%
- Rhabdomyosarcoma (Soft Tissue Tumors): 7%
- Germ Cell Tumors: 6%
- Neuroblastoma and Other Peripheral Nerve Cell Tumor: 5%
- Bone Tumors: 5%
- Kidney Tumors (including Wilms Tumor): 4%
- Liver Tumors (including Hepatoblastoma): 2%
- Retinoblastoma: 2%
- Other: 1%

Characteristics

- Rare (4-5 x lower incidences than in adults)
- Different treatment strategies (minimize long-term sequelae)
- Different pathogenetic mechanisms (developmental, MAPK pathway vs. RTK, onc-fusions, epigenetic regulation)

>100 subtypes of pediatric solid tumors

REVIEW

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

Sofian M. Pfister^{1,2}, Miguel Reyes-Mugica³, Isako K.C. Chan⁴, Henrik Hastie⁵, Alexander I. Lazar⁶, Sabrina Rios⁷, Araya Ferrer⁸, Jason A. Jarambowski⁹, Kristy Patricia Jones¹⁰, D. Ashley Hill¹¹, Thomas S. Jacques¹², Peter Wesseling¹³, Charles H. Lopez-Torres¹⁴, Andreas von Deimling¹⁵, Christian P. Koels¹⁶, Ian A. Cree¹⁷, and Kris Alaggio¹⁸

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 arising 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 created in a multidisciplinary approach, incorporating morphologic, IHC, and molecular characteristics. The volume is organized according to organ sites and provides a single, state-of-the-art compilation of pediatric tumor types. A special emphasis was placed on "biomarkers" which variably recapitulate the morphologic/molecular features 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 to a molecularly driven classification systematically taking recent discoveries in pediatric tumor genetics into account.

INTRODUCTION

Why Pediatric Tumors Need Separate Classification

Childhood cancers are fundamentally different in nature from those occurring in adults. Despite being extremely

rare, they represent the second or third most common cause of cancer-related death in children (1). In contrast to malignancies in adults, which are mostly of epithelial origin, many childhood cancers are

highly diverse. Childhood cancer incidence (2,3), including brain, bone, and soft tissue, is increasing (4,5). The WHO Classification of Pediatric Tumors (6) is a multidisciplinary effort, incorporating morphologic, IHC, and molecular characteristics. The volume is organized according to organ sites and provides a single, state-of-the-art compilation of pediatric tumor types. A special emphasis was placed on "biomarkers" which variably recapitulate the morphologic/molecular features of organs from which they originate.

Corresponding Author: Sofian M. Pfister, MD, Department of Pediatrics, University of Colorado Denver, Aurora, CO, USA. Email: sofian.pfister@ucdenver.edu

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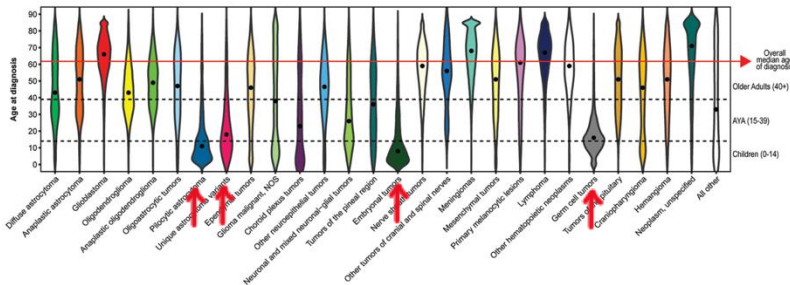
CANCER DISCOVERY | 3

Pfister, SM et al; Cancer Discov. 2022; PMID:PMC9401511

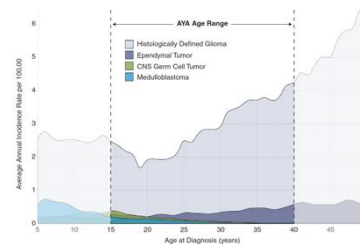
Adult versus Pediatric Brain Tumors Incidence & Subtypes

1 Distribution of age at diagnosis by selected primary brain tumors

Price et al; Neuro-oncology 2024; <https://doi.org/10.1093/neuonc/noae145>



2 Overview of the relative incidence of brain tumors with age



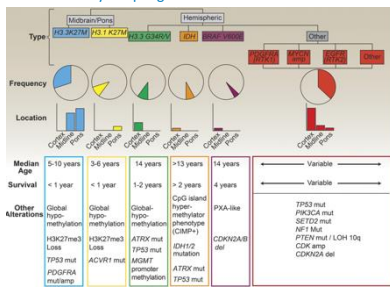
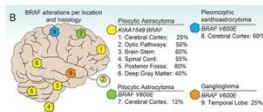
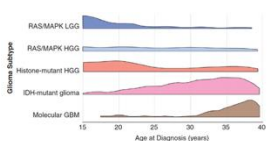
Lim-Fat et al; Neuro-oncology 2025, PMID:PMC11726256

