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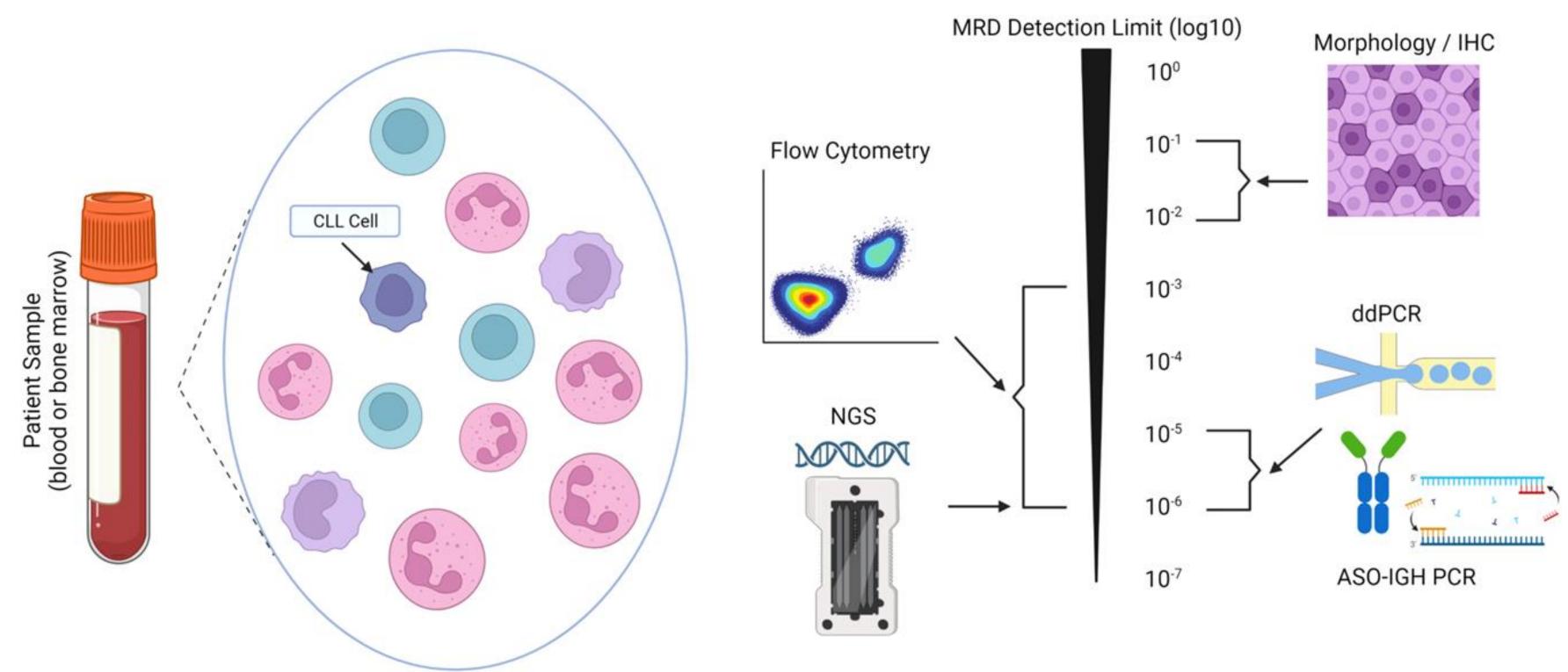


MRD in CLL

- Il Raggiungimento di uno stato di MRD negativa (uMRD) si è dimostrato un potente fattore prognostico, correlato a più lunga progressione libera da malattia e sopravvivenza globale.
- Negli ultimi anni le tecniche di NGS si sono affermate come valido strumento per la valutazione della MRD.



MRD in CLL: tecnologie utilizzate



This image was created with Biorender

Joanna M. Rhodes, Carlos A. Lopez, Jacqueline C. Barrientos; MRD-directed therapy in CLL: ready for prime time?. *Hematology Am Soc Hematol Educ Program* 2023; 2023 (1): 413–420.





Metodologie a confronto

Methodology	Strengths	Weaknesses		
Multiparameter Flow Cytometry	Fast; Absolute Quantification; Information at a cellular level; Wide availability	Variable antigen expression could lead to false negative results; High grade of expertise needed; Medium sensitivity with less than 8-colours		
Allele-Specific Oligonucleotide PCR	High sensitivity	Time-consuming in the design of patient-specific primers; Requirement for optimal DNA quality and quantity		
Digital PCR	Absolute quantification; High sensitivity; Avoids PCR inhibitors due to compartmentalization of target sequences	Lack of standardization No possibility to find new variants Allele-specific design		
Next-Generation Sequencing or High-Throughput Sequencing	High Sensitivity (>10 ⁻⁶); Patient-specific primers not necessary; Versatility	Lack of standardization; High degree of bioinformatics expertise; Expensive		

Sánchez R, Ayala R, Martínez-López J. MinimalResidualDiseaseMonitoring with Next-Generation Sequencing Methodologies in HematologicalMalignancies. Int J Mol Sci. 2019 Jun 10;20(11):2832.

