In situ Biomarker analysis in cancer Immunotherapy : development of quantitative multiplex IHC

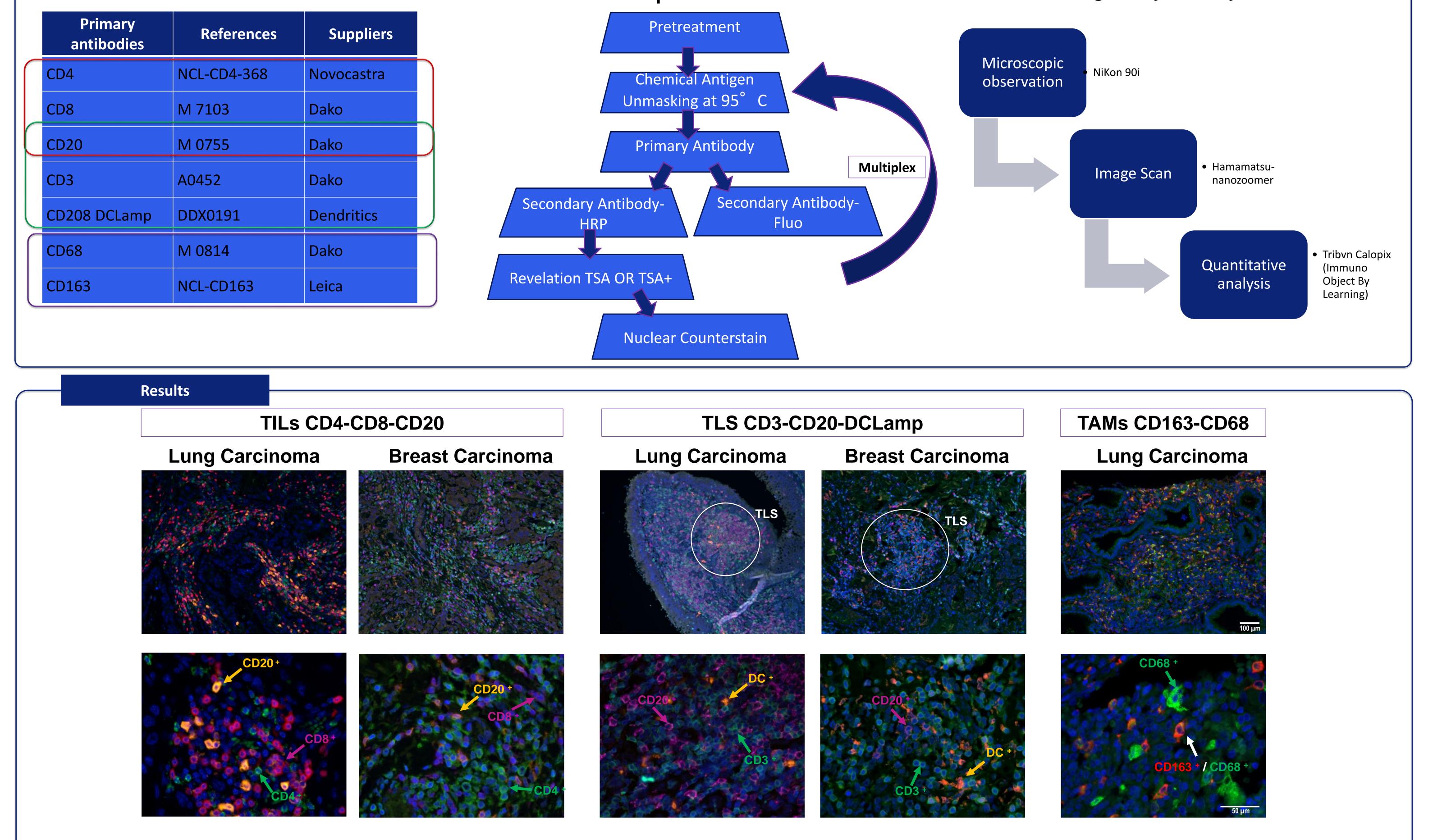
S. Cochin, N. Kehrer, C. Reymann, L. Barraud

Objectives

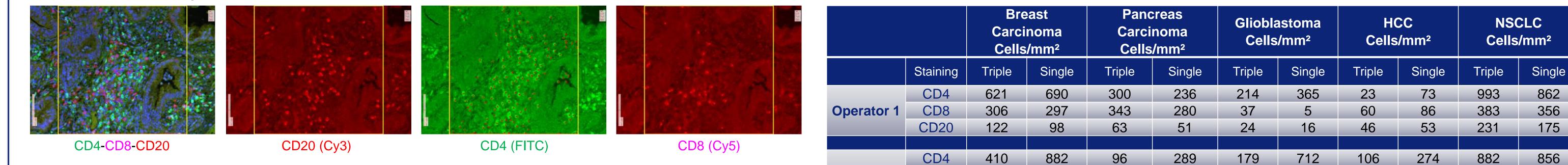
transgene

There is now growing evidence that the immune contexture influences cancer progression and clinical outcome of patients. The tumors microenvironment (TME) is the bed of cancer progression and the target of increasing drugs in development. The objective is to develop and partially validate multiplex IHC panels to analyze the human immune TME. The developed panels enabled to analyze the ymphocyte compartment (TIL), the macrophage status (M1 versus M2) and the presence of Tertiary Lymphoid Structures (TLS) in clinical sample. Multiplex method and quantitative analyses allowed to obtain maximum information about TME in precious clinical sample (biopsies).

	Material and Method		
Single and Multi-stainings were performed using OPAL system from Perkin Elmer on FFPE Human tumor section : Breast carcinoma, Lung carcinoma			
Ass	essment of three panels :	TILs (Tumors Infiltrating Lymphocytes): CD4-CD8-CD20	
		TLS (Tertiary Lymphoid Structure) CD3-CD20-DCLamp	
		TAMs : (Macrophages differentiation) CD163-CD68	
	Used Antiboo	ies IHC process OPAL	Image Analysis assay



Quantitative analysis on TILs



Validation process (ongoing on TIL)

- Objectives
 - Multiplex versus simplex labelling

First observations (raw data) : Variability induced by methods:

468

147

399

39

30

130

24

- Between simplex and multiplex

371

136

380

168

CD8

CD20

Operator 2

- Quantification method via Calopix
 - Repeatability : repeat process by one operator
 - Reproducibility : repeat process by two operators

- ROA definition : tumor versus parenchyma/necrosis...
- Tumor type: difficult for HCC, easier for NSCLC ...

Conclusion and Next Steps

Development of new *in situ* biomarkers is essential to understand the influence of TME on tumor progression for immunotherapy. The assessment of multiplex IHC panels allows the immune "phenotyping" of this TME. In addition, we start to develop a validated process to quantify these biomarkers Next step :

- Complete validation of quantification: increasing the repeatability and the reproducibility data (including a third operator)
- Describe an harmonized quantification process for future analysis
- Other markers :
- tumor architecture via Pan Cytokeratin , CD31, MHC I...
- Innate immunity : NK, Neutrophil, regulatory cells Treg, Marker M1 : iNOS



400 Boulevard Gonthier d'Andernach - Parc d'Innovation - CS80166 67405 Illkirch Graffenstaden Cedex - France



136

306

401

252

69

38

312

148