3 Incidence of different subtypes of glioma, medulloblastoma, and ependymoma

Glioma

Low-grade glioma are the most common

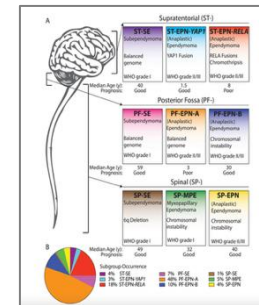
Pediatric high-grade glioma are rare but devastating and carry unique genetic alterations



Medulloblastoma

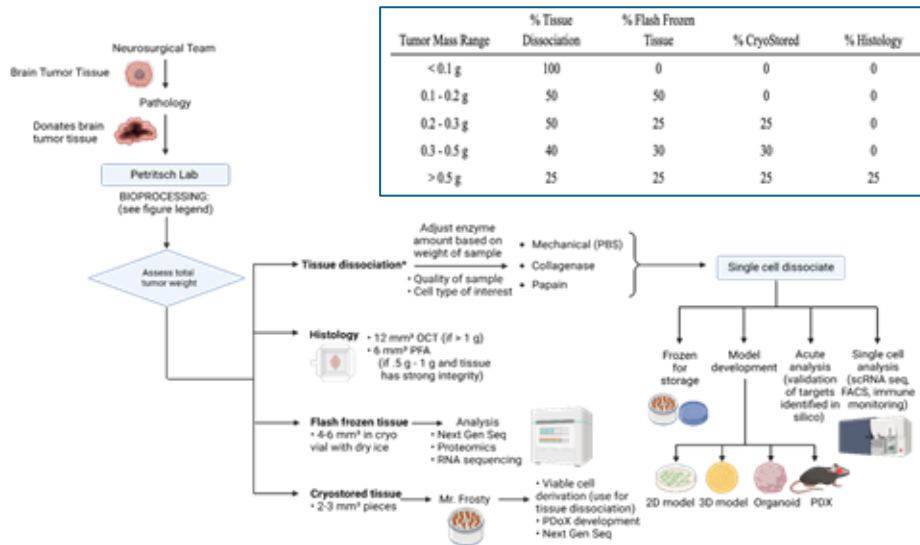
Subgroup	WNT	SHH	Group 3	Group 4
Incidence	10%	30%	35%	
Subtype	WNT α	SHH α	SHH β	SHH δ
Gender	♂ ♀	♂ ♀	♂ ♀	♂ ♀
Subtype proportion	2	5	10	40
Age	3-17	>10	0-3	0-3
Metastases	9%	21%	20%	33%
5 year survival	97%	100%	70%	90%
Copy Number Changes	6	TP53 del, CDKN2A del, PTEN loss	Balanced genome	10q22.3, 11q13.2, 17p11.2
Other events	TP53 mutation, PTEN loss, CDKN2A del	TP53 mutation, PTEN loss, CDKN2A del	TP53 mutation, PTEN loss, CDKN2A del	TP53 mutation, PTEN loss, CDKN2A del
Histology	Classic, LCA(w/m)	Desmoplastic, Nodular	Extensive nodularity	LCA

Ependymoma



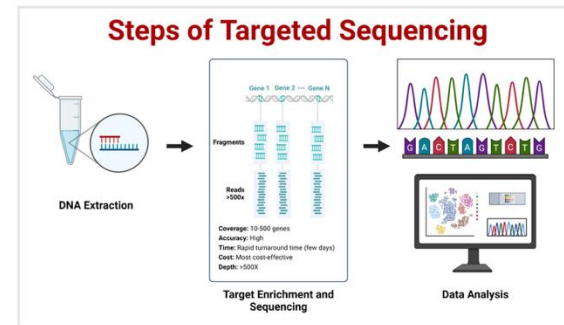
Pediatric Cancer Model Development - Workflow

Established Bioprocessing Workflow in our laboratory



Tumor Mass Range	% Tissue Dissociation	% Flash Frozen Tissue	% CryoStored	% Histology
<0.1 g	100	0	0	0
0.1 - 0.2 g	50	50	0	0
0.2 - 0.3 g	50	25	25	0
0.3 - 0.5 g	40	30	30	0
>0.5 g	25	25	25	25

