



Perspectives of Clonal Evolution at the Single-Cell Level

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Disclosures

- Freeline: Consultancy
- Novalgen: Patents, Royalties and Consultancy
- Jannsen: Speakers Fees
- AstraZeneca: Speakers Fees
- Beigene: Advisory Board, Travel
- Abbvie: Travel and Honorarium
- Electra: Advisory Board
- Takeda: Travel

Single Cell Analysis in CLL

- Longstanding & Established
- Exponential development of technology & bioinformatics
 - Increasing numbers of cells
 - Increasing numbers of parameters



Single Cell

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Which part of the cell to study?





Nagler & Wu. 2023; Campillo-Marcos et al. 2021; Ren et al. 2018

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Accurate Determination of CLL Phylogeny

• Patient with del(13q), del(17p), MYH1 and TP53 mutations

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Wang et al. 2017

Linking Mutations to Transcriptional State

• Consequence of SF3B1 mutations



Wang et al. 2016

Epigenetic states at a single cell level

Reduced representation bisulfite sequencing



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Gaiti et al. 2019

Epigenetic states at a single cell level

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Gaiti et al. 2019

- Mitochondrial genome 16.6kb
- Increased mutation rate
- Large numbers of mitochondria/cell
- In vivo clone tracking







Ludwig et al. 2019; Lareau et al. 2021





Penter & Gohil et al. 2021

• Multiple resistance mutations targeting same agent in mutually exclusive clones



Thompson et al. 2022

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Thompson et al. 2022



Thijjsen et al. 2022



Thijjsen et al. 2022



Dissecting the Tumour Microenvironment



Sun et al. 2022

Dissecting the Tumour Microenvironment



Sun et al. 2022

Immune dysfunction in CLL







Purroy et al. 2022

Immune dysfunction in CLL



Purroy et al. 2022

Insights into Richter Transformation



Nadeu et al. 2022

Conclusions

- High Dimensional Single Cell Approaches provide unparalleled insights into leukemic and immune cells in CLL
- Ongoing advances will drive further understanding of CLL biology, response and resistance
- Spatial transcriptomics will provide the next tranche of insights