

Vaccine Large Language Model (VaxLLM)

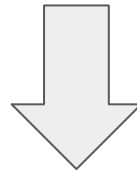
Presenter: Laurel (Xingxian) Li

12/09/2024

Motivation

The Challenge

- Development of database (ex. ontologies) needs manual annotation
- Rapid increase in scientific literature overwhelms manual annotation efforts.
- Annotated information is essential to advance vaccine development



Urgent need to automate the vaccine annotation process from resources

VIOLIN database

- vaccine research data curation and storage
- include vaccine candidates developed against various pathogens



<https://violinet.org/>

B. abortus DNA vaccine pcDNA-SOD

Vaccine Information

- **Vaccine Ontology ID:** [VO_0000018](#)
- **Type:** DNA vaccine
- **Antigen:** *B. abortus* Cu/Zn Superoxide dismutase ([Munoz-Montesino et al., 2004](#)).
- **SodC from *B. abortus* strain 2308 gene engineering:**
 - **Type:** DNA vaccine preparation
 - **Description:** *B. abortus* sodC was subcloned into the expression vector pcDNA3 ([Munoz-Montesino et al., 2004](#)).
 - **Detailed Gene Information:** Click [Here](#).
- **Vector:** pcDNA3 ([Munoz-Montesino et al., 2004](#))
- **Preparation:** Recombinant plasmid pBAII-3, containing the gene for *B. abortus* Cu-Zn SOD (*sodC*), and its own promoter, was initially obtained from a pUC9 genomic library of *B. abortus* strain 2308. A 1.1-kb fragment containing the *sodC* gene and its promoter sequences was excised and ligated into the expression vector pcDNA3 downstream of the cytomegalovirus promoter. The resulting plasmid was designated pcDNA-SOD. A colony of *E. coli* containing pcDNASOD was cultured and used for large-scale plasmid DNA isolation. The DNA was resuspended in PBS at a final concentration of 1 mg/ml. The pcDNA-SOD plasmid construct was verified by restriction digestion and by sequencing of the complete insert ([Munoz-Montesino et al., 2004](#)).
- **Description:** Cu-Zn superoxide dismutase (SOD) is one of the protective immunogens of *Brucella abortus*. Intramuscular injection with a plasmid DNA carrying the SOD gene (pcDNA-SOD) was able to induce a protective immune response in mice ([Munoz-](#)

LLM Annotation Focus:

using PubMed abstract for itemized vaccine annotation

1. Vaccine type (live attenuated, recombinant, etc)
2. Vaccine antigen
3. Vaccine formulation (antigen, protein, gene, adjuvant)
4. Host species use as a specific lab animal model
5. Experiment used to investigate the vaccine

From Baseline Llama 3 to VaxLLM

Data: PMID: 38257995 {Abstract chunk cut here}

Vaccine Introduction: Efforts are underway to develop an improved brucellosis vaccine, and possibly a universal vaccine.

Live attenuated

Vaccine Type: Mutant vaccines (znBM-lacZ and znBM-mCherry)

Vaccine Antigen: Brucella melitensis *Brucella melitensis* mutants znBM-lacZ znBM-mCherry

Vaccine Formulation: Not specified in the provided text

Host Species Used as Laboratory Animal Model: Mice

Experiment Used to investigate the vaccine:

1. In vitro infection of bone-marrow-derived macrophages and human TF-1 myeloid cells with znBM-lacZ and znBM-mCherry mutants.
2. Mice were vaccinated once by the intraperitoneal route with either znBMZ or B. abortus S19 vaccine, followed by assessment of splenic brucella colonization and T cell immunity.

