Paola Tacchetti IRCCS Azienda Ospedaliero-Universitaria di Bologna Istituto di Ematologia "Seràgnoli" Terapie di reindirizzamento delle cellule immuni effettrici: Anticorpi bispecifici anti-GPRC5D e FcRH5 con approvazione FDA/EMA ed in fase di sviluppo (e combinazione con altri agenti)

30-31 gennaio 2024 BOLOGNA, Royal Hotel Carlton

Disclosures of Paola Tacchetti

Company name	Research support	Employee	Consultant	Stockholder	Speakers bureau	Advisory board	Other
Janssen						x	Honoraria
BMS/Celgene						x	Honoraria
Amgen						x	Honoraria
Sanofi						x	Honoraria
GSK						x	Honoraria
Abbvie							Honoraria
Takeda							Honoraria
Pfizer							Honoraria

Non-BCMA Targets BsAbs

GPRC5D

- GPRC5D is a member of the G protein-coupled receptor family with an unknown function
- It is highly expressed on malignant PCs, as well as hard keratinized structures (hair shaft, nail, and central region of the tongue)

GPRC5D

Talquetamab Forimtamig TALQUETAMAB FDA: RRMM ≥4 prior tp, TCE EMA: RRMM ≥3 prior tp, TCE

FcRH5

- FcRH5 is a surface protein in the Ig superfamily
- It is expressed only in B cells, with increasing expression in mature B cells and plasma cells
- FcRH5 is involved in proliferation and isotype expression

FcRH5

Cevostamab

Atamaniuk J, et al. Eur J Clin Invest 2012; Smith EL, et al. Sci Transl Med 2019; Li J et al. Cancer Cell 2017; Chari A, et al. N Engl J Med 2022; Carlo-Stella C et al. ASH 2022; Trudel S, et al. Blood 2021

Talquetamab, GPRC5D × CD3 T-Cell Redirecting BsAb: MonumenTAL-1



MonumenTAL-1 ph 1/2 st

Adults with measurable MM Phase I: progression on or intolerance to all established therapies; ECOG PS 0–1 Phase II: ≥3 prior lines of therapy that included a PI, an IMiD, and an anti-CD38 antibody: ECOG PS 0–2

Multicenter, open-label phase I/II trial
 Primary endpoint (phase II): ORR

*Previous anti-BCMA therapy allowed; T-cell redirection therapy naive.

Talquetamab 0.4 mg/kg SC QW*

Talquetamab 0.8 mg/kg SC Q2W*

■ Secondary endpoints (phase II): DOR, ≥ VGPR rate, ≥ CR, sCR rate, TTR, PFS, OS, MRD, safety

RRMM patients after a median of 5 PL, 74% (@0.4mg/kg) / 69% (@0.8mg/kg) triple-class refractory

TALQUETAMAB FDA: RRMM ≥4 prior tp, TCE EMA: RRMM ≥3 prior tp, TCE

Characteristic	0.4 mg/kg SC QW (n=143)	0.8 mg/kg SC Q2W (n=145)	Prior TCR (n=51)
Age (years), median (range)	67.0 (46-86)	67.0 (38–84)	61.0 (38–78)
Male, n (%)	78 (54.5)	83 (57.2)	31 (60.8)
Bone marrow plasma cells ≥60%,ª n (%)	17 (12.3)	32 (22.7)	8 (17.0)
Extramedullary plasmacytomas≥1, ^b n (%)	33 (23.1)	37 (25.5)	16 (31.4)
High-risk cytogenetics, ^c n (%)	41 (31.1)	37 (28.9)	18 (40.9)
ISS stage, ^d n (%)			
I	62 (43.4)	64 (44.4)	24 (47.1)
II	53 (37.1)	45 (31.3)	18 (35.3)
III	28 (19.6)	35 (24.3)	9 (17.6)
Prior lines of therapy, median (range)	<mark>5 (2–13)</mark>	<mark>5 (2–17)</mark>	<mark>6 (3–15)</mark>