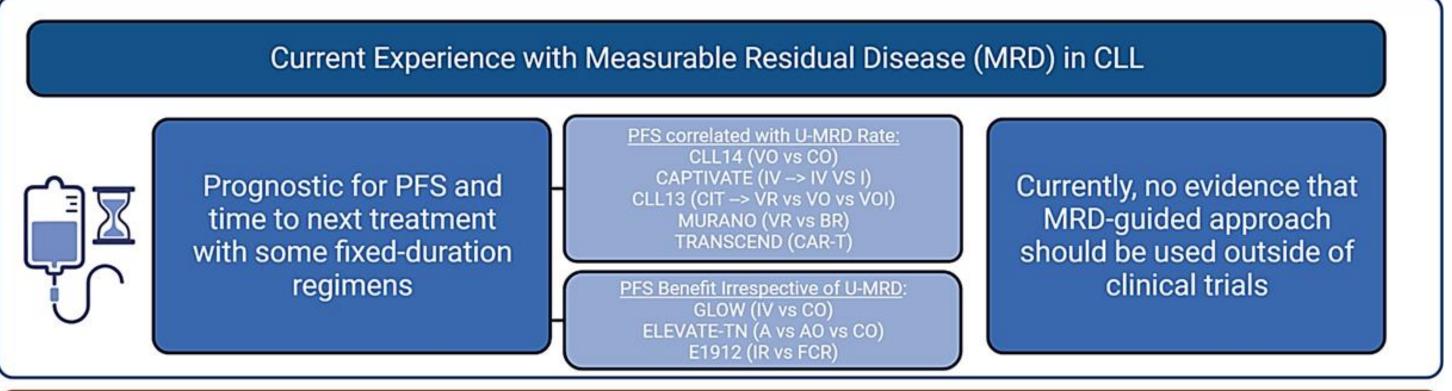


MRD-directed therapy in CLL: stato dell'arte

- Studi condotti durante trattamento con BTKis non hanno messo in evidenza PFS e OS correlata allo stato MRD
- Successivamente, nel trattamento con BCL-2 inibitori (+obinotuzumab/rituximab), la valutazione dello stato MRD si è reso fondamentale nella pratica clinica.
- Studi come CLL14 hanno dimostrato una percentuale significativa di pazienti che ottengono uMRD dopo trattamento.
- Gruppo di lavoro CLL-AVEN su caratteristiche biologiche e outcome in pazienti affetti da CLL trattati con target agents (valutazione MRD in NGS at EOCT, EOT, Follow-up ogni 6 mesi)



MRD-directed therapy in CLL: stato dell'arte



Key Questions to be Answered for "Prime Time" use of MRD in CLL Timing for testing? Factors Confounding "Ideal" U-MRD Target Ideal compartment to test? Disease with high-risk genomics (del17p, Method for Testing? IGHVu) may achieve initial U-MRD but still relapse, affecting prognostication Aggressive Genetics (Relapse) Testing too early or with too insensitive a Which MRD endpoints to method might underestimate utility of target to guide treatment 'Early" U-MRD U-MRD to predict PFS duration? **U-MRD** "Late" U-MRD Some continuous regimens (e.g. FLAIR) can take longer to achieve UMRD Which patients benefit most Time after Treatment from MRD-adapted treatment? This image was created with Biorender

Joanna M. Rhodes, Carlos A. Lopez, Jacqueline C. Barrientos; MRD-directed therapy in CLL: ready for prime time?. *Hematology Am Soc Hematol Educ Program* 2023; 2023 (1): 413–420.

MRD: Measurable Disease; CLL: Chronic Lymphocytic Leukemia; PFS: Progression-Free Survival; VO: venetoclax + obinutuzumab; CO: chlorambucil + obinutuzumab; IV: Ibrutinib + Venetoclax; I: Ibrutinib; CIT: Chemoimmunotherapy; VR: Venetoclax + Rituximab; VOI: venetoclax + obinutuzumab + ibrutinib; BR: Bendamustine + rituximab; CAR-T: Chimeric antigen receptor T-cell Therapy; CR: Complete Remission; U-MRD: undetectable MRD; A: acalabrutinib; AO: acalabrutinib + obinutuzumab; IGHVu: IGHV unmutated



IMMUNOTERAPIA E FARMACI BIOLOGICI NEL TRATTAMENTO DEI LINFOMI E LLC: ATTUALITÀ E FUTURO



National Comprehensive NCCN Guidelines Version 3.2025 NCCN Cancer Network®

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Chronic Lymphocytic Leukemia/Small Lymphocytic Lymphoma