Llama 3 8B direct usage for paper PMID 38257995

Incorrect or missing information

VaxLLM

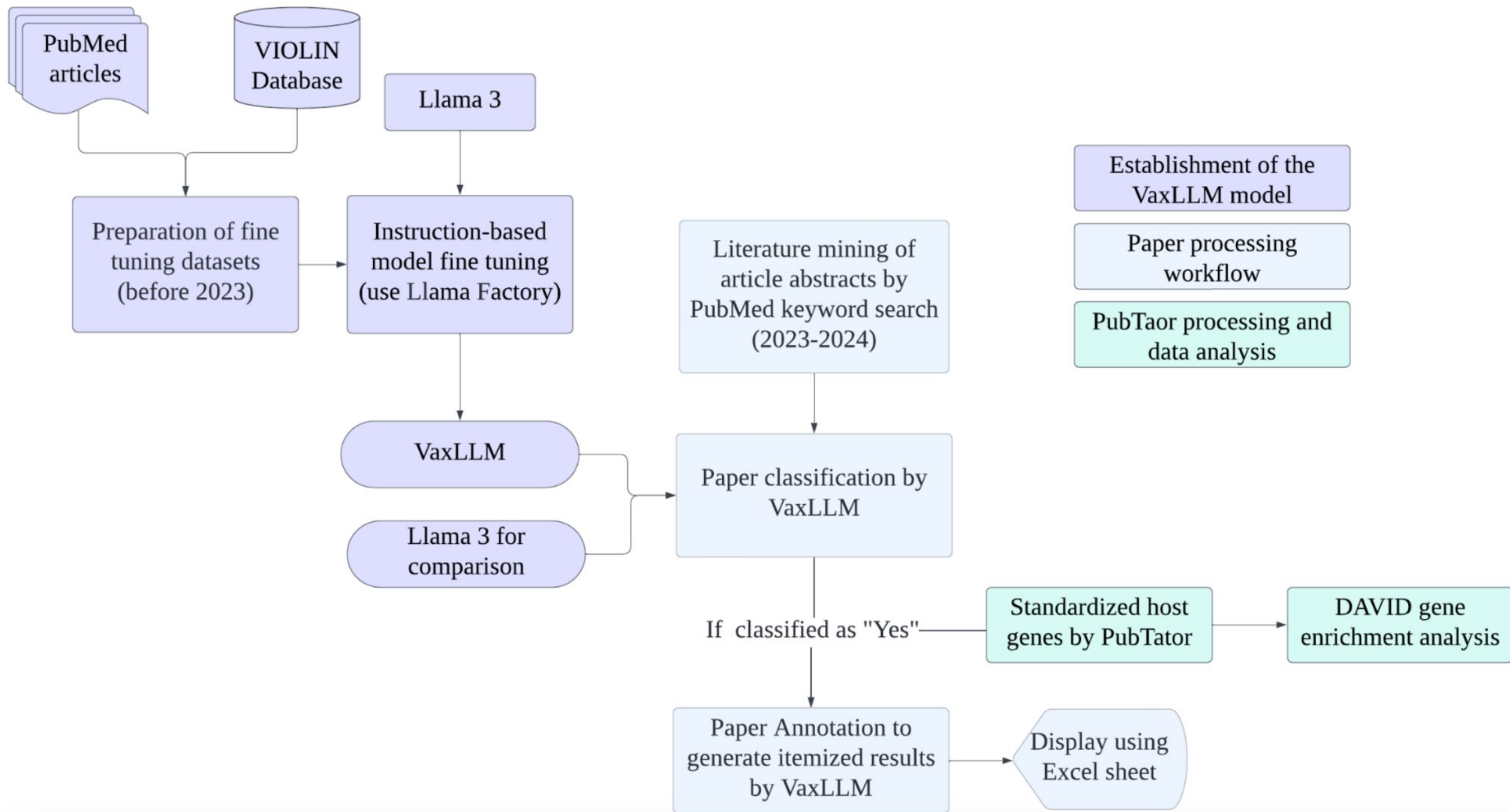
- A fine-tuned Large Language Model derived from Llama3-8B

The whole project is to develop a pipeline to:

perform **classification** and **standardized annotation** of *Brucella* vaccine articles using VaxLLM

Classification Focus:

filter the relevant articles containing specific vaccine formulation



Development of VaxLLM - Model fine tuning

Let the model know what is correct annotation → VIOLIN !



