



Istituto Nazionale di Fisica Nucleare

GAP

(towards) Genome Adjusted Particle therapy

RL – Giorgio Baiocco



Informazioni generali

Sezioni Coinvolte:

- TIFPA (RN)
- LNL
- RM2/RM3
- PV/CNAO
- TO
- CT

Durata del progetto:
2026-2028

FTE 2026 sezione PV : 2.15

Anagrafica di Pavia 2026:

Giorgio Baiocco	PA	50
Isabella Guardamagna	TecLab	50
Leonardo Lonati	AdR	50
Cecilia Riani	PhD	25
Pullia Marco	CNAO	10
Facoetti Angelica	CNAO	10
Charalampopoulou Alexandra	CNAO	10
Carnevale Federica	CNAO	10

Goal del progetto

Obiettivi

- **Integrate genomic data into radiobiological models** to develop a framework towards **Genomic Adjusted Particle Therapy (GAP)**.
- **Identify gene expression signatures** modulated by different radiation qualities (protons, carbon ions, and X-rays) using next-generation sequencing (NGS).
- **Develop predictive models** that combine genomic, microdosimetric, and radiobiological data to predict treatment outcomes.
- **Characterize radiation fields** using advanced new microdosimeters (MUSICA, DIODE) and reference microdosimeters (miniTEPC) to ensure precise radiation field correlation with biological effects at differential spatial scales.
- **Evaluate biological responses** across multiple cell lines (lung and pancreatic, healthy and tumor) using 2D and 3D assays, including apoptosis, senescence, and extracellular vesicle release.

Obiettivi specifici

- INFN-Pavia section is in charge of:
 - experimental measurements on cell-culture media collected from samples from TIFPA (healthy and tumor lung cell lines) and CNAO (healthy and tumour pancreas cell lines) to isolate and characterize EV's
 - RNA extraction and RNE-seq (external service) to obtain data for bioinformatic analysis
 - Radiobiological measurements with carbon ions at the CNAO facility (starting from the 2° year) on pancreatic cells cultures in 2D and spheroids

GARD: genomic adjusted radiation dose

Genomic analysis at 2Gy survival fraction

Table 1. SF2 values for 48 cell lines in the database

Cell line	Recorded SF2	Cell line	Recorded SF2
BREAST_HS578T	0.79	COLON_COLO205	0.69
BREAST_MDAMB231	0.82	COLON_HCC-2998	0.44
COLON_HCT116	0.38	COLON_HT29	0.79
COLON_HCT15	0.4	COLON_KM12	0.42
COLON_SW620	0.62	MELAN_LOXIMVI	0.68
LEUK_CCRFCEM	0.185	MELAN_M14	0.42
LEUK_HL60	0.315	MELAN_MALME3M	0.8
LEUK_MOLT4	0.05	MELAN_SKMEL28	0.74
MELAN_SKMEL2	0.66	MELAN_SKMEL5	0.72
NSCLC_A549ATCC	0.61	MELAN_UACC257	0.48
NSCLC_H460	0.84	MELAN_UACC62	0.52
NSCLC_HOP62	0.164	NSCLC_EKVX	0.7
NSCLC_NCIH23	0.086	NSCLC_HOP92	0.43
OVAR_OVCAR5	0.408	OVAR_OVCAR3	0.55
RENAL_SN12C	0.62	OVAR_OVCAR4	0.29
BREAST_BT549	0.632	OVAR_OVCAR8	0.6
BREAST_MCF7	0.576	OVAR_SKOV3	0.9
BREAST_MDAMB435	0.1795	PROSTATE_DU145	0.52
BREAST_T47D	0.52	PROSTATE_PC3	0.484
CNS_SF268	0.45	RENAL_7860	0.66
CNS_SF539	0.82	RENAL_A498	0.61
CNS_SN19	0.43	RENAL_ACHN	0.72
CNS_SN175	0.55	RENAL_CAKI1	0.37
CNS_U251	0.57	RENAL_UO31	0.62

Abbreviation: SF2 = survival fraction at 2 Gy.

Linear regression on selected genes
RSI (Radiation sensitivity index)

$$\begin{aligned} \text{SF2} = & - 0.0098009^* \text{AR} + 0.0128283^* \text{cJun} \\ & + 0.0254552^* \text{STAT1} - 0.0017589^* \text{PKC} \\ & - 0.0038171^* \text{RelA} + 0.1070213^* \text{cABL} \\ & - 0.0002509^* \text{SUMO1} - 0.0092431^* \text{CDK1} \\ & - 0.0204469^* \text{HDAC} - 0.0441683^* \text{IRF1} \end{aligned}$$