Table 1: Undetectable MRD Rates for BCL2i-Containing Regimens

Disease Setting	Trial	Regimen	No. of Patients	Patient Characteristics	Median Follow-up	Undetectable MRD (S10 ⁻⁴ , uMRD4)	Method used for MRD detection	
	CLL14 ²⁸	Venetoclax + obinutuzumab (VenO)	216	≥65 years		EOT+2: 75% (blood)	ASO-PCR;	
	(Phase III)	Chlorambucil + obinutuzumab	216	(CIRS >6; CrCl <70 mL/min)	65 months	EOT+2: 33% (blood)	MRD-flow; NGS	
	CAPTIVATE ¹² (Phase II; Time-limited cohort)	Venetoclax + ibrutinib	159	≤70 years; ECOG PS 0–1	28 months	EOT+3: 77% (blood); 60% (BM)		
	CAPTIVATE ¹² (Phase II; MRD cohort)	Venetoclax + ibrutinib (3 cycles of lead-in ibrutinib followed by 12 cycles of ibrutinib + venetoclax)	164	≤70 years; ECOG PS 0–1 (Prerandomization)		75% (blood); 68% (BM)	MRD-flow (8-color flow cytometry)	
		Venetoclax + ibrutinib	32	≤70 years; ECOG PS 0–1	31 months	69% (blood); 66% (BM)		
Previously		Ibrutinib	31	(Randomization; uMRD not confirmed)		45% (blood); 42% (BM)		
Untreated CLL/SLL	GLOW (Phase III) ¹⁷	Venetoclax + ibrutinib	106	≥65 years or <65 years	34 months	EOT+3: 55% (blood); 52% (BM)	NGS	
CLUGEE		Chlorambucil + obinutuzumab	105	who also had CIRS >6 or CrCl <70 mL/min		EOT+3: 39% (blood); 17% (BM)		
	GAIA/CLL13 ²⁸ (Phase III)	Venetoclax + ibrutinib + obinutuzumab	231		39 months	15 months: 92% (blood); 78% (BM)	MRD-flow (4-color flow cytometry)	
		VenO	229	≤65 years or >65 years		15 months: 87% (blood); 73% (BM)		
		Venetoclax + rituximab (VenR)	237	[without del(17p) or TP53 mutation]		15 months: 57% (blood); 43% (BM)		
		Chemoimmunotherapy (FCR ≤65 years; BR >65 years)	229	,		15 months: 52% (blood); 37% (BM)		
	AMPLIFY ²⁸	Venetoclax + acalabrutinib	Median age: 61 ve			EOT: 45% (blood); EOT+3: 38% (blood)		
		VenO + acalabrutinib	286	[without del(17p) or	41 months	EOT: 95% (blood); EOT+3: 94% (blood)	NGS; MRD-flow	
		FCR or Bendamustine + rituximab (BR)	290	TP53 mutation])		EOT: 73% (blood); EOT+3: 78% (blood)		
Relapsed or Refractory CLL/SLL		VenR	194	≥18 years; ECOG PS 0–1:	36 months	62% (blood)	ASO-PCR and/or MRD-flow (4-color flow cytometry)	
	MURANO (Phase III) ⁸²	BR	195	adequate bone marrow, liver, and kidney function		13% (blood)		
	CLARITY ⁶⁵ (Phase II)	Ibrutinib + venetoclax	53	Median age: 64 years ECOG PS 0-2	21 months	53% (blood); 36% (BM)	MRD-flow	



MRD-directed therapy in CLL: ruolo dell' NGS

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Discussion

RESPONSE DEFINITIONS AFTER TREATMENT FOR CLL/SLL^a

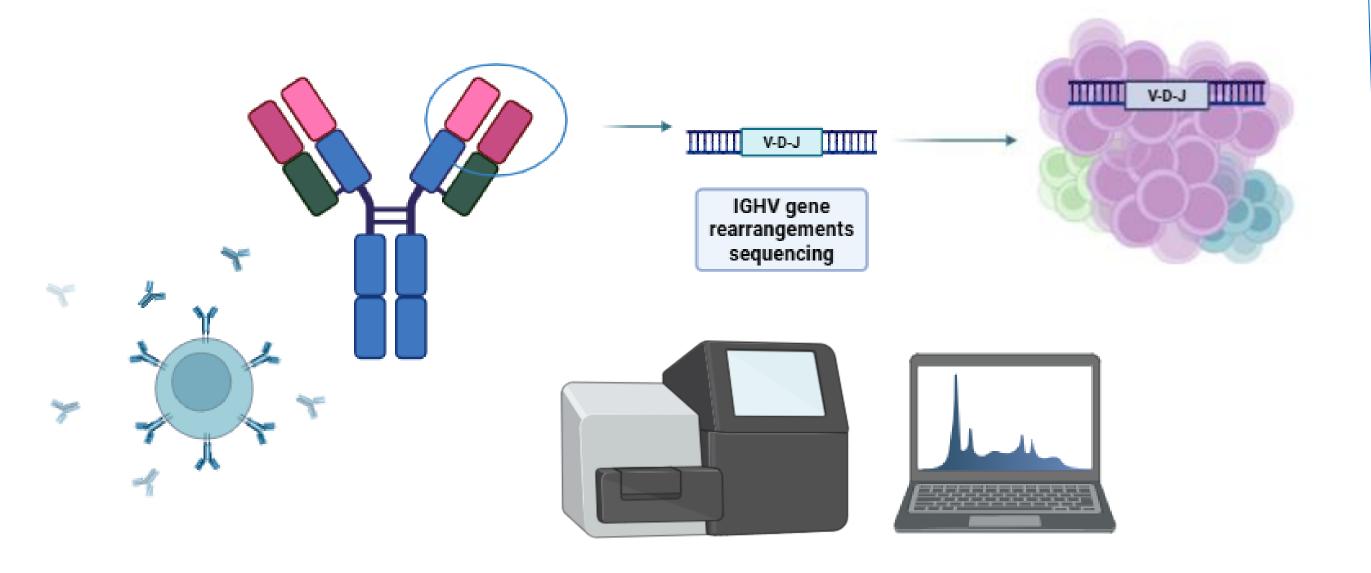
Minimal Residual Disease (MRD) Assessment:

- Evidence from clinical trials suggests that undetectable MRD in the peripheral blood after the end of time-limited treatment is an important predictor of efficacy. e,f,g,h,i
- Allele-specific oligonucleotide polymerase chain reaction (ASO-PCR) and six-color flow cytometry (MRD flow) are the two validated methods used for the detection of MRD at the level of 10⁻⁴ to 10⁻⁵. Next-generation sequencing (NGS)-based assays have been shown to be more sensitive, thus allowing for the detection of MRD at the level of 10⁻⁶. I,m,n
- MRD evaluation should be performed using an assay with a sensitivity of 10⁴ according to the standardized European Research Initiative on CLL (ERIC)
 method or standardized NGS method.
 - MRD su campioni BM o PB
 - L'analisi NGS richiede un campione pre-trattamento
 - La valutazione della MRD può essere utile nella pratica clinica per fornire informazioni dettagliate sulla PFS dopo il trattamento, ma non è ad oggi raccomandata nelle decisioni terapeutiche.

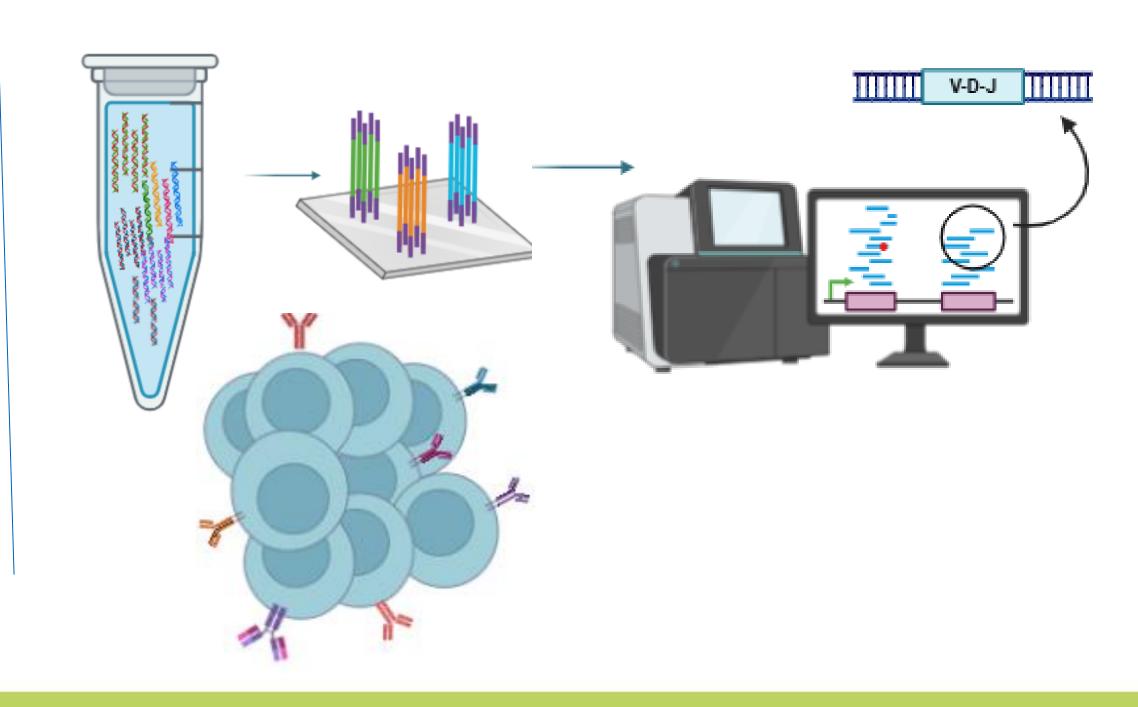


MRD-directed therapy in CLL: workflow NGS

Esordio:



Rivalutazione MRD:



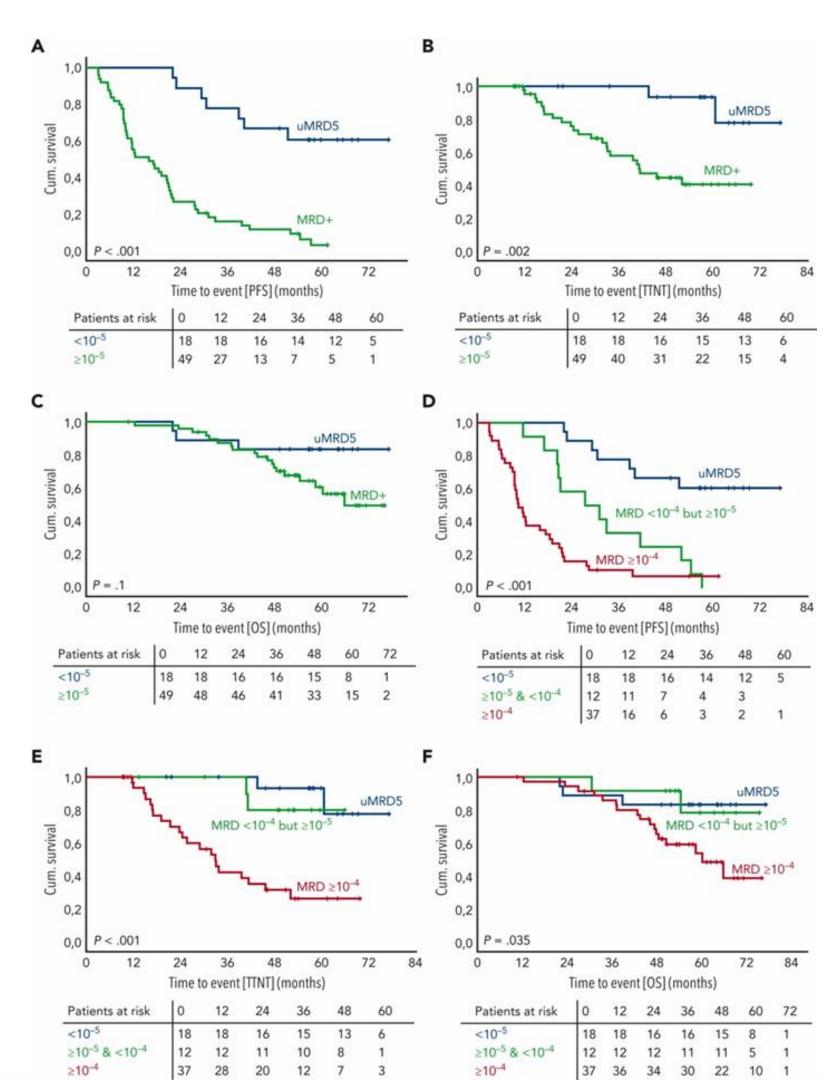


Vantaggi della tecnologia NGS:

sensibilità

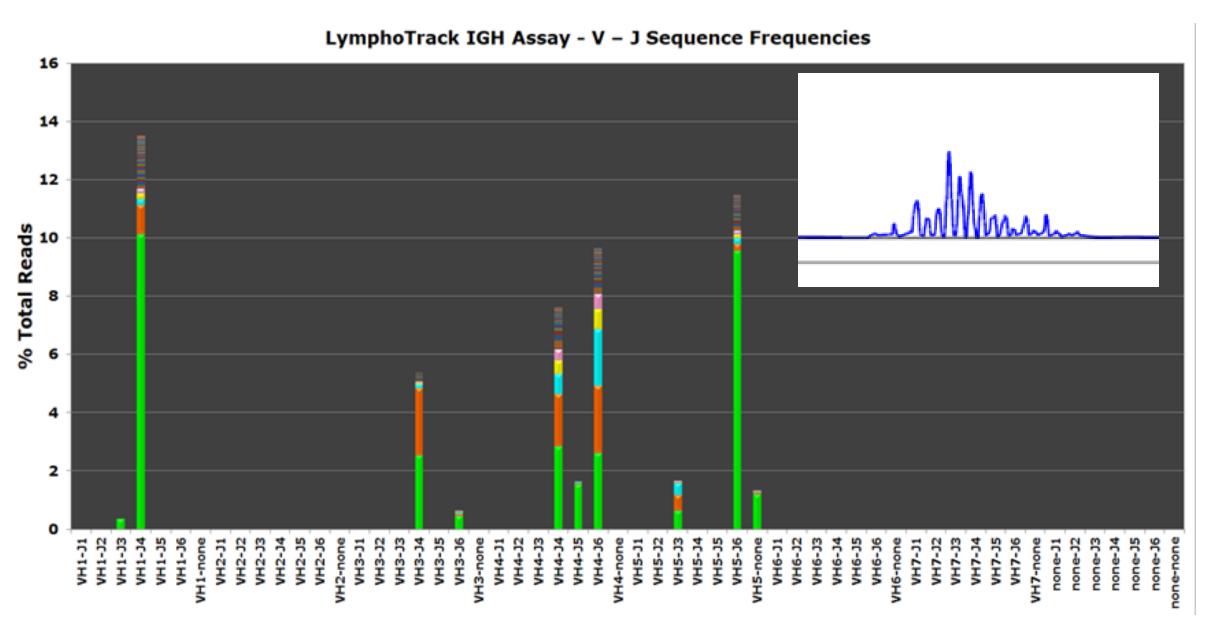
- uMRD <10⁻⁵ è associata a maggiore PFS e intervallo al successivo trattamento.
- 10⁻⁴>MRD≥10⁻⁵ presentano PFS inferiore rispetto a uMRD<10⁻⁵, ma superiore al gruppo con MRD>10⁻⁴.
- I risultati supportano l'utilità clinica del test NGS IGHV leader-based.

Paul J. Hengeveld et al.; Detecting measurable residual disease beyond 10⁻⁴ by an IGHV leader-based NGS approach improves prognostic stratification in CLL. *Blood* 2023; 141 (5): 519–528