Classification training data:

- Positive Examples: 50 scientific articles abstracts (From VIOLIN reference)
- Negative Examples: 100 articles abstracts (From PubMed before 2023)

Annotation training data:

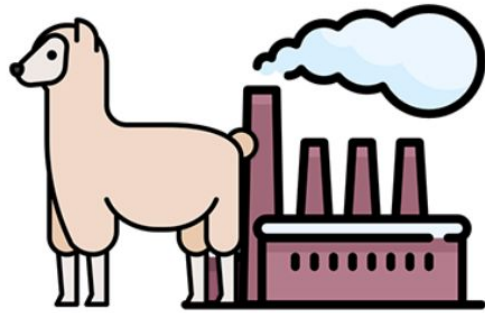
- The standardized annotation from VIOLIN

Training data format

Alpaca Format:

- **Instruction:** our instruction (e.g., "Is this article about a *Brucella* vaccine?..." or "Extract the following details...").
- **Input:** The abstract of the article.
- **Output:** The classification result (Yes/No) or detailed vaccine annotation

Fine-tuning process: Llama Factory



LLaMA-Factory





Easy and Efficient LLM Fine-Tuning



<https://github.com/hiyouga/LLaMA-Factory>









A package for fine-tuning of hundreds of LLMs

adopted LoRA, 4 bits for VaxLLM, T4 GPU

 **Xingxian123/VaxLLM**   like 1

 Question Answering  Transformers  Safetensors llama text-generation llama-factory  text-generation-inference

 Inference Endpoints  License: mit

 **Model card**  Files  Community  Settings   Train  Deploy  Use this model

 Edit model card

 **Gated model** You have been granted access to this model

Model Card for Model ID

VaxLLM (Vaccine Large Language Model) is a fine-tuned Llama-3 model to automatically perform the classification and annotation of vaccine-related articles, using Brucella vaccines as a case study.

Downloads last month
17



 **Safetensors** 

Model size 8.03B params Tensor type BF16 

Inference Examples 

PubMed literature Mining

PubMed Advanced Search Builder

[User Guide](#)

Add terms to the query box

Date - Publication to **AND**

Query box

Search

- from 2023 to 2024
- "*Brucella* vaccine" term keyword search
- identify 148 papers

(Brucellosis vaccine) AND (("2023/01/01"[Date - Publication] : "3000"[Date - I] **Search**

[Advanced](#) [Create alert](#) [Create RSS](#) [User Guide](#)

Sort by:

145 results

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Prompt Engineering

1. Vaccine Classification:

Using the following data: '**{Abstract information}**', is this article about a brucella vaccine? To classify an article as being about a brucella vaccine, you must successfully extract at least some information about the vaccine formulation. This includes details such as the antigen, protein, gene, adjuvant, or vaccine platform mentioned in the abstract.

2. Vaccine Annotations:

Extract the following details using the given data: '**{Abstract information}**':
Vaccine Introduction, Vaccine Antigen, Vaccine Type, Vaccine Formulation, Host Species Used as Laboratory Animal Model, Experiment Used to investigate the vaccine
Ensure each response is based solely on the provided data. Ensure the response is formatted as follows:

Response:

Vaccine Introduction:

Vaccine Type:

Vaccine Antigen:

Vaccine Formulation:

Host Species Used as Laboratory Animal Model:

Experiment Used to investigate the vaccine:

Classification results

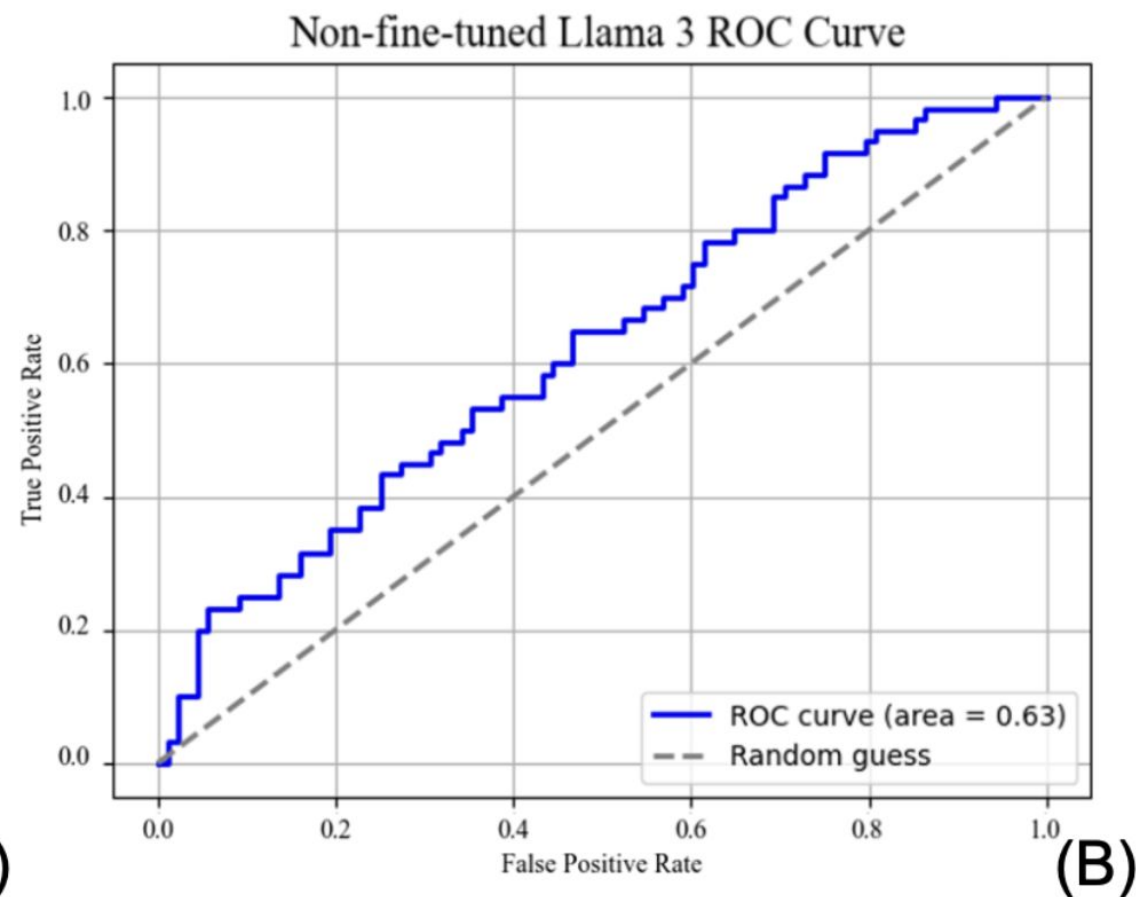
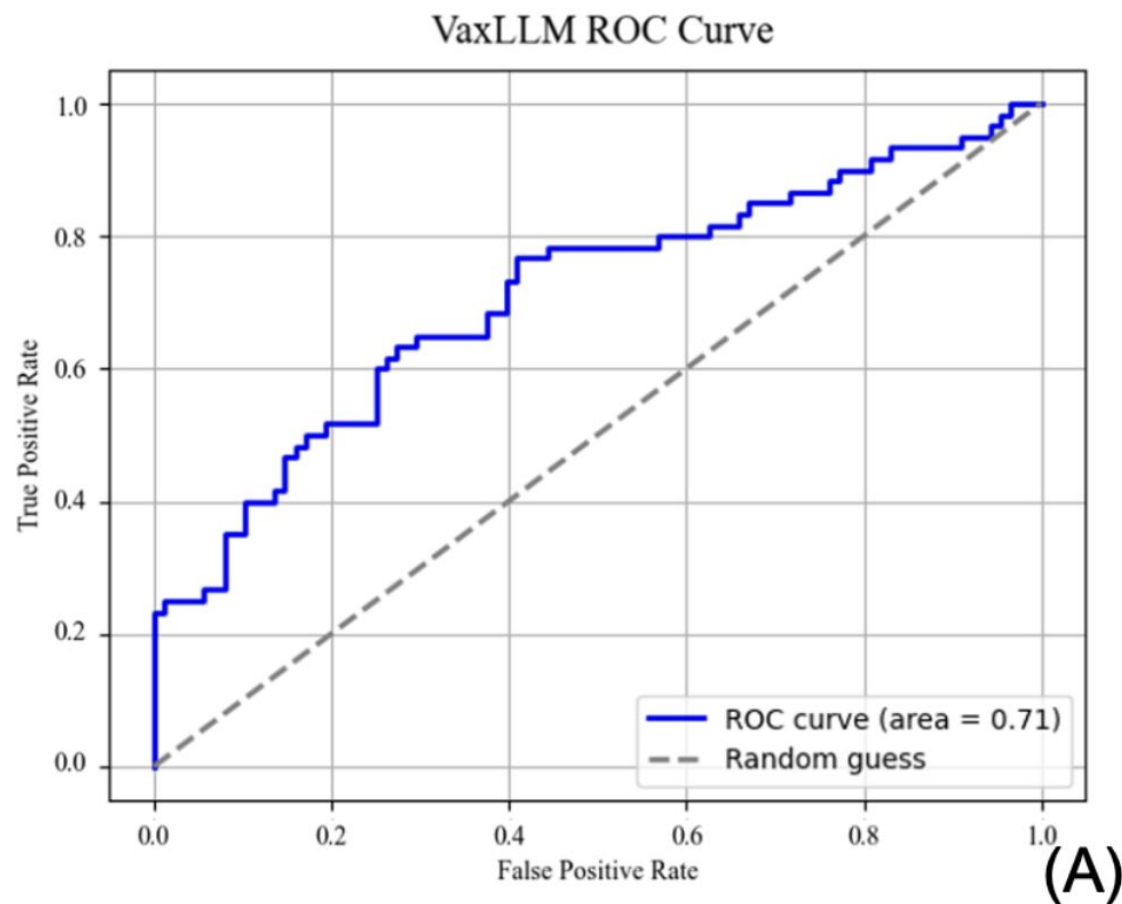
VaxLLM after finetuning

True Label	Predicted Label	
	Predicted No	Predicted Yes
Actual No	84	7
Actual Yes	0	60

Direct usage from Llama 3

True Label	Predicted Label	
	Predicted No	Predicted Yes
Actual No	60	29
Actual Yes	3	56

Classification results



VaxLLM annotation output

Data: PMID: 38257995 {Abstract chunk cut here}

Vaccine Introduction: This live attenuated vaccine is based on *Brucella melitensis* mutants znBM-lacZ (znBMZ) and znBM-mCherry (znBM-mC), designed to confer systemic immunity against virulent *B. melitensis* 16M challenge.

Vaccine Type: Live attenuated vaccine.

Vaccine Antigen: *Brucella melitensis* mutants znBM-lacZ (znBMZ) and znBM-mCherry (znBM-mC).

Vaccine Formulation: The vaccine consists of the *Brucella melitensis* mutants znBM-lacZ (znBMZ) and znBM-mCherry (znBM-mC).

Host Species Used as Laboratory Animal Model: Mice.

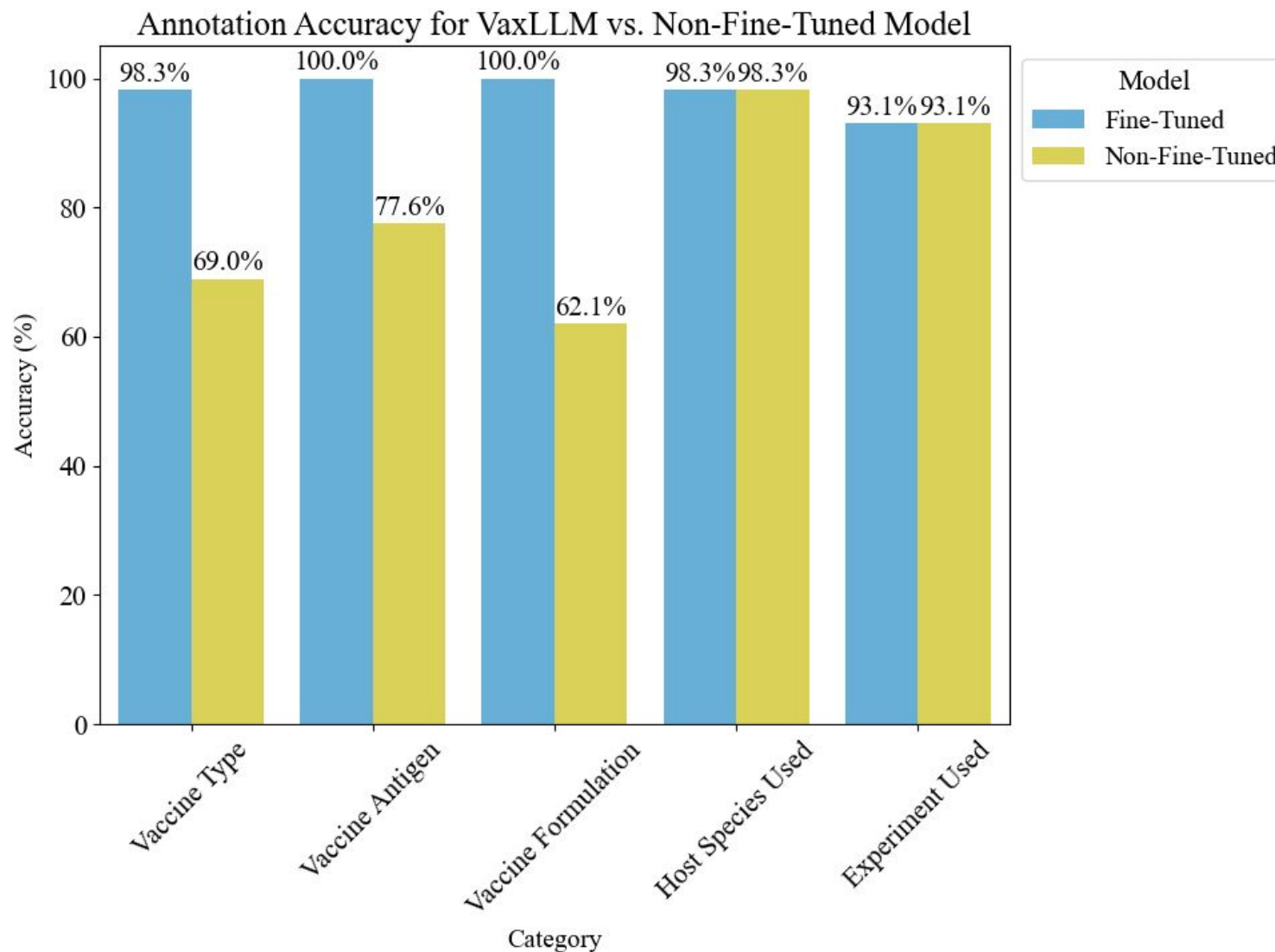
Experiment Used: Bone-marrow-derived macrophages and human TF-1 myeloid cells were infected with both mutants to assess their attenuation. Mice were vaccinated once by the intraperitoneal route with either znBMZ or *B. abortus* S19 vaccine, and their immune response was evaluated by measuring IFN- γ ⁺ CD4⁺ and CD8⁺ T cells, as well as CD4⁺ and CD8⁺ effector memory T cells.

VaxLLM usage for paper PMID
38257995

correct information

VaxLLM annotation accuracy

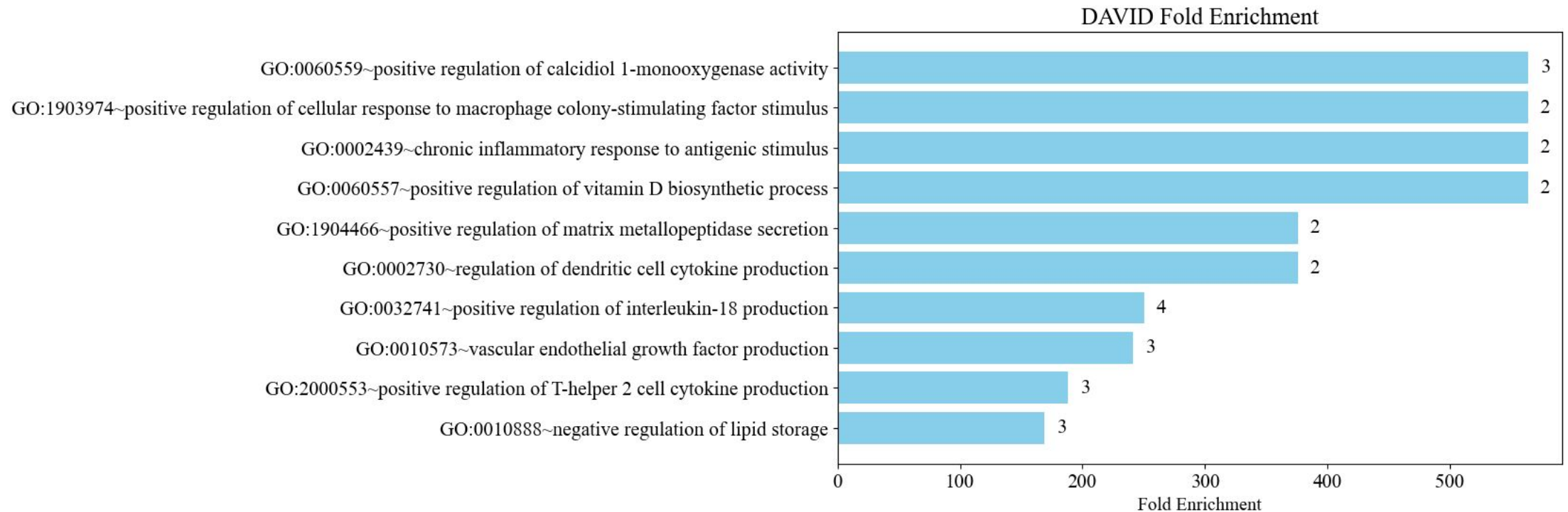
Overall
accuracy:
97.9%



Potential downstream use case

Brucella vaccines data-analysis

- demonstrate here as gene enrichment analysis



PubTator usage

•PubTator — uses Name Entity Recognition (NER) to extract the following information:

- Genes (both antigen and host genes)
- Protein
- Chemicals, if relevant
- Species mentioned

Pubtator result example

```
37515088|t|Characterization of Brucella abortus Mutant A19mut2, a Potential DIVA Vaccine Candidate
37515088|a|BACKGROUND: Brucella abortus is the main causative agent for bovine brucellosis. B. a
37515088      111      129      Lipopolysaccharide      Chemical      MESH:D008070
37515088      280      298      Brucella infection      Disease MESH:D002006
37515088      336      354      lipopolysaccharide      Chemical      MESH:D008070
37515088      356      359      LPS      Chemical      MESH:D008070
37515088      674      691      acetyltransferase      Gene      20468107
37515088      798      814      O-polysaccharide      Chemical      -
37515088      880      883      LPS      Chemical      MESH:D008070
37515088      1328     1337     IFN-gamma      Gene      15978
37515088      1342     1346     IL10      Gene      16153
37515088      1418     1421     LPS      Chemical      MESH:D008070
```

Gene enrichment analysis

- VaxLLM classify 60 articles as yes
- Run the Pubtator of all these articles
- Identify and extract the gene list of 37 standardized for DAVID gene enrichment analysis.

Meaning of VaxLLM

- Automates the annotation of *Brucella* vaccine literature.
- High accuracy and standardized outputs support database integration.
- May realize the integration of thousands of vaccine information in VIOLIN

Future direction and limitation

- Use *Brucella* vaccines as demo to text methodology, may implement more vaccines
- Full text paper Vs. abstract
- may provide standardization (ex. map to ontology)

Data Availability

VaxLLM data and sample code:

<https://github.com/xingxianli/VaxLLM>

Model availability:

<https://huggingface.co/Xingxian123/VaxLLM>

Q & A

