

TraIN:

**Translating knowledge of
cell-to-cell communication molecules
from Immunology to Neuroscience
with RNAseq data**

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Team Lead

Lucia Guerri → Mom of scientific idea

The Professionals

Miranda Darby → Mentor and Review Committee

Amanda Bell → Dataset magician

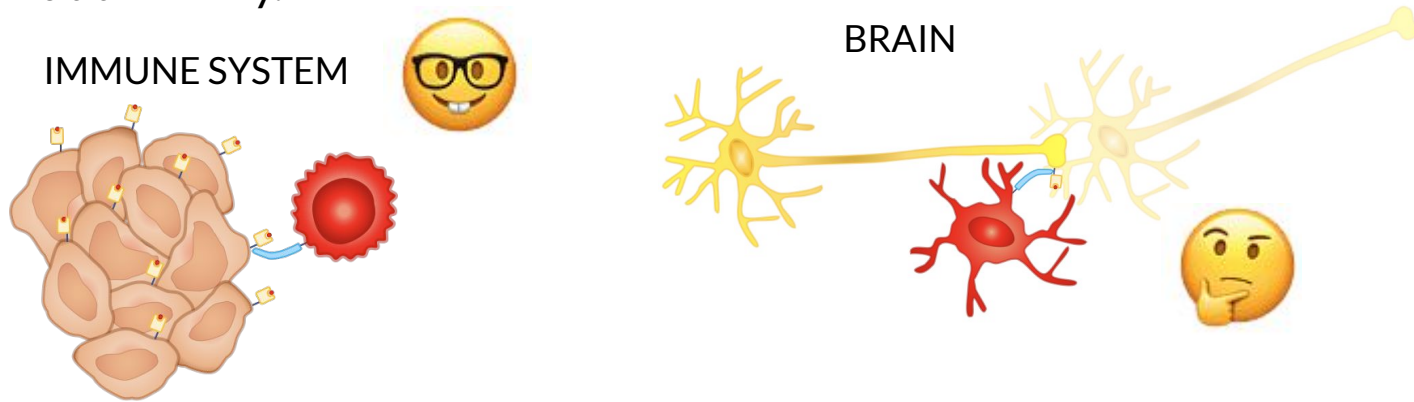
Saba Nafees → Mathematics magician & Writer

Jingwen Gu → Stats magician & Sysadmin

Van Truong → Data & Scientific Illustrator magician

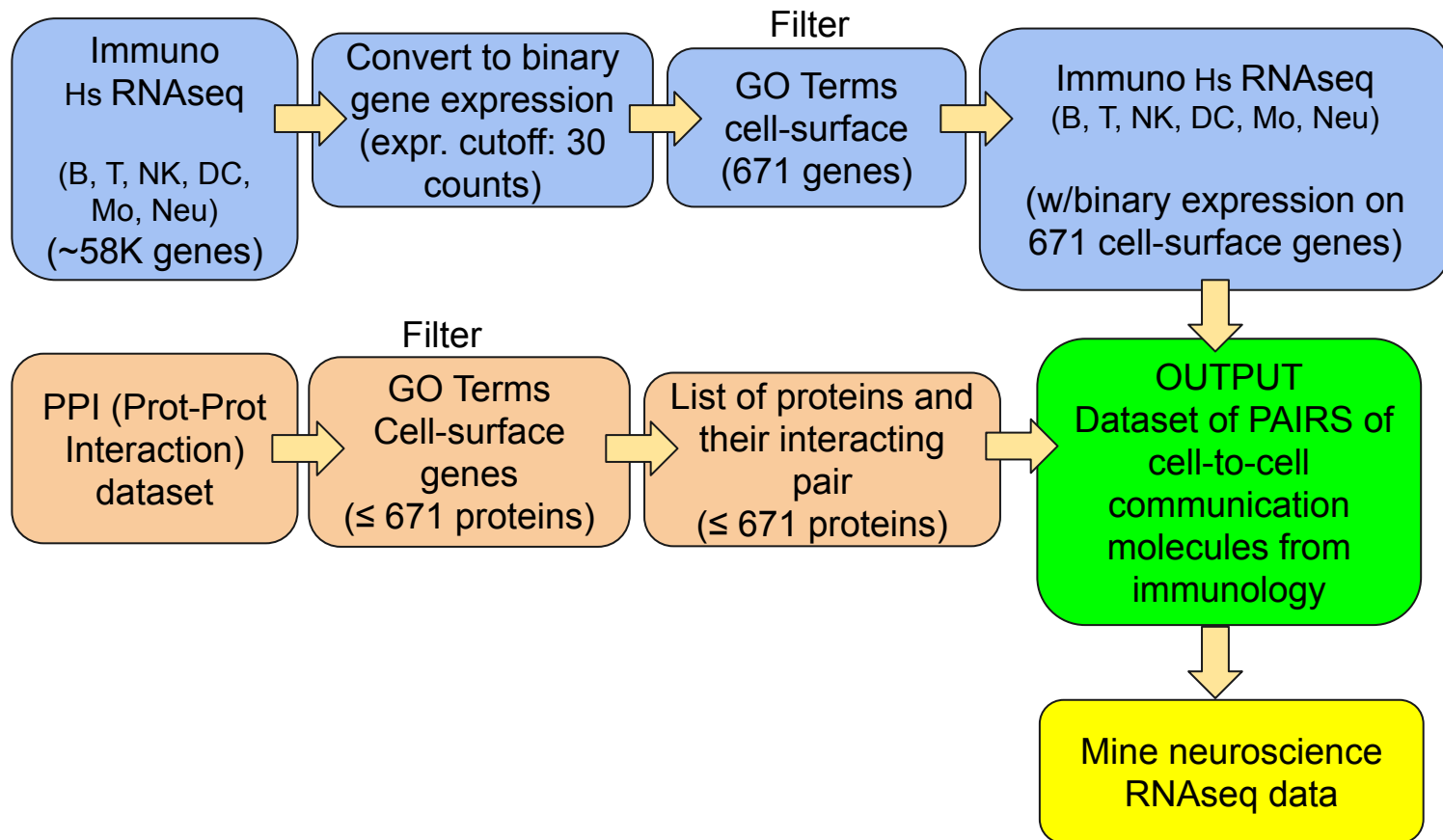
WHY?

- The study of cell-to-cell communication molecules has been particularly strong in immunology, while remaining largely understudied in brain.
- The same pairs of communication molecules are employed by several tissues throughout the body.



- By translating knowledge of cell-to-cell communication molecules (and their conserved intracellular signaling) from immunology to neuroscience with RNAseq data, we can accelerate hypothesis generation in neuroscience.

WORKFLOW:






GOALS:

Hackathon:

Day 1:

- ✓ • **Brainstorm** about best **strategies** to address the scientific question (“Translate knowledge of cell-to-cell communication molecules from immunology to neuroscience”)
- ✓ • **Break down** the theoretical project into concrete technical pipeline
- ✓ • Identify best RNAseq human and murine **datasets**
- ✓ • Define **pipeline** to generate a **database** of “**PAIRS of cell-to-cell communication molecules from immunology**”

Day 2:

-    • Generate a **database** of “**PAIRS of cell-to-cell communication molecules from immunology**”
- ✓ • **Brainstorm** about
 - **Scoring system** for supervised pipeline
 - **Machine-learning** for partially-unsupervised tool (adapt WGCNA?)

Day 3:

-    • Generate working prototype of **supervised pipeline**

Stretch:

- Generate working prototype of **machine-learning partially-unsupervised tool**