Characteristic	0.4 mg/kg SC QW (n=143)	0.8 mg/kg SC Q2W (n=145)	Prior TCR (n=51)
Exposure status, n (%)			
Triple-class ^e	143 (100)	145 (100)	51 (100)
Penta-drug ^f	105 (73.4)	101 (69.7)	40 (78.4)
BsAb	NA	NA	18 (35.3) ^g
CAR-T therapy	NA	NA	36 (70.6) ^h
BsAb + CAR-T therapy	NA	NA	3 (6.0)
Refractory status, n (%)			
Triple-class ^e	106 (74.1)	100 (69.0)	43 (84.3)
Penta-drug ^f	42 (29.4)	34 (23.4)	21 (41.2)
To last line of therapy	134 (93.7)	137 (94.5)	31 (60.8)

Chari A, et al. N Engl J Med 2022; Schinke CD et al. EHA 2023

MonumenTAL-1: Efficacy



Outcome	0.4 mg/kg SC QW (n=143)	0.8 mg/kg SC Q2W (n=145)	Prior TCR (n=51)
mFU, mo	18.8	12.7	14.8
12-mo DOR rate in patients with ≥CR, %	78.9	90.5	80.5
mPFS, mo (95% CI)	7.5 (5.7–9.4)	14.2 (9.6−NE) ^ь	5.1 (3.4–12.3)
12-mo PFS rate, %	34.9	54.4	38.1
12-mo OS rate, %	76.4	77.4	62.9

MonumenTAL-1:1

- >71% ORR across QW and Q2W TCR naïve cohorts
- <u>65% ORR in the prior TCR cohort</u>

ASH23 update:²

- ORR of 73% and mDOR of >1 year in the post-CAR-T setting
- 52% response in patients exposed to prior bispecific antibodies

MonumenTAL-1: Efficacy and Safety of Patient Subgroups From MonumenTAL-1

TABLE 2: ORR in high-risk subgroups

	0.4 mg/kg QW								
Outcome	Overall (N=143)	Age ≥75 years (n=21)	Renal impairment (n=40)	High-risk cytogenetics (n=41)	ISS stage III (n=28)	EMD (n=33)	Triple-class refractory (n=106)		
mFU, mo	18.8	18.7	19.5	19.2	18.5	18.4	18.7		
ORR, n (%)	106 (74.1)	15 (71.4)	26 (65.0)	29 (70.7)	18 (64.3)	16 (48.5)	77 (72.6)		

	0.8 mg/kg Q2W							
Outcome	Overall (N=145)	Age ≥75 years (n=32)	Renal impairment (n=45)	High-risk cytogenetics (n=37)	ISS stage III (n=35)	EMD (n=37)	Triple-class refractory (n=100)	
mFU, mo	12.7	11.9	13.0	12.5	13.3	12.1	12.8	
ORR, n (%)	104 (71.7)	24 (75.0)	30 (66.7)	28 (75.7)	21 (60.0)	16 (43.2)	69 (69.0)	

mFU, median follow-up.

FIGURE 2: Outcomes among select high-risk subgroups in Q2W cohort



FIGURE 3: ORR in prior BCMA subgroups



*Some patients received >1 prior BCMA therapy, leading to differences in total patient count.

Talquetamab vs Real-World Physician's Choice of Therapy: Comparative Efficacy in Patients With TCE RRMM

Data sources

- Individual patient-level data from MonumenTAL-1 were included for pts who received sc talquetamab 0.4 mg/kg QW or 0.8 mg/kg Q2W by a data cut-off of January 2023
- An external control group was created from eligible pts in the Flatiron database who met MonumenTAL-1 eligibility criteria by a data cut-off of July 2022

