## **Supplementary information**

# Genomic predictors of response to PD-1 inhibition in children with germline DNA replication repair deficiency

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**Supplementary Fig.S1.** KM estimates of progression-free and overall survival for **(a)** all tumours, **(b)** CNS tumours, **(c)** all solid tumours (CNS and non-CNS). **(d)** Progression free survival for non-CNS solid tumours versus CNS tumours (p=0.01; 2-sided). The log-rank test was used to compare between the groups.





**Supplementary Fig.S2.** Univariable analysis of clinical predictors for **(a)** response and **(b)** survival following ICI at recurrence for replication-repair deficient cancers. Univariable logistic regression, estimated through generalized estimating equations, was fitted to assess association between each clinical variable and response. Univariable Cox regression model with robust standard errors was fitted to assess association between each clinical variable and response.



Supplementary Fig.S3. Biomarkers for response to ICI in replication-repair deficient cancers. (a) Spearman's rank correlation coefficient for dependent and independent variables. (b, c) Effects size (95% confidence intervals) for univariable analyses for the independent predictors for (b) response (assessed using univariable logistic regression) and (c) survival (assessed using univariable Cox regression model). (d-g) Dual role of SNVs and MS-indels in predicting outcomes: Multivariable analyses for (d) response (using logistic regression) and (e) survival (using Cox regression model) showing effects size and 95% confidence intervals. Kaplan-Meier (KM) curves for tumors with (f) both high and (g) both low SNVs and MS-indels (cut offs at the median values). All p-values are 2-sided.



**Supplementary Fig.S4. (a)** SNVs outnumber MS-indels in tumors with MMRD+PPD (Wilcoxon-Mann-Whitney test). Each bar represents the SNV count (black) and MS-indel (grey) for each individual tumor with identifiers mentioned on the x=axis. (b) Overall survival for MMRD tumors by SNVs/Mb (>=median). (c-f) Response and survival for MS-indels in coding microsatellites in MMRD+PPD versus MMRD tumors. (g-h) Response and survival for neoantigens generated by MS-indels in coding microsatellites in MMRD+PPD versus MMRD tumors. for survival, median values were utilized to stratify into 'high' and 'low' groups. For all box-plots for responders and non-responders, data are represented as median +/- interquartile range. For statistical significance in comparing responders and non-responders, the Wilcoxon-Mann-Whitney test was used. Survival analysis was performed using the Kaplan-Meier method, and the log-rank test was used to compare groups. All p values are 2-sided.

Predictive Genomic Biomarkers



Supplementary Fig.S5. Predictive Genomic and Immune Biomarkers for CNS versus non-CNS solid tumors (excluding haematological malignancies). (a) SNVs and (b) MS-indel burden were similar for CNS and non-CNS solid tumors. Responders and non-responders stratified by (d) SNV and (e) MS-indel burden and tissue type (Green- Non-CNS solid tumors, blue- CNS tumours). (e) CD8 and (f) PDL1 were significantly higher for non-CNS solid tumors. Responders had higher levels of (g) CD8 T-cell infiltration and (h) PD-L1 staining as compared to non-responders in all tumours (Green- Non-CNS solid tumors, blue- CNS tumours). Representative images from non-CNS solid tumours, all of which demonstrated (i) high CD8 and (j) PD-L1 expression (20X). All immunohistochemistry was analysed by two independent pathologists. For all box-plots for responders and non-responders, data are represented as median +/- interquartile range. For statistical significance in comparing responders and non-responders, the Wilcoxon-Mann-Whitney test was used. All p values are 2-sided.



**Supplementary Fig.S6. (A)** TCGA glioma dataset demonstrated 34 tumors (6.8%) with TMB >5 mutations/Mb. **(B)** Extreme mutation burden in replication-repair deficient (RRD) glioma responders (n=21) as compared to non-responders (n=10) and adult hypermutant gliomas (n=34). **(C)** High MS-indel burden in replication-repair deficient (RRD) gliomas (responder: n=21; non-responder: n=10) as compared to adult hypermutant gliomas (n=34). For statistical significance in comparing responders and non-responders, the Wilcoxon-Mann-Whitney test was used. All p values are 2-sided.



**Supplementary Fig.S7.** Concordance between immunohistochemical analyses and interpretation between two independent pathologists (CH/NA and OK) shown using **(a)** Spearman's rank correlation and **(b)** Bland-Altman plots.

(Red dotted lines: 95% confidence intervals)



Supplementary Fig.S8. Immune inference of tumor microenvironment at baseline and at flare. Deconvolution analyses of tumor samples of two patients (a) P31 and (b) P33 using CIBERSORT\_Absolute and EPIC (CD8+ T-cells are highlighted in red). (c) and (d) Same tumors demonstrating the actual abundance of all cell populations.



