



Annual Reporting for Faculty Supported Research Centres and Networks

All Centres (provisional Centres; McGill Centres), Research groups and Networks that receive funding from the Faculty of Medicine are required to provide two components of reporting:

1. an Annual Report of Activities and Outcomes (see below),
2. a Financial Statement (see attached Excel document).

The reporting period is May 1, 2019 – April 30, 2020.

Deadline: Monday, June 15th 2020

Please send both documents to Faculty of Medicine's Research Office (riac.med@mcgill.ca).

Continued support from the Faculty is contingent on:

1. the receipt of the reporting documents on time,
2. the evaluation of reported activities by the Faculty's Committee for Oversight of Research Units (CORU),
3. the availability of Faculty funds.

Your strong engagement in the Faculty's mission for continued research excellence and financial stewardship is truly appreciated.

Annual Report of Activities and Outcomes

Please respect the page limits, where indicated, or the report will be returned.

(The accepted font is Times New Roman or Calibri regular 11 pts)

1. Name of the Unit: **McGill Cystic Fibrosis Translational Research Centre (CFTRc)**

2. Director's contact information:

Director: **Dr. John Hanrahan**
john.hanrahan@mcgill.ca
514-398-8320

Co-Director (Clinical): **Dr. Larry Lands**
larry.lands@mcgill.ca
514-412-4444

Associate Director: **Dr. David Thomas**
david.thomas@mcgill.ca
514-398-2973

Manager: **Dr. Annick Guyot**
annick.m.guyot@mcgill.ca
514-398-4323 x1

3. If the Unit is a **Senate-approved** McGill Research Centre, indicate date of approval:

October 8th, 2015

4. Mission Statement of the Unit:

The Cystic Fibrosis Translational Research Centre (CFTRc) focuses primarily on finding a cure for cystic fibrosis, and raising awareness in the general public and research communities. The CFTRc provides a platform for basic CF research and the development of therapies targeting the basic defect that underlies cystic fibrosis and other protein trafficking diseases. The centre is composed of a network of dedicated, passionate researchers who work in close relation with other CF organizations such as Cystic Fibrosis Canada.

5. Number of Unit members:

12 regular members and 18 associate members

6. Number of members affiliated with McGill's Faculty of Medicine:

11 regular members and 8 associate members

7. Unit's website:

URL: <http://www.mcgill.infoca/cftrc>

Note: The website needs to feature the following:

- all sources of funding support (including the Faculty of Medicine's logo),
- the List of Members and their institutional affiliation with appropriate links,
- the activities supported by the Unit

- all previous Annual Reports.

8. Summary of past year's **goals and objectives** of the Unit. (limit: ½ page)

- Use bullet points or numbered lists and be as quantitative as possible.
- Indicate unforeseen changes, opportunities or difficulties.

The goal of the CFTRc remains to accelerate the development of a cure for cystic fibrosis. To achieve this goal it serves as a base for scientific networking/collaboration, training, and provides resources needed to carry out mechanistic studies of CF disease and develop new therapeutics. More specifically, the CFTRc provides:

- a stimulating training environment for students and postdocs that is enhanced by collaboration with other centres, international conferences and workshops, and seminars by leading researchers.
- a structure that facilitates liaison with Cystic Fibrosis Canada, the US Cystic Fibrosis Foundation, UK CF Trust, and the European CF Society, and which can promote the awareness of CF through outreach activities and publicity.
- training on, and access to, state-of-the-art core equipment for fundamental CF research, including electrophysiology, large scale (bioreactor) culture of mammalian cells for biochemical studies, spectro-fluorometry, and imaging of protein trafficking in live cells using confocal and TIRF fluorescence microscopy.
- access to Ussing chamber setups (18 channels) for epithelial electrophysiology, an automated patch clamp (QPatch, see publication by Billet et al., 2017 for details), and expertise in assay development, screening and hit validation.
- pulmonary cells isolated from the lungs of CF patients undergoing transplantation and from non-CF donors. The Primary Airway Cell Biobank (PACB; a core facility operating within the CFTRc) isolates and distributes lung cells, tissue and biofluid samples to researchers across Canada and internationally. As a biobank certified by the International Society of Biological and Environmental Repositories (ISBER), the PACB reaches potential users in Canada and internationally. It is promoted by CF Canada (<https://www.cysticfibrosis.ca/our-programs/research/core-facilities/pacb>), CF Foundation (<https://www.cff.org/Research/Researcher-Resources/Tools-and-Resources/CF-Foundation-Biorepository/>), the UBC Office of Biobank Education and Research Biobank Certification Program (<https://biobanking.org/biobanks/view/218>), and the Association of Biomolecular Resource Facilities (ABRF) core marketplace (<https://abrf.org/abrf-core-marketplace>). The PACB is a member of the Canadian Network of Scientific Platforms (CNSP), a pan-Canadian network of professionals working on technical, managerial and administrative levels in scientific platforms (i.e. core facilities and technology resources). The network represents diverse technologies and institutions across the country and its mandate is to raise awareness and increase the value of scientific platforms, educate research personnel, set and promote best practices, enhance communication and cooperation within the research community, and promote interaction with industry. See: cnsprcps.ca The PACB also biobanks cells from donors with other respiratory diseases including asthma, COPD, emphysema, IPF and has seen requests for these increase during the pandemic.

Challenges: The PACB is no longer funded by Cystic Fibrosis Canada. We are currently seeking other sources of support to ensure a stable base budget. For stability we need to cover 1.5 FTE salaries, materials and supplies, and equipment maintenance. At present there is a shortfall of about \$60K, which we try to make up through research contracts with companies. CFC does provide up to \$10,000 to the CHUM to subsidize our tissue procurement, however we do not manage those funds and cannot provide a budget for them. That allocation for lung tissue procurement in 2019 was used up after ~7 months. Fortunately Drs. Brochiero and Ferrero (transplant surgeon) kindly continued to supply the PACB with samples without receiving further CFC funding.

9. **Major achievements** enabled by the support obtained from the Faculty. **(limit: 1 page)**

(see Appendix for suggested metrics)

Note: We care to evaluate how the Unit is doing as a whole greater than the sum of its talents. For this reason, do not list achievements from a single PI member of the Unit. Instead, please report only the achievements from the coordinated efforts of at least two PI members of the Unit.

- Use bullet points
- Insert an appendix for publications, grants, etc. if necessary.
- **CFTRc has leveraged funds** from the Faculty to obtain additional support since its inception. Funds have been raised from Vertex Pharmaceuticals Inc. for seminars and training enrichment. These ended 31 Dec 2019 however we will apply for renewal after the pandemic situation is resolved. CFTRc has also leveraged funds from the Faculty by attracting contract research projects with companies based in Quebec, USA, UK and Germany.
- **CFTRc annual symposium:** co-organized and co-sponsored by CFTRc in collaboration with CF Canada (Quebec) took place on May 9, 2019 at the Hotel Mortagne in Boucherville. It was held on back-to-back days in conjunction with the CFC (Quebec) clinical meeting, which brings together caregivers from across Québec. There were 2 renowned international speakers, one from Hong Kong (Dr. Lap Chee Tsui, OC, leader of the team that cloned the CF gene) and one from the USA (Dr. Ray Frizzell, a pioneer in epithelial physiology who established the mechanism of airway Cl⁻ secretion). Other excellent presentations were given by CFTRc members PIs and trainees from McGill University, RI-MUHC, CR-CHUM, and University of Sherbrooke. The event was attended by about 80 researchers and the Chief Scientific Officer of CF Canada John Wallenburg, and it was sponsored by Vertex Pharmaceuticals, Horizon Pharma and Mylan. The program booklet is attached (**appendix 1A**) along with photos from the event (**appendix 1B**). This CFTRc-organized symposium helped launch the CFC annual fundraising campaign in May.
- **CFTRc seminar series:** cutting edge seminars were organized through the Centre and sponsored by Vertex Pharmaceuticals.
 - *Dr Lap-Chee Tsui, GBM, O.C., O. Ont., Founding president of the Hong Kong Academy of Sciences, President of Victor and William Fung Foundation, was co-sponsored by Vertex and the CFTRc and integrated into the annual CF symposium. Dr. Tsui led the team that identified the gene for cystic fibrosis by positional cloning (chromosome jumping and walking). He provided an overview of the cloning of CFTR on its 30th anniversary, and his perspective on the future of CF research. Poster attached (**appendix 1C**)
 - *Dr Ray Frizzell, Professor, Department of Pediatrics and Cell Biology, Director of Cystic Fibrosis Research Center, University of Pittsburgh School of Medicine, was co-sponsored by Vertex and the CFTRc and integrated into the Physiology Department seminar series. He described a new pathway for stimulation of CFTR biogenesis by the SUMO E3 ligase PIAS4. Poster attached (**appendix 1D**)
 - *Dr. Subash Sad, Professor, Department of Biochemistry, Microbiology and Immunology, Faculty of Medicine, University of Ottawa, was co-sponsored by Vertex and the CFTRc and integrated into the Physiology Department seminar series. His presentation dealt with immunological responses in CF. Poster attached (**appendix 1E**)
 - *Dr. Bob Scholte, Professor, Department of Pediatric Pulmonology and Department of Cell Biology, ERASMUS Medical Center, Rotterdam, The Netherlands, was cosponsored by Vertex and the CFTRc and integrated into the Physiology Department seminar series. His presentation described recent functional studies using intestinal organoids. Poster attached (**appendix 1F**)

- **Journal Club:** This year we have established a journal club in which members of different labs take turns presenting key articles in their research area that may interest the entire group. One goal is to provide trainees with program enrichment by increasing their knowledge of the literature and by enabling them to practice presenting research to others. Such meetings increase communication between members and might lead to new collaborations.
- **Visiting scientists hosted by the Unit:** The CFTRc hosted:
 - *Dr. Andrea Yool, a visiting professor from University of Adelaide, Australia (July 2019)
 - *Dr. Yen Li, a visiting post-doc from University of Saskatchewan (October 2019)
- **Governance:** There have been informal meetings of groups of CFTRc members almost weekly during the past year. The fourth annual CFTRc meeting was held on Thursday, May 9, 2019 in Boucherville. It was attended by 14 members (10 from McGill, 3 from UdeM, 1 from Sherbrooke). The agenda is attached (**appendix 1G**)
- **What/how much additional Funding was generated that was directly attributable to activities of the Centre?**

Activities of the CFTRc were directly responsible for generating:

 - *\$67,316 in revenue by the PACB. These funds were used to support 1 FTE (Carolina Martini) who runs the primary airway cell biobank.
 - *\$6,043 in microscope user fees. These funds were used to support part time microscope maintenance and training support from the ABIF for CFTRc members
 - *\$35,000 from Vertex Pharmaceuticals over 2.5 years to support a CFTRc seminar series. Seminars were integrated into departmental seminar series to increase benefit to the McGill community.
- **Trainee travel Awards:** four travel grants were awarded to trainees (3 graduate students and 1 postdoc from Brown, Brochiero, Moraes and Lukacs labs, respectively), so they could attend conferences in Canada, USA and Germany to present their work (**letters of approval appendix 1H**).
- **Evidence of collaboration within the CFTRc:**
 - Co-supervised students and postdoctoral trainees-**
 - Claire Brown and John Orlowski:**
Alyssa Dai (Honours Anatomy and Cell Biology student and one 396 project).
 - Emmanuelle Brochiero and Ryszard Grygorczyk:**
Ju Jing Tan (PhD Student), director (R. Grygorczyk) and co-supervisor E. Brochiero
 - John Hanrahan and Paul Wiseman:**
Asmahan AbuArish (Research Associate) and Kamila Mustafina (PhD Student)
 - John Hanrahan and Larry Lands:**
Mark Turner (PDF)
 - John Hanrahan and David Thomas:**
Yukiko Sato (PhD student)
 - Mark Tewfik and Don Sheppard:**
Sarah Khalife (MSc student)
 - MarcTewfik and Simon Rousseau:**
Thereza Queiroga (PDF)

Other examples of integration between members of the CFTRc-

Emmanuelle Brochiero has active collaborations with **André Cantin**, **Dao Nguyen** and **Simon Rousseau** and manages the supply of tissues to the PACB. She shares a Team grant (\$30k) from the Respiratory Health Research Network with **Roger Levesque**. She is also Co-PI on a grant application with **Larry Lands** that is currently under review at the Canadian donation and transplantation program (\$30k). **André Cantin** and PhD student **Mégane Lebel** collaborate with **Emmanuelle Brochiero** on the isolation of lung fibroblasts to study invadosome structures as a marker of fibrosis (manuscript in preparation). This project has recently been awarded funding for 2 years by Boehringer-Ingelheim. **David Thomas is co-PI on a CIHR grant with John Hanrahan and is developing an innovative project on acquired cystic fibrosis with Astra Zeneca, an**

international pharmaceutical company, the Max-Planck Lead Discovery Centre, McGill and the CQDM. The scientific programme is well advanced and they have recently been awarded funding under the CQDN SynergiQc programme. Kalle Gehring holds a CFI grant with Gergely Lukacs and they collaborate on the application of HDX-MS to membrane proteins and complexes. **Chris Moraes** collaborates with **Dao Nguyen** on mechano-bactericidal nanopillars (ms under review) with **John Hanrahan** on stiffness tunable air liquid interface epithelial culture substrates (recently published *Lab on a Chip*, 19, 2786 – 2798, 2019)

- **Collaborations with other units:**

- *with other McGill groups: Centre de Recherche en Biologie Structurale (CRBS), Meakins Christie lab, MUHC-Research Institute

- *with groups outside McGill: networking with the FRQ-S Réseau en santé respiratoire, E-rare consortium based at the Leibniz Research Labs for Biotechnology and Artificial Organs (LEBAO) in Hannover, Germany and other E- rare consortium partners in Rotterdam, Genoa, Lisbon, Toronto and Naples

- **Publications: appendix 1I:** list of publications with collaboration between at least 2 center members

- **Communications:**

<http://www.mcgill.ca/cftrc>

<https://www.mcgill.ca/lifesciencescomplex/facilities/cystic-fibrosis-translational-research-centre-platforms>

<https://navigator.innovation.ca/en/facility/mcgill-university/cystic-fibrosis-translational-research-center-cftrc>

10. **New Members** who joined the Unit in the past year and their **institutional affiliation(s)**.

A complete list of members is on the CFTRc web site. Jörg Fritz has begun a collaboration with Samantha Gruenheid on CF and they are planning a joint application to the US CF Foundation. Jörg would like to join the CFTRc as an associate member and this will be voted on at the next CFTRc meeting.

11. Members who have **left the Unit** over the reported year.

No members have left during the past 12 months.

12. State how the current and forecasted activities of your Unit align with the Education or Research mission (Strategic Research Plan) of the Faculty of Medicine and/or other Faculties at McGill
(limit: ½ page):

The activities of the CFTRc are fully consistent with the strategic research plan and mission of the Faculty of the Medicine. The centre aligns with Theme 4 of the McGill Strategic Research Plan to “Advance Biomedical and Health Sciences for Healthy Populations” and supports this vision in several ways. First, it provides a structure that promotes collaboration between members, as indicated by the many joint publications and co-supervised trainees. Second, it provides an international footprint that is recognized in Europe and the USA. For example the PACB has attracted customers from Northern Ireland, England, Japan and several centres in the USA and interaction with groups at the Wyss Institute at Harvard and Biology Dept at Stanford during the past year. The CFTRc enhances the international profile of the group members stimulates innovative research programs. Third, it provides an enhanced training environment for postdocs and students. During the past year it attracted visiting international students from the UK (Amy Walker, UCL) and France (Aurelie Liege, U Poitiers), in large part because the “whole is greater than the sum of its parts”. Finally, the emphasis on translational research is greatly enhanced by the resources and expertise in the CFTRc, in that it has helped members attract funds from industry. As contract work is completed for various companies (ERAD, Verona Pharma, Engelhard), new opportunities emerge (EnGene, Laurent Pharma, Feldan Therapeutics) in large part because CFTRc is known as a translational

research centre. This collaboration with industry also maximizes the benefits from past CFI investments that provided the infrastructure for the CFTRc (approx. \$5M). The centre is of strategic importance not only for CF research but also for research in other areas related to e.g. protein folding and long QT syndrome type 2(Lukacs), alpha1 antitrypsin deficiency (Thomas), COPD (Thomas and Hanrahan), asthma (Martin, Lands) and COVID19 (Hanrahan).

13. Explain why support from the Faculty of Medicine continues to be crucial to the operations of the Unit **(limit: ½ page)**:

Support from the Faculty of Medicine is essential for continued operation of the centre. They are used to pay a part time Administrative manager (Dr. Annick Guyot), a highly successful, seminar series sponsored by Vertex, which enhances the seminar programs of several departments by bringing in national and international speakers. These seminars are training enrichment for postdocs and students and also introduce visiting speakers to the talent at McGill. Annick is the main coordinator of the research day for the annual CF symposium that CFTRc organizes for CF Canada (Quebec). She assembles the scientific organizing committee of CFTRc members (L. Lands, D. Nguyen, J. Hanrahan in 2019) and coordinates the venue and many other logistics with staff at Fibrose Kystique Quebec. She also manages the finances of the centre, equipment repairs, and handles the distribution of travel awards that are distributed based on merit after review by a standing committee of 3 PIs. It is especially important to have an excellent administrator with strong organizational skills in a centre that has members located in many departments (Biochemistry, Microbiology and Immunology, Physiology, Experimental Medicine), faculties (Medicine, Science, Engineering) and institutions (RI-MUHC, U de Montreal, U Sherbrooke, UBC).

14. List action items that the Unit has taken or will consider taking in the next year towards growth and sustainability of its operations **(limit: ½ page)**

- COVID-19 has been a major disruption for our centre in 2020. For safety reasons we will replace the in-person executive board meeting with a virtual meeting and will schedule it in September 2020.
- The 2020 CF Research symposium which was scheduled for Thursday April 30th, 2020 at the Hotel Omni Mont-Royal in Montreal had to be postponed due to the lockdown. It is tentatively rescheduled for Dec 3 2020 as the hotel had space then, however it probably need to be moved online, therefore we have reduced our budget request for this meeting. Two international speakers that were coming to speak on Apr. 30 (Alice Prince, Columbia Univ. Med. Center, NY and Jay Kolls, Director, Center for Translational Research in Infection and Inflammation, Tulane School of Medicine, New Orleans) have agreed to present on Dec. 3 or whenever the symposium can be held, and also agreed to give their talks online if necessary.
- CFTRc journal club will be held on zoom during the fall semester for safety reasons. If this format works well we may keep it indefinitely as it would be more convenient for the CFTRc members situated at the MUHC and other institutions.
- The current manager of the primary airway cell biobank of the CFTRc, Carolina Martini, will leave the centre in Dec 2020 to begin graduate studies at U de Montreal. We have recently recruited a replacement (Jessica de la Torre) who is being trained to take over. In addition to training, the overlap period will be used to carryout contract research for companies and work with CFTRc collaborators. We are requesting some salary for Jessica while Carolina is still at McGill so that she can be trained to run the facility.
- We discussed the imminent retirement of Jacopo Mortola at the CFTRc annual meeting in May 2019 and decided that we will encourage the department and faculty to recruit a respiration physiologist whose research interests would fit with the CFTRc. The most recent cyclical review recommended hiring in this area to build on the strengths that already exist in the department.

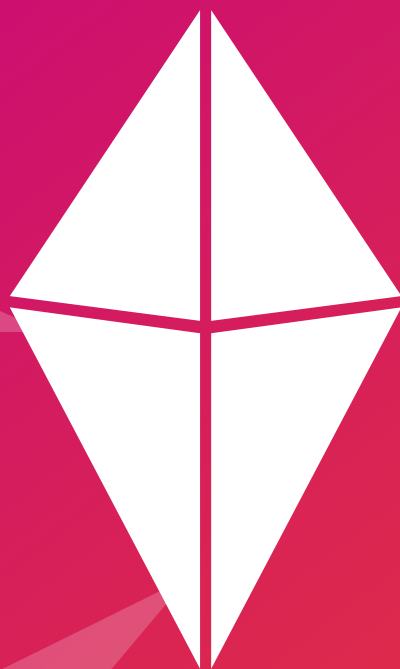
- due to the high cost of renewing the service contract on the CFTRc laser scanning microscope, we have implemented modest user fees (we had them initially, then eliminated them to encourage usage of the equipment). The microscope is now heavily used by CFTRc members and by other users in the McGill community.

15. Provide suggestions about how the Faculty could do better to support the Unit and research efforts in general (e.g., centralized data repositories, institutional data management plans, support for software developments, guidance for adopting open-science practices, simplification of administrative procedures, etc.) **(no page limit but please be specific and unleash your creativity!)**

Sponsoring workshops and other opportunities for networking between centers at McGill would be useful. For example, many members of the CFTRc are interested in airway inflammation and infection in CF. Inflammation and infection is also a focus of the complex traits group yet there is little formal communication between centres despite some common membership. There is more communication between the CFTRc and RI-MUHC/Meakins Christie lab and the CFTRc research day has been held there in the past.

In the attached (Excel) **Year-End Financial Report** please detail:

1. Expenditures of funding provided by the Faculty of Medicine and other sources, towards meeting the objectives of the Unit,
2. Any in-kind contributions provided to the Unit by other partners and sponsors,
3. Projected budget for the coming year (including request to the Faculty of Medicine).



UN SOUFFLE DE COLLABORATION

*Symposium québécois des
chercheurs et cliniciens FK*



Cystic Fibrosis
Fibrose kystique
Canada

Jeudi 9 et vendredi 10 mai 2019
Hôtel Mortagne



I would like to extend a warm welcome to the 2019 CFC-CFTRc symposium « An Era of Collaboration». This symposium will bring us up to date on CF research being carried out here in Quebec and hopefully will foster collaboration between different labs with common interests and complementary expertise. It also provides us with an opportunity to invite outstanding international speakers including Dr. Lap Chee Tsui, lead researcher responsible for cloning the CF gene, and Dr. Ray Frizzell, who has made many seminal contributions to our understanding of epithelial chloride transport. We are looking forward to their talks and to other exciting presentations throughout the day on diverse topics.

