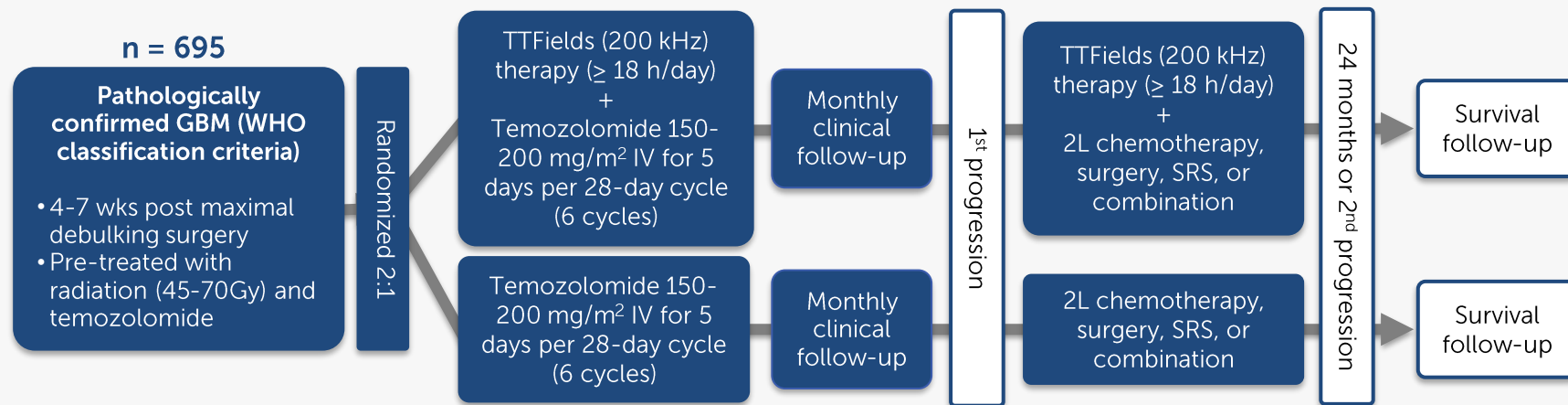




tumor treating fields clinical evidence

EF-14 Phase 3 pivotal trial evaluated Optune Gio + TMZ in 695 patients with ndGBM



Start date: June 2009
Primary completion: December 2016
Study completion: March 2017
Study sites: 83 (global)

Primary endpoint:

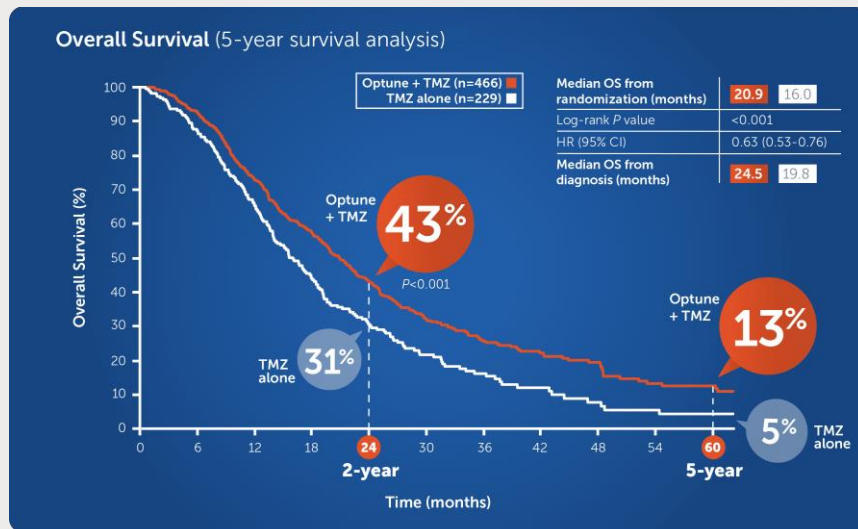
- Progression-free survival

Secondary endpoints:

- Overall survival

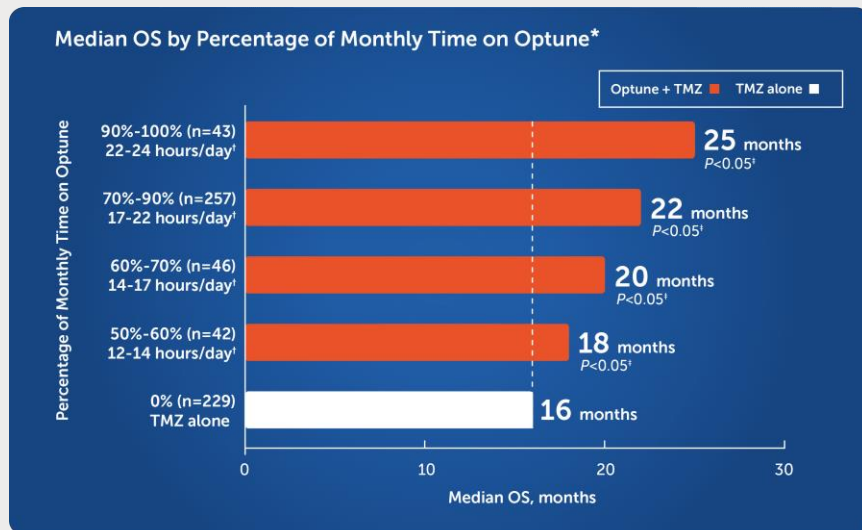
in ndGBM, Optune Gio + TMZ provided an unprecedented long-term survival benefit

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more time on Optune Gio predicted increased significant survival benefit

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29.3%
vs. 4.5%

5-YEAR PROBABILITY
OF SURVIVAL WITH
90% COMPLIANCE
(n=43) VS SURVIVAL
WITH TMZ ALONE

Journal of Neuro-Oncology (2019) 141:467–473
https://doi.org/10.1007/s11068-019-03027-y

CLINICAL STUDY



Increased compliance with tumor treating fields therapy is prognostic for improved survival in the treatment of glioblastoma: a subgroup analysis of the EF-14 phase III trial

S. A. Toms¹ · C. K. Kins² · G. Nicholas³ · Z. Raza⁴

Received: 30 August 2019 / Accepted: 27 November 2019 / Published online: 1 December 2019
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Abstract

Background Tumor treating fields (TTFields) is a non-invasive, antineoplastic therapy. In the EF-14 phase III trial in newly diagnosed glioblastoma, TTFields plus temozolomide (TTFields+TMZ) improved progression-free (PFS) and overall survival (OS) versus TMZ alone. Previous data indicate a ≥ 75% daily compliance improves outcomes. We analyzed compliance data from TTFields+TMZ patients in the EF-14 study to correlate TTFields compliance with PFS and OS and identify potential lower boundaries for compliance with improved clinical outcomes.

Methods Compliance was assessed by usage data from the NovoTTF-100A device and calculated as percentage per month of TTFields delivery. TTFields+TMZ patients were segregated into subgroups by percent monthly compliance. A Cox proportional hazard model controlled for sex, extent of resection, MGMT methylation status, age, region, and performance status was used to investigate the effect of compliance on PFS and OS.

Results A threshold value of 50% compliance with TTFields+TMZ improved PFS (HR 0.76, 95% CI 0.47–1.05) and OS (HR 0.67, 95% CI 0.45–0.96) versus TMZ alone with improved outcomes as compliance increased. At compliance > 90%, median survival was 24.9 months (25.7 months from diagnosis) and 5-year survival rate was 29.3%. Compliance was independent of gender, extent of resection, MGMT methylation status, age, region and performance status (HR 0.76; $p = 0.031$; OS at compliance: 3.75% vs. 2.9%).

Conclusion A compliance threshold of 50% with TTFields+TMZ correlated with significantly improved OS and PFS versus TMZ alone. Patients with compliance > 90% showed extended median and 5-year survival rates. Increased compliance with TTFields therapy is independently prognostic for improved survival in glioblastoma.

Keywords Glioblastoma · Tumor treating fields · Compliance · Monthly usage

Introduction

Glioblastoma (GBM) is the most common and aggressive adult brain tumor, accounting for 56% of all gliomas and 15% of all primary brain tumors with an annual incidence in the United States that increases with age—ranging from 0.2 per 100,000 in 0–19 year old population to the highest rate of 25.3 per 100,000 in the 75–94 year old population [1]. Glioblastoma remains incurable with a median survival of only 15 months overall [2]. The previous standard treatment for newly diagnosed GBM include maximally safe surgical resection followed by radiation therapy (RT) and adjuvant temozolomide (TMZ) chemotherapy [3].

Tumor treating fields (TTFields) are a unique treatment modality [4, 5] for GBM that effects rapidly dividing glioma cells through the action of low-intensity, intermediate

© S. A. Toms

* See the "Supplementary Materials" link on the journal website.

¹ Department of Neurosurgery, Western Sydney Medical School of Western Sydney University, Penrith, NSW, Australia

² Saint National University, Reading, South East, UK

³ Ontario Hospital Research Institute, Ottawa, ON, Canada

⁴ Tel Aviv Medical Center, Tel Aviv, Israel

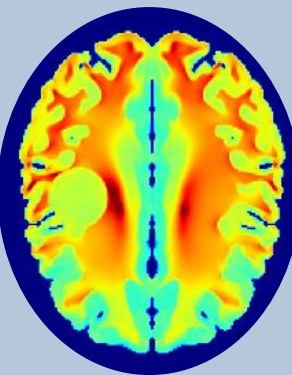


higher TTFields therapy dose can lead to increased efficacy

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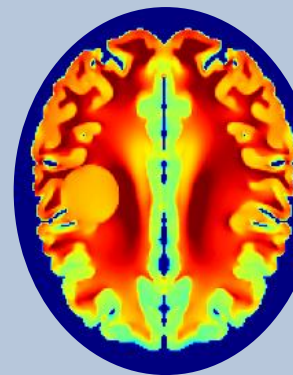


EXISTING ARRAYS AP channel, 1,364 mAmps



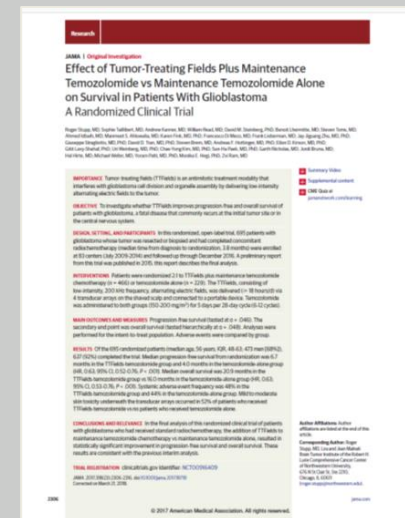
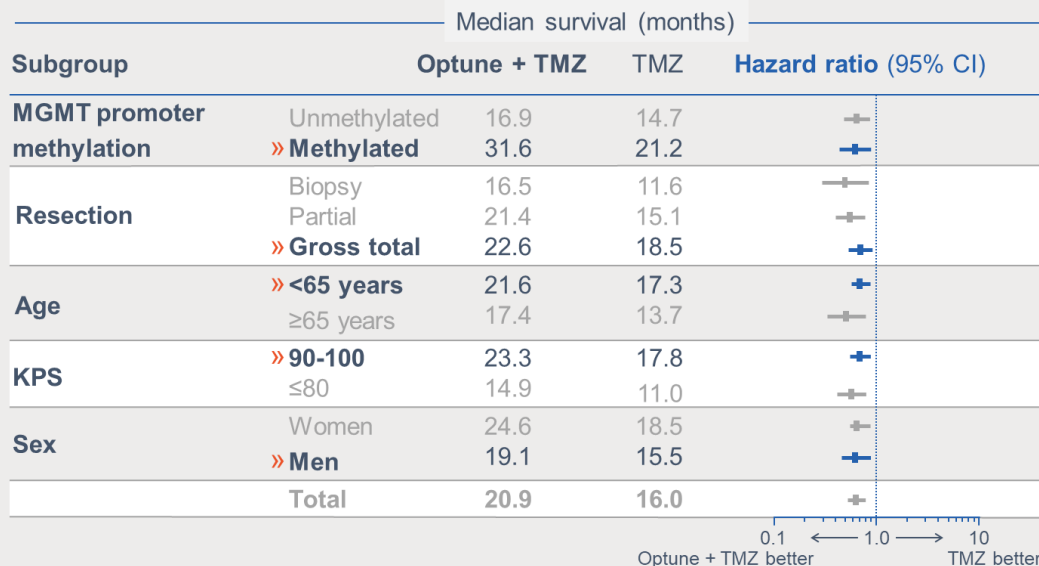
VS.