### c P33

T-cell Receptor	Clone Count	aaSeqCDR3	IEDB	VDJdb	McPAS-TCR
TRB	1414	CASSQGGWDR_PTAYQETQYF	Not reported	Not reported	Not reported
TRB	1034	CATSVQGYNEQFF	Not reported	Not reported	Not reported
TRB	785	CASSPGGLTTGNTIYF	Not reported	Not reported	Not reported

**Supplementary Fig.S9. (a)** Sequencing reads for two patients at diagnosis and at time of flare (Data are presented as mean values +/- SEM). **(b)** TRB richness and diversity measures at flare compared to baseline for P31 (sequencing had reached saturation for both samples at pre-therapy and at flare). **(c)** Complementary-determining regions (CDR3) sequences of the major three clonotypes detected at flare in P33 (TRB: T cell receptor beta).



Supplementary Fig.S10. Immune responses in peripheral blood at flare. (a) 4-1BB+ CD4+ T-cells in blood from responders without flare, non-responders and flare. For the box-plots, data are represented as median +/- interquartile range. For statistical significance in comparing responders and non-responders, the Wilcoxon-Mann-Whitney test was used. All p values are 2-sided. (b) TRB clonotypes were tracked in 3 patients at baseline, time of flare, and following ICI continuation. Each box represents a distinct TRB clonotype. Clonotypes detected multiple time-points are color-coded and tracked in serial samples over time.

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### Flow Cytometry Gating Strategy