	Talq	uetamab QW vs R	WPC	Talquetamab Q2W vs RWPC			
Outcome/analysis	Median, mo	HR (95% CI)	P value	Median, mo	HR (95% CI)	P value	
PFS							
Primary analysis	7.5 vs 4.0	0.55 (0.44-0.69)	<0.0001	14.2 vs 4.0	0.40 (0.31-0.53)	<0.0001	
Fully adjusted model	7.5 vs 4.2	0.56 (0.45-0.71)	<0.0001	14.2 vs 4.0	0.41 (0.31-0.54)	<0.0001	
TTNT							
Primary analysis	9.1 vs 5.1	0.59 (0.47-0.74)	<0.0001	13.3 vs 5.1	0.45 (0.35-0.59)	<0.0001	
Fully adjusted model	9.1 vs 5.1	0.60 (0.48-0.77)	<0.0001	13.3 vs 5.0	0.46 (0.36-0.61)	<0.0001	
OS							
Primary analysis	NR vs 16.5	0.56 (0.40-0.78)	0.0007	NR vs 15.9	0.48 (0.33-0.70)	0.0002	
Fully adjusted model	NR vs 16.8	0.58 (0.41-0.83)	0.0029	NR vs 17.5	0.50 (0.34-0.75)	0.0008	

FIGURE 1: Key patient eligibility criteria

eria	MonumenTAL-1 (QW [n=143] and Q2W [n=145]) and RWPC (n=1169) ^a						
Patient eligibility crit	• Triple-class exposed • ≥3 prior LOT • Progression ≤12 months after last therapy • No prior receipt of T-cell redirection therapy, such as CAR-T or bispecific antibodies	• ECOG performance status ≤2 • Hemoglobin ≥8 g/dL • Creatinine clearance ≥40 mL/min/1.73 m ²					

*RWPC cohort included 629 patients who received 1169 treatment regimens across all eligible lines of therapy. CAR, chimeric antigen receptor.

TABLE 1: Treatment regimens in the RWPC cohort $\ensuremath{^a}$

Treatment regimen	Frequency, n (%) ^b (n= 1169)
Pomalidomide, elotuzumab, dexamethasone	56 (4.8)
Pomalidomide, daratumumab, dexamethasone	46 (3.9)
Clinical study drug	43 (3.7)
Carfilzomib, dexamethasone	42 (3.6)
Carfilzomib, cyclophosphamide, dexamethasone	36 (3.1)
Carfilzomib, dexamethasone, pomalidomide	32 (2.7)
Belantamab mafodotin-blmf	23 (2.0)
Bortezomib, selinexor, dexamethasone	23 (2.0)
Elotuzumab, lenalidomide, dexamethasone	22 (1.9)
Daratumumab, dexamethasone	21 (1.8)
Selinexor, dexamethasone	21 (1.8)
Daratumumab, dexamethasone, lenalidomide	19 (1.6)
Pomalidomide, dexamethasone	19 (1.6)
Bortezomib, daratumumab, dexamethasone	18 (1.5)
Clinical study drug, dexamethasone	18 (1.5)
Daratumumab/hyaluronidase-fihj, dexamethasone, pomalidomide	16 (1.4)

 $^{\circ}$ Only treatment combinations used in ≥ 16 patients are presented. $^{\circ}$ Percentages are calculated with the number of treatment regimens received by the 629 patients in the RWPC cohort set as the denominator (n=1169).

MonumenTAL-1: Safety

Most common AEs
included CRS,
infection, dysgeusia,
and skin/nail toxicity

 5 patients discontinued due to skin-related AEs and dysgeusia

	Most Common AEs,	n = 143		n = 145		n = 51	
dysgeusia, /nail toxicity	N (%)	Any Grade	Grade ≥ 3	Any Grade	Grade ≥ 3	Any Grade	Grade ≥ 3
	Hematologic						
s ued due to	Anemia	66 (44.8)	45 (31.5)	66 (45.5)	40 (27.6)	25 (49.0)	14 (27.5)
ted AEs	Neutropenia	50 (53.0)	44 (30.8)	41 (28.3)	32 (22.1)	28 (54.9)	27 (52.9)
eusia	Thrombocytopenia	39 (27.3)	29 (20.3)	43 (29.7)	27 (18.6)	19 (37.3)	15 (29.4)
	Nonhematologic						
	CRS	113 (79.0)	3 (2.1)	108 (74.5)	1 (0.7)	39 (76.5)	1 (2.0)
CRS, dysgeusia, nail toxicity s ued due to ced AEs leusia On-target, off-tumor effects	Infection	84 (58.7)	28 (19.6)	96 (66.2)	21 (14.5)	37 (72.5)	14 (27.5)
	Dysgeusia	103 (72.0)	-	103 (71.0)	-	39 (76.5)	-
On-target,	Skin related	80 (55.9)	0	106 (73.1)	1 (0.7)	35 (68.6)	0
effects	Nail related	78 (54.5)	0	78 (53.8)	0	32 (62.7)	0
	Rash related	57 (39.9)	2 (1.4)	43 (29.7)	8 (5.5)	18 (35.3)	2 (3.9)
	Weight decrease	59 (41.3)	3 (2.1)	60 (41.4)	8 (5.5)	15 (29.4)	0

TCR-Naive, Q2W Dose

Prior TCR

TCR-Naive, QW Dose













Chari A, et al. N Engl J Med 2022; Schinke CD et al. EHA 2023; Purcell K, et al. IMS 2023. Oral NSO-07

Management Considerations for Dermatologic Toxicities Associated With Talquetamab

A single-center experience of dermatologic AEs in patients from MonumenTAL-1

	Total (N=24)		0.4 mg/l (n	kg SC QW =8)	0.8 mg/kg SC Q2W (n=16)	
Incidence of AEs in Mount Sinai patients, n (%)	Any grade	Grade 3/4	Any grade	Grade 3/4	Any grade	Grade 3/4
Skin (dry skin and exfoliation)	21 (88)	1 (4)	6 (75)	0	15 (94)	1 (6)
Rash	11 (46)	8 (33)	3 (38)	1 (13)	8 (50)	7 (44)
Nail (thinning and peeling)	14 (58)	0	4 (50)	0	10 (63)	0

Skin, rash, and nail AEs were generally not painful

Most dermatologic non-nail Aes (dry skin/exfoliation and rash AEs) resolved (63–86%)

Timing of AEs	Median time to onset	Median time to resolution
Skin (dry skin and exfoliation)	4.6 weeks	7.1 weeks
Rash	6.6 weeks	3.2 weeks
Nail (thinning and peeling)	15.0 weeks	16.4 weeks

Dry skin	Heavy moisturizers	Pruritus	Loratadine 10 mg PO daily for 3–5 days post talquetamab dose and triamcinolone 0.1% cream BID	AEs	Patient education
Hand and/or foot peeling	Ammonium lactate 12% lotion to soles and palms BID	Injection site reaction		Skin(dry skin and exfoliation) and rash	 Short, lukewarm showers, and moisturize afterward and throughout the day Alert team of any rashes or redness around injection area immediately Alert team of any secondary (fungal, bacterial) skin infections immediately
Nail thinning and peeling	Nail hardeners, topical vitamin E oil, and triamcinolone 0.025% ointment	Body rash/drug rash	Above plus consider methylprednisolone taper and betamethasone 0.05% cream BID	Nail	 Apply heavy moisturizer to cuticles and keep nails/cuticles short and clean Alert team of any signs of a fungal infection (thickened or discolored nails)

Consider dose HOLD for other grade 3 dermatologic AEs

Purcell K, et al. IMS 2023. Oral NSO-07

MonumentTAL-1: Efficacy and Safety of Less Frequent/Lower Intensity Dosing of Talquetamab



100 79.2 (19/24) 80 SCR 25.0 % 60 Patients, CR VGPR >VGPR 40 29.2 PR 75.0 20 4.2

Prospective cohorts with change in AE status after switch vs matched cohort without dose reduction^a