NEW ARRAYS AP channel, 1,685 mAmps



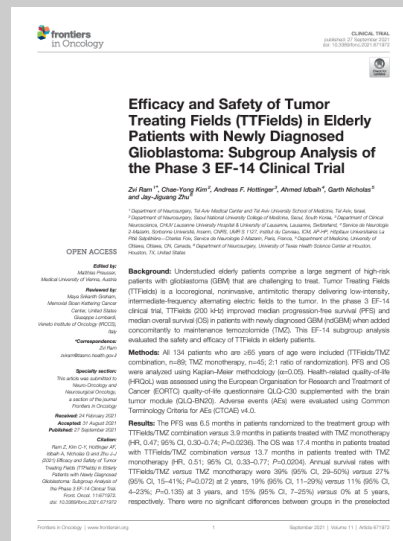
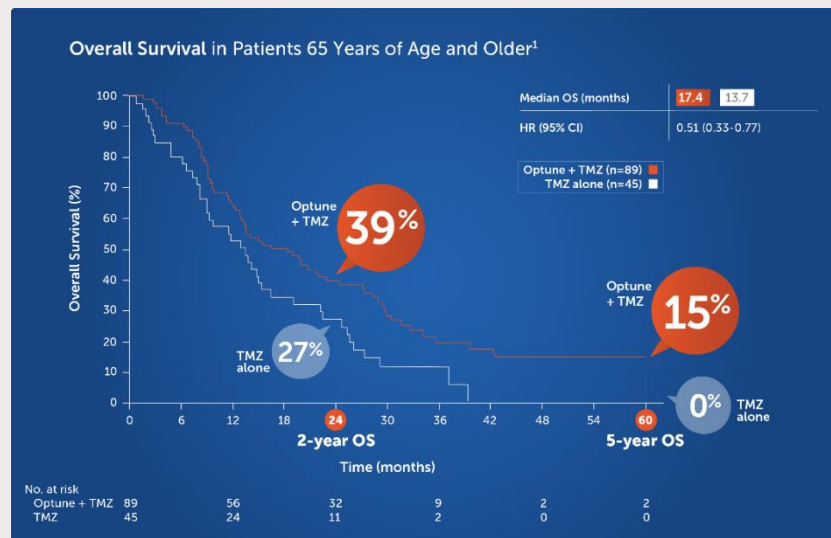
all analyzed subgroups experienced a benefit when adding Optune Gio to TMZ

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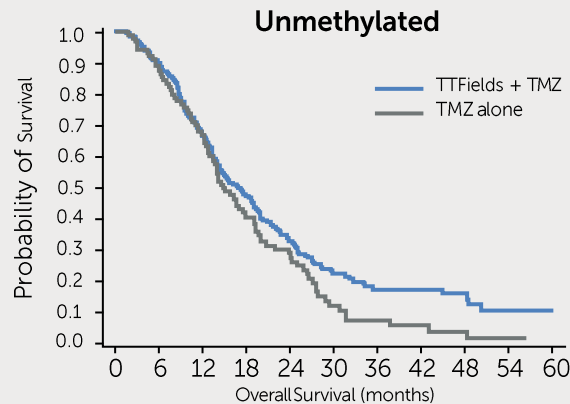


Optune Gio was associated with increased survival in patients 65 years and older

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	TTFields + TMZ (n = 137)	TMZ Alone (n = 77)
Median OS, months	31.6	21.2
Range, months	21.1–48.5	12.3–37.9
HR (95% CI) ¹	0.62 (0.43–0.88)	

	TTFields + TMZ (n = 209)	TMZ Alone (n = 95)
Median OS, months	16.9	14.7
Range, months	9.7–28.2	9.8–24.8
HR (95% CI) ¹	0.66 (0.49–0.85)	



Optune Gio has a strong safety profile with no significant increase in serious AEs compared with TMZ alone

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Incidence of grade 3/4 AEs occurring in ≥5% of patients during 5 years of follow-up	Optune + TMZ (n=456) %	TMZ alone (n=216) %
≥1 AE	48	44
Blood and lymphatic system disorders Thrombocytopenia	13 9	11 5
Gastrointestinal disorders	5	4
Asthenia, fatigue, and gait disturbance	9	6
Infections	7	5
Injury, poisoning, and procedural complications (falls and medical device site reaction)	5	3
Metabolism and nutrition disorders (anorexia, dehydration, and hyperglycemia)	4	5
Musculoskeletal and connective tissue disorders	5	4
Nervous system disorders Seizures	24 6	20 6
Respiratory, thoracic, and mediastinal disorders (pulmonary embolism, dyspnea, and aspiration pneumonia)	5	5

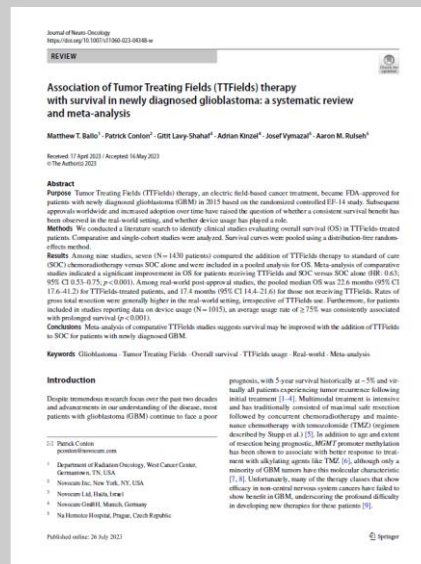
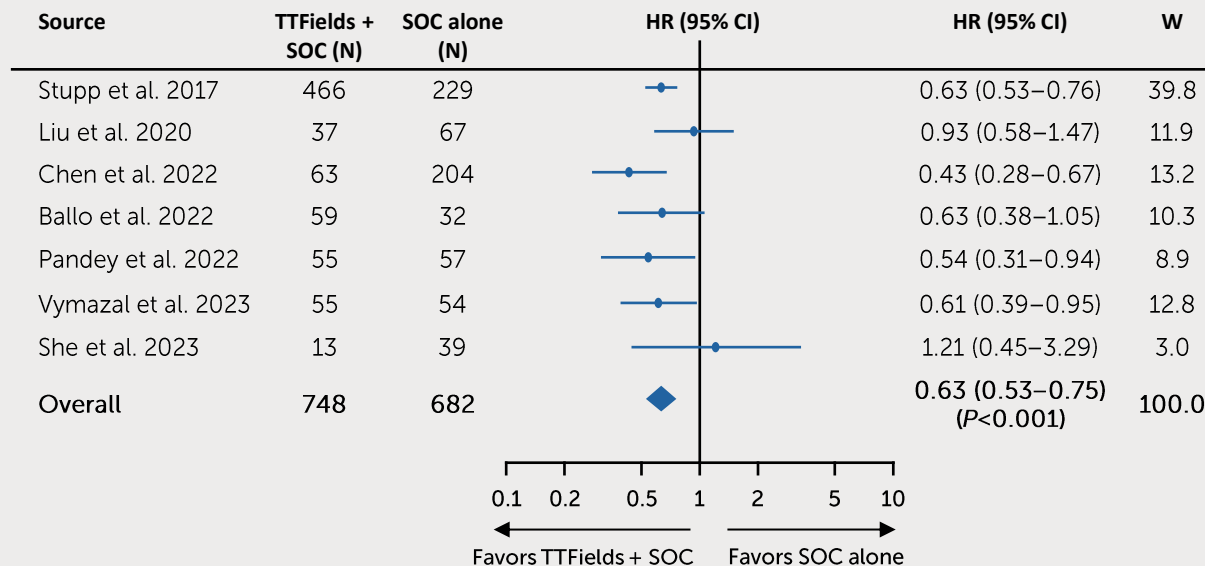




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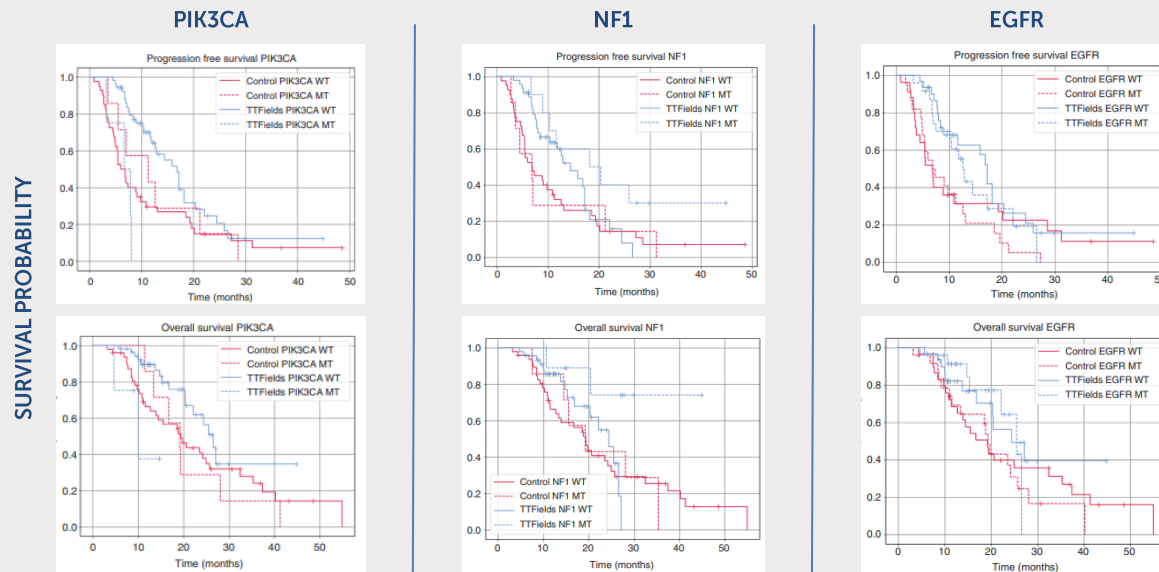
meta-analysis in ndGBM showed significant improvement in OS, and usage $\geq 75\%$ consistently prolonged survival, corroborating pivotal trial data

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TTFields therapy provide consistent activity for patients with GBM irrespective of molecular alterations

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Neuro-Oncology Advances

Volume 4, Issue 1, January-December 2022 | <https://doi.org/10.1093/advances/abz008> | Advance Access Date 27 June 2022

Molecular alterations associated with improved outcome in patients with glioblastoma treated with Tumor-Treating Fields

Manjaj Pandey, Joanne Xie, Sandeep Mittal, Si Zeng, Michelle Saul, Santosh Kesari, Amir Azadi, Herbert Newton, Karina Deniz, Katherine Ladner, Ashley Sunmual, W. Michael Kim, and Enli Lou*

West Cancer Center and Research Institute, Memphis, Tennessee, USA (M.P.); Carol Life Sciences, Phoenix, Arizona, USA (J.X.); J.D. M.D., M.B.A., W.M.A.; Virginia Tech Carilion School of Medicine, Roanoke, Virginia, USA (S.Z.); Pacific Neuroscience Institute, Saint John's Cancer Institute, Santa Monica, California, USA (M.S.); Arizona Oncology, Phoenix, Arizona, USA (K.D.); Neuro-Oncology Center, Advent Health Cancer Institute, Orlando, Florida, USA (H.N.); Division of Hematology, Oncology and Transplantation, Mason Cancer Center, University of Minnesota, Minneapolis, Minnesota, USA (A.A.); E.L. E.L. Cancer Center Institute, Charlotte, North Carolina, USA (A.L.)