Vantaggi della tecnologia NGS: caratterizzazione cloni multipli



Total cour	201 387									
Rank -	Sequence -	Length	Merge coulx	V-gene 💌	J-gene	▼ % total read ▼	Cumulative 0	Mutation rate to partial V-gene (%	In-frame (Y/N	No Stop codon (Y/I▼
1	GATCCTCTTTTTGC	478	14030	IGHV1-3_01	IGHJ4_02	13.51	13.51	0.68	Υ	Y
2	CTCGCCCTCCTCC	472	12780	IGHV5-51_01	IGHJ6_02	12.30	25.81	0.00	Y	Y
3	TCCTGGTGGCAG	461	4461	IGHV4-4_02	IGHJ4_02	4.29	30.10	7.77	Y	Υ
4	TCCTGCTGGTGG(471	3912	IGHV4-39_01	IGHJ6_02	3.76	33.87	0.67	Υ	Y
5	TCCTGCTGGTGG(468	3601	IGHV4-39_01	IGHJ4_02	3.47	37.34	2.68	Υ	Υ
6	TCCTCCTGGTGGC	455	3572	IGHV4-4_02	IGHJ6_02	3.44	40.77	1.01	Υ	Y
7	GGTTTTCCTTGTTC	475	3114	IGHV3-74_01	IGHJ4_02	2.99	43.77	2.03	Υ	Υ
8	GGTTTTCCTCGTT	482	2875	IGHV3-33_01	IGHJ4_02	2.76	46.53	1.35	Υ	Y
9	TCCTCCTGGTGGC	450	2360	IGHV4-34_01	IGHJ6_02	2.27	48.80	0.00	Υ	Y
10	TCCTCCTGGTGGC	448	1771	IGHV4-34_01	IGHJ5_02	1.70	50.51	0.00	N	N



Vantaggi della tecnologia NGS: analisi dei subcloni

- LLC è considerata come patologia monoclonale, meno del 5% dei pazienti presenta riarrangiamenti multipli (Plevova et. al,2014)
- Grazie alla tecnologia NGS si possono identificare più facilmente diverse sequenze clonali all'interno del campione di esordio.
- Attualmente manca una standardizzazione in merito al follow-up MRD dei campioni con riarrangiamenti clonali multipli.



Controlli di qualità



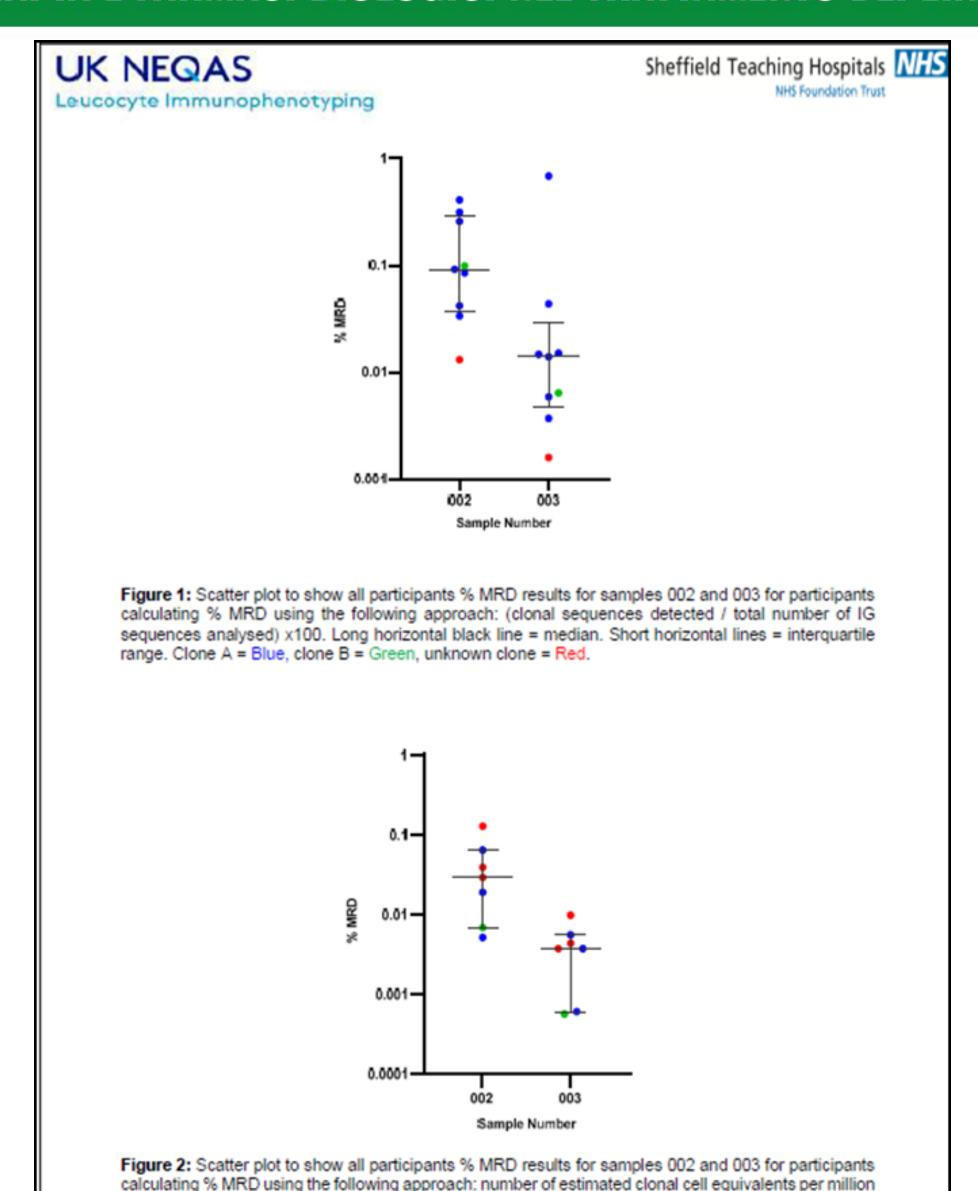


Measurable Residual Disease for Lymphoid Neoplasms by Molecular Methods (Not Accredited)

- Disponibile controllo qualità UK NEQAS dal 2023 (Not Accredited)
- Composto da 2 campioni MRD rilevabile (soglie decrescenti) e 1 campione MRD non rilevabile.
- Nell' anno 2024 è stata evidenziata la necessità di standardizzare il LoD e la soglia di positività MRD. Raccomandano il raggiungimento di sensibilità 10⁻⁴ – 10⁻⁵ per i saggi in NGS.