### Supplementary Fig. S11. Gating strategy for flow cytometry utilized in the index study.

P38 10	P37 IQ	P35 IQ	10	P35 IQ	P34 IQ	P33 IQ	P22 IQ	P31 IQ	P30 10	P23 10	0	P28 10	10	P27 10	P26 10	P25 10	P24 IQ	P23 10	P22  0	P21 IQ	P20 10	P19 IQ	P18 IQ	P17 IQ	10	Pt6 IQ	P15 IQ	P14 IQ	P13 IQ	P07 10	P05 10	Q	P05 10	P04 10	P03 IQ	P02 IQ	10	P01 IQ	Ratient id Stud
20 7	10 27.4	21 13.6	13	14 125	06 13.2	12 28	18 11	99 91	27 12.9	38 16.9	3	26 10.8	34	35 11	23 11.6	44 93	45 7.9	65	32 27.6	46 93	19 10.6	43 9	40 14	33 30.9	07	08 13.8	37 5	17 16	23 16	22 83	39 16.8	01	02 16.7	24 24	11 12.6	04 3	41	42 15	yID (years)
M Ly	F CM	M CM		F CM	F CM	F CM	M Ly	F CM	M CM	M Ly		M CM		F CM	FLy	M CM	M CM	F CM	F CM	M CM	M CM	F CM	M Ly	M P		M CM	F CM	M CM	M P	M CM	FLy		M CM	M Lynch(	F CM	M CM		F CM	97 Sex Gen
nah MSH	MRD MS+	MRD MSF		MRD MS+	MRD PMS	MRD MS+	nah PMS	MRD PMS	MAD BW8	nah MU-		MRD MS+		MRD MS+	ndt MSE	MAD PMS	MRD PMS	MRD PMS	MAD PMS	MAD BWS	MRD PMS	MRD MSF	nah MSH	190 POL		MRD PMS	MRD PMS	MRD PMS	POL POL	MRD PMS	nch MU-		MRD MS+	(dganic) PMS2, P	MAD WSF	MAD BWS		MAD PMS	mine Gen
Ŕ	6	5		6	8	2	ю	ю	ю	-		2		<del>1</del> 3 Т	2	ю	ю	ю	ю	ю	183	5	2	'n		13	13	ю	h	13	=		8	WSH5	5	ю	_	ю	6
Anaplastic Astrocytoma	Gliddestone	Anaplastic Astrocytoma	Glioblastoma	Anaplastic Astrocytoma	<b>GNSEmbryonal turnor</b>	Glioblastoma	lleopacal Signet Cell Carcinoma	Giddastoma	Acute Myeloid Leukemia	Colorectal Adenocal cinoma	Giobastoma	Colorectal Adancear cinoma	Glioblastoma	Cell Acute Lymphoblastic Lymphoma	Gliddiastoma	Gliddastoma	Giobastoma	Gliddastoma	Gliddastoma	Glioblastoma	Gliddestone	Gliddlastoma	Glioblastoma	Gliddestone	Colorectal Adenocar cinoma	Glioblastoma	Gliddestone	Gliddlastoma	Colorectal Adenocar cinoma	Glioblastoma	Colorectal Adancear cinoma	Anaplastic Astrocytoma	Colorectal Adancear cinoma	Colorectal Adenocar cinoma	Colorectal Adenocal cinoma	Gliddastoma	Gliddastoma	Colorectal Adenocal cinoma	Tumour Type
Surgery+ Adjuvant treatment	Surgery + Adjuvant treatment	Surger y+ Adjuvant treatment	Sugery+ Adjuvant/reatment	Surgery + Adjuvant treatment	Surger y+ Adjuvant treatment	Surgery+ Adjuvant treatment	Surgery+ Adjuvant/reatment	Sugery+ Adjuartheatment	Chemotherapy	Sugery+ Adjuvanthreatment	Sugery+ Adjuvant treatment	Surgery only	Sugery+ Adjuvant/reatment	Chamotharapy	Surgery+ Adjuvant/reatment	Sugery+ Adjuvanthreatment	Sugery+ Adjuvant treatment	Surgery+ Adjuvant treatment	Sugery+ Adjuvanthreatment	Surgery+ Adjuvant treatment	Surgery + Adjuvant treatment	Surger y+ Adjuvant treatment	Surgery+ Adjuvant treatment	Surgery + Adjuvant treatment	Surgery only	Surgery+ Adjuvant treatment	Surgery + Adjuvant treatment	Surger y+ Adjuvant treatment	Surgery+ Adjuvant treatment	Surgery+ Adjuvant treatment	Surgery only	Surgery+ Adjuvant/reatment	Surgery only	Surger y + Adjuvant treatment	Surgery+ Adjuvant/reatment	Surger y+ Adjuvant treatment	Surger y+ Adjuvant treatment	Surgery+ Adjuvant/reatment	Previous treatment Ag
85	28.1	14.4		13.5	14.1	31	12.3	97	13.2	17.2		11.3		11.3	12.1	95	82	7.1	27.8	95	11.3	92	14.7	31.2		14.5	55	16.5	17.5	92	16.8		17	24.2	13	45		15.9	e at ICI (years)
No	No	No	No	No	No	No	Yes	No	Not applicable	Yes	No	No	No	Not applicable	No	No	No	No	Yes	No	No	No	No	No	Yes	No	No	Yes	Yes	No	No	No	No	Yes	Yes	No	No	No	Diseminated disease
Nivdunab	Nivdunab	Nivolumab	Nivolumab	Nivdunab	Nivolumab	Nivdunab	Nivolumab	Nivdumab	Nivdunab	Rembrolizumab	Nivdumab	Nivdunab	Nivolumab	Nivolumab	Nivdumab	Nivdunab	Nivdumab	Nivdumab	Nivdunab	Nivdunab	Nivdunab	Nivolumab	Nivdunab	Rembrolizumab	Pembrolizumab	Rembrolizumab	Nivdunab	Nivolumab	Nivolumab	Nivdunab	Rembrolizumab	Rembrolizumab	Rembrolizumab	Rembrolizumab	Rembrolizumab	Nivdunab	Nivdumab	Nivolumab	ICI agent
Progressive disease	Progressive disease	Progressive disease	No central review	No central review	Progressive disease	Progressive disease	Progressive disease	Progressive disease	Not applicable	Progressive disease	Progressive disease	Nomessurable disease	Progressive disease	Not applicable	Progressive disease	Completer esponse	Completer expanse	Partial response	No central review	Stable di sesse	Progressive disease	Stable di sease	Stable di sease	Partial response	Partial response	Progressive disease	Partial response	Completer espanse	Completer esponse	Stable di sease	Partial response	No central review	Nomessurable disease	Partial response	Completer esponse	Partial response	Completer esponse	Stable di sesse	N893.