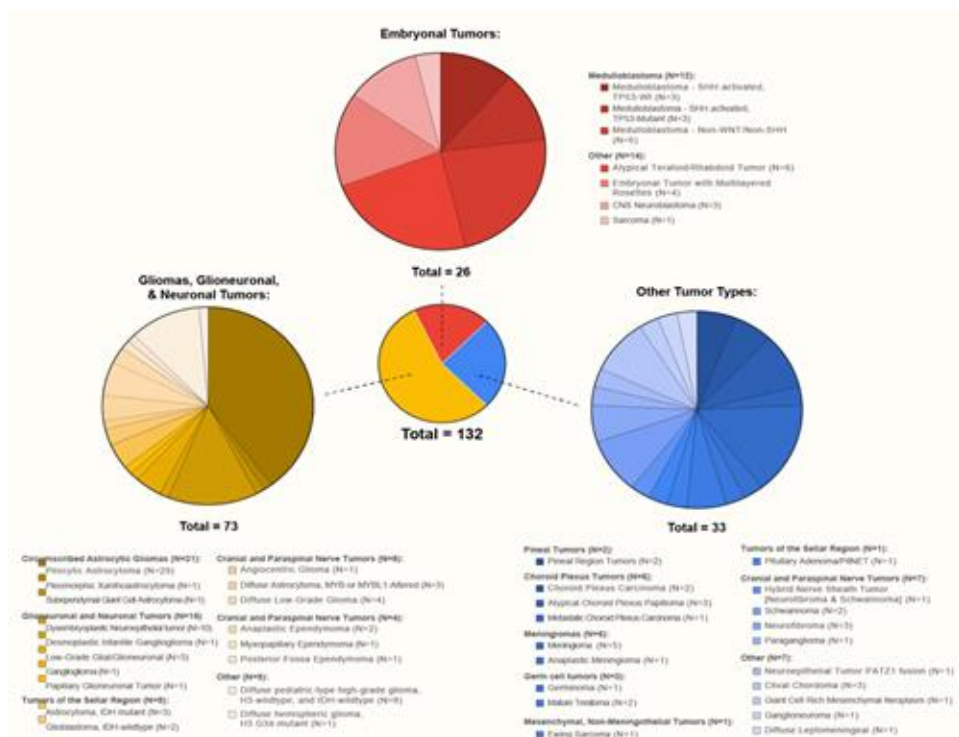
Established Mutation Identification



Stanford Actionable Mutation Panel ~250 genes, UCSF 500 Caris (WES)

An established biobanking workflow with standardized targeted sequencing of patient Tumors to identify recurrent mutations, including oncogenes

Representative Cohort of Pediatric Brain Cancer Types at Stanford Pediatrics CMDC



No. of Sellar Cases	Tumor Type	Gene/Molecular Profiles Characteristically Altered
3	Astrocytoma, IDH-mutant	IDH1, IDH2, ATRX, TP53, CDKN2A/B
2	Glioblastoma, IDH-wildtype	IDH-wildtype, TERT promoter, chromosomes 7/10, EGFR
2	Diffuse astrocytoma, MIB-1 or MIB1 altered	MIB1, MIB1
1	Angiocentric glioma	MIB1
4	Diffuse low-grade glioma, MAPK pathway-altered	FGFR1, BRAF
1	Diffuse hemispheric glioma, H3 G34-mutant	H3 G34, TP53, ATRX
9	Diffuse pediatric-type high-grade glioma, H3-wildtype and IDH-wildtype	IDH-wildtype, H3-wildtype, PDGFRA, MYCN, EGFR (methylome)
30	Pilo-cytic astrocytoma	KIAA1549-BRAF, BRAF, NF1
1	Pleomorphic xanthoastrocytoma	BRAF, CDKN2A/B
1	Subependymal giant cell astrocytoma	TSC1, TSC2
2	Choroid glioma	PRKCA
1	Ganglion cell tumors	BRAF
10	Dysmaturational neuroepithelial tumor	FGFR1
1	Papillary glioneuronal tumor	PRKCA
1	Diffuse leptomenigeal glioneuronal tumor	KIAA1549-BRAF fusion, 3p (methylome)
3	Supratentorial ependymoma	ZFTA, RELB, YAP1, MAML2
1	Posterior fossa ependymoma	H3 K27me3, EZH2 (methylome)
6	Medulloblastoma, SHH-activated	TP53, PTCH1, SUFU, SMO, MYCN, GLI2 (methylome)
6	Medulloblastoma, non-WNT/non-SHH	MYC, MYCN, PRDM8, KDM6A (methylome)
1	Atypical teratoid/oligo-dendrogloma	SMARCB1, SMARCA4
5	Embryonal tumor with multilayered rosettes	C19orf6, DICER1
3	CNS-neuroblastoma, FOXR2-activated	FOXK2
6	Meningeomas	NF2, AKT1, TRAF7, SMO, PIK3CA, KLF4, SMARCB1, BAP1 in subtypes; H3K27me3, TERT promoter, CDKN2A/B in CNS WHO grade 1