*Corresponding Author: Enli Lou, MD, PhD, FACP, Associate Professor of Medicine, Division of Hematology, Oncology and Transplantation, University of Minnesota, Mayo West Code 460, 430 Delaware Street SE, Minneapolis, MN 55455, USA (enli@mc.man.unc.edu)

Abstract

Background. The genomic and overall biologic landscape of glioblastoma (GB) has become clearer over the past 2 decades, as predictive and prognostic biomarkers of both de novo and transformed forms of GB have been identified. The oral chemotherapeutic agent temozolomide (TMZ) has been integral to standard-of-care treatment for nearly 2 decades. More recently, the use of non-pharmacologic interventions, such as application of alternating electric fields, called Tumor-Treating Fields (TTFields), has emerged as a complementary treatment option that increases overall survival (OS) in patients with newly diagnosed GB. The genomic features associated with improved or lack of response to TTFields are unknown.

Methods. We performed comprehensive genomic analysis of GB tumors resected from 58 patients who were on to receive treatment using TTFields, and compared results to 57 patients who received standard treatment without TTFields.

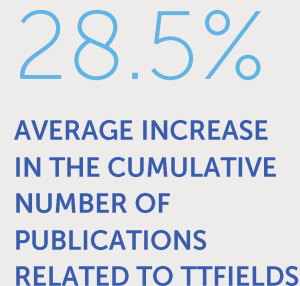
Results. We found that molecular driver alterations in NF1, and wild-type PIK3CA and epidermal growth factor receptor (EGFR), were associated with increased benefit from TTFields as measured by progression-free survival (PFS) and OS. There were no differences when stratified by PFS status. When NF1, PIK3CA, and EGFR status were combined as a Molecular Survival Score, the combination of the 3 factors significantly correlated with improved OS and PFS in TTFields-treated patients compared to patients not treated with TTFields.

Conclusions. These results shed light on potential driver and passenger mutations in GB that can be validated as predictive biomarkers of response to TTFields treatment, and provide an objective and testable genomic-based approach to assessing response.

Key Points

- Alterations in NF1 were associated with increased benefit from TTFields.
- Wild-type PIK3CA and EGFR also aligned with increased benefit from this approach.
- These results provide insight into molecular differences that can be validated to tailor treatment.

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[illegible]



The Journal of Clinical Investigation RESEARCH ARTICLE

Tumor Treating Fields dually activate STING and AIM2 inflammasomes to induce adjuvant immunity in glioblastoma

Dongjiang Chen,¹ Sen B. Le,¹ Tanun E. Hutchinson,¹ Anda-Alexandra Calinescu,¹ Mathew Sebastian,¹ Dan Jin,¹ Tianyi Liu,¹ Ashley Chlasek,¹ Maryam Raheman,¹ and David D. Tsan¹

Tumor Treating Fields (TTFields), an approved therapy for glioblastoma (GBM) and malignant melanoma, employ noninvasive application of low-intensity, intermediate frequency, alternating electric fields to disrupt the mitotic spindle, leading to chromosome missegregation and apoptosis. Emerging evidence suggests that TTFields may also induce inflammation, however, the mechanism underlying this property and whether it can be harnessed therapeutically are unclear. We have previously shown that TTFields induce the formation of protein aggregates in the form of amyloid-like fibrillar clusters that are primarily recruited and activated by macrophages – cyclic AMP response element binding protein (CREB) and interleukin 2 (IL-2) and their cognate α 2-macroglobulin of interferon gamma (STING) and α 1-antitrypsin (A1AT) immunofluorescence profiles are predominantly pericytic, type 1, and type 2 macrophages, respectively. In this study, we have used a murine model of TTFields-treated GBM to determine the antitumor immune response intensity and a cellular response to the electric field. In a STING- and A1AT-dependent manner, using single-cell and bulk RNA sequencing of purified blood mononuclear cells, we detected robust post-TTFields activation of adaptive immunity in patients with GBM and a T1906-based murine model and identified a gene signature that is predictive of improved survival. These findings suggest that the mechanism of action of TTFields may include a strategy using TTFields as cancer immunotherapy in GBM and potentially other solid tumors.

Introduction

Introduction
Glioblastoma (GBM) is the most common and lethal brain cancer in adults and one of the least immunogenic tumors (1). Recent work has revealed striking immune dysregulation and functional impairment in patients with GBM. Resident systemic T lymphocytes infiltrate the GBM microenvironment, but the majority of GBM tumors also possess a profoundly immunosuppressed or cold tumor microenvironment (TME), characterized by scarce tumor-infiltrating lymphocytes (TILs) and an abundance of inhibitory myeloid cells (2). Understanding these barriers presents a long, multi-faceted, immune-mediated tumor control. To "heat up" the cold GBM TME, recent efforts have focused on tumor cell–immune pathways with mixed results, such as dendritic cell (DC) vaccines (3), adoptive T cell transfer (4), and checkpoint-inhibiting cytokine mAb, or disrupting BTLA integrity to control tumor-specific cytotoxic T lymphocytes (CTLs) (4). However, it remains a challenge to leverage a direct, active role of tumor cells in the TME to elicit an anti-tumor immune response.

[illegible]

Submitted: March 3, 2021; **Accepted:** February 16, 2022; **Published:** April 15, 2022.
Reference information: J Clin Invest. 2022;132(8):e1545256.
<https://doi.org/10.1172/JCI1545256>.

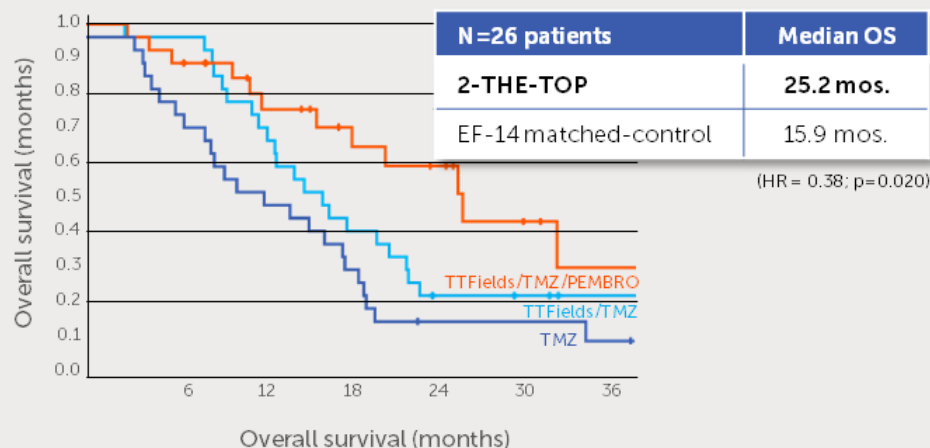
there is early evidence of efficacy in newly diagnosed GBM patients when TTFields therapy is added to immune checkpoint inhibitors

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Overall Survival

2-THE-TOP single arm study vs. external controls



WFNOS 2022 Top 10 Session 2 / March 26 (Sat), 10:15-11:30



Phase 2 study of pembrolizumab plus TTFields plus temozolomide in patients with newly diagnosed glioblastoma (2-THE-TOP)

David Tran, Ashley Ghaweddin, Dongjiang Chen, Maryam Rahmani
Department of Neurosurgery, Division of Neuro-Oncology, University of Florida, United States

Background: Emerging data indicate that TTFields, the new anti-mitotic treatment for GBM, stimulate immunity via the type-1 interferon (T1IFN) pathway of STING and ASK2 inflammasomes. Thus, we hypothesize that TTFields synergize with immune checkpoint inhibitors to induce anti-tumor immunity in GBM.

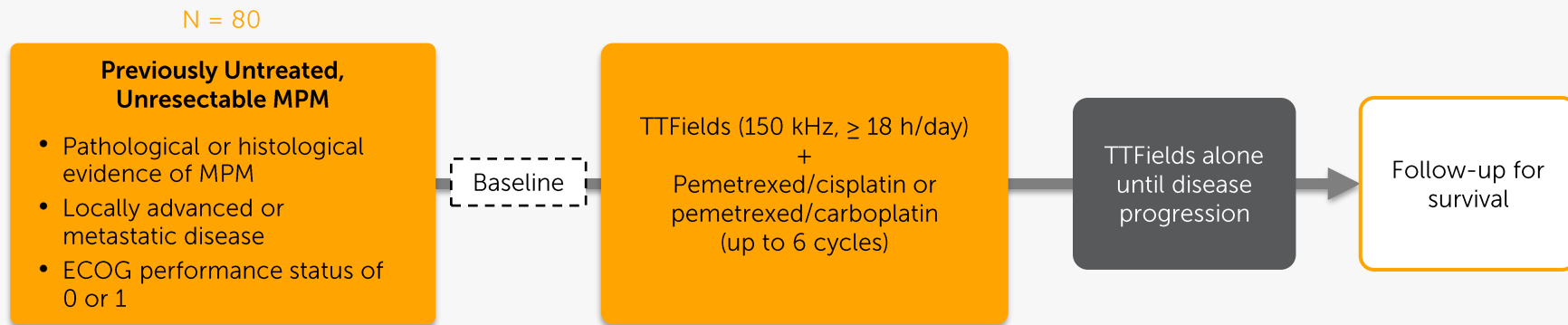
Methods: We conducted a phase 2 study combining peritumoral TTFields and maintenance TMZ in 24 patients with newly diagnosed GBM (nGBM). To distinguish immune effects of TTFields from pembrolizumab, TTFields was started at cycle 1 of TMZ and pembrolizumab (200 mg Q3 weeks) at cycle 2. The primary endpoint was PFS vs the historical control of TTFields plus TMZ (IAMA18-2306-2316) and immune signatures of TTFields and pembrolizumab by single-cell genomics of PBMCs. Secondary endpoints included toxicity and OS.

Results: As of 09/24/2021, 24 patients with a median age of 60 years were enrolled. Fourteen (58%) had biopsy only or subtotal resection. Nineteen (79%) had unresected MGMT and 1 (4%) had no MGMT. The median follow-up was 18 and 14.2 months for PFS and OS, respectively. Thirteen (54%) were progression free and 16 (67%) were alive. Of 22 patients with follow-up of 6 months, the median PFS was 3.11 vs 6.7 months in the control. Six (27%) patients with measurable tumors have achieved partial to complete objective response. We sequenced 16,704 PBMCs at 12 patients before and after TTFields and detected robust gene TTFields T cell activation in 11 of 12 patients via the T1IFN regulatory which was strongly correlated with T1IFN clinical expansion (Spearman coefficient $r=0.8$, $p=0.004$). Importantly, we defined a T cell-based gene signature of TTFields effects on T1IFN clinical expansion. The most common serious adverse events were thrombocytopenia, seizure, and metabolic disturbance in 4 (17%), 3 (13%), and 2 (8%) patients, respectively.

Conclusion: The triple combination is well tolerated and shows early evidence of efficacy in nGBM patients. Survival and molecular data will be updated.