Yes	No	Yes	No	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	No	Yes	Yes	Yes	No	No	Yes	Yes	No	Yes	Yes	No	No	No	Yes	No	Yes	No	No	No	Yes	No	No	No	Yes	No	No	Progression
No	Ch/	P40	No	Ch/	P40	No	No	No	No	No	No	No	bilimumab	P40	No	No	No	bilimumab	No	No.	Iplimunab + Re-inadiation + Targated ag	P40	No	Ch/	No	Re-irradiation	Ch/	Iplimunab + Re-irradiation + Targeted ag	No	No	Ch/	bilimumab	No	No	No	bilinumab	No	No	Treatment in addition to PD1 ICI
Non-responder	Flare and stopped	Flare and stopped	Responder	Non-responder	Non-responder	Flare and stopped	Flare and stopped	Flare and stopped	Non-responder	Flare and stopped	Flare and stopped	Responder	Flare and stopped	Non-responder	Non-responder	Responder	Responder	Flare and response	Non-responder	Flare and response	nt Non-responder	Responder	Responder	Responder	Flare and response	Non-responder	Responder	nt Responder	Responder	Responder	Responder	Non-responder	Responder	Responder	Responder	Flare and response	Responder	Responder	Final Tumor Response
Dead	Dead	Dead		Dead	Dead	Dead	Dead	Dead	Dead	Dead		Dead		Dead	Dead	Aive	Aive	Dead	Dead	Aive	Dead	Aive	Aive	Aive		Dead	Aive	Dead	Alve	Aive	Aive		Aive	Alive	Aive	Alve		Aive	Patient status
MMRD	MMRD+PPO	MMRD	MMRD+PPD	MMRD	MMRD+PPD	MMRD+PPD	DBWW	MMRD+PPD	MMRD	DBWW	MMRD	MMRD	MMRD	MMRD	MMRD	Not available	MMRD+PPD	MMRD+PPO	Not available	MMRD+PPO	MMRD+PPO	MMRD+PPD	Not available	MMRD+PPO	MMRD+PPD	MMRD+PPD	MMRD+PPO	MMRD+PPD	MMRD+PPD	MMRD+PPD	MMRD	MMRD	Not available	MMRD	MMRD+PPD	MMRD+PPD	MMRD+PPO	MMRD+PPO	Tumor RRD sta
1478	54341	1406	25175	1499	9859	19 533	6520	34116	1404	5785	1008	2035	12254	810	1995	Not available	13950	39642	Not available	12123	16810	61414	Notavailable	23641	17070	31 690	22.483	14544	21 195	20330	1172	3993	Notavailable	3995	23433	35396	22510	24349	tus Total necantigen
200	18481	351	9797	434	2001	8223	654	10224	197	608	257	403	43)4	289	48	Not available	11542	11522	Not available	4993	9933	22341	Not available	11502	221	10331	7414	4213	8333	6301	207	786	Not available	673	16417	10227	7395	8233	Total nonsynonymous variants
127	7046	157	3355	173	1085	3058	0	4000	8	0	103	159	2820	140	191	Notavalable	4933	4187	Notavalable	1993	3999	8230	Not available	4309	150	4134	2876	142	2816	2391	85	0	Not available	0	6214	-4211	3409	3993	Total synonymous variants
827	37497	744	19902	883	533	17020	2737	21090	523	2785	623	1025	11771	647	1284	Not available	22234	24186	Not available	10295	19553	49310	Not available	23133	8334	20793	14739	10203	18579	17228	497	1725	Not available	2820	3469	21488	16242	17805	Total Mutations (SNVs and Indes)
1254	748.34	14.88	398.04	17.16	110.76	340.4	54.74	421.8	10.46	55.9	12.55	20.5	235.42	12.94	25.68	Notavailable	454.68	483.72	Notavailable	205.92	391.16	918.2	Notavailable	492.76	133.68	415.86	235.98	204.18	373.58	344.95	994	345	Notavailable	56.4	6832.18	429.76	324.84	352.12	TMBMb
501	37210	699	19537	787	5254	16376	1531	20490	41	1305	501	791	11832	646	871	Not a valiable	22590	22222	Not a valiable	10001	18555	45395	Not a vail able	22927	3310	20198	14351	9229	18572	13789	409	1673	Not a vail able	1808	33537	20881	15767	18530	Total SNVs
10.02	7442	13.33	392.74	1574	105.08	327.52	33.62	4092	8.22	26.1	10.02	1582	233.84	12.92	17,42	Not a valiable	451.8	494.44	Not a valiable	201.02	371.1	911.92	Not a valiable	498.54	66.2	403.95	287.02	184.53	371.44	275.38	8.18	33.45	Not a valiable	32.16	670.74	417.62	315.34	330.6	SWM
125	27	75	265	71	284	644	1205	63)	112	1490	127	234	79	_	413	Not av all able	644	954	Not av all able	255	1003	314	Not av all able	211	3524	995	443	9 <del>9</del> )	1077	3459	88	52	Not av all able	1212	82	607	43	1076	Total Indels
252	514	1.5	5.3	1.42	568	12.88	24.12	126	224	238	254	468	158	0.02	826	Notavailable	12.88	19.28	Notavailable	5.9	20.05	628	Notavailable	422	72.48	11.9	895	196	214	69,18	1.76	104	Notavailable	24.24	18.44	12.14	9.5	21.52	Indels/Mb
122	993	101	178	98	1226	829	458	516	â	545	130	194	95	å	195	Not available	199	959	Not available	158	a	480	Notavalable	201	1370	227	668	332	174	1423	123	1984	Notavalable	393	335	342	612	832	Total MS-indels

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# Table S1. Summary of clinical features, outcomes and biomarkers (n=45 tumors in 38 patients)