



Description of the set of the se

de Greffe Pulmonaire (COLT)



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CLAD : mécanismes



Royer PJ et al., transplantation 2016

COLT

Research of early predictive factors of CLAD Towards 4P medicine ?



COLT



M Reynaud-Gaubert

COLT / SysCLAD Story



Dysfonction Chronique du Greffon:

Classification of adjudicated patients after 3 years post transplant



• Nov 2017: 1st adjudications at 5 years: 250 patients « stable » at 3 years

MMP9 Story

Airway epithelial cells exposed to allogeneic T cells produce MMP9 through a CCL2/CCR2 pathway: implications for chronic lung allograft dysfunction.



Airway epithelial cells exposed to allogeneic T cells produce MMP9 through a CCL2/CCR2 pathway: implications for chronic lung allograft dysfunction.



Pain M, Royer PJ, AJT 2016

Airway epithelial cells exposed to allogeneic T cells produce MMP9 through a CCL2/CCR2 pathway: implications for chronic lung allograft dysfunction.



A cut-off value of 139 ng/ml of MMP-9 allowed the prediction of BOS at 6 months with 94% sensitivity and 73% specificity

Pain M, Royer PJ, AJT

Regulatory T cells in transplantation

- Regulation of inflammation : asthma, autoimmune diseases, ...
- Growing interest in studying T regs in organ transplantation : operational tolerance, acute rejection, clinical trials.

The number of FoxP3+ cells in transbronchial lung allograft biopsies does not predict bronchiolitis obliterans syndrome within the first five years after transplantation

Dorrit Krustrup^a, Martin Iversen^b, Torben Martinussen^c, Hans Henrik L. Schultz^b and Claus B. Andersen^a

Dynamics of Human Regulatory T Cells in Lung Lavages of Lung Transplant Recipients

David C. Neujahr,^{1,5} Adriana C. Cardona,² Onome Ulukpo,² Mark Rigby,² Andres Pelaez,¹ Allan Ramirez,¹ Anthony A. Gal,³ Seth D. Force,⁴ E. Clinton Lawrence,¹ Allan D. Kirk,² and Christian P. Larsen²

REGULATORY CD4+CD25+ T CELLS IN THE PERIPHERAL BLOOD OF LUNG TRANSPLANT RECIPIENTS: CORRELATION WITH TRANSPLANT OUTCOME

FEDERICA MELONI,^{1,5} PATRIZIO VITULO,¹ ALESSIA MARONE BIANCO,¹ ENRICA PASCHETTO,¹ MONICA MOROSINI,¹ ALESSANDRO CASCINA,¹ IOLANDA MAZZUCCHELLI,² LAURA CIARDELLI,³ TIBERIO OGGIONNI,¹ ANNA MARIA FIETTA,¹ ERNESTO POZZI,¹ AND MARIO VIGANÒ⁴

Lung function early after lung transplantation is correlated with the frequency of regulatory T cells

Tomoyuki Nakagiri · Gregor Warnecke · Murat Avsar · Stefanie Thissen · Bianca Kruse · Christian Kühn · Petra Ziehme · Ann-Kathrin Knöfel · Nodir Madrahimov · Meinoshin Okumura · Yoshiki Sawa · Jens Gottlieb · André R. Simon · Axel Haverich · Martin Strüber

Decreased Percentage of CD4⁺FoxP3⁺ Cells in Bronchoalveolar Lavage From Lung Transplant Recipients Correlates With Development of Bronchiolitis Obliterans Syndrome

Sangeeta M. Bhorade,^{1,5} Hong Chen,¹ Luciana Molinero,² Chuanhong Liao,³ Edward R. Garrity,¹ Wickii T. Vigneswaran,⁴ Rebecca Shilling,¹ Anne Sperling,¹ Anita Chong,⁴ and Maria-Luisa Alegre²

IMMUNOBIOLOGY AND GENOMICS

T Regulatory Cells in Stable Posttransplant Bronchiolitis Obliterans Syndrome

Emilie Mamessier,¹ Anne-Marie Lorec,² Pascal Thomas,³ Monique Badier,³ Antoine Magnan,^{1,4} and Martine Reynaud-Gaubert¹

• No consensus about the link between Tregs and BOS occurrence.