Keywords: TTFields immunotherapy; pembrolizumab; STING; single cell analysis

STELLAR Phase 2 trial evaluated TTFields therapy + pemetrexed and cisplatin or carboplatin in MPM



Start date: February 2015
Primary completion: April 2018
Study completion: April 2018
Study sites: 13 (Europe)

Primary endpoints:

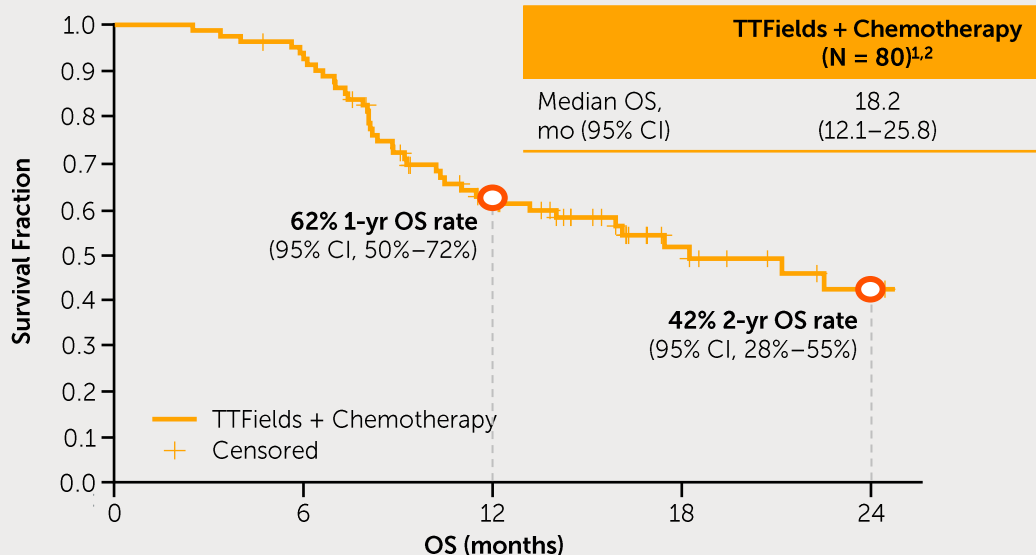
- OS

Secondary endpoints:

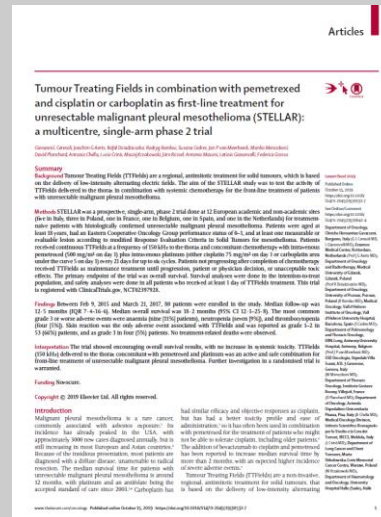
- PFS, ORR (modified RECIST criteria for MPM), safety

MPM patients who used Optune Lua first line achieved 18.2 months median OS

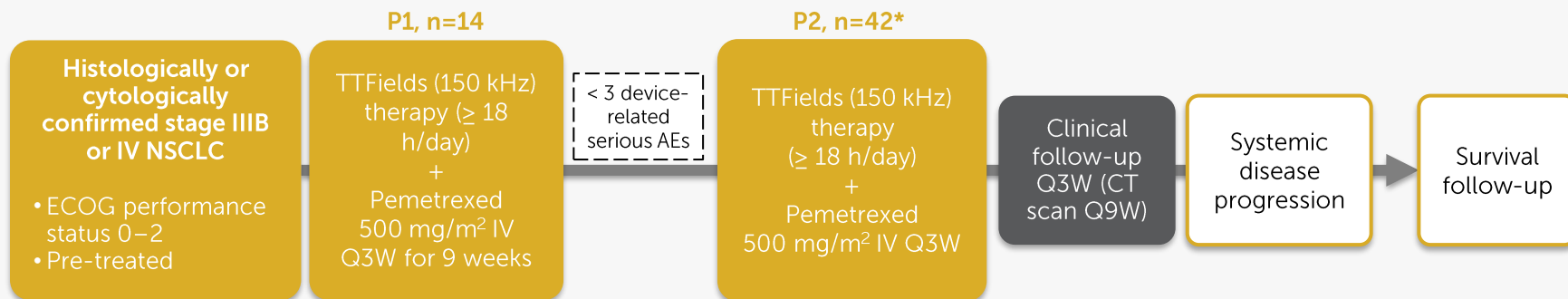
FOR MORE INFORMATION, USE THE QR CODE:



Adapted from Ceresoli GL et al. 2019



EF-15 Phase 2 trial evaluated TTFields therapy + pemetrexed in NSCLC



Start date: May 2008

Primary completion: July 2011

Study completion: July 2011

Study sites: 4 (Switzerland)

Primary endpoints:

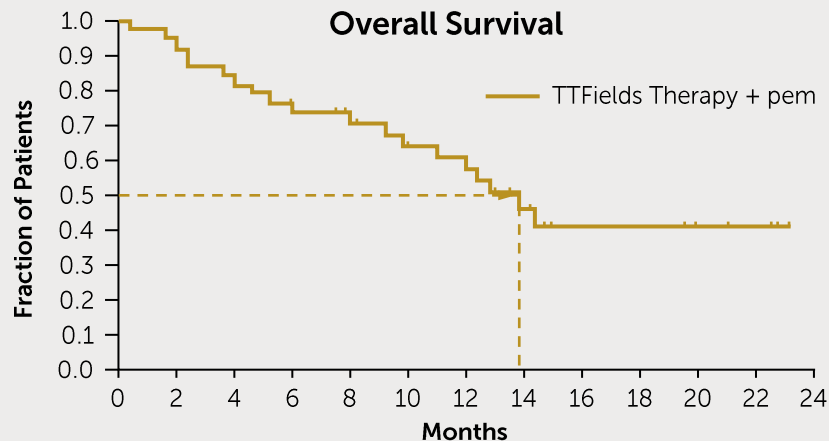
- Device related toxicity (P1), Time to in-field progression (P2)

Secondary endpoints:

- OS, ORR, time to systemic progression, safety

TTFields therapy together with pemetrexed improved disease control within the treatment field in second line NSCLC

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Median in-field PFS

Median PFS

Median OS

1yr Survival

TTFields + Pemetrexed

6.5 mo

5.0 mo

13.8 mo

57.0%

Pemetrexed alone

n/a

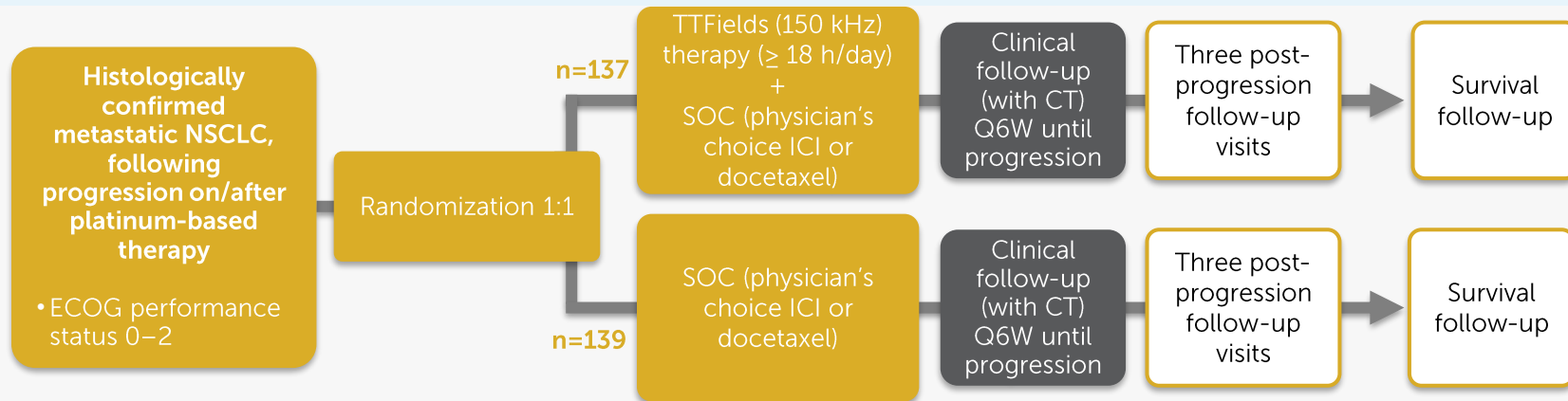
2.9 mo

8.3 mo

29.7%



LUNAR Phase 3 trial evaluated TTFields therapy + SOC in metastatic NSCLC, post-platinum



Start date: December 2016

Primary completion: December 2022

Study completion: December 2022

Study sites: 124

Primary endpoints:

- OS

Secondary endpoints:

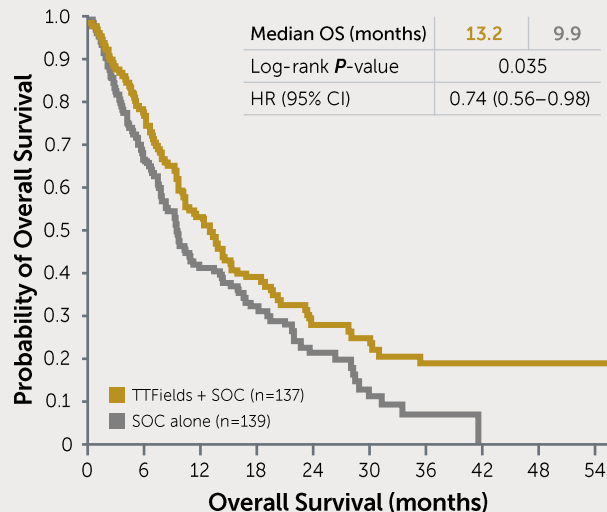
- OS (by cohort), PFS, ORR, QoL, safety

TTFields therapy together with either standard of care therapies or immune checkpoint inhibitor improved overall survival in second-line NSCLC

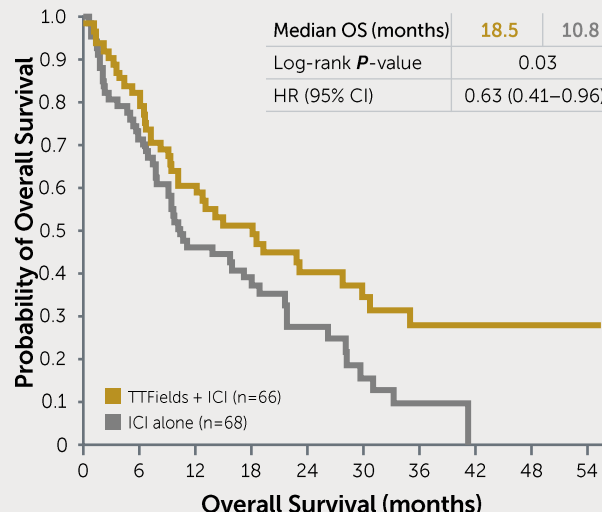
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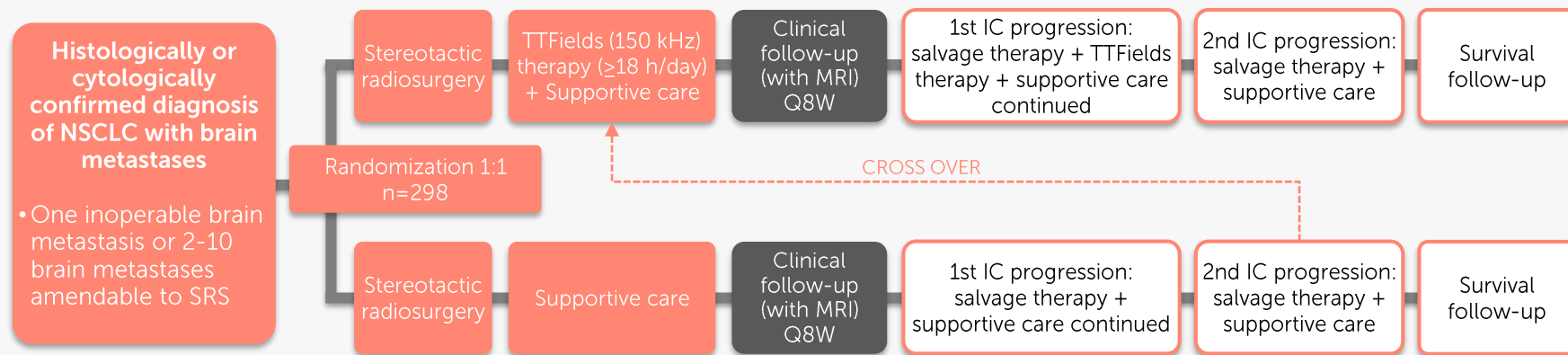
Overall survival (ITT population)