Trend toward improved resolution of GPRC5D-related AEs, except weight loss

	Prospective (n=19)
Median follow-up, mo (range)ª	13.2 (4.0+–16.1)
Median PFS, mo (95% Cl)ª	13.2 (8.8–NE)
12-mo PFS rate, % (95% Cl)ª	50.1 (27.9–68.7)
Median DOR, mo (95% Cl)	NE (8.3–NE)

- In the 0.8 mg/kg Q2W registrational cohort (n=145)^{1,b}
 - ORR: 71.7%
 - Median PFS: 14.2 mo (95% CI, 9.6-NE)
 - 12-mo PFS rate: 54.4%
 - Median DOR: NE (95% CI, 13.0–NE)

Forimtamig: anti GPRC5DxCD3 Bispecific antibody



Phase 1/2 study D15 D1 D1 D8 C1 C2 C3+ \cdots 7 7 14 14 $days^*$ $days^*$

Common (≥20%) hematologic and non-hematologic AEs by Grade



51 (iv) and 57 (sc) pts, 5 median prior LOT

- 70% TCR
- 20% prior anti-BCMA
- 30% EMD

Response rate across all tested target doses (IV: 18–10,000 μ g; SC 30–7200 μ g) in efficacy-evaluable patients*



Cevostamab: anti FcRH5xCD3 Bispecific antibody







Common (≥15%) hematologic and non-hematologic AEs in all patients by Grade[‡]





mDOR: 11.5 mo

Cevostamab: durable responses off therapy



Non-BCMA Bispecific Antibodies

	Anti-GPRC5d Talquetamab		Anti-GPRC5D Forimtamig		Anti-FCRH5 Cevostamab	
Patients (n)	143 T-cell redirecting therapy naïve (ADC allowed)	145 T-cell redirecting therapy naïve (ADC allowed)	51 Prior anti-BCMA TT (CARs/BsAb) allowed	51 Prior anti-BCMA TT (CARs/BsAb) allowed	57 Prior anti-BCMA TT (CARs/BsAb) allowed	161
Dosing schedule	405 μg/Kg SC QW	800 µg/Kg SC Q2W	5-1600 μg/Kg SC	18-10000µg/Kg IV Q2-3W	1200-7200 µg/Kg IV Q2-3W	20-198 mg IV Q3W
Prior LoT	5	5	6	5	4	6
TCR/Penta-ref (%)	74/29	69/23	84/41	62/36	72/42	85/68
ORR/≥CR (%) ORR prior BCMA (%)	74.1/33.6	71.7/38.7	64.7/35.3 75% prior CAR-T 44.4% prior BsAbs	71/35 50	64/25 55	56.7/8.9
PFS DoR OS	7.5 79% at 12m (≥CR) 76% at 12m	14.2 90% at 12m (≥CR) 76% at 12m	5.1 63% at 12m (≥CR) 80% at 12m	NR	NR	NR 11.5 m NR
Toxicity	CRS overall is reported in 74-80% of patients and no more tan 3% G3-4 Infections overall are reported in 46-73% of patients and nearly 20% G3-4 Dysgeusia is reported in 70-75% of patients Skin/Nails toxicity reported in 56-70% of the patients					

Schinke et al- ASCO 2023; Chari et al. NEJM 2023; Carlo-Stella et al. ASH 2022; Trudel et a ASH2021; Harrison et al . IMS 2023

Talquetamab + Daratumumab: TriMM-2 Ph 1b study



Talquetamab + Pomalidomide: MonumenTAL-2 Ph 1b study





ORRs were consistent across patient subgroups

- 100% (3/3) in CAR-T–exposed pts in the QW cohort (no patients had CAR-T exposure in Q2W)
- 100% (5/5 in QW, 3/3 in Q2W) in poma-exposed pts in both cohorts
- 50% (1/2 in QW) and 67% (2/3 in Q2W) in pts with EMD
- 80% (4/5 in QW) and 75% (3/4 in Q2W) in pts with HRCg