- ~90% of cases yield tissue for research
- 45 subtypes of brain cancer and 12 subtypes of non-CNS cancers captured over a 4-year period
- Longitudinal collection is crucial to capture diverse, rare cancers

Pediatric Cancer Types for HCMI

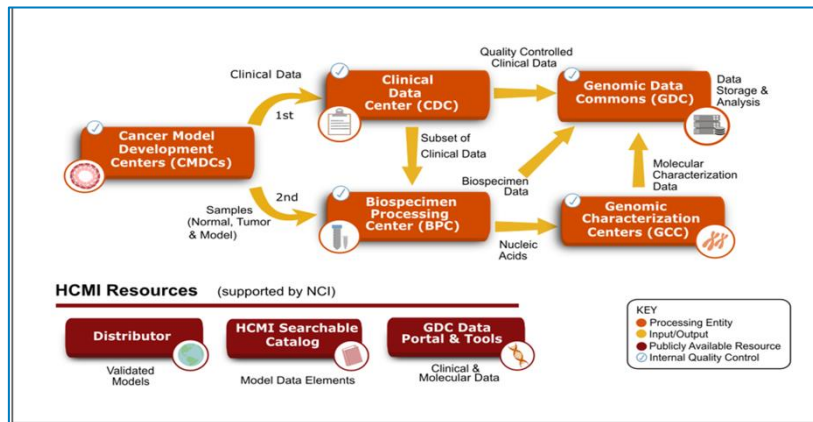
▪ Pediatric central nervous system (CNS) solid tumors

- Diffuse Midline Glioma (DMG), H3-K27-mutant
- Diffuse High-Grade Glioma (HGG), MAPK activated, H3 and IDH wildtype & IDH-mutant
- Recurrent Lower Grade Glioma
- Medulloblastoma
- Ependymoma

▪ Non-CNS pediatric solid tumors

- Malignant Rare Soft Tissue and Bone sarcoma
- Wilms' Tumor
- Neuroblastoma
- Hepatoblastoma

▪ Other rare cancer types and subtypes as feasible.



(Source: <https://ocg.cancer.gov/programs/hcmi/nci-cancer-model-development>)

60-70% success rate for model development

27 subtypes of brain cancer and 4 of non-CNS cancer were captured in models

Challenges: small tumor size

Planning and management oversight by the team at Frederick National Laboratory for Cancer Research, Leidos Biomedical Research, Inc.

This project has been funded in part with federal funds from the Childhood Cancer Data Initiative (CCDI), National Cancer Institute, National Institutes of Health, Task Order numbers 75N91020F00035, under contract no. 75N91019D00024.

Stanford Pediatric CMDC

Emon Nasajpour
Dena Panovska
Ruolun Wei
Claudia K. Petritsch

Neurosurgery Pediatric Clinic

Laura Prolo
Kelly Mahaney
Cormac Maher
Westly Phillips
Gabriella Morton

Monje Lab Rapid Autopsy Program

Michelle Monje
Isabella Chau
Ann-Helen Liljensten
Lijun Li

Bass Center Tissue Bank (BCTB)

Sheri Spunt
Raya Saab

Min Huang

Neuropathology

Hannes Vogel
John Newman
Jeff Nirschl
David Solomon

Valley Childrens Hospital Fresno

Natalie Limoges
Patricia Clerkin
Melissa Martinez

National Cancer Institute

Daniela S. Gerhard
Justin Benavidez
Cindy W. Kyi
Julyann Perez-Mayoral
Eva Tonsing-Carter
Jean-Claude Zenklusen
Louis Staudt
CCDI: Subhashini Jagu,

BPC: Sarah Coppens

CDC: Megan Stine

ATCC: Carolina Lucchesi

Frederick National Lab Leidos Biomedical

Research Inc: Rachana Agarwal, Conrado Soria
The HCI Network



This project has been funded in part with federal funds from the Childhood Cancer Data Initiative (CCDI), National Cancer Institute, National Institutes of Health, Task Order numbers 75N91020F00035, under contract no. 75N91019D00024.

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Q&A

Join Us at Our Next Webinar

CCDI Pediatric, Adolescent, and Young Adult Rare Cancer Study

Monday, April 27, 2026
12:00–1:00 p.m. ET

Come learn about the newly launched study from Mary Frances Wedekind, D.O., Associate Research Physician at NCI!

Learn more and register at events.cancer.gov/ccdi/webinar

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Questions? Email us at:
NCIChildhoodCancerDataInitiative@mail.nih.gov

Thank you for attending!



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