Overall survival (ICI-treated patients)



METIS Phase 3 trial evaluated TTFields therapy + supportive care in NSCLC brain metastases, following SRS



Start date: October 2016
Primary completion: March 2023
Study sites: 125

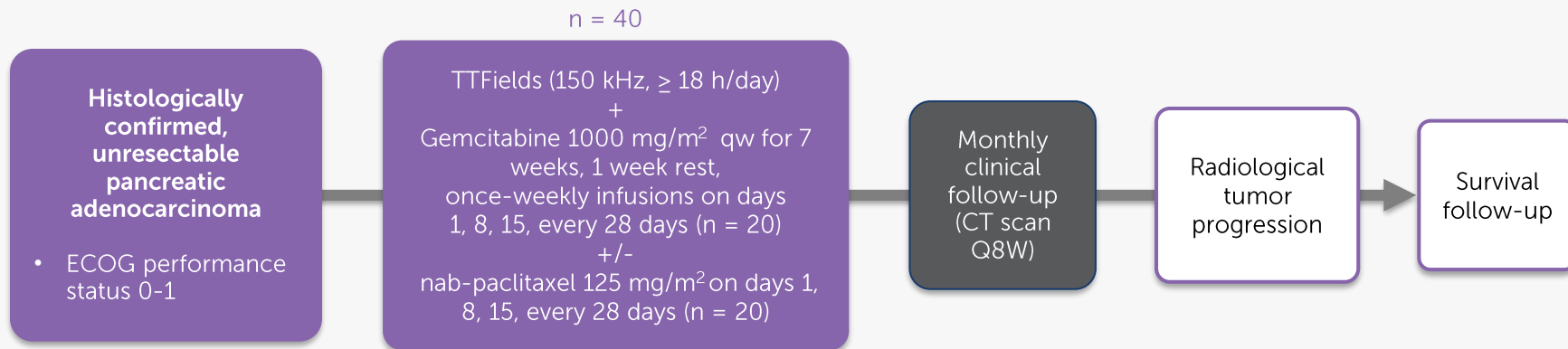
Primary endpoints:

- Time to intracranial progression

Secondary endpoints:

- Time to neurocognitive failure, OS, radiological response rate, time to 2nd intracranial progression, time to 1st and 2nd progression by cohort (1-4 metastases, 5-10 metastases), rate of intracranial progression at two-month intervals, time to distant progression, rate of cognitive decline, neurocognitive failure-free survival, quality of life, adverse events

PANOVA phase 2 trial evaluated TTFields therapy + gemcitabine +/- nab-paclitaxel in pancreatic cancer



Start date: Nov 2013

Primary completion date: Dec 2017

Study completion date: Dec 2017

Study sites: 6 (Europe)

Primary endpoint:

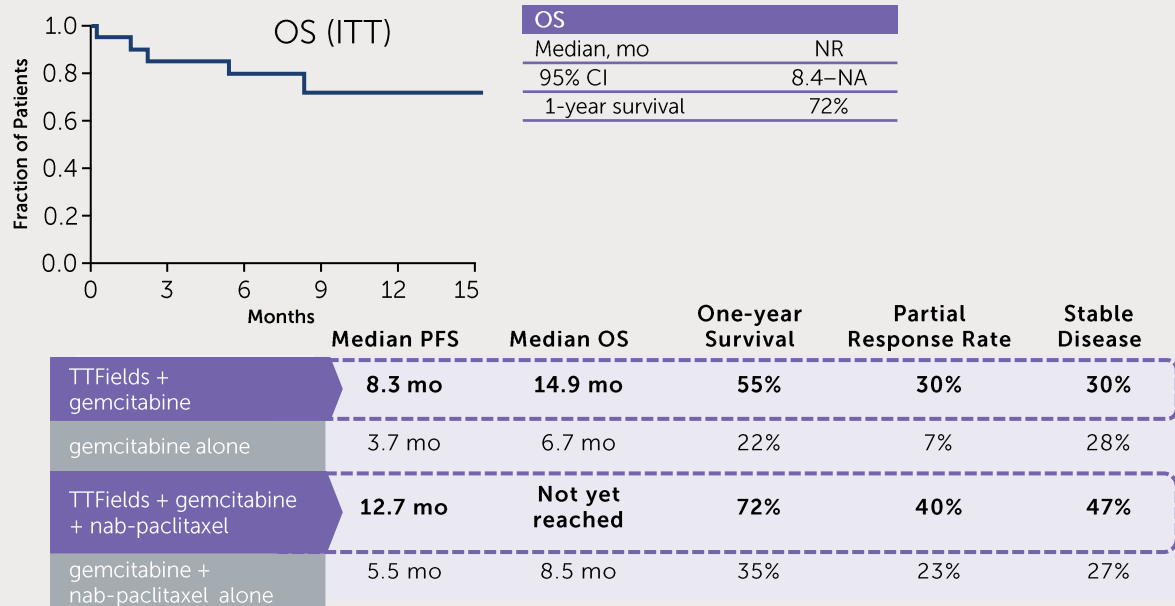
- Safety

Secondary endpoints:

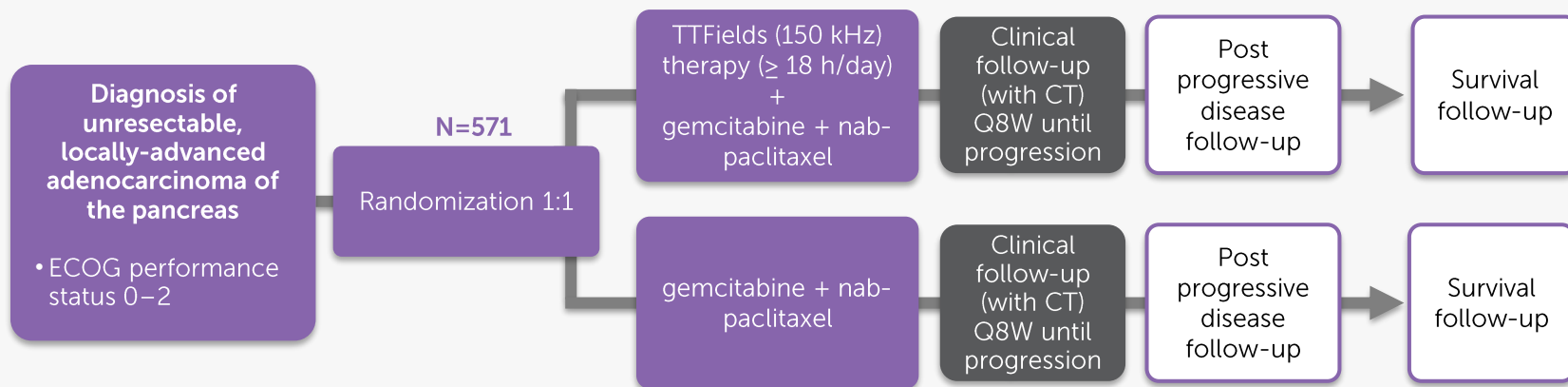
- TTFields monthly usage, PFS, OS

TTFields therapy together with chemotherapy were well tolerated for patients with advanced pancreatic cancer

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PANOVA-3 Phase 3 trial evaluated TTFields therapy + gemcitabine + nab-paclitaxel in unresectable, locally advanced pancreatic cancer



Start date: February 2018
Primary completion: October 2024
Study completion: October 2024
Study sites: 199

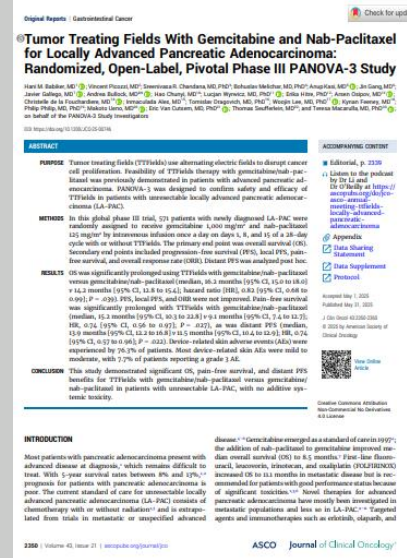
Primary endpoints:

- OS

Secondary endpoints:

- PFS, local PFS, ORR, 1-year survival rate, QoL, pain-free survival, puncture-free survival, resectability rate, safety

**MET PRIMARY
ENDPOINT**

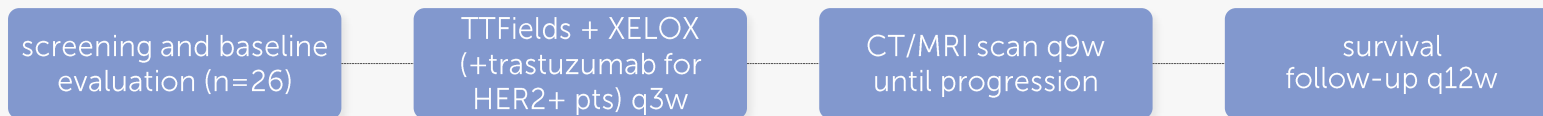


encouraging response rate and durability signals
in EF-31 phase 2 gastric cancer trial

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EF-31 PHASE 2 PILOT TRIAL DESIGN¹



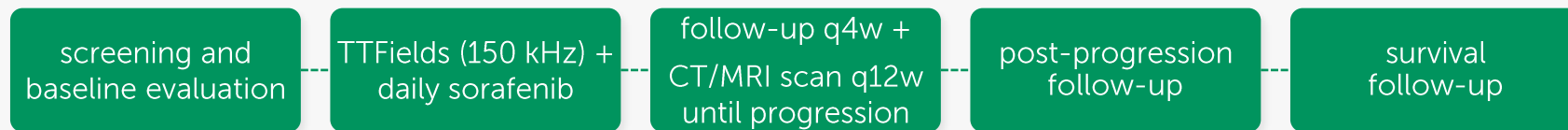
	OBJECTIVE RESPONSE RATE	MEDIAN PROGRESSION-FREE SURVIVAL	DURATION OF RESPONSE	ONE-YEAR SURVIVAL
TTFields + chemotherapy	50%	7.8mo	10.3mo	72%
SOC chemotherapy ²	41-45%	6.9mo	6.9mo	48%

encouraging signals in liver cancer despite poor prognosis and low treatment exposure in HEPANOVA phase 2 trial

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HEPANOVA PHASE 2 PILOT TRIAL DESIGN²



76%

DISEASE CONTROL RATE
(n=21)

VS. 43% CONTROL³

9.5%

OBJECTIVE RESPONSE RATE
(n=21)

VS. 4.5% CONTROL

91%

DISEASE CONTROL RATE

18%

OBJECTIVE RESPONSE RATE

patients that received ≥ 12 wks of TTFields (n=11)