35 pts, 3 median prior LOT

- 31% (0.4) / 21% (0.8) TCR
- 25% (0.4) / 0% (0.8) prior anti-BCMA TCRed tp
- 31% (0.4) / 16% (0.8) poma exposed

TEAE (≥20%), n (%)	All patients (N=35)		
	Any Grade	Grade 3/4	
Hematologic TEAEs			
Neutropenia	22 (62.9)	19 (54.3)	
Anemia	13 (37.1)	9 (25.7)	
Thrombocytopenia	10 (28.6)	7 (20.0)	
Thrombocytopenia	10 (28.6)	7 (20.0)	

TEAE (≥5%), n (%)	All patients (N=35)		
	Any Grade	Grade 3/4	
Infections	28 (80.0)	8 (22.9)	
Pneumonia	8 (22.9)	5 (14.3)	
Upper respiratory tract infection	8 (22.9)	1 (2.9)	
COVID-19	6 (17.1)	1 (2.9)	
Oral candidiasis	3 (8.6)	0	
Urinary tract infection	3 (8.6)	1 (2.9)	
Influenza	2 (5.7)	0	
Respiratory syncytial virus infection	2 (5.7)	1 (2.9)	
Rhinovirus infection	2 (5.7)	0	
Sinusitis	2 (5.7)	0	

TEAE (≥25%), n (%)	All patients (N=35)			
	Any Grade	Grade 3/4		
Nonhematologic TEAEs				
Taste related ^a	30 (85.7)	N/A		
Infections	28 (80.0)	8 (22.9)		
CRS	26 (74.3)	1 (2.9)		
Skin related ^b	26 (74.3)	2 (5.7)		
Nail related ^c	24 (68.6)	0		
Dry mouth	19 (54.3)	0		
Fatigue	19 (54.3)	5 (14.3)		
Pyrexia	13 (37.1)	1 (2.9)		
Nausea	12 (34.3)	0		
Diarrhea	10 (28.6)	0		
Hypokalemia	10 (28.6)	2 (5.7)		
Back pain	9 (25.7)	1 (2.9)		
Headache	9 (25.7)	1 (2.9)		

Matous J et al. ASH 2023

Teclistamab and Talquetamab: First Combination of Bispecific Antibodies to Target 2 Distinct Myeloma Antigens – RedirecTT-1 phase 1b study









October 25, 2021





Extramedullary disease

	All dose levels (N=35)	Tec 3.0 mg/kg + tal 0.8 mg/kg Q2W (N=11)
Median DOR, ^f months (95% Cl)	12.9 (4.17–NE)	NE (4.17–NE)
Median PFS, ^g months (95% Cl)	6.1 (2.5–9.9)	<mark>9.9</mark> (2.4–NE)

January 2022

■ PR ■ VGPR ■ CR ■ sCR

Cohen et al. ASCO 2023

JNJ-79635322 Is a Potential First-in-Class Trispecific Antibody Targeting BCMA, GPRC5D, and CD3

- Dual antigen targeting may enhance tumor response by circumventing tumor heterogeneity and antigen loss and improving potency due to antigen binding avidity
- JNJ-79635322 (JNJ-5322) is an IgG1 trispecific antibody that binds to CD3 on T cells and BCMA and GPRC5D on MM cells



- In vitro, JNJ-5322 induced potent and dose-dependent cytotoxicity with concomitant T-cell activation only in myeloma cell lines that expressed one or both target proteins (BCMA, GPRC5D)
- JNJ-5322 also induced CD138⁺ plasma cell depletion when tested using patient-derived myeloma bone marrow mononuclear cells in a co-culture assay
- In vivo, JNJ-5322 induced potent antitumor activity in models that expressed one or both target proteins

A phase 1 dose-escalating study of JNJ-5322 in patients with RRMM is ongoing (NCT05652335)