GARD: genomic adjusted radiation dose

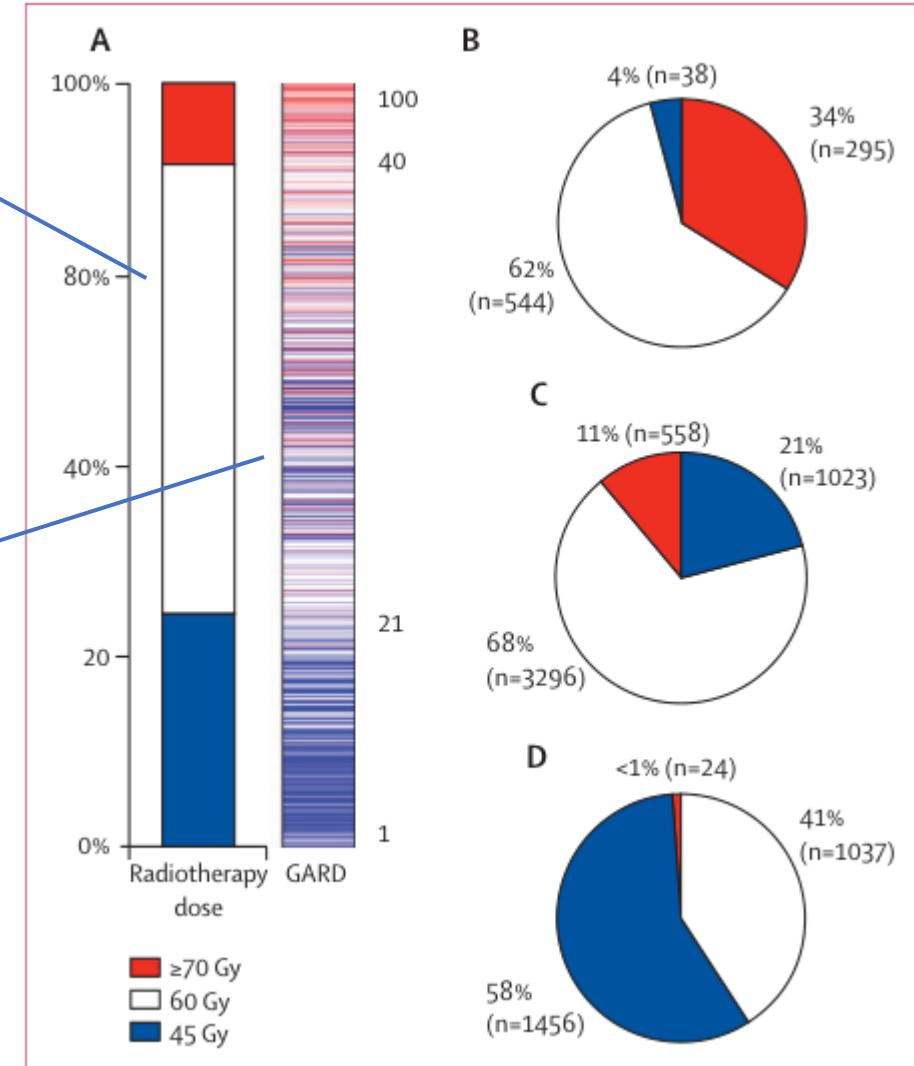
Radiosensitivity
adjusted by patient
specific genomic
profiling

$$S = e^{-nd(\alpha + \beta d)}$$

$$\alpha_g = -\frac{\ln RSI}{nd} - \beta d, \quad (1)$$

Dose received by
patient

Dose **should have**
been received by
patient according to
GARD



Preventivi

2026

	missioni	missioni_sj	consumo	altri_cons	inventario	spservizi	spservizi_sj	totali	totali_sj
LNL	1.5	1.5	2					3.5	1.5
PV				7		7		14	
ROMA2	3		1.5					4.5	
TIFP				14.5			5	14.5	5
TO	4		1.5	19.5	4			29	
Totali	8.5	1.5	5	41	4	7	5	65.5	6.5

INFN-Pavia section

- Consumabili per misure citometria e kit reagenti per isolamento vescicole e estrazione RNA, siero per colture cellulari di prova
- Acquisto di linee cellulari (sana e tumorale) per esperimenti di radiobiologia
- Sequenziamento RNA da ditta esterna, compreso shipment e costi di data delivery (~24 campioni)

Previsione di budget

2027

- Consumabili: 61 k€;
- Missioni: 14 k€;
- Instruments: 7 k€;
- Service: 2.5 k€ [sj]
- Total: 82 + 2.5 k€ [sj]

2028

- Consumabili: 47 k€;
- Total: 47 k€

Total request for year 1-2-3:

- Y1 - 65.5 k€ + 6.5 k€ [sj]
- Y2 - 82 + 2.5 k€ [sj]
- Y3 - 47 k€
- **TOTAL = 194.5 + 9 [sj] k€**

INFN-Pavia section

	I anno	II anno	III anno	Totale	GROUP
consumabili citometria, kit reagenti per isolamento vescicole e estrazione RNA	4	4		8	PV
RNA-SEQ da ditta compresa spedizione e costi di data delivery	7	7		14	PV
linee cellulari	3			3	PV
consumabili per cell culture e misure radiobiologiche presso CNAO		15	25	40	PV e PV-CNAO
NGS			10	10	PV-CNAO
	14	26	35	75	