2025-2026 anticipated clinical development milestones

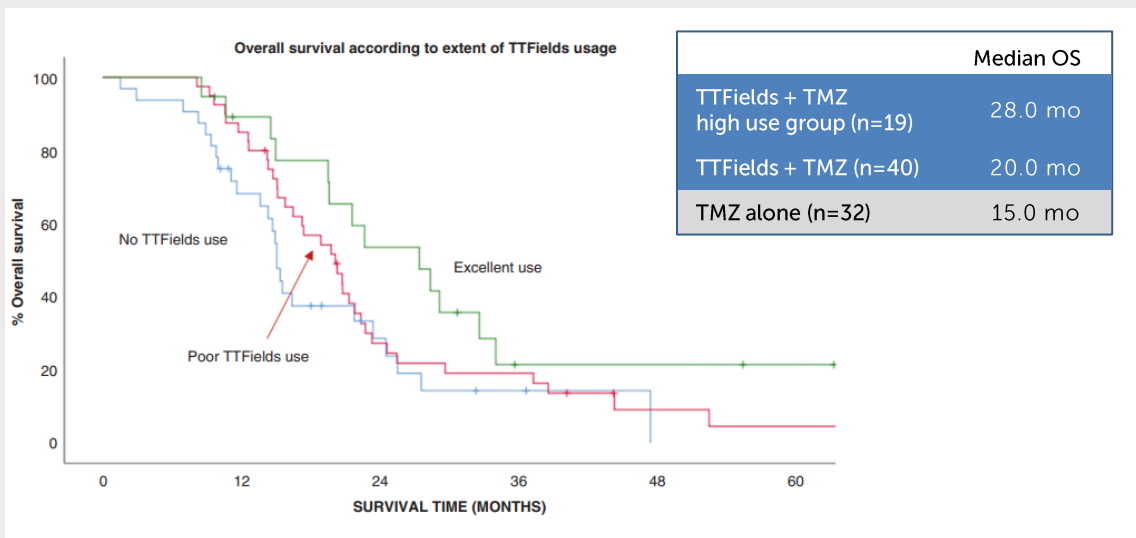
	TRIAL	TTFIELDS +	PHASE 2	PHASE 3	APPROVED
CNS indications	glioblastoma	EF-14	TMZ		✓
		TRIDENT	TMZ + radiation	DATA IN 2026	
		KEYNOTE D58	TMZ + pembrolizumab	enrolling	
	brain metastases	EF-11	monotherapy (recurrent GBM)		✓
		METIS	monotherapy	SUBMISSION IN 2025	
torso indications	non-small cell lung cancer	LUNAR	docetaxel or PD-L1 inhibitor (2L)		✓
		LUNAR-2	pembrolizumab + platinum (1L)	enrolling	
		LUNAR-4	pembrolizumab (2L retreatment)	enrolling	
	mesothelioma	STELLAR	pemetrexed + cisplatin/carboplatin		✓
	pancreatic cancer	PANOVA-3	nab-paclitaxel + gemcitabine (LAPC)	SUBMISSION IN 2025	
		PANOVA-4	atezolizumab + nab-paclitaxel + gemcitabine (MPC)	DATA IN 2026	



real-world and
routine clinical care
study evidence

real-world evidence showed ndGBM median overall survival extension by over 12 months in the high use TTFields group

FOR MORE
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THE QR CODE:



Neuro-Oncology Advances

4111-1-6-2022 | <https://doi.org/10.1093/advances/101> | Advance Access date 19 September 2022

Determinants of tumor treating field usage in patients with primary glioblastoma: A single institutional experience

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Abstract

Background. Determinants of tumor treating fields (TTFields) usage in patients receiving combined modality therapy for primary CNS solid-type glioblastoma are currently unknown.

Methods. Ninety-one patients underwent maximal debulking surgical resection, completed external beam radiotherapy with concurrent temozolomide (TMZ), and initiated adjuvant TMZ with or without TTFields. We performed a retrospective analysis of patient, tumor, and treatment-related factors that affected TTFields usage.

Results. We identified three TTFields usage subgroups: 32 patients that declined TTFields, 40 patients that started, but had monthly compliance of less than 75% or used it for less than 2 months, and 19 patients who used TTFields for 2 or more months and maintained average monthly compliance greater than 75%. With 26.5 months median follow-up for surviving patients, the 1- and 3-year actuarial overall survival for all patients was 80% and 19%, respectively. On multivariate analysis TTFields use ($P = .03$), extent of surgical resection ($P = 0.02$), and MGMT methylation status ($P = .01$) were significantly associated with overall survival. TTFields usage was explored as a continuous variable and higher average usage was associated with longer overall survival ($P = .03$). There was no relationship between patient, tumor, or treatment-related factors and a patient's decision to use TTFields.

Conclusions. No subgroup of patients was more or less likely to initiate TTFields therapy and no subgroup was more or less likely to use TTFields as prescribed. The degree of TTFields compliance may be associated with improved survival independent of other factors.

Key Point

• It is reasonable to offer all patients with primary glioblastoma TTFields therapy as we could not identify a group that was more or less likely to discontinue therapy or unable to initiate therapy. Patients benefit from TTFields regardless of tumor or patient characteristics.

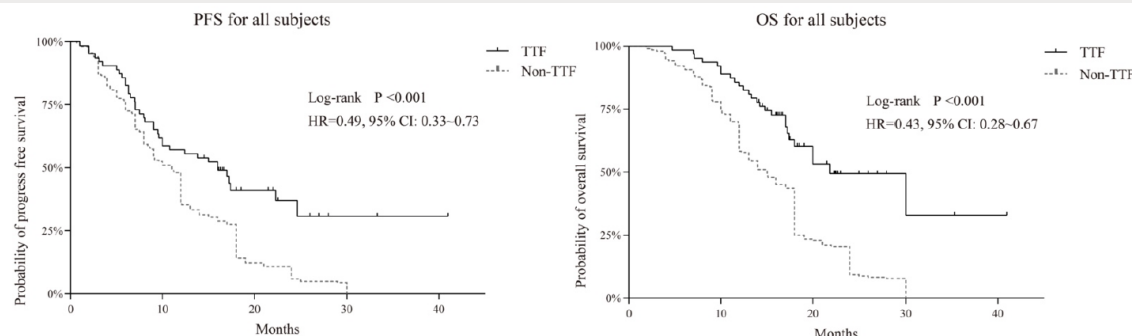
Glioblastoma is the most common and aggressive primary malignant brain tumor diagnosed in adults and has a poor prognosis, with only 15–20% of patients living past 5 years following diagnosis. Even with the best standard of care, consisting of maximal safe surgical resection, radiation therapy, and temozolomide (TMZ) chemotherapy, median overall survival has historically been only 14.6 months.^{1,2}

Tumor treating fields (TTFields) represent a novel therapy in the treatment of glioblastoma. TTFields deliver low intensity, intermediate-frequency (200 kHz) alternating electric fields

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real-world evidence validates EF-14 with statistically significant improvement in PFS and OS in Chinese patients with ndGBM

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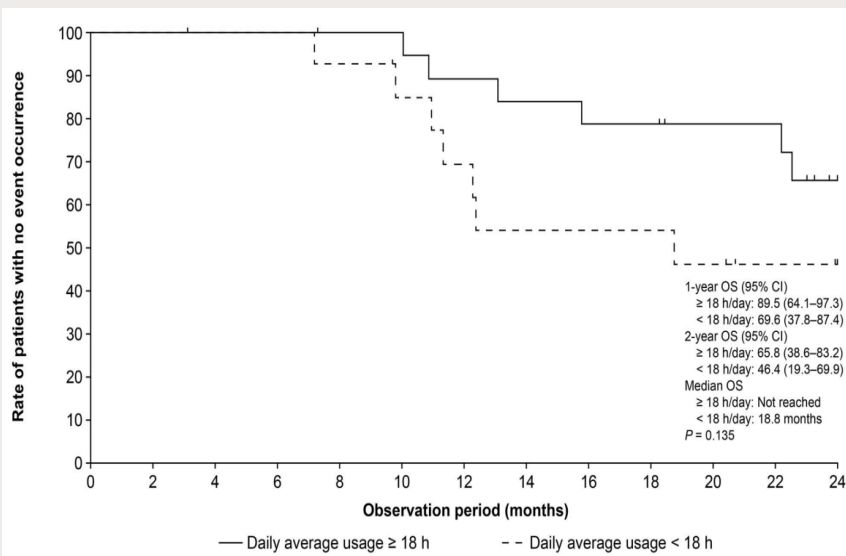


	Median OS	Median PFS
TTFields + TMZ (n=63)	21.8 mo	16.0 mo
TMZ alone (n=204)	15.0 mo	11.0 mo



post-approval study supports safety and efficacy profile of TTFields therapy in ndGBM Japanese patients, validating EF-14 improved survival rates

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	1-year survival	2-year survival
TMZ alone ¹	65%	31%
TTFields + TMZ (n=14)	77.9%	53.6%
TTFields + TMZ high use group (n=21)	89.5%	65.8%

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Original Article

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Original Article

Safety and efficacy of tumour-treating fields (TTFields) therapy for newly diagnosed glioblastoma in Japanese patients using the Novo-TTF System: a prospective post-approval study

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Abstract

Background: Tumour-treating fields (TTFields) therapy is a non-invasive, anti-cancer treatment. Efficacy and safety of tumour-treating fields therapy in adults with newly diagnosed glioblastoma were demonstrated in the pivotal phase 2 EF-14 study (NCT00708050). Here, we report post-approval data of tumour-treating fields therapy in Japanese patients with newly diagnosed glioblastoma.

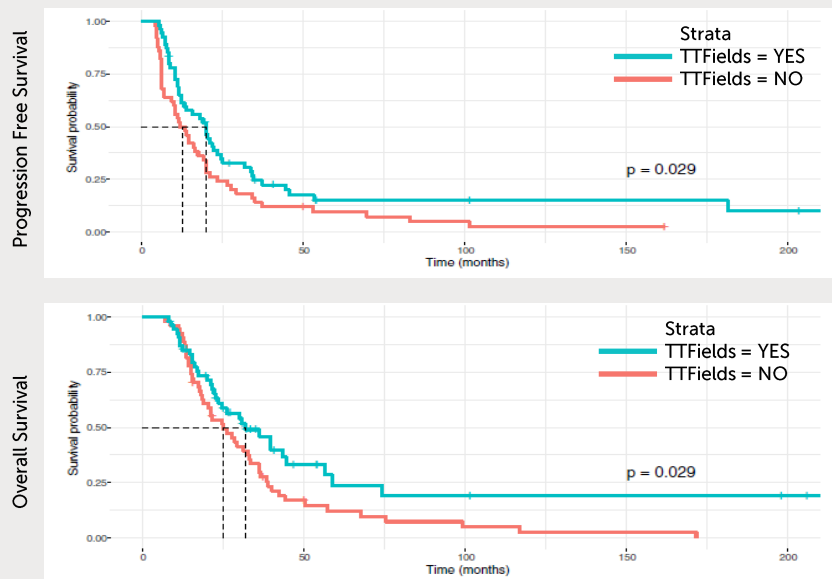
Methods: Unselected post-marketing surveillance data from Japanese patients with newly diagnosed glioblastoma treated with tumour-treating fields therapy (December 2016–June 2020) were retrospectively analysed. The primary endpoints were skin, neurological and psychiatric adverse events. The secondary endpoints were 1- and 2-year overall survival rates, and the 6-month progression-free survival. Adverse events were analysed using MedDRA v24.0. The overall survival and progression-free survival were assessed using the Kaplan–Meier survival analysis (log-rank testing). The Cox proportional hazard regression analyses were also performed.