IMMUNOTERAPIA E FARMACI BIOLOGICI NEL TRATTAMENTO DEI LINFOMI E LLC: ATTUALITÀ E FUTURO





total nucleated cells detected x100. Long horizontal black line = median. Short horizontal lines =

interquartile range. Clone A = Blue, clone B = Green, unknown clone = Red.

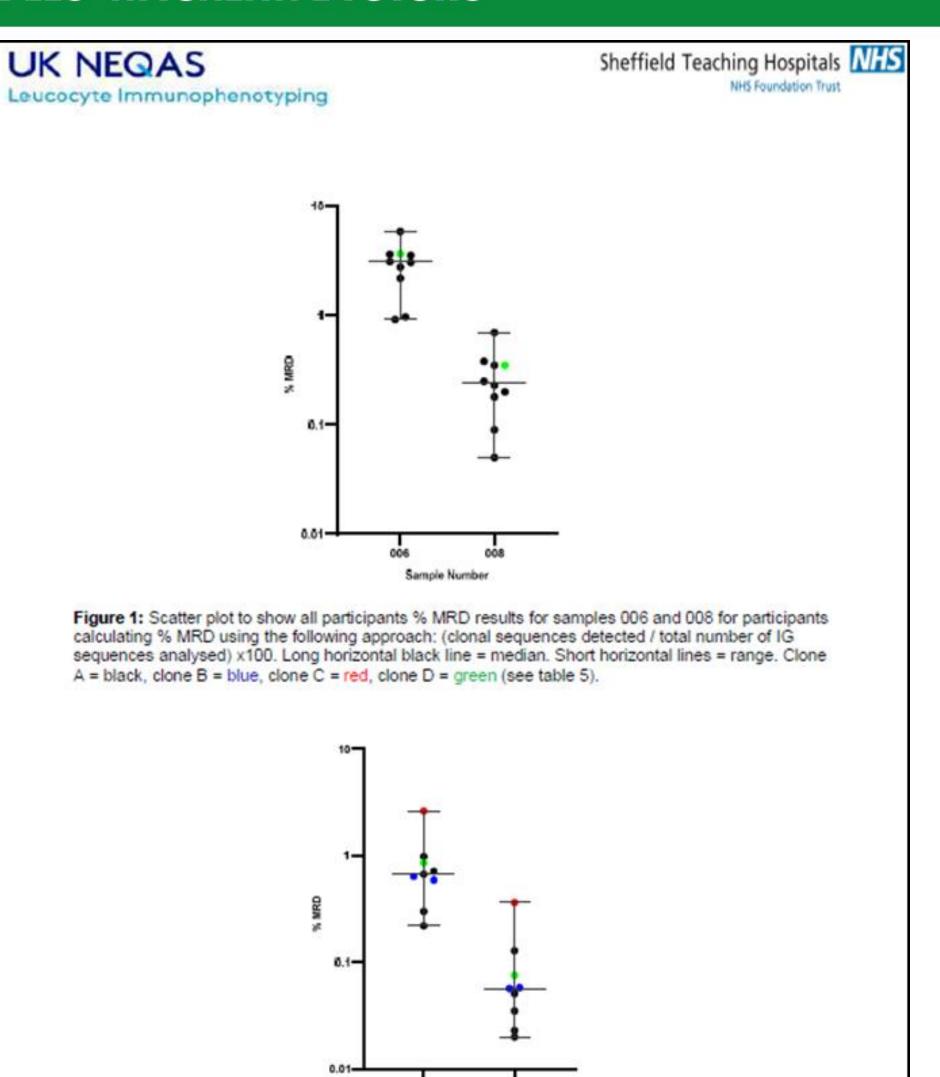


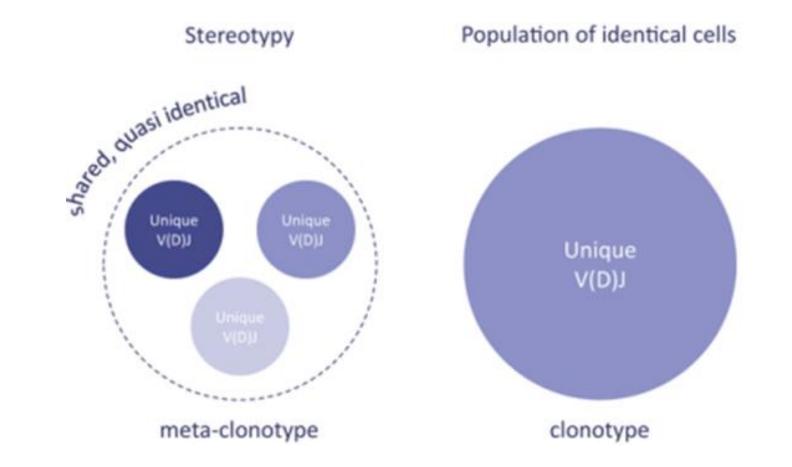
Figure 2: Scatter plot to show all participants % MRD results for samples 006 and 008 for participants calculating % MRD using the following approach: number of estimated clonal cell equivalents per million total nucleated cells detected x100. Long horizontal black line = median. Short horizontal lines = range. Clone A = black, clone B = blue, clone C = red, clone D = green (see table 5).