Regulatory T cells profile



Regulatory T cells profile



0.0

SP

1.1 nTregs FoxP3low CD45RA+

1.3 mTregs FoxP3hi CD45RA-

1.2 nonTregs FoxP3low CD45RA-

and Shimon Sakaguchi1,7,8,*

Conclusion

• Increase of Tregs proportion with a memory phenotype early after TP for patients who will declare a BOS in the five years.



• New potential biomarker of the BOS occurrence, which could help to manage CLAD after lung transplantation in the next decades.

Circulating CD9+ B cells: biomarker of long-term BOS-free survival after Lung Transplantation (94/100)

Carole Brosseau ^{1,2,5,6+}, Maxim Durand ^{1,2,3+}, Eugénie Durand ^{1,2}, Richard Danger ^{1,2}, Jennifer Loy ^{5,6}, Aurore Foureau ^{5,6}, Mélanie Chesneau ^{1,2}, Philippe Lacoste ^{5,6}, Pierre-Joseph Royer ^{5,6}, Adrien Tissot^{1,2,3,5,6}, Antoine Roux⁵, Martine Reynaud-Gaubert⁶, Romain Kessler⁷, Sacha Mussot⁸, Claire Dromer⁹, Olivier Brugière¹⁰, Jean François Mornex¹¹, Romain Guillemain¹², Marcel Dahan¹³, Christiane Knoop¹⁴, Christophe Pison¹⁵, Laurent Nicod¹⁶, Antoine Magnan ^{5,6\$}, Sophie Brouard ^{1,2,4\$} & COLT and SysCLAD consortia*.





Months

Months



Blood Gene Expression Predicts

Bronchiolitis Obliterans Syndrome Appearance

After Lung Transplantation

Richard Danger*, Pierre-Joseph Royer*,

Damien Reboulleau, Eugénie Durand, Jennifer Loy, Adrien Tissot, Philippe Lacoste, Antoine Roux, Martine Reynaud-Gaubert, Carine Gomez, Romain Kessler, Sacha Mussot, Claire Dromer, Olivier Brugière, Jean-François Mornex, Romain Guillemain, Marcel Dahan, Christiane Knoop, Karine Botturi, Christophe Pison, Angela Koutsokera, Laurent P. Nicod,

Sophie Brouard*, Antoine Magnan* and the COLT and SysCLAD Consortia







Blood gene expression associated with BOS

• Identification of differential genes (t-statistic from R limma package; p.value <5% and fold change >1.5)



• Enrichment of down-regulated and immune-related genes in PRED versus STA comparison

Gene ontology analysis:



number of genes within the enriched ontology

Prediction of BOS: independent validation

• Validation of genes associated with BOS appearance







POU2AF1: POU class 2 associating factor 1 *BLK*: B lymphoid tyrosine kinase *TCL1A:* T-cell leukaemia/lymphoma 1A

Prediction of BOS: independent validation



TCL1A: T-cell leukaemia/lymphoma 1A

Blood Gene Expression Predicts Bronchiolitis Obliterans Syndrome Appearance After Lung Transplantation

Conclusion

- Identification of 3 genes as predictive biomarkers of BOS
- Whole blood and qPCR: non-invasive and compatible with clinical settings
- Suggests a role of B cells in BOS mechanisms

patent: EP16306125.2 work submitted



Validation of these 3 genes in a large prospective cohort



Conclusions: Biomarqueurs issus de COLT et SysCLAD

- MMP9
- Treg
- B transitionnels CD9 +
- Pou2AF1, BLK, TCL1A

A valider en pratique clinique

Score composite à construire

SysCLAD consortium

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COLT Consortium

Flow Cytometry Platform



















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