There will also be several interesting flash presentations from graduate students and postdocs and we encourage them to discuss their work with other participants including speakers to benefit from their insights and feedback.

This symposium is an annual event for the CF Translational Research centre (CFTRc), a virtual CF centre that was formally established in the fall of 2015. It is based at McGill but includes many other institutions. Its main goal is to accelerate the development of a cure for CF by providing access to state-of-the-art equipment, CF cells and other materials, by promoting collaboration between laboratories, and by organizing symposia, workshops, seminars, and travel awards for trainees to present their work at conferences and obtain feedback.

This is my opportunity to acknowledge the tremendous efforts of Dr. Annick Guyot in putting together this symposium. I also thank Cystic Fibrosis Canada for their helpful collaboration and our sponsors for making the symposium possible.

I wish you all an interesting and productive meeting.

A handwritten signature in black ink, reading "John Hanrahan". The signature is fluid and cursive, with the first name "John" being more prominent than the last name "Hanrahan".

Dr. John Hanrahan
Director of the CFTRc



Retrospective on CF research and future directions

Lap-Chee Tsui

*GBM, O.C., O. Ont.,
Founding President of the Hong Kong Academy of Sciences,
President of Victor and William Fung Foundation*

Briefly, I would go over the key steps that led to the identification of the Cystic Fibrosis Transmembrane Conductance Regulator gene and what we learned from the genetics of the CF disease. The gene identification has allowed better definition of the basic defect in CF, deeper understanding of its pathophysiology and research into new strategies to treat the patients in the past 30 years. Further advances in CF disease management continue to require close collaborations among scientists and medical service providers, and, even patients themselves.

📖 Prof. the Honourable Tsui Lap-Chee, world-renowned molecular biologist, is currently Founding President of the Hong Kong Academy of Sciences, President of Victor and William Fung Foundation, Director of Qiushi Academy for Advanced Studies and Master of Residential College in International Campus of Zhejiang University, and University of Toronto's Emeritus University Professor. He was the 14th Vice Chancellor of The University of Hong Kong. Prior to his appointment at HKU, Prof. Tsui was Geneticist-in-Chief and Head of the Genetics and Genomic Biology Program of the Research Institute at The Hospital for Sick Children in Toronto. He is world renowned for his research work in human genetics and genomics. He has also made significant contributions to the study of the human genome, especially the characterization of chromosome 7, and identification of additional disease genes. He has over 300 peer-reviewed scientific publications and 65 invited book chapters. He is the recipient of many national and international prizes, including the 2018 Warren Alpert Foundation Prize. His other awards include 16 honorary doctoral degrees from prestigious universities around the world.

Synergistic rescue of $\Delta F508$ and CFTR2 mutation functional expression defects by structure-guided corrector combinations

Guido Veit¹, Haijin Xu¹, Elise Dreano², Radu G Avramescu¹, Miklos Bagdany¹, Lenore K Beitel¹, Ariel Roldan¹, Mark A Hancock¹, Cecilia Lay³, Wei Li³, Katelin Morin³, Sandra Gao³, Puiying A Mak³, Edward Ainscow³, Anthony P Orth³, Peter McNamara³, Aleksander Edelman², Saul Frenkiel³, Elias Matouk³, Isabelle Sermet-Gaudelus², William G Barnes³, Gergely L Lukacs¹

1. McGill University, Montréal, Canada; 2. Institut Necker-Enfants Malades (INEM) - INSERM U1151, Paris, France; 3. Genomic Institute of the Novartis Research Foundation, San Diego, USA.

The most common cystic fibrosis mutation, $\Delta F508$ in nucleotide binding domain 1 (NBD1), impairs CFTR-coupled domain folding, plasma membrane (PM) expression, function, and stability. Robust $\Delta F508$ -CFTR correction is achieved by stabilization of NBD1, the interfaces between NBD1 and membrane-spanning domains (MSDs), as well as NBD2, the former two representing primary conformational defects, established by using combination of genetic and pharmacological means. Thus, a rationally designed, structure-guided corrector strategy may require the combination of type I correctors supporting the NBD1-MSD1 and NBD1-MSD2 interface formation, type II correctors targeting NBD2, and type III correctors stabilizing the NBD1 domain.

VX-809 (lumacaftor), the first approved corrector, exhibits a type I mechanism and in combination with the gating potentiator VX-770 (ivacaftor) provides only modest clinical benefit to patients carrying two copies of the $\Delta F508$ mutation.

Here we report the identification of compounds for all three corrector types in a screen of ~600,000 small molecules by monitoring the PM expression of the HRP-tagged $\Delta F508$ in CFBE41o- (CFBE) epithelia. Compounds that increased $\Delta F508$ PM densities both in the presence and absence of VX-809 contained type III correctors, compounds that required VX-809 included type II correctors, and compounds exhibiting redundancy with VX-809 encompassed type I correctors. The mechanisms of action (MOAs) of correctors were determined by domain-interrogation and domain-specific binding assays, competition with reference compounds, and Δ NBD2-CFTR PM density measurements. While type I-III correctors alone displayed only modest correction, combination of correctors from all three classes synergistically increased the $\Delta F508$ PM density and function in CFBE by up to ~9 fold in comparison to VX-809 alone, augmented the mutant ER maturation and the abundance of the complex-glycosylated form, promoted the peripheral stability, and largely normalized the single channel function of $\Delta F508$. These results correlated well with CFTR gain-of-function in human bronchial epithelia and human nasal epithelia from CFTR ^{$\Delta F508/\Delta F508$} patients. Corrector combinations lead to ~50% of wild-type-level correction in human nasal and bronchial epithelia and in mouse nasal epithelia. Likewise, corrector combinations were effective against rare missense mutations in various CFTR domains, probably acting via structural allostery, suggesting a mechanistic framework for their broad application.

This study provides proof of principle for synergy screening to identify correctors with distinct MOAs, which, when used in structure-guided combinations, achieve therapeutically relevant correction levels of $\Delta F508$ and other processing mutants.

Synergistic inhibition of CFTR dependent chloride secretion by urban air pollution particulate matter and oxidative stress in airway epithelial cells

Victor Dumitru^{1,2}, Jie Liao^{1,2}, Premkumari Kumarathasan³, Renaud Vincent³, John W. Hanrahan^{1,2}

1. Department of Physiology, McGill University; 2. McGill Cystic Fibrosis Translational Research centre (CFTRc); 3. Environmental Health Science and Research Bureau, Health Canada

The FDA recommends that clinically relevant drug-drug interactions be defined during drug development and assessed in prospective clinical studies prior to filing a new drug application. Consequently many interactions of approved CFTR modulators with other drugs are known. By contrast, the interactions of CFTR modulators with environmental factors have not been studied although they could impact efficacy. For example rescued F508del-CFTR is downregulated by lower concentrations of cigarette smoke extract compared to wild-type CFTR (1). Urban air pollution particulate matter (PM) induces ER stress and the unfolded protein response (2) and elevates oxidative stress that may contribute to the cellular response to PM (3), however the effect on rescued F508del-CFTR after corrector treatment is not known. To examine the effects of PM and oxidative stress individually and in combination on airway epithelial cells, comparing responses of non-CF and CF cells. Polarized CFBE cell monolayers and primary HBE cells mounted in Ussing chambers were used to measure chloride secretion. qPCR was performed on targets associated with oxidative stress. PM (standard reference material 1648) from the National Institute of Standards and Technology was used as PM. CFTR-dependent secretion was unaffected by overnight exposure to PM alone but was strongly inhibited when cells were treated with both PM and the oxidant stressors tert-butylhydroquinone or hydrogen peroxide. CFBE cells expressing F508del-CFTR that had been partially rescued by the corrector drug VX-809 were more susceptible to PM + oxidant stress than cells expressing WT-CFTR. mRNA transcript levels for the antioxidant enzymes glutathione synthetase, superoxide dismutase 2, and catalase were elevated by oxidant exposure as expected, however this response was also impaired in cells expressing F508del-CFTR and further reduced by the presence of PM + oxidant stress. Immunoblots of primary human bronchial epithelial cell lysates revealed a slight decrease in CFTR protein expression, although a reduction in basolateral transport also contributed to the acute inhibition of CFTR-dependent secretion by oxidants. We conclude that oxidant stress and PM cause synergistic inhibition of airway secretion and antioxidant defenses are compromised in CF cells. Finally, much of this inhibition of CFTR-dependent secretion may occur at the basolateral membrane.

The dual phosphodiesterase 3/4 inhibitor RPL554 stimulates rare class III and IV CFTR mutants

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Over 2,000 mutations have been reported in the *cftr* gene, many of which cause disease but are rare and have no effective treatment. Thus there is an unmet need for new, preferably mutation-agnostic, therapies for cystic fibrosis (CF). Phosphodiesterase inhibitors (PDEis) are one potential class of therapeutics as they have been shown to elevate intracellular cAMP levels and stimulate CFTR-dependent secretion in human airway epithelia, however the number of people of CF that could be helped by PDEi's remains to be determined. Recently, we demonstrated that an inhibitor of human PDE3 and PDE4, RPL554 (Verona Pharma), was able to stimulate the class IV CFTR mutant R117H CFTR endogenously expressed in well-differentiated primary human bronchial epithelial cells. Therefore, we sought to assess whether RPL554 could also stimulate other class IV CFTR mutants and explored its effects on class III CFTR mutants for the first time. Fisher Rat Thyroid cells transduced with lentiviruses to stably express R334W or T338I CFTR (class IV) or S549R or G551D CFTR (class III) were used to study regulation by RPL554. We found that RPL554 elevates intracellular [cAMP] leading to a potentiation of forskolin-stimulated R334W, T338I, G551D and S549R CFTR when used either alone or in combination with the CFTR modulators VX809 and VX770. Furthermore, we obtained biochemical evidence that VX809 can increase the cell surface expression of T338I, G551D and S549R CFTR, which correlated with enhanced cAMP-stimulated activity. Together, our findings strengthen the therapeutic potential as RPL554 as an anti-CF therapy for CF patients with class III/IV mutations and expand its scope as a drug that can benefit numerous CF patients.



Airway smooth muscle, airway hyperresponsiveness and cystic fibrosis

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Cystic fibrosis is associated with an “asthma” phenotype in a high proportion of affected persons. Although primarily an epithelial disease CFTR deficiency in other tissues may be implicated in significant ways in the clinical expression of disease. We have focused on airway smooth muscle as a possible contributor to the pathogenesis of the asthma phenotype. Airway smooth muscle expresses CFTR and involvement in the contractile properties of the muscle has been demonstrated, primarily through the study of cultured smooth muscle cells. We have examined the airways for evidence of remodeling of the muscle and have confirmed greater muscle mass in airways from CF affected subjects, in agreement with previous studies. The intrinsic biomechanical properties of the muscle ex vivo were not different muscle harvested from previously healthy subjects, showing similar maximal force development on exposure to methacholine and normal dynamic properties when stimulated either electrically or with methacholine. However, the impairment of relaxation to β -agonist stimulation was shown after pre-treatment with interleukin-13, a T 2 cytokine associated with CF, compared to control muscle. In addition, we have found enhanced proliferation of CF smooth muscle in vitro compared to non-CF control cells. The mechanism is unclear but is associated with a smaller of proportion of cells in the G0/G1 phase of the cell cycle, suggesting an acceleration of entry to the S phase. An enhanced expression of myosin light chain kinase was also demonstrated but the significance of this finding is not clear.