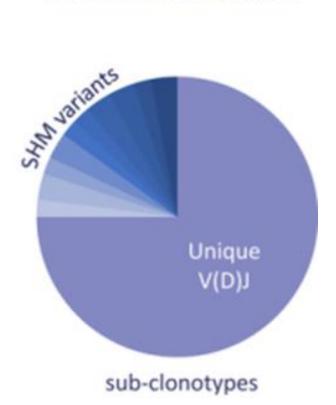


Prospettive future della tecnologia NGS

Diversità intraclonale

- L'accumulo di SHM dovuto alla continua stimolazione antigenica porta alla diversificazione delle sequenze BcR rilevabili nel repertorio immunitario sequenziato.
- Lo studio del repertorio può dare informazioni utili in merito all'evoluzione della malattia e alle stimolazioni antigeniche in corso.
- Attualmente determinabile tramite programmi bioinformatici di creazione alberi filogenetici delle famiglie dei geni IgH (ClonalTREE, GCTree, GLaMST, IgTree, Mtree et al.)





Intraclonal diversification

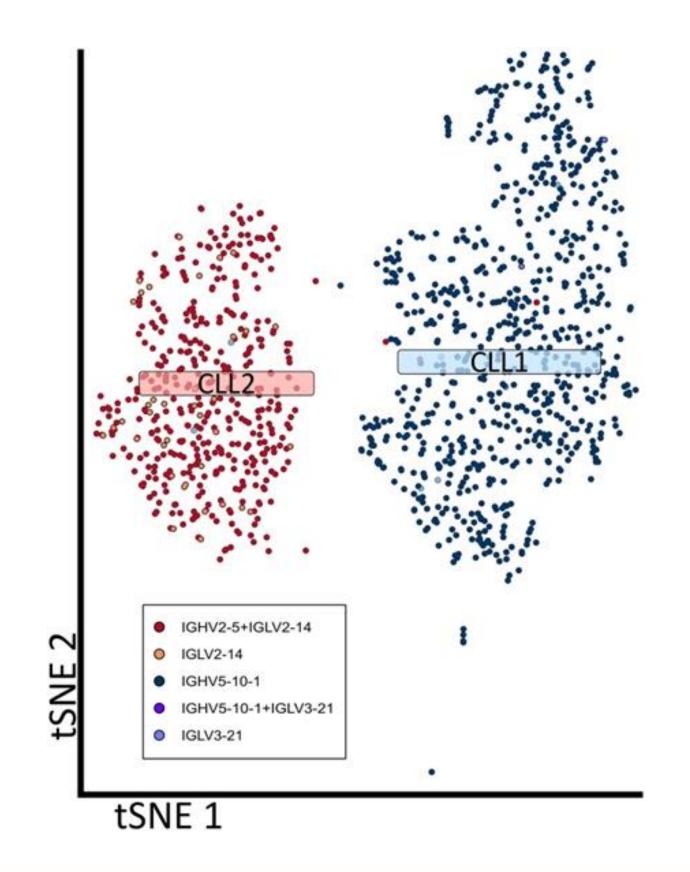
Clonotype definitions for immunogenetic studies: proposals from the EuroClonality NGS Working Group June 2023
Leukemia 37(8):1-3



Prospettive future della tecnologia NGS

Single-cell analysis

- Permette di identificare l'evoluzione e l'etrogeneità clonale ed eventuali nuovi target terapeutici.
- Nella CLL permetterebbe una migliore distinzione tra pool di cellule biclonali o bialleliche (Dampmann et al.,2024)



"T-distributed stochastic neighbor embedding (tSNE) displaying IG gene heavy (HV) and light chain (LV) usage of the CLL cell populations as determined by single-cell RNA analysis. Cells using IGHV5/IGLV3-21 were assigned to cluster CLL1, cells using IGHV2/IGLV2-14 to cluster CLL2. Cells are colored in lighter shades (orange, lilac, light blue) if only one heavy or one light chain was found in these cells".

Dampmann, M., Kibler, A., von Tresckow, J., Reinhardt, H. C., Küppers, R., & Budeus, B. (2024). Single-cell analysis of a bi-clonal chronic lymphocytic leukemia reveals two clones with distinct gene expression pattern. *Leukemia & Lymphoma*, 66(4), 744–752.



Conclusioni

- La valutazione MRD con tecnologia NGS rappresenta oggi uno strumento promettente per la valutazione della risposta al trattamento a durata fissa nella LLC.
- Si necessita standardizzazione in merito al LoD e agli endpoint di valutazione della MRD.
- In futuro la metodica si presenta come un valido strumento per espandere le conoscenze in merito alla cinetica del repertorio immunitario nella LLC.



Grazie per l'attenzione