Results: Forty patients with newly diagnosed glioblastoma were enrolled (82.5% male; median age 59 years; median baseline Karnofsky Performance Scale score 90). The most common tumour-treating fields therapy-related adverse event was beneath-army local skin reaction (10% of patients). The adverse events were mostly mild to moderate in severity. Neurological disorders were observed in 2.5% patients (one patient reported dysphagia). No psychiatric disorders were reported. The 1- and 2-year overall survival rates were 73.9% (95% CI 65.6–80.3) and 53.6% (95% CI 40.7–66.7%), respectively. The 6-month progression-free survival was 72.5% (95% CI 62.1–82.9%). These survival

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long term study of ndGBM patients, covering 18-year period, confirms TTFields' positive effect on PFS and OS

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	median PFS	median OS
TTFields + TMZ (n=55)	19.75 mo	31.67 mo
TMZ alone ¹	12.45 mo	24.80 mo

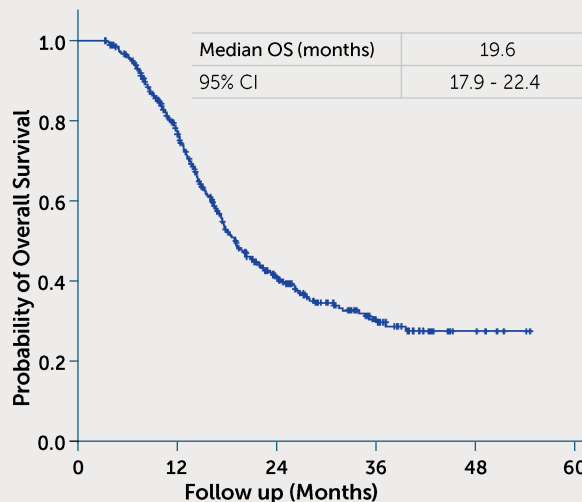


TIGER study of routine clinical care in German ndGBM patients corroborates overall survival and safety outcomes from EF-14

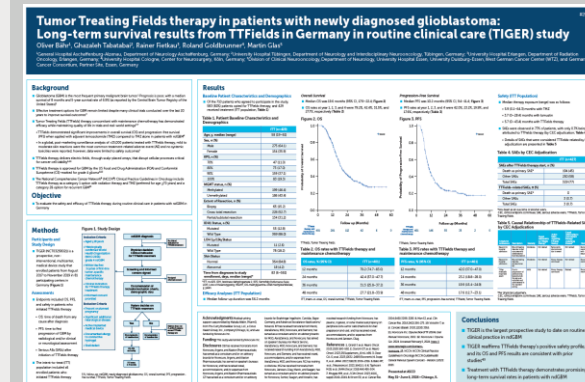
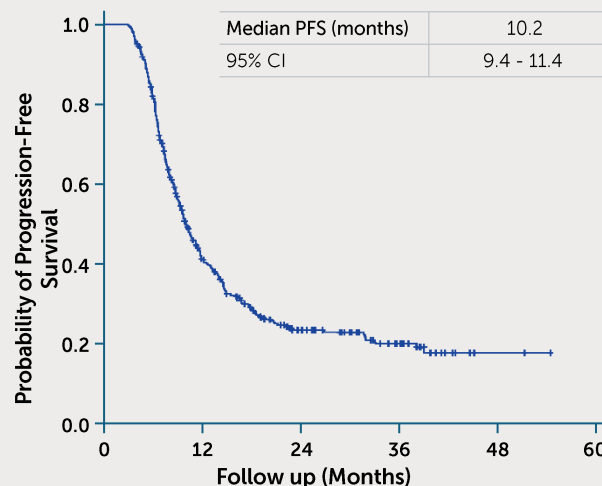
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Overall Survival



Progression-Free Survival



review article identifies TTFields therapy as one of few factors driving increased overall survival in GBM patients since the 2005 Stupp-protocol

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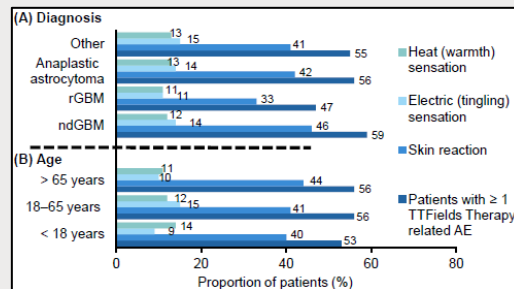
Neff et al.

Characteristic	N	HR ¹	95% CI ¹	p-value
Age (years)	19,414	1.02	1.02, 1.03	<0.001
Sex				
Female	8,046	—	—	reference
Male	11,368	1.10	1.07, 1.14	<0.001
Elkhauser Comorbidity Score	19,414	1.01	1.01, 1.01	<0.001
Tumor-Treating Fields (ever)				
No	16,353	—	—	reference
Yes	3,061	0.77	0.73, 0.80	<0.001
Received radiation or radiosurgery (ever)				
No	7,370	—	—	reference
Yes	12,044	0.88	0.85, 0.91	<0.001
Bevacizumab (ever)				
No	15,741	—	—	reference
Yes	3,673	0.85	0.82, 0.88	<0.001

In this commercially insured dataset, TTFields improved OS to a greater extent (HR=0.77) vs. Bevacizumab (HR=0.85) or Radiation use (HR=0.88)

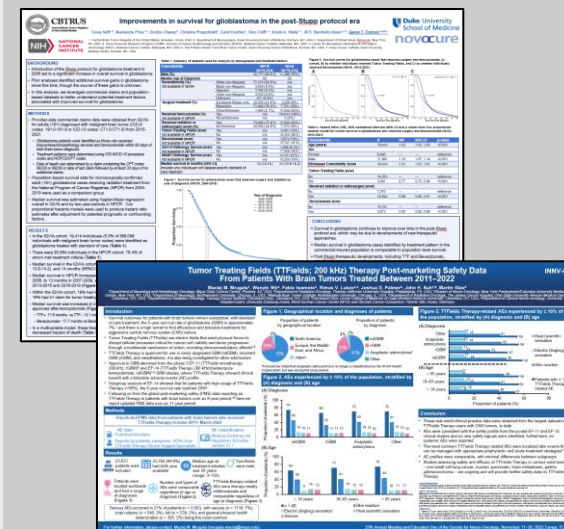
TTFields subset n=3,061 over 6 years

Mrugala et al.



AEs were consistent with the safety profile from the pivotal EF-11 and EF-14 clinical studies

n=23,822 over 11 years





tumor treating fields
mechanism of
action appendix

patients with aggressive solid tumors often face suboptimal survival outcomes, despite advancements in treatment modalities

These outcomes are due to diverse treatment challenges, including:



Therapeutic tumor resistance



Drug-to-drug interactions



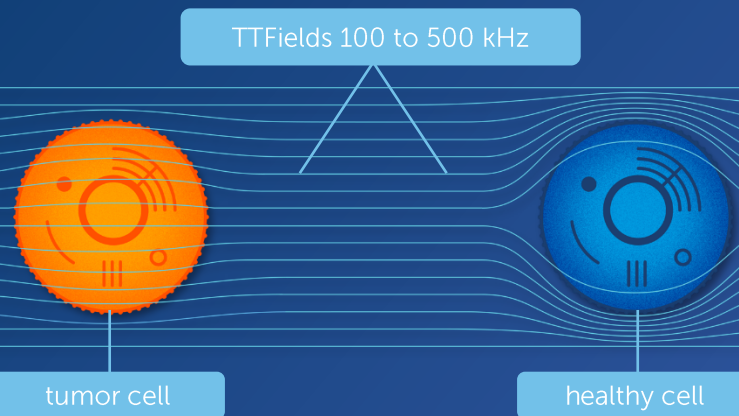
Additive systemic toxicities

With a poor survival outlook, physicians and patients need additional treatment strategies

Tumor Treating Fields (TTFields) are electric fields that exert physical forces to kill cancer cells via a variety of mechanisms



TTFields spare healthy cells because they have different properties than cancer cells across a range of tumor types

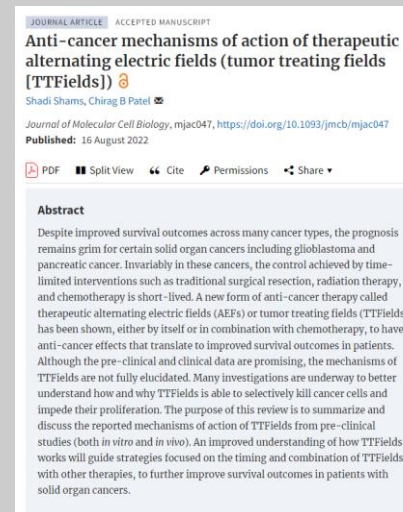


a growing body of evidence supporting multiple mechanisms of action

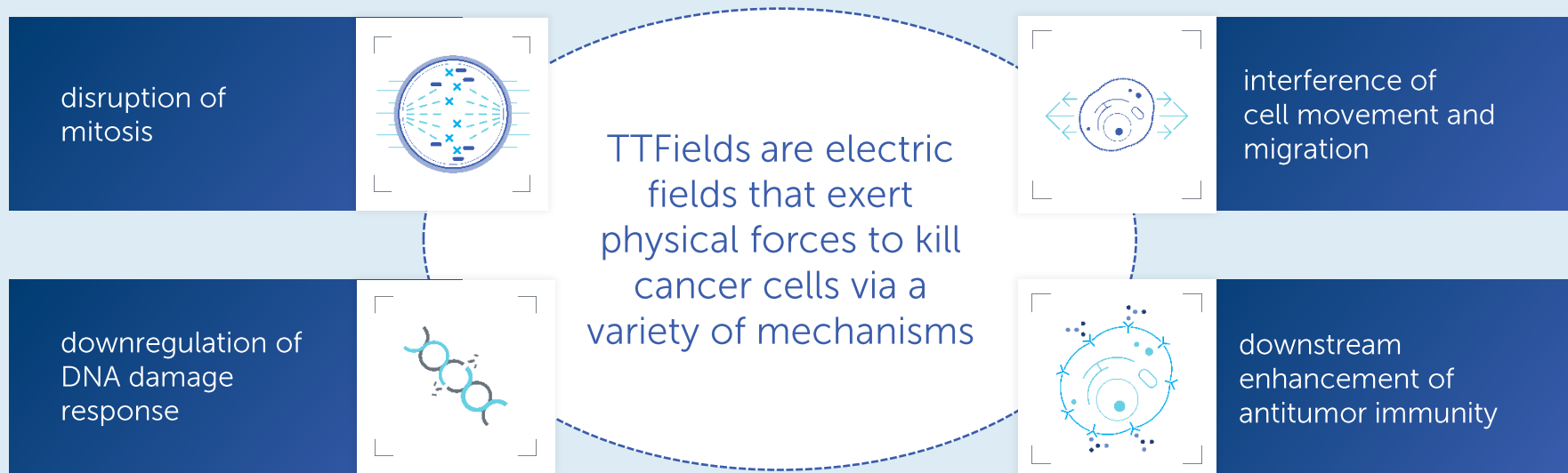
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- Preclinical research has shown interference with cancer cell motility and migration, activation of anti-tumor immunity, downregulation of genes important for DNA damage repair, and other potential mechanisms
- May demonstrate enhanced effects across solid tumor types when used with chemotherapy, radiotherapy, immune checkpoint inhibition, or PARP inhibition in preclinical models

