



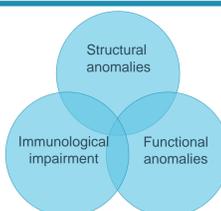
Novel imaging-supported trisomic mouse model of RSV infection for immunological studies of respiratory infections in the context of Down syndrome

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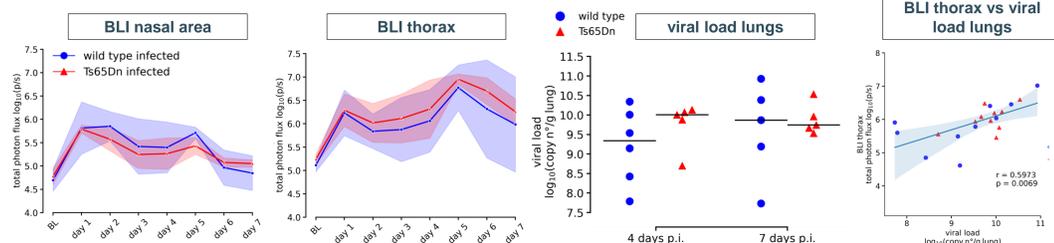
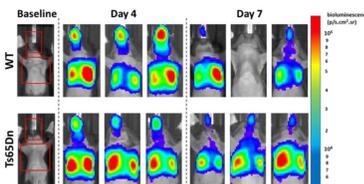
Importance of the study

- Individuals with Down syndrome are prone to severe and recurrent lower respiratory tract infections with RSV [1].
- Down syndrome-related structural, functional, and immunological anomalies contribute to infection severity [2].
- We characterized an imaging-based human RSV Ts65Dn Down syndrome mouse model, to longitudinally visualize the infection, pathology, and immune response.



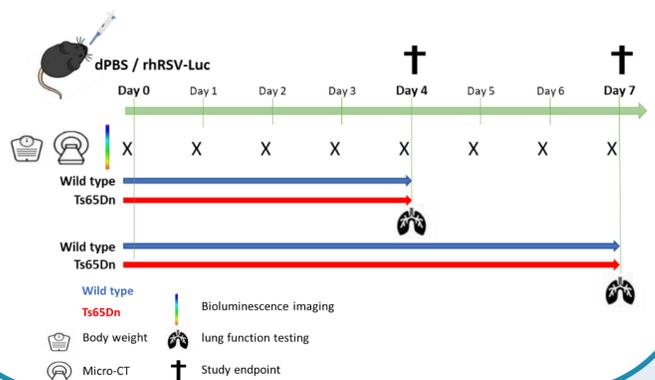
Infection development

- BLI enables **non-invasive spatiotemporal whole-body imaging** of RSV infection development.
- Non-invasive **BLI results correlated** with invasive **RT-qPCR** results.
- Ts65Dn and wild-type had **similar infection development**.
- Two wild-type mice** showed signs of viral clearance on BLI (BLI signal \leq baseline) 7 days post-infection.



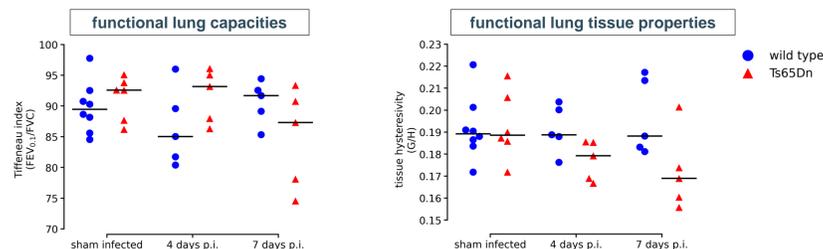
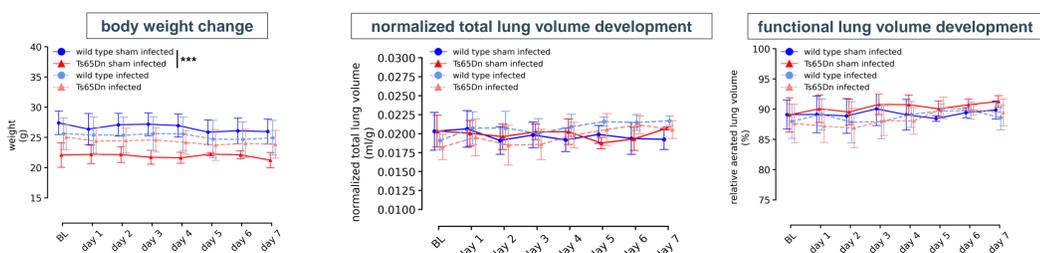
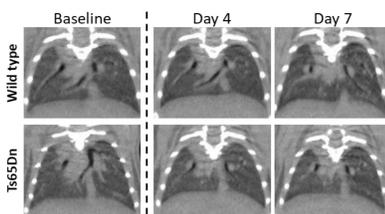
Methodology

- 5-6 weeks old Ts65Dn mice and euploid littermates
- Intranasally challenged with a **human recombinant RSV luciferase encoding strain** [3]
- Daily **non-invasive imaging** of disease progression
- Infectious development: **bioluminescent imaging (BLI)**
- Pathology development: **micro-computed tomography (μ CT) + automated delineation** of functional lung volumes (convolutional neural network).
- Endpoint measurements:** viral load lungs (RT-qPCR), invasive lung function measurement (FlexiVent), bronchoalveolar lavage (BAL) cell count, flow cytometry of the lungs and spleen.



Pathology development

- Infected mice had **no signs of pneumonia**.
- Functional lung volumes** were **unaffected** during the course of the infection.
- Lower initial body weight** of sham-infected Ts65Dn mice corresponded with **lower total lung volumes**.
- Non-invasive μ CT results corresponded with invasive **lung function capacity and tissue property** measurements (Tiffeneau index and tissue hysteresivity).



Conclusion

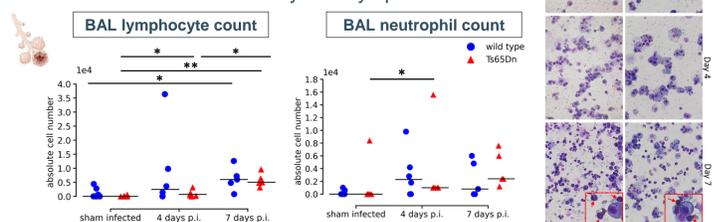
- We reported the **first multimodal-imaging supported animal model** to study lower respiratory tract infections in the context of Down syndrome.
- The **Ts65Dn mouse model** showed similar infection and pathology development as wild-type mice with signs of a **less efficient viral clearing and humoral response**.
- Our model has **potential for prophylactic and immunological studies** in the context of lower respiratory tract infections in Down syndrome.

References:

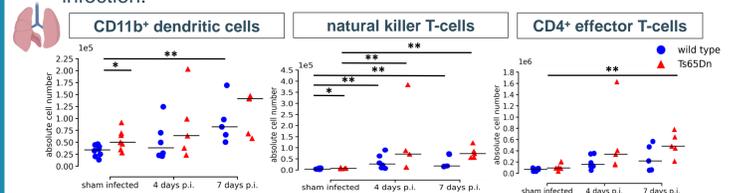
- Sánchez-Luna M, Medrano C, Lirio J, the RISK-21 Study Group. Down syndrome as risk factor for respiratory syncytial virus hospitalization: A prospective multicenter epidemiological study. *Influenza Other Respi Viruses* 2017; 11: 157–164.
- K. L. Colvin and M. E. Yeager, 'What people with Down Syndrome can teach us about cardiopulmonary disease', *European Respiratory Review*, vol. 26, no. 143, Mar. 2017, doi: 10.1183/16000617.0098-2016.
- M.-A. Rameix-Welti et al., 'Visualizing the replication of respiratory syncytial virus in cells and in living mice', *Nat Commun*, vol. 5, no. 1, Art. no. 1, Oct. 2014, doi: 10.1038/ncomms6104.

Immunological response

- RSV infection resulted in increased **lymphocyte recruitment in the airways**.
- Ts65Dn mice** had increased **neutrophil recruitment in the airways** 4 days post-infection.



- Sham-infected Ts65Dn mice** had increased **pro-inflammatory CD11b⁺ dendritic cells** and natural killer T-cells (NKT) numbers.
- During infection **both genotypes** had upregulated **NKT-cell numbers** and an efficient **CD8⁺ effector T-cell response**.
- An increased CD4⁺ effector T-cell response in **Ts65Dn mice**, suggests a more **extensive genotype-related T-cell response** following infection.



- Absence of myeloid response in the spleen of Ts65Dn mice, suggest a **local myeloid response in Ts65Dn lungs**.
- Wild-type mice** had increased **B-cell and CD8 T-cell numbers** 4 days post-infection, resulting in genotype related CD4/CD8 T-cell differences.
- Wild-type mice** had a more **effective humoral response** and a more **efficient viral clearing**.