Combination strategies of Non-BCMA BsAbs in Ph 1-2 clinical trials

Study	Agents	
Combination with other anti-myeloma agents		
NCT04108195_TriMM-2 (Phase 1)	Teclistamab + daratumumab; Talquetamab + daratumumab; then \pm pomalidomide	
NCT05050097_MonumenTAL-2 (Phase 1b)	Talquetamab + carfilzomib; Talquetamab + carfilzomib + daratumumab; Talquetamab + lenalidomide; Talquetamab + lenalidomide + daratumumab; Talquetamab + pomalidomide	
NCT05338775_TriMM-3 (Phase 1)	Teclistamab + PD-1 inhibitor; Talquetamab + PD-1 inhibitor	
NCT06055075 (Phase 1)	Forimtamig + carfilzomib; Forimtamig + daratumumab	
Combination of 2 bispecific molecules targeting various MM antigens		
NCT04586426 (Phase 1, Part 2: Dose expansion cohort)	Talquetamab + Teclistamab; Talquetamab + Teclistamab + daratumumab	
NCT05927571 (Phase 1)	Cevostamab + Elranatamab	

Ongoing phase 3 studies with Non-BCMA BsAbs as treatment of RRMM at early relapse or NDMM

Study	Regimen	Condition
MonumenTAL-3 NCT05455320	Tal-Dara vs Tal-Dara-Poma vs DaraPd	RRMM ≥ 1 prior LOT (including len and PI)
MajesTEC-7 NCT05552222	Tec-Dara-Len vs DaraRd Tal-Dara-Len vs DaraRd	NDMM ineligible or not intended for ASCT

Article

nature medicine

Article

https://doi.org/10.1038/s41591-023-02491-5

Mechanisms of antigen escape from BCMAor GPRC5D-targeted immunotherapies in multiple myeloma

- Genomic events on the *TNFRSF17* locus were identified in 8 out of 16 patients (50%) who progressed after prior anti-BCMA/GPRC5D CAR-T and/or TCE therapies.
- BCMA extracellular domain mutations leading to functional loss of epitopes were demonstrated at the time of PD in 5 out of 14 patients treated with BCMA-targeting TCE. These events were possibly driven by the longitudinal selective therapeutic pressure exerted by TCE in comparison to the transient immune selection post-CAR T.
- **Biallelic mutations of the** *GPRC5D* **locus after GPRC5D-targeting TCE** were found in 4 patients, suggesting that under the immune-therapeutic pressure GPRC5D can be more readily lost than *TNFRSF17* in MM cells.

Cancer Cell

The pre-existing T cell landscape determines the response to bispecific T cell engagers in multiple myeloma patients

- Single-cell TCR tracing identifies conserved T cell responses to TCEs in humans
- Clonal expansion of effector CD8+ T cells is an immunological driver of TCE therapy
- Naive T cells require additional MHC class I signal and differentiate upon TCE activation
- The abundance of exhausted CD8+ clones predicts response failure in multiple myeloma



Friedrich MJ al, Cancer Cell 2023

Lee H et al, Nature Medicine 2023

Thanks!

Seràgnoli Institute of Hematology

Michele Cavo



Myeloma Research Unit

Clinical Research Unit

Elena Zamagni Paola Tacchetti Lucia Pantani Katia Mancuso Ilaria Rizzello Chiara Sartor Miriam Iezza Marco Talarico Flavia Bigi Michele Puppi Ilaria Sacchetti Enrica Manzato Simone Masci Roberta Restuccia

Lab of Cytogenetics Nicoletta Testoni Giulia Marzocchi

Lab of Molecular Biology

Carolina Terragna Marina Martello Enrica Borsi Vincenza Solli Andrea Poletti Silvia Armuzzi Gaia Mazzocchetti Ilaria Vigliotta Barbara Taurisano Ignazia Pistis

Data Management

Simona Barbato Giorgia Lazzarini Francesca Trombetta Alessandra Scatà Nicola Francesco Parisi Nicola Paprusso Federica Di Camillo