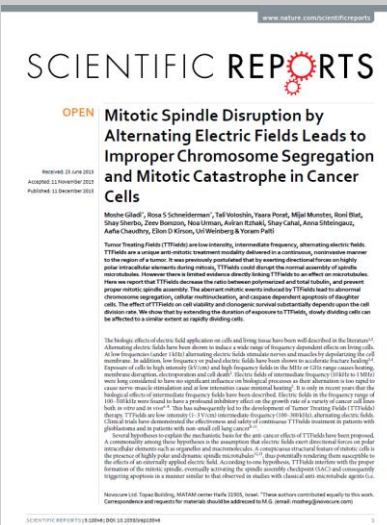
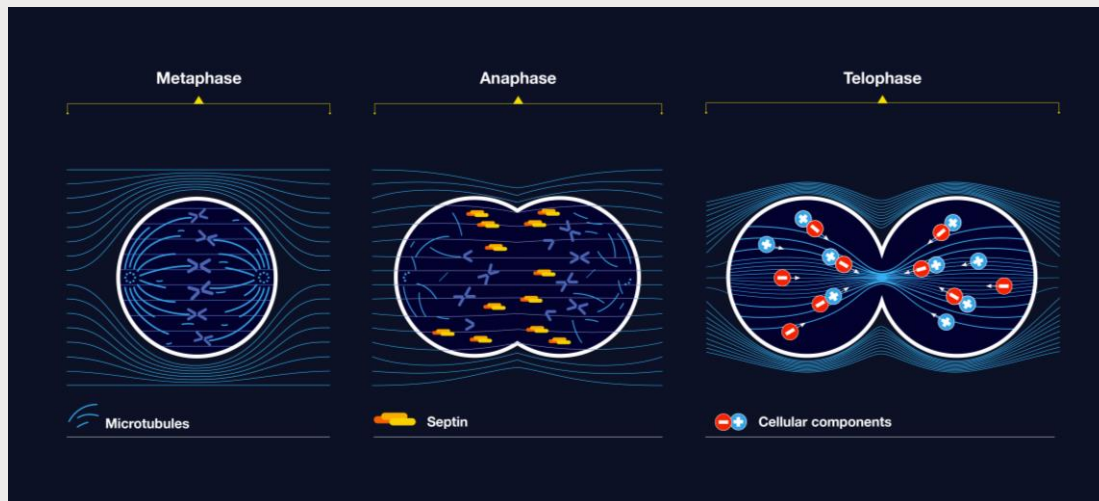


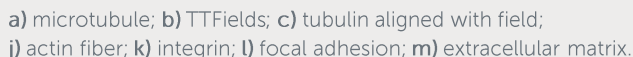
Tumor Treating Fields have multiple, distinct mechanisms of action



TTFields have been shown to disrupt mitosis in cancer cells by exerting physical forces on their polar components

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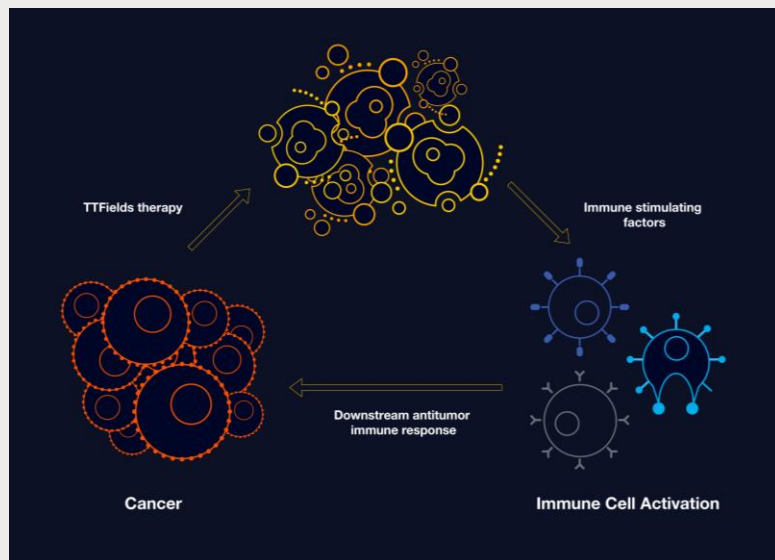
A model illustrating the mechanism by which TTFields modulates cancer cell motility.

- (1) Microtubules are required to specify the direction of cell movement. GEF-H1 catalytic activity is downregulated through microtubule binding.
- (2) TTFields exert directional forces on polar tubulins leading to their alignment in the direction of the field. This, in turn, leads to the reorganization of the microtubule network resulting in changes in the abundance of microtubules and initiation of the GEF-H1/RhoA/ROCK signaling pathway
- (3) to increase actin bundling
- (4) and formation of focal adhesions,
- (5) which disrupt cell polarity and migration directionality.



TTFields-mediated cell disruption activates the immune system and triggers a downstream antitumor cell response

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TTFields induces downstream immunogenic cell death, including release of DAMPs (damage-associated molecular patterns)

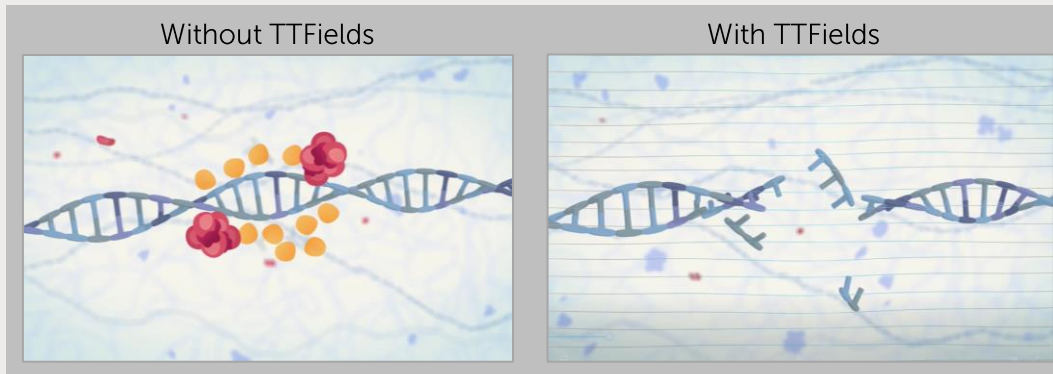


TTFields downregulate genes important for DNA damage repair

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- TTFields disrupt DNA damage repair in cancer cells by downregulating genes that are part of the well-known FA-BRCA pathway^{1,2}



OPEN

Tumor-treating fields elicit a conditional vulnerability to ionizing radiation via the downregulation of BRCA1 signaling and reduced DNA double-strand break repair capacity in non-small cell lung cancer cell lines

Narasimhan Kumar Karanam¹, Kalyanman Srinivasan¹, Lingshuo Ding¹, Brock Blain¹, Debabrata Saha¹ and Michael D Story^{1,2}

The use of tumor-treating fields (TTFields) has revolutionized the treatment of recurrent and newly diagnosed glioblastoma (GBM). TTFields are low-intensity, intermediate frequency alternating electric fields that are applied to tumor regions and cells using non-invasive arrays. The predominant mechanism by which TTFields are thought to kill tumor cells is the disruption of mitosis. Using five non-small cell lung cancer (NSCLC) cell lines we found that there is a variable response to cell proliferation and cell killing between these NSCLC cell lines that was independent of cell status. TTFields treatment increased the G2M population, with a concomitant reduction in S-phase cells followed by the appearance of a sub-G1 population indicative of apoptosis. Temporal changes in gene expression during TTFields exposure were evaluated to identify molecular signaling changes underlying the differential TTFields response. The most differentially expressed genes were associated with the cell cycle and cell proliferation pathways. However, the expression of genes found within the BRCA1-DNA damage response were significantly downregulated (P < 0.05) during TTFields treatment. DNA double-strand break (DSB) repair foci increased when cells were exposed to TTFields as did the appearance of chromosome type aberrations, suggesting an interplay mechanism responsible for cell death involving DNA repair. Exposing cells to TTFields immediately following ionizing radiation resulted in increased chromosome aberrations and a reduced capacity to repair DNA DSBs, which were likely responsible for at least a portion of the enhanced cell killing seen with the combination. These findings suggest that TTFields induce a state of 'BRCAness' leading to a conditional susceptibility resulting in enhanced sensitivity to ionizing radiation and provide a strong rationale for the use of TTFields as a combined modality therapy with radiation or other DNA-damaging agents.

Cell Death and Disease (2021) 12, 40711 | doi:10.1038/s41419-021-11671-3

Lung cancer is the second most prevalent cancer and the leading cause of cancer-related death in the United States.^{1,2}

Non-small cell lung cancer (NSCLC) is the most prevalent type, accounting for ~85% of new cases.^{3,4} A plethora of treatment options, including surgical resection, chemotherapy, radiation therapy and immunotherapy,^{5,6} have improved outcomes for patients with stage I and II NSCLC, but ~30% and 50%, respectively, relapse despite the use of these options. 5-year survival rates for patients with late stage IIIA, IIIB and IV are 10%, 5% and 5%, respectively, with limited hope for improved outcomes.^{7,8}

TTFields have been shown to be synergistic with conventional therapies to increase survival rates. The advent of Tumor-Treating Fields (TTFields), a novel physical treatment modality, has been effective for the treatment of solid, BRCA1-mediated primary and recurrent tumors.^{9,10} TTFields modulate gene expression and alter a transcriptionally regulated intermediate frequency (100–200 kHz) alternating electric field across the tumor space.¹¹ TTFields create a heterogeneous intracellular environment that induces a dielectrophoretic movement of polar molecules towards the region of higher field intensity, effectively preventing cytoskeletal and other critical biochemical functions.¹² As such, TTFields perturb and target cancer cells through the inhibition of cell proliferation, effectively causing non-cleaving chromosomal, in addition, TTFields do not abrogate nerves and muscle because of their high frequency, and do not perturb heart function or breathing.^{13,14} The combination of TTFields and chemotherapy has been shown to be synergistic, leading to improved outcomes in recurrent and newly diagnosed glioblastoma (GBM).^{15,16}

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TTFields is a highly versatile first-in-class treatment modality

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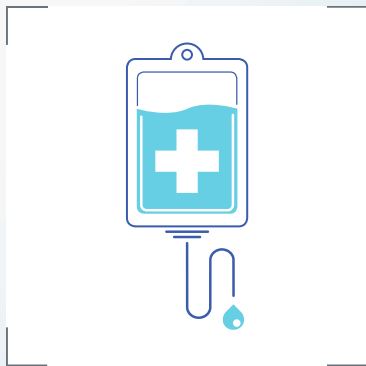


- TTFields therapy has significant potential for broad applicability across solid tumor types and lines of therapy
- Investigation of TTFields therapy is ongoing across clinical trials in multiple tumor types
- In approved indications, TTFields therapy is well tolerated, suggesting a low risk of additive systemic toxicity when used with other cancer treatment modalities

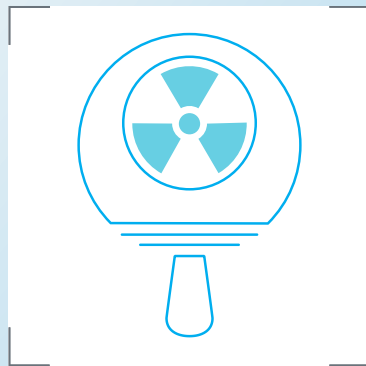


TTFields therapy can be added to cancer treatment modalities in approved indications

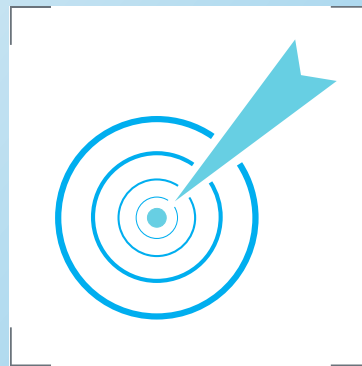
TTFields **demonstrate enhanced effects** across multiple solid tumor types, when used concomitantly with each of the following:



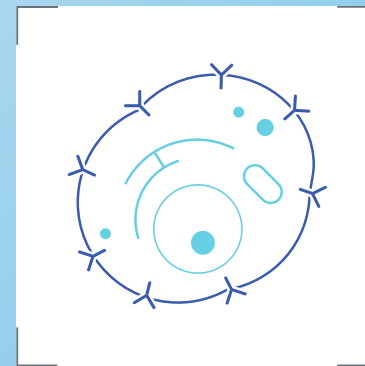
Chemotherapy



Radiation therapy (RT)



Targeted therapies



Immuno-oncologic (IO) agents