In conclusion, CF airway smooth muscle shows distinct characteristics that may favour its remodeling in vivo and the inflammatory environment, associated with T 2 cytokine expression may lead to resistance to relaxation to β -agonist stimulation.

Fenretinide mimics CFTR-induced correction of DHA/AA imbalance and blocks LPS-induced MUC5AC overexpression without affecting MUC5B

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Cystic fibrosis (CF) is the most common genetic disease in Caucasians. CF manifests through the accumulation of mucus in the lungs, which serves as the fertile soil for the growth of microorganisms, leading to recurrent infections and ultimately lung failure. Mucus in CF patients consists of DNA from dead neutrophils and mucins produced by goblet cells. MUC5AC mucin is responsible for the pathological plugging of the airways whereas MUC5B has a protective role against bacterial infection. Therefore, decreasing the level of MUC5AC without affecting MUC5B would be a desirable mucoregulatory treatment.

We demonstrated that pre-treating mice with fenretinide in a chronic model of *P. aeruginosa* lung infection efficiently prevents the accumulation of mucus. Fenretinide prevented lipopolysaccharide-induced increase of MUC5AC gene expression, without affecting the level of MUC5B in the lung goblet cell line. Furthermore, fenretinide treatment efficiently reversed pro-inflammatory imbalance of fatty acids by increasing the levels of docosahexanoic and decreasing the level of arachidonic acid in lung epithelial cell line and primary leukocytes derived from CF patients.

Mechanism of action of the synergistic combination tomatidine-aminoglycoside against *Staphylococcus aureus* virulent and persistent phenotypes

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Background: *Staphylococcus aureus* (SA) can adopt two phenotypes, prototypical (WT) and its small-colony variant (SCV). These two phenotypes are often recovered from the lungs of cystic fibrosis (CF) patients and are frequently co-isolated with *Pseudomonas aeruginosa* (PA). Aminoglycosides such as tobramycin or gentamicin (AMG) were shown efficient to control and reduce PA prevalence to 40% in patients with CF. Although PA is one of the most deleterious pathogens in CF, prophylaxis targeting PA over the past few years did not profoundly modify the overall number of exacerbations per year or the number of days of hospitalization per exacerbation per year based on the CF registry data. On the other hand, *S. aureus* recovery rate from patients increased to more than 53% of the patients. Furthermore, reports indicate that co-isolation of both SA (MRSA or SCVs) and PA worsen patients' health. Tackling all three pathogens seems imperative. Tomatidine (TO), a phytomolecule extracted from tomatoes, exerts a strong bactericidal activity on the *S. aureus* SCV phenotype (minimal inhibitory concentration [MIC] of 0.06 µg/mL) and is part of the novel steroidal alkaloid antibiotic class. Moreover, when TO is combined to an aminoglycoside, the combination shows a strong synergistic activity against WT SA (AMG MIC of 0.06 µg/mL). We recently determined that the molecular target of TO was the ATP synthase subunit c (*atpE*) and that TO reduced ATP production in *S. aureus*. We report here how TO, with AMG, exerts its bactericidal activity against both the WT and SCV phenotypes of SA.

Methods: Since TO affects the bacterial ATP synthase, we measured the membrane potential. Bacteria in broth were incubated with various concentrations of antibiotics (TO or TO-AMG). Bacteria were then washed in PBS, the fluorophore DiOC₂ was added and incubated for 30 min before flow cytometry. To assess the production of reactive oxygen species (ROS), bacteria were suspended in broth and incubated 2h at 35°C before the addition of 10 µM H₂DCFDA for 1h. Bacteria were then washed and transferred to a 96-well plate containing broth and antibiotics (ciprofloxacin as a control). Fluorescence was measured (λ exc 494_{nm}, λ emi 521_{nm}) over a 13-h period. AMG uptake was measured using a Texas-Red tagged AMG with the addition of TO. All results are reported as a percentage of that measured for WT without antibiotic.

Results: TO reduced WT membrane potential in a dose-dependent manner and reached a low of 35% at the highest doses (≥8 µg/mL). On the other hand, SCV membrane potential, which was about 10% of that of WT, further dropped to about ≤2% at very low TO concentrations (≥0.0035 µg/mL). This was also accompanied by 2 times more ROS production than that seen in the no antibiotic control. Besides, there was no difference in the membrane potential of WT when comparing the effect of AMG to that of TO-AMG. However, the combination TO-AMG generated 2.5 times more ROS compared to that caused by AMG alone. TO also increased AMG uptake by more than 1.5 times.

Conclusions: TO is able to reduce the membrane potential of both SA WT and SCV phenotypes, but the membrane potential only dropped to a critical level in the SCV. Also, significant ROS are only produced in SCV, which are highly susceptible to TO. Similarly, only the TO-AMG combination generated significant ROS production in the WT and increased AMG uptake, which explains the strong synergy with aminoglycosides. The TO-AMG combination may represent a novel therapeutic paradigm for lung infections in CF patients, targeting both *S. aureus* phenotypes (WT and SCV) and *P. aeruginosa* all together.

Despite antagonistic activities *in vitro*, *Pseudomonas aeruginosa* enhances *Staphylococcus aureus* colonization in a murine lung infection model

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Background: A defective muco-ciliary clearance of bacteria in cystic fibrosis (CF) patients results in recurrent pulmonary infections. *S. aureus* (SA) and *P. aeruginosa* (PA) are the most prevalent lung pathogens in CF and are frequently co-isolated. Their co-infection is associated with a worse clinical outcome, as noted by decreased lung functions and frequent pulmonary exacerbations. However, despite their co-occurrence, PA and SA prototypic strains exhibit antagonism *in vitro*: PA reduces SA growth and induces the small-colony variant (SCV) phenotype. Therefore, we attempted to better understand the apparent conflict between the *in vitro* observations and the high SA-PA co-occurrence in CF. We previously described clinical SA-PA co-isolates not displaying such an antagonism. The present study compares the colonization of various strains, including reference, clinical co-isolates or virulence-attenuated mutants, in a murine co-infection model.

Methods: Growth kinetics were followed for co-cultures of five SA-PA clinical pairs isolated from adult CF patients. Selective plates were used to determine viable counts for each species. In other experiments, SA was spread over an agar plate where a PA spot was applied in the centre to visualize SA SCV formation around PA. *In vivo* interactions were characterized using a mouse lung infection model. Intra-tracheal inoculations of SA, PA or SA-PA pairs were performed, and infections developed for 24h. Lung homogenates were plated on selective media allowing CFU counts of either SA or PA. Inflammation was assessed by myeloperoxidase (MPO) quantification and expression of two known receptors for cellular adhesion of SA, ICAM-1 and ITGA-5, were measured by RT-qPCR.

Results: Growth kinetics showed that some PA antagonized their SA co-isolate with a drop of $\sim 3.6 \log_{10}$ CFU/mL initiated after 8h of co-culture. However, other pairs did not interact negatively showing equivalent growth for SA in mono- and co-cultures. The agar co-culture model revealed the formation of SA SCVs in pairs where PA antagonized SA growth kinetics, while some non-antagonistic pairs did not result in the formation of SCVs. Paradoxically, in the infection model, SA colonization was significantly higher in SA-PA co-infections, even for pairs showing antagonism *in vitro*. In fact, SA colonization was most enhanced in co-infections with antagonistic PA. The SA colonization increased up to $2.00 \log_{10}$ in co-infections compared to their respective SA mono-infections ($P < 0.05$). PA did not benefit from the co-infection showing equivalent colonization of lung tissues in presence or absence of SA ($P > 0.05$). Upon compiling all results from 200 co-infections (35 PA and 10 SA strains, including clinical, reference and mutant strains), SA colonization was found to be proportional to PA colonization. Different virulence-attenuated mutants for both species were evaluated in the co-infection model, but no virulence or regulatory genes could specifically be associated to this phenomenon. The level of inflammation was measured via MPO quantification but was not correlated to the promotion of SA colonization. However, RT-qPCR of ICAM-1 and ITGA-5 showed that PA significantly increased their expression, both in mono- or co-infections with SA.

Conclusion: The observation that lung colonization by SA is improved in presence of PA may explain the frequent co-infections by these pathogens in CF and may contribute to their combined detrimental effect on patient health. The more effective SA colonization in presence of PA could involve an increase of its cell surface receptors by PA.



Susceptibility of CFTR-mutant mice to *C. rodentium* infection: a new model to study the role of CFTR in the gut?

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Although commonly associated with a respiratory phenotype and chronic lung infection, Cystic Fibrosis (CF) is a multi-organ disease with manifestations in multiple organ systems, including the gastrointestinal tract. CF is caused by mutations in the cystic fibrosis transmembrane conductance regulator (CFTR), which is located in the membrane of various epithelial tissues and immune cells and plays a critical role in ion and fluid homeostasis. Mouse models of CF generally display little lung pathology, but do possess an overt intestinal phenotype, characterized by obstruction, inflammation, and bacterial overgrowth. To probe the intestinal phenotype of CF mutant mice, we infected Δ F508 CFTR mice (Cftrtm1Eur) and *Cftr*^{-/-} mice with *Citrobacter rodentium*, a Gram-negative intestinal mouse pathogen. We found that CFTR mutant mice were highly susceptible to infection with *C. rodentium*, displaying high levels of mortality and atypical localization and greater colonization of bacteria than WT littermate mice. Detailed immunophenotyping revealed differences in the intestinal immune status of CFTR mutant mice at steady state and following infection. Further experiments using conditional knockout and gut-corrected mouse models suggest that loss of CFTR expression in the intestinal epithelium is not sufficient to cause *C. rodentium* infection susceptibility. We propose that *C. rodentium* infection provides a new model system to study the role of CFTR in the gut which may be relevant to intestinal disease observed in CF patients.



Correction of the CFTR3849+10kb C>T mutation using a CRISPR-Cas9 NHEJ strategy delivered by receptor-targeted nanocomplexes

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Introduction: Cystic Fibrosis is an autosomal recessive disorder caused by mutations in the CFTR gene. The 10th most common mutation, 3849+10kb C>T, generates a cryptic splice site, resulting in the formation of a pseudoexon containing a PTC, producing a truncated version of the protein. CRISPR/Cas9 allows for precise targeting of mutations by a guide RNA targeting molecule followed by double strand DNA cleavage by Cas9 nuclease. However, delivery of the CRISPR components into cells and target organs remains a challenge.

Aims: Our approach is to deliver CRISPR with a non-viral nanoparticle, previously described for in vivo DNA and siRNA delivery. These nanoparticles comprise peptide and lipid components, which package nucleic acids and target their delivery to epithelial cells. Gene editing of airway epithelial cells is permanent, therefore repeated delivery with these non-immunogenic nanoparticles could be performed to reach a sufficient level of genetic correction.

We designed pairs of Cas9 guide RNAs to create targeted double-stranded breaks in CFTR either side of the mutation, resulting in high efficiency excision via non-homologous end-joining repair, when tested in a mini-gene assay in HEK293T cells. Experiments were then repeated in bronchial cells isolated from a patient homozygous for this mutation to confirm functional restoration of the CFTR protein channel.

Methods: Primary CFBE cells were lentivirally transduced with the *BMI-1* proto-oncogene to expand proliferative potential for the course of experiments. Pairs of gRNAs were complexed with Cas9 protein and formulated with our lipid-based nanoparticles for transfection. 48 h post-transfection, genomic DNA was isolated, and cells were re-seeded for repeat transfections. Cells were then expanded for ALI culture. After 5 weeks differentiation on ALI, CFTR mRNA was analysed via qRT-PCR, and protein expression analysed by Ussing Chamber.

Results: After one transfection, a DSB efficiency of 26% was achieved in primary CFBE cells, as measured by Inference of CRISPR Edits (ICE) software, and T7 endonuclease assay. Of this, 10% had the expected 187 bp excision, indicative of successful cutting by both guides. After four repeat nanoparticle transfections, a DSB efficiency of 82% was achieved, 61% of sequences having the expected excision. This was able to restore CFTR mRNA expression and, importantly, CFTR channel function as measured by Ussing Chamber.

Conclusion: This approach could be used to correct aberrant splicing signals in several other CF mutations. Moreover, this targeted gene excision strategy may also be applicable in the study many other genetic disorders where deep-intronic mutations have been identified as a disease cause.



***S. aureus* and *P. aeruginosa* infections in cystic fibrosis: beyond bacteria, to the host response of CFTR-targeting therapies**

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Progressive lung damage due to chronic inflammation and bacterial infections, especially with *P. aeruginosa* and *S. aureus*, remains the first cause of morbidity and mortality in cystic fibrosis (CF) patients. Although novel CFTR-targeting therapies have recently emerged, their efficiency remains limited and variable among patients. There is now evidence, including from our laboratory, that *P. aeruginosa* infection down-regulates wt-CFTR and restrains the functional rescue of F508del-CFTR by CFTR correctors. Our previous work also highlighted a link between CFTR function, bacterial infections and epithelial repair mechanisms after injury. Indeed, we discovered that the capability of CF airway epithelia to heal is less efficient than in healthy subjects, most likely due to the basic CFTR defect and bacterial infections. We then discovered that CFTR rescue with correctors enhances epithelial repair in non-pathogenic conditions; this effect is however dampened in the presence of *P. aeruginosa* infection.

Our general objective is to 1) better define the impact of *S. aureus* and *P. aeruginosa* on the response of host epithelial cells to CFTR-targeting therapies and 2) identify therapeutic strategies to improve CFTR rescue and epithelial repair, despite the presence of *S. aureus* or *P. aeruginosa* infection. To achieve our goals, we are using primary cultures of airway epithelial cells (AEC) collected from non-CF and CF patients, exposed to bacterial exoproducts (from *P. aeruginosa* or *S. aureus*) and treated with CFTR modulators (Orkambi (VX-809 corrector + VX-770 potentiator) or Symdeco (VX-661 + VX-770)). CFTR-F508del rescue (CFTR maturation and currents) as well as repair processes (migration/lamellipodia dynamics, cytoskeletal organization, wound healing rates) were then assessed. The effect of various *P. aeruginosa* strains on CFTR was also evaluated in mouse lung tissues collected after chronic *P. aeruginosa* airway infection.

Our data first indicated that the effects of *P. aeruginosa* strains vary as a function of their genotypic/phenotypic characteristics. Whereas an Early strain (intermittent acute infection) severely dampened CFTR expression *in vitro* and *in vivo*, an engineered mutant of the *lasR* gene (Early Δ lasR) and a late isolate (from the same patient, when chronic infections are established) did not alter CFTR. We also discovered that interfering with bacterial quorum sensing (QS) with a QS inhibitor (QSI, HDMF) prevented the deleterious effect of *P. aeruginosa* on CFTR rescue by correctors. We then demonstrated that *P. aeruginosa* and *S. aureus* exoproducts significantly impaired airway epithelial repair processes. The deleterious impact of *P. aeruginosa* was prevented by QS inhibition. Our work also showed that the Symdeco or Orkambi combinations elicited a greater beneficial effect, than the corrector alone, on the repair of airway epithelia from F508del/F508del and heterozygous (F508del+another class II mutation). The effects of Orkambi or Symdeco were however dampened in the presence of *P. aeruginosa* or *S. aureus* exoproducts, indicating that a complementary approach is required to efficiently restore epithelial integrity. Interestingly, *P. aeruginosa* did not affect K⁺ channels, which are widely expressed in airway epithelial tissue, including in progenitor cells. Moreover, a treatment with Orkambi or Symdeco, combined with K⁺ channel activators, greatly improved airway integrity, despite the presence of infection. Such strategies, targeting *P. aeruginosa* (with QSI) and both CFTR and K⁺ channels (with modulators) would deserve further investigation to enhance the efficiency of treatments in cystic fibrosis.

Étude des biomarqueurs prédictifs associés à l'ischémie de reperfusion et à la dysfonction primaire du greffon suite à une transplantation pulmonaire chez les patients atteints de fibrose kystique

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Malgré l'amélioration des traitements pour les patients atteints de fibrose kystique (FK), la transplantation pulmonaire demeure la seule option de survie en phase respiratoire terminale. Toutefois, malgré l'amélioration des méthodes de préservation du greffon, des techniques chirurgicales, des soins péri-opératoires et de l'immunosuppression, le taux de survie à 5 ans (+/- 65%) des greffés pulmonaires reste insuffisant. La dysfonction primaire du greffon (DPG), se développant dans les 72h après la transplantation, est la première cause de décès en période péri-opératoire. Elle est aussi associée à des risques accrus d'infection, de syndrome de détresse respiratoire, et à plus long terme, de rejet chronique et de taux de survie réduits à 5 et 10 ans. Il a été établi que la lésion d'ischémie/reperfusion (I/R) du greffon représente un risque majeur de la DPG. À ce jour, il n'existe aucun traitement pharmacologique efficace pour la DPG. Il est donc crucial d'identifier de nouvelles cibles thérapeutiques.

Nous avons donc proposé l'hypothèse que la dysfonction de l'épithélium alvéolaire joue un rôle majeur dans la pathophysiologie de l'I/R et de la DPG suite à la transplantation pulmonaire chez les patients souffrant de FK.

Les objectifs spécifiques du projet, alliant une composante cellulaire, un volet *in vivo* (EVLP chez le cochon) et translationnel chez les patients transplantés sont de : 1) démontrer l'importance de marqueurs d'intégrité, de dommages et de fonctionnalité de l'épithélium alvéolaire, associés à l'I/R *in vitro* et *in vivo*; 2) valider ces marqueurs prédictifs dans le développement et la sévérité de la DPG chez les transplantés pulmonaires.

En utilisant un protocole mimant l'ischémie/reperfusion (I/R), à partir de culture primaire de cellules épithéliales alvéolaires de rats, nous avons une diminution de l'expression d'un canal sodique (ENaC) jouant un rôle primordial dans la fonction de clairance liquidienne alvéolaire, une baisse de l'expression d'une protéine impliquée dans les jonctions serrées (ZO-1) et de la résistance transépithéliale (RTE), indiquant une altération de l'intégrité alvéolaire. De plus, la capacité de réparation suite aux lésions est affectée.

Grâce à un modèle porcin avec inflammation induite au LPS nous avons pu observer un dommage tissulaire, un œdème pulmonaire, une réaction inflammatoire exacerbée ainsi qu'une baisse de l'expression d'ENaC, qui ne peut être reversée par la procédure de perfusion *ex-vivo* du poumon (EVLP).

Parmi les 53 patients FK transplantés pulmonaires que nous avons recrutés, 22 ont développés par la suite une DPG de grade 2 ou 3 dans les 72h suivant la transplantation pulmonaire. Nos résultats préliminaires, à partir de prélèvements faits lors de ces transplantations, indiquent une réponse inflammatoire, un dommage alvéolaire et une baisse de l'expression d'ENaC et ZO-1 déjà présents dans les greffons des donneurs, parmi les patients greffés ayant par la suite développés une DPG, comparé aux receveurs sans développement de la DPG.

Les résultats de nos études *in vitro* et *in vivo* ainsi que du volet translationnel chez l'humain, indiquent une altération de l'épithélium alvéolaire en lien avec l'I/R et la DPG. Compte tenu du dommage précoce observé dans le greffon donneur, notre but est maintenant d'identifier les caractéristiques phénotypiques du donneur (cause de décès, âge, sexe, intervention pré-collecte, etc.) associées au développement de la DPG chez les receveurs. Ces données seront cruciales pour développer de nouvelles stratégies thérapeutiques visant à améliorer la survie des transplantés pulmonaires.

Les canaux potassiques : nouvelles stratégies thérapeutiques dans la réparation de l'épithélium des voies aériennes fibrose kystique en présence d'exoproduits infectieux de *S. aureus*

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Introduction: Chez les patients atteints de fibrose kystique (FK), la présence d'infection/inflammation chroniques entraîne une atteinte pulmonaire progressive, qui est la cause principale de morbidité et mortalité. Bien que la prévalence des pathogènes respiratoires soit variable d'un patient à l'autre et en fonction de son âge, la bactérie *Staphylococcus aureus* (SA) prédomine chez les jeunes patients, tandis que les infections à *Pseudomonas aeruginosa* (PA) deviennent plus fréquentes chez les patients adultes. Nous avons précédemment démontré que l'infection, par des exoproduits sécrétés par PA, altère la réparation de l'épithélium des voies aériennes FK ainsi que l'efficacité des correcteurs de CFTR. Notre but est de développer des stratégies efficaces favorisant la réparation de l'épithélium des voies aériennes FK en ciblant les canaux potassiques K⁺(K_VLQT1 et K_{ATP}) et le canal CFTR (Symdeco[®]) en présence d'exoproduits de SA.

Méthodes: Nous avons collecté les cellules épithéliales des voies aériennes humaines FK (hAEC) à partir des poumons provenant de patients FK transplantés homozygotes F508del/F508del, mutation la plus fréquente. La vitesse de réparation a été évaluée en réalisant des tests de blessure à partir de cultures primaires d'hAEC. Les cellules ont été soumises à des traitements pharmacologiques : un activateur de K_VLQT1 (ML-277, 4 µM) et/ou de K_{ATP} (pinacidil, 50 µM); et/ou des modulateurs de CFTR utilisés en clinique (Symdeco[®] : combinaison du correcteur Tezacaftor VX-661, 4 µM et du potentiateur Ivacaftor VX-770, 100 nM) en présence ou en absence d'exoproduits bactériens de SA.

Résultats: Nous avons tout d'abord mis en évidence que les canaux potassiques K_VLQT1 et K_{ATP}, étaient exprimés dans les hAEC FK et que leurs expressions n'étaient pas affectées suite à une exposition d'exoproduits bactériens de SA. Ensuite, nous avons pu également confirmer l'effet délétère de l'infection par SA sur la réparation épithéliale FK. Nous avons pu noter un faible effet stimulateur du Symdeco sur la réparation épithéliale dans des conditions infectieuses. Toutefois, la vitesse de réparation est nettement améliorée par les activateurs ML-277 et pinacidil. De façon intéressante, un effet synergique est observé avec des co-traitements aux activateurs des canaux K⁺ (ML-277 et pinacidil) et modulateurs de CFTR (VX-661 + VX-770), malgré la présence d'exoproduits de SA.

Conclusion: Nos résultats démontrent que des traitements combinés avec des modulateurs des canaux CFTR et K⁺ pourraient ainsi être une stratégie efficace pour favoriser la réparation de l'épithélium respiratoire FK et ce malgré la présence d'infection bactérienne. Cette stratégie pourrait être non seulement testée sur des cellules issues de patients FK homozygotes pour la mutation F508del mais également sur d'autres classes de mutations.

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***Pseudomonas aeruginosa* evasion of neutrophil antibacterial functions in early cystic fibrosis lung infection**

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Pseudomonas aeruginosa (PA) is the predominant pathogen which causes chronic lung infection in patients with cystic fibrosis (CF). Failure to clear PA by the host innate immune responses leads to persistent PA infections in the CF airways, which has been associated with lung function decline and worse clinical outcomes. Although inhaled tobramycin treatment has demonstrated efficacy in improving eradication of PA in CF children with new onset infections, 40% patients still failed inhaled tobramycin eradication therapy. The adequate recruitment and antibacterial functions of neutrophils (PMNs) upon initial PA infection is likely a key step required for successful PA eradication even when antibiotics are used. Interestingly, microbiological analysis from clinical trials of PA eradication therapy suggested that certain PA bacterial phenotypes (loss of flagellar and pilus-mediated motilities, mucoidy or wrinkly colony morphology) are associated with PA eradication failure, but results vary between studies and the mechanism remains to be determined. Thus, we hypothesize that PA isolates that persist after inhaled tobramycin in CF patients with initial PA infection elicit impaired PMN mediated antibacterial functions. To examine this, we tested a collection of PA strains isolated from initial PA infection from CF children followed at the Sick Kids as part of an “Early PA eradication” study. We compared *in vitro* phagocytosis and intracellular bacterial killing by PMN-like cells (differentiated HL-60) in response to persistent clinical PA isolates (N = 10 persistent patients with 18 isolates) vs eradicated clinical PA isolates (N = 32 eradicated patients with 53 isolates). So far, we observed a significantly lower PMN phagocytosis ($p < 0.01$) and intracellular bacterial killing ($p < 0.05$) of persistent PA compared to eradicated PA. To identify which bacterial phenotypes have impacts on PMN antibacterial functions, we compared various bacterial phenotypes (Type IV pilus mediated twitching and flagellum mediated swimming motility, overproduction of alginate or Psl exopolysaccharides, biofilm production, pyocyanin and protease secretion) using univariate and multivariable regression. We found that PMN phagocytosis is significantly associated with twitching motility ($r = 0.26$, $p < 0.01$) and mucoidy ($r = -0.28$, $p < 0.01$). Furthermore, our preliminary data in a subset of PA isolates, persistent PA ($n = 7$) produce a remarkably higher levels of Psl ($p < 0.01$) compared to eradicated PA ($n = 7$). In order to determine whether Psl produced by CF clinical PA isolates plays a role in PMN phagocytosis, we compared the effect of an anti-Psl monoclonal antibody (mAb) Psl0096 in *in vitro* PMN phagocytosis of persistent and eradicated PA with high and low Psl production respectively. We observed that PMN phagocytosis of PA isolates was significantly increased in a Psl dependent manner by the treatment of Psl0096 compared to its isotype control. Our results to date therefore suggest that Psl may be a determinant which significantly contributes to impaired PMNs phagocytosis in addition to mucoidy and lack of twitching, and may be associated with PA eradication failure in CF patients. We will extend our preliminary findings by first validating our *in vitro* results using PA clinical isolates from another independent patient cohort. We will further test the relevance of our *in vitro* findings in a murine pulmonary infection model for *in vivo* PMN phagocytosis and bacterial clearance in response to persistent and eradicated PA isolates.



Predicting and tuning the dynamics of microbial evolution to delay drug resistance

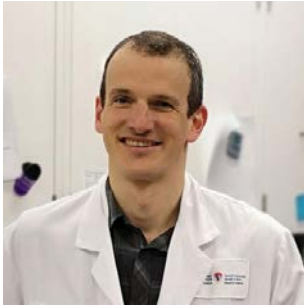
Adrian Serohijos

Département de Biochimie, Centre Robert-Cedergren en Bioinformatique et Génomique, Université de Montréal

The rise of antimicrobial resistance (AMR) has compromised our ability to manage chronic diseases, such as cystic fibrosis. To circumvent AMR, we need to discover new antibiotics. However, this will not be enough. We also need to conserve the efficacy and maximize the duration of usefulness of both current and future antibiotics. Antibiotic resistance will always arise, but if we are able to predict how they evolve, we can delay their emergence. At the fundamental level, the evolution of AMR, like any other biological system, is governed by processes at several scales of biological organization—molecular, cellular, organismal, and population level. Although the immediate effects of mutations are on the properties of proteins or fitness of individual cells, the eventual evolutionary success of these mutations is affected by population dynamics.

In this talk, I describe our efforts to develop a quantitative and predictive model of the emergence of AMR by bridging these multiple scales in bacterial evolution. First, we determined the comprehensive “mutational landscape” of gene targets that are crucial target of antimicrobials. We find that the overall survival of mutations that eventually become clinical isolates is strongly determined by the fitness of the bacteria in the presence of the drug (the “resistance level”) and by the fitness of the bacteria without the drug (the “fitness cost”). Using simulation and population genetics theory, we then predict the survival probability of these mutations under different selective regimes defined by drug concentration and population structure of the bacteria. These predictions are validated by lab evolution and data from clinical isolates. Altogether, by driving the bacterial populations into a region of the landscape where they are unable to grow, we can delay drug resistance.

www.serohijoslab.org



The potential roles of the Aryl-Hydrocarbon Receptor in the host defense response of airway epithelial cells exposed to *Pseudomonas aeruginosa*

Perrine Bortolotti, Lucie Roussel, Dao Nguyen, Carolyn J. Bagloli and
Simon Rousseau

Meakins-Christie Labs, RI-MUHC, Montréal, Canada

Pseudomonas aeruginosa are gram-negative bacteria that frequently infect the lungs of cystic fibrosis (CF) patients. This bacterium is highly responsive to changes in its environment, resulting in the expression of a diverse array of bacterial genes that may contribute to host-pathogen interactions. *P. aeruginosa* is well-known to induce neutrophilic inflammation via the activation of Toll-Like Receptors (TLRs). Recently, it was shown that pyocyanin, a phenazine produced by *P. aeruginosa*, binds to the aryl hydrocarbon receptor (AhR), leading to neutrophilic inflammation. AhR is a ligand-dependent transcription factor involved in the regulation of innate immunity, which also has non-genomic functions. Understanding the role of the AhR in the *P. aeruginosa*-induced epithelial infection and its subsequent immune response could help better understand the complexity of CF lung disease. In this presentation, the differential contribution of AhR and TLRs to neutrophilic inflammation will be presented as well as a potential role of AhR in the *P. aeruginosa* “detoxifying” response.

Understanding how pathoadaptation of *Pseudomonas aeruginosa* in cystic fibrosis alters host-microbe interactions in the airway epithelium

Lisa Hennemann, E. Faure, P. Bortolotti, S. Rousseau, Dao Nguyen

McGill University, Montreal, Canada

Chronic *Pseudomonas aeruginosa* (PA) infections occur in 60-80% of all adult Cystic Fibrosis (CF) patients and are associated with an accelerated lung function decline and increased mortality. Due to host and bacterial factors that allow PA to rapidly adapt to the host environment and cause impaired bacterial clearance, PA is nearly impossible to eradicate once it has established a chronic infection. PA infections cause progressive lung disease by inducing an exuberant and non-resolving lung inflammation. While PA is traditionally considered an extracellular pathogen, our lab has recently demonstrated the presence of intracellular PA within airway epithelial cells (AECs) of CF lung explant tissues, indicating the possible presence of an intracellular reservoir *in vivo*. In this project, we investigate how chronic PA pathoadaptations modulate host-microbe interactions in the CF airway, including intracellular survival in airway epithelial cells, a novel mechanism of PA persistence. We hypothesize that genetic adaptations commonly observed in PA isolates from chronic infections promote increased lung inflammation and an intracellular PA lifestyle which promote bacterial persistence.

Our first aim is to analyze the impact of LasR (major quorum sensing transcriptional activator in PA) loss of function on ICAM-1, an important pro-inflammatory mediator expressed by AEC in response to PA extracellular diffusible bacterial products and intracellular bacteria. Preliminary results show that expression of membrane-bound ICAM-1 in AEC is upregulated in response to *lasR* mutant filtrates compared to filtrates from wild-type PA and that this upregulation enhances neutrophil binding to AECs *in vitro*, suggesting that it might promote neutrophilic lung inflammation in the CF airway.

Our second aim is to examine whether LasR and Type 3 secretion system (T3SS) loss of function, two common pathoadaptations, modulate internalization of PA into AECs as well as its intracellular persistence *in vitro*. So far, we have observed that *lasR* mutation promotes internalization into CFBE $\Delta F508$ cells, an immortalized cell line homozygous for the most common CFTR mutation observed in CF patients, as well as primary CF AECs. We have further found that loss of T3SS injectisome function enhances bacterial persistence in AECs for up to 120h, and this may be unrelated to T3SS-induced cytotoxicity or secreted effectors.

In conclusion, our results suggest that loss of LasR might increase lung inflammation and thus contribute to CF lung function decline. Furthermore, loss of LasR and T3SS function, two phenotypic adaptations commonly observed in PA strains from chronic CF infections, promote an intracellular PA lifestyle, potentially allowing bacteria to avoid clearance and persist. Consequently, further examination of the involved mechanisms might provide insight into how common pathoadaptations contribute to disease progression and PA persistence.

Future experiments will further explore how the T3SS modulates PA intracellular persistence and whether our ICAM-1 *in vitro* data translates to increased membrane-bound ICAM-1 induced by *lasR* mutants compared to wild-type PA *in vivo*.

The opposing effect of cigarette smoke on CFTR activity is dependent on the channel's phosphorylation state

Aiswarya Premchandrar, Andrea Schnur, Miklos Bagdany and Gergely L Lukacs

Departments of Physiology and Biochemistry, McGill University, Montreal, Canada

Defective CFTR function is a major risk factor in CF, COPD, and other lung diseases. Chronic cigarette smoke (CS) exposure, a leading cause of COPD, inhibits CFTR activity at transcriptional, biochemical, and functional levels. The organic components of CS, the condensate (CSC), however, transiently activate the resting CFTR channel in respiratory epithelia. The comparable phospho-occupancy of ten PKA consensus sites, after CSC and forskolin exposure, determined by affinity-enriched tandem mass spectrometry, suggest that PKA stimulation causes the channel activation. This is accomplished by subcompartmentalised elevation of cAMP concentration due to inhibition of the MRP4, a cAMP export pump, which is a constituent of the CFTR macromolecular signalling complex at the cell surface. In sharp contrast, CSC reversibly inhibits the phosphorylated CFTR channel activity *in vivo* and in phospholipid bilayers, without altering its phospho-occupancy and cell surface expression. Put together, we posit that CS may also elicit a dual acute effect on CFTR in the airways; an acute activation of the resting channel that enhances the protective mucociliary clearance efficiency, and a subsequent channel inactivation that contributes to the COPD lung pathology.

Role of the proteasome in the biosynthetic arrest of SLC26A9 by F508del-CFTR

Yukiko Sato^{1,3}, Renaud Robert^{1,3}, David Thomas^{2,3'}, John Hanrahan^{1,3}

1. Department of Physiology, McGill University, Montréal, Canada; 2. Department of Biochemistry, McGill University, Montréal, Canada; 3. McGill Cystic Fibrosis Translational Research center

Introduction: Currently available drugs that correct F508del-CFTR misfolding provide only modest clinical benefit for most CF patients, therefore alternative anion channels and other potentially druggable targets are being explored. SLC26A9 is a constitutively active anion channel expressed in human airways that modifies the severity of CF airway disease in patients with the G551D-CFTR mutation. The functional expression of complex glycosylated SLC26A9 is reduced in cells that express F508del-CFTR mediated in part by a PDZ- and CAL-dependent mechanism. SLC26A transporters also interact with the regulatory (R) domain of WT-CFTR through its Sulfate Transporter AntiSigma factor antagonist (STAS) domain, however the role of this interaction in SLC26A9 biosynthetic arrest is uncertain. Indeed, it is not known if the interaction with immature F508del-CFTR in the endoplasmic reticulum (ER) contributes to the retention and proteasomal degradation of SLC26A9.

Aim: to understand the interaction between SLC26A9 and CFTR and its impact on SLC26A9 expression.

Methods: BHK cells overexpressing wild-type (WT) or F508del-CFTR and parental BHK cells lacking CFTR were transiently transfected with SLC26A9 cDNA. SLC26A9 protein was quantified in lysates by immunoblotting and at the plasma membrane by cell surface biotinylation. SLC26A9 levels were assessed in well-differentiated CF and non-CF primary human bronchial epithelial cells (pHBEs) by immunofluorescence staining and confocal microscopy.

Results: Total and plasma membrane SLC26A9 expression were both lower in BHK cells when co-expressed with F508del-CFTR than when expressed alone or with WT-CFTR. Similar results were obtained when well-differentiated CF and non-CF primary human bronchial epithelial cells (pHBEs) were studied by confocal imaging. Since F508del-CFTR misfolding leads to its retention in the ER and subsequent proteasomal degradation, we examined the effects of proteasome inhibitors on SLC26A9 degradation. Inhibiting the proteasome increased SLC26A9 immunofluorescence in F508del-CFTR homozygous pHBE cells but not in non-CF pHBEs, suggesting there is enhanced proteasomal degradation of SLC26A9 with F508del-CFTR in pHBEs. This difference in proteasome inhibitor sensitivity was not observed in non-epithelial cells overexpressing SLC26A9 with F508del-CFTR or WT-CFTR. Apical SLC26A9 levels increased in pHBEs and in non-epithelial cells when F508del-CFTR was partially corrected using low temperature or VX-809, and this rescue was mimicked by co-transfecting cells with WT-CFTR. The rate of SLC26A9 degradation was measured when expressed alone or with WT-CFTR or F508del-CFTR. In the presence of the protein synthesis inhibitor cycloheximide, degradation of immature SLC26A9 was enhanced in F508del-expressing cells suggesting there is CFTR-dependent, proteasome-mediated degradation of SLC26A9 at the ER.

Conclusions: These results suggest that ER retention of F508del-CFTR and SLC26A9 leads to premature degradation of both proteins and SLC26A9 biosynthetic arrest is due in part to an interaction between them. Disrupting this interaction will make SLC26A9 an exciting therapeutic target for most CF patients.

Support: Studentships from CF Canada, and Fonds de recherche du Québec – Santé to YS, and grants from the Canada Foundation for Innovation to DYT and CF Canada to JWH.



Divergent functions and selective interactions of the Hsp70 chaperone system with CFTR

Jason C. Young

McGill University, Montreal, Canada

Chaperones assist the folding and quality control degradation of the CFTR channel, both its wild-type form and the misfolded $\Delta F508$ cystic fibrosis mutant. The Hsc70/Hsp70 chaperone promotes folding, but also ubiquitination by the E3 ligase CHIP, leading to ER-associated degradation, or internalization and lysosomal degradation from the plasma membrane. We addressed the overall role of Hsc70/Hsp70 and co-chaperones DNAJA1 and DNAJA2 in cells. Unexpectedly, the net effect of Hsp70 was to suppress CFTR trafficking. While DNAJA1 supported biosynthetic folding, DNAJA2 specifically enhanced ER-associated degradation through Hsp70 and CHIP. Excess Hsp70 also promoted CFTR degradation, but this occurred through the lysosomal pathway and required CHIP but not Hsp90 chaperone complex formation. Notably, the Hsp70 inhibitor MKT077 enhanced levels of mature CFTR and $\Delta F508$ -CFTR, by slowing turnover and allowing delayed maturation, respectively. MKT077 also boosted the channel activity of $\Delta F508$ -CFTR when combined with the corrector compound VX809. To further understand the divergent effects of chaperones, we screened a synthetic peptide library to identify binding sites within CFTR for Hsc70, DNAJA1 and DNAJA2. While many sites were shared between Hsc70 and DNAJA2, DNAJA1 binding was more restrictive. Moreover, the sites mapped to regions critical for the folding of CFTR. These results suggest how the chaperones act in coordination on these structurally labile regions for different outcomes. Modulation of the chaperone system may offer ways to relieve the misfolding phenotype.



Behavior of the CF disease modifier: SLC26A9

Raymond A. Frizzell and Carol A. Bertrand

Department of Pediatrics, University of Pittsburgh School of Medicine, Pittsburgh, PA 15224

The diversity in CF pathology makes the SLC26A family of anion transporters and ion channels attractive candidates to investigate as interacting partners with CFTR due to their diverse anion transport functions, tissue distributions and reciprocal regulation. Genome-wide association studies (GWAS) have implicated SLC26A9, originally identified in the lung, as a modifier of several CF-associated pathologies, including meconium ileus (1), CF-related diabetes (2), and prenatal exocrine pancreatic damage (3). Recently, GWAS-identified a non-coding, single nucleotide polymorphism in SLC26A9 (A9) as a modulator of the airway response to potentiator VX-770 in CF patients with the gating mutation G551D CFTR (4).

We found previously that primary, human bronchial epithelia (HBE) from non-CF donors exhibit constitutive anion secretion attributable to A9 (5); however, this secretory process was absent in HBE from CF donors. Therefore, we asked whether changes in A9 constitutive activity could be attributed to a loss of CFTR trafficking, and what role PDZ interactions played. HEK293 cells co-expressing A9 with the trafficking mutant F508del CFTR exhibited a significant reduction in constitutive current compared to cells co-expressing A9 with either WT or G551D CFTR, and the expression of A9 at the plasma membrane was reduced. A9 interacted with NHERF-1 and CAL, and its interaction with both PDZ proteins significantly increased with co-expression of WT CFTR. However, expression with F508del CFTR only increased A9's interaction with CAL, and core-glycosylated F508del CFTR co-immunoprecipitated endogenous CAL. Mutation of A9's PDZ motif restored its constitutive activity when co-expressed with F508del; also, correcting F508del CFTR trafficking in CF HBE with VX-809 restored A9 activity. Thus, CFTR can modify A9's interactions with PDZ-domain proteins in different cellular compartments along the protein maturation pathway (6). A9 has the potential to restore anion secretion to CF airway epithelia as it is farther along the maturation pathway than F508del CFTR, thus having therapeutic implications.

- (1) Sun L, Rommens JM et al. Multiple apical plasma membrane constituents are associated with susceptibility to meconium ileus in individuals with cystic fibrosis. *Nat Genet* 44: 562, 2012.
- (2) Blackman SM et al. Genetic modifiers of cystic fibrosis-related diabetes. *Diabetes* 62: 3627, 2013.
- (3) Miller MR et al. Variants in Solute Carrier SLC26A9 Modify Prenatal Exocrine Pancreatic Damage in Cystic Fibrosis. *J Pediatr*, 166: 1152, 2015.
- (4) Strug LJ, et al. Cystic fibrosis gene modifier SLC26A9 modulates airway response to CFTR-directed therapeutics. *Hum Mol Genet*, 25: 4590, 2016.
- (5) Bertrand CA et al. SLC26A9 is a constitutively active, CFTR-regulated anion conductance in human bronchial epithelia. *J Gen Physiol*, 133: 421, 2009.
- (6) Bertrand CA et al. The CFTR trafficking mutation F508del inhibits the constitutive activity of SLC26A9. *Am J Physiol Lung Cell Mol Physiol*, 312: L912-L925, 2017.

Appendix 1B



Appendix 1B: pictures taken at the CF symposium held on May 9, 2019 in Boucherville.





Retrospective on CF research and future directions

Lap-Chee Tsui

*GBM, O.C., O. Ont.,
Founding President of the Hong Kong Academy of Sciences,
President of Victor and William Fung Foundation*

Briefly, I would go over the key steps that led to the identification of the Cystic Fibrosis Transmembrane Conductance Regulator gene and what we learned from the genetics of the CF disease. The gene identification has allowed better definition of the basic defect in CF, deeper understanding of its pathophysiology and research into new strategies to treat the patients in the past 30 years. Further advances in CF disease management continue to require close collaborations among scientists and medical service providers, and, even patients themselves.

📖 Prof. the Honourable Tsui Lap-Chee, world-renowned molecular biologist, is currently Founding President of the Hong Kong Academy of Sciences, President of Victor and William Fung Foundation, Director of Qiushi Academy for Advanced Studies and Master of Residential College in International Campus of Zhejiang University, and University of Toronto's Emeritus University Professor. He was the 14th Vice Chancellor of The University of Hong Kong. Prior to his appointment at HKU, Prof. Tsui was Geneticist-in-Chief and Head of the Genetics and Genomic Biology Program of the Research Institute at The Hospital for Sick Children in Toronto. He is world renowned for his research work in human genetics and genomics. He has also made significant contributions to the study of the human genome, especially the characterization of chromosome 7, and identification of additional disease genes. He has over 300 peer-reviewed scientific publications and 65 invited book chapters. He is the recipient of many national and international prizes, including the 2018 Warren Alpert Foundation Prize. His other awards include 16 honorary doctoral degrees from prestigious universities around the world.



Guest Speaker

DR. RAYMOND FRIZZELL

University of Pittsburgh School of Medicine
Department of Cell Biology and Director of Cystic Fibrosis
Research Center

Title:

*“Different SUMO paralogs determine
the fate of CFTR: biogenesis vs.
degradation”*

Date: Friday, May 10, 2019

Time: 11:00 a.m.

**Place: Room 1034, 10th floor
McIntyre Medical Sciences Bldg.
3655 Promenade Sir William Osler
Montreal, QC H3G 1Y6**

Refreshments will be served before the seminar
Enquiries: 514-398-4318

SEMINAR ANNOUNCEMENT

Cystic Fibrosis Translational Research
CENTRE
de Recherche Translational sur la Fibrose Kystique



Guest Speaker:

DR. SUBASH SAD

**Department of Biochemistry, Microbiology and
Immunology, Faculty of Medicine, University of Ottawa**

Title:

***“Revenge of the Zombies: Lessons
learned from cystic fibrosis”***

Date: Friday, October 18, 2019

Time: 11:00 a.m.

**Place: Room 1034, 10th floor
McIntyre Medical Sciences Bldg.
3655 Promenade Sir William Osler
Montreal, QC H3G 1Y6**

SEMINAR ANNOUNCEMENT

Cystic Fibrosis Translational Research
CENTRE
de Recherche Translational sur la Fibrose Kystique



Guest Speaker:

DR. BOB SCHOLTE

**Erasmus Medical Center, Department of Pediatric
Pulmonology and Department of Cell Biology
Rotterdam, The Netherlands**

Title:

***“Therapy of Cystic Fibrosis, when, what
target, and how much is enough?”***

Date: Friday, November 8, 2019

Time: 11:00 a.m.

**Place: Room 1034, 10th floor
McIntyre Medical Sciences Bldg.
3655 Promenade Sir William Osler
Montreal, QC H3G 1Y6**



CFTRc Annual Meeting
9 May 2019
Hotel Mortagne – Boucherville

AGENDA

- 1) Governance
- 2) Membership
- 3) Training
- 4) Platforms: PACB and Imaging
- 6) CF symposium 2020
- 7) CF seminar series: Vertex sponsored
- 8) Report for the Faculty of Medicine
- 9) AOB



Thursday, May 16th 2019

NaotoSoya
McGill University
Physiology

Objet: CFTRC Travel Allowance

Dear Naoto,

Your application for a CFTRC Travel Allowance has been considered and it is my pleasure to advise that you have been awarded a Travel Allowance to attend the 2nd International Conference on HDX-MS (Banff, Alberta) from May 21-24 2019. The CFTRC will be reimbursing you for a maximum amount of \$1,500. Your supervisor, Dr. Lukacs, is responsible for paying any additional costs for your participation at this meeting.

Please accept our good wishes for the success of your research.

Yours sincerely,

John W. Hanrahan
Professor, Physiology
Director, *Cystic Fibrosis Translational Research Centre*

c.c. Dr. Gergely Lukacs

We require the following info for our files. Please complete and return by email to the attention of Dr. Annick Guyot. Thank you.

Accept this award: ☒ Yes ☐ No

Decline this award. Reason (optional):

Signature
Date May 16, 2019



Wednesday, July 10th 2019

Caroline Landry
CRCHUM
Biomedical Sciences

Objet: CFTRc Travel Allowance

Dear Caroline,

Your application for a CFTRc Travel Allowance has been considered and it is my pleasure to advise that you have been awarded a Travel Allowance to attend The 2019 Canadian Transplant Summit and The CDTRP (Banff, Alberta) from October 15-19 2019. The CFTRc will be reimbursing you for a maximum amount of \$1,500. Your supervisor, Dr. Brochiero, is responsible for paying any additional costs for your participation at this meeting.

Please accept our good wishes for the success of your research.

Yours sincerely,

John W. Hanrahan
Professor, Physiology
Director, *Cystic Fibrosis Translational Research Centre*

We require the following info for our files. Please complete and return by email to the attention of Dr. Annick Guyot. Thank you.

Accept this award:

☒ Yes

☐ No

Decline this award. Reason (optional)

Signature

Date

Caroline Landry
July 19th 2019

c.c. Dr. Emmanuelle Brochiero



Tuesday, June 11th 2019

Sonya Kouthouridis
McGill University
Chemical Engineering

Objet: CFTRc Travel Allowance

Dear Sonya,

Your application for a CFTRc Travel Allowance has been considered and it is my pleasure to advise that you have been awarded a Travel Allowance to attend the Summer Biomechanics, Bioengineering and Biotransport Conference (Seven Springs, USA) from June 25-28 2019. The CFTRc will be reimbursing you for a maximum amount of \$1,500. Your supervisor, Dr. Moraes, is responsible for paying any additional costs for your participation at this meeting.

Please accept our good wishes for the success of your research.

Yours sincerely,

John W. Hanrahan
Professor, Physiology
Director, *Cystic Fibrosis Translational Research Centre*

c.c. Dr. Chris Moraes

We require the following info for our files. Please complete and return by email to the attention of Dr. Annick Guyot. Thank you.

Accept this award: ☒ Yes ☐ No

Decline this award. Reason (optional):

Signature

Date

June 12th, 2019



Friday, September 13th 2019

Abira Rajah
McGill University
Physiology

Objet: *CFTRC Travel Allowance*

Dear Abira,

Your application for a CFTRC Travel Allowance has been considered and it is my pleasure to advise that you have been awarded a Travel Allowance to attend The Seeing is Believing: Imaging the Molecular Processes of Life Symposia (Heidelberg, Germany) from October 9-12 2019. The CFTRC will be reimbursing you for a maximum amount of \$1,500. Your supervisor, Dr. Brown, is responsible for paying any additional costs for your participation at this meeting.

Please accept our good wishes for the success of your research.

Yours sincerely,

John W. Hanrahan
Professor, Physiology
Director, *Cystic Fibrosis Translational Research Centre*

We require the following info for our files. Please complete and return by email to the attention of Dr. Annick Guyot. Thank you.

Accept this award:

☒ Yes

☐ No

Decline this award. Reason (optional):

Signature

Date *Sept. 16, 2019*

c.c. Dr. Claire Brown

Appendix 1I

Publications co-authored

1. The *Pseudomonas aeruginosa* Population among Cystic Fibrosis Patients in Quebec, Canada: a Disease Hot Spot without Known Epidemic Isolates.
Jeukens J, Freschi L, Kukavica-Ibrulj I, Emond-Rheault JG, Allard C, Barbeau J, **Cantin A**, Charette SJ, Déziel E, **Malouin F**, Milot J, **Nguyen D**, Popa C, Boyle B, **Levesque RC**.
J Clin Microbiol. 2019 May 24;57(6):e02019-18. doi: 10.1128/JCM.02019-18.
Impact Factor: 4.959
2. Agonists that stimulate secretion promote the recruitment of CFTR into membrane lipid microdomains.
Abu-Arish A, Pandžić E, Kim D, Tseng HW, **Wiseman PW**, **Hanrahan JW**.
J Gen Physiol. 2019 Jun 3;151(6):834-849. doi: 10.1085/jgp.201812143.
Impact Factor: 4.258
3. Pendrin Mediates Bicarbonate Secretion and Enhances Cystic Fibrosis Transmembrane Conductance Regulator Function in Airway Surface Epithelia.
Kim D, Huang J, Billet A, Abu-Arish A, Goepp J, Matthes E, **Tewfik MA**, **Frenkiel S**, **Hanrahan JW**.
Am J Respir Cell Mol Biol. 2019 Jun;60(6):705-716. doi: 10.1165/rcmb.2018-0158OC.
Impact Factor: 4.34
4. Bioactive Thymosin Alpha-1 Does Not Influence F508del-CFTR Maturation and Activity.
Armirotti A, Tomati V, Matthes E, Veit G, Cholon DM, Phuan PW, Braccia C, Guidone D, Gentzsch M, **Lukacs GL**, Verkman AS, Galletta LJV, **Hanrahan JW**, Pedemonte N.
Sci Rep. 2019 Jul 16;9(1):10310. doi: 10.1038/s41598-019-46639-1.
Impact Factor: 4.011
5. Paxillin S273 Phosphorylation Regulates Adhesion Dynamics and Cell Migration through a Common Protein Complex with PAK1 and β PIX.
Rajah A, Boudreau CG, Ilie A, Wee TL, Tang K, Borisov AZ, **Orlowski J**, **Brown CM**.
Sci Rep. 2019 Aug 7;9(1):11430. doi: 10.1038/s41598-019-47722-3.
Impact Factor: 4.011
6. Culture-Dependent Bioprospecting of Bacterial Isolates From the Canadian High Arctic Displaying Antibacterial Activity.
Marcolefes E, Leung T, Okshevsky M, McKay G, Hignett E, Hamel J, Aguirre G, Blenner-Hassett O, Boyle B, **Lévesque RC**, **Nguyen D**, **Gruenheid S**, Whyte L.
Front Microbiol. 2019 Aug 9;10:1836. doi: 10.3389/fmicb.2019.01836.
Impact Factor: 4.259
7. Hsp70 and DNAJA2 limit CFTR levels through degradation.
Kim Chiaw P, Hantouche C, Wong MJH, Matthes E, Robert R, **Hanrahan JW**, Shrier A, **Young JC**.
PLoS One. 2019 Aug 13;14(8):e0220984. doi: 10.1371/journal.pone.0220984.
Impact Factor: 2.776
8. Cystic Fibrosis: Proteostatic correctors of CFTR trafficking and alternative therapeutic targets.
Hanrahan JW, Sato Y, Carlile GW, Jansen G, **Young JC**, **Thomas DY**.
Expert Opin Ther Targets. 2019 Aug;23(8):711-724. doi: 10.1080/14728222.2019.1628948.
Impact Factor: 4.621

9. Magnetic microboats for floating, stiffness tunable, air-liquid interface epithelial cultures.
Chandrasekaran A, Kouthouridis S, Lee W, Lin N, Ma Z, Turner MJ, **Hanrahan JW, Moraes C.**
Lab Chip. 2019 Sep 7;19(17):2786-2798. doi: 10.1039/c9lc00267g.
Impact Factor: 6.914

10. A High-Throughput Image Correlation Method for Rapid Analysis of Fluorophore Photoblinking and Photobleaching Rates.
Sehayek S, Gidi Y, Glembockyte V, Brandão HB, François P, **Cosa G, Wiseman PW.**
ACS Nano. 2019 Oct 22;13(10):11955-11966. doi: 10.1021/acsnano.9b06033.
Impact Factor: 13.903

11. Mutation-specific dual potentiators maximize rescue of CFTR gating mutants.
Veit G, Da Fonte DF, Avramescu RG, Premchandrar A, Bagdany M, Xu H, Bensinger D, Stubba D, Schmidt B, **Matouk E, Lukacs GL.**
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