

Result creation

Create result

CREATE A NEW RESULT

NAME

1

DESCRIPTION

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SUBMITCANCEL

1. Enter the name of the result.
2. Enter the description of the result (add and format text, images, videos, links...). The description should be a few paragraphs long. All relevant references should be added directly at the end of the section. See help section for references formatting guidelines.

Result information

BALF PROTEOME VS. LUNG TRANSCRIPTOMICS ANALYSIS

[← Back to study](#)

RESULT INFORMATION ⓘ

[Edit ↗](#)

NAME

BALF PROTEOME VS. LUNG TRANSCRIPTOMICS ANALYSIS

DESCRIPTION

We show examples below of single node NPAs (so-called backbone NPAs) of typical inflammatory mediators and their corresponding BALF analyte measurements for the macrophage activation-related entities IL-6 and tumor necrosis factor (TNF) and for the epithelial proinflammatory signaling-related entities IL-1R and VCAM-1. The trends of a marked enhancement of backbone NPAs in the 3R4F group and their decrease upon cessation or switching (left panel) were consistently reflected by the amounts of corresponding proteins in the BALF (right panel, also shown as changes relative to sham). Interestingly, the NPA approach was more sensitive than the BALF measurement, highlighting weak perturbations of IL-6- and TNF signaling in the pMRTP group and residual perturbations of IL-6, TNF α , IL-1R, and VCAM-1 in the cessation/switching groups at month 7 that were not reflected in the BALF measurements.

1. Edit the result details. The description should be a few paragraphs long and can contain images, videos, links... All relevant references should be added directly at the end of the section.

Result creation

Results elements

RESULT ELEMENTS

Protocols 1	Datasets 2	Endpoints 3	Tested sample type 4	Diseases & pathways 5
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Select the protocols to be associated with this result. You can associate protocols that have been created separately (through your dashboard).

PROTOCOLS
None

1. Select the protocols to be associated with this result. You can associate protocols that have been created separately (through your dashboard).
2. Select the datasets to be associated with this result. You can associate datasets that have been created separately (during the creation/update of a study).
3. Select the endpoint(s) that has/have been measured to obtain this result. If the result's associated endpoint is not listed below, contact the admin to get it added. It is important to thoroughly select the endpoint as they will be used as filters when searching on the platform.
4. Select the sample type(s) that has/have been tested to obtain this result. If the result's associated sample type is not listed below, contact the admin to get it added. It is important to thoroughly select the sample type(s) as they will be used as filters when searching on the platform.
5. Select the diseases and pathways to be associated with this result. If the result's associated disease or pathway is not listed below, contact the admin to get it added. It is important to thoroughly select the disease(s) and pathway(s) as they will be used as filters when searching on the platform.