

PhenOM Inference API User Guide

This user guide summarizes how to use the PhenOM Inference API based on the latest spec.

Base URL & Reference

- **API base URL**

`https://phenom-api.om1.com`

Authentication

- **Auth method:** OAuth2 Bearer token (Auth0)
- **Header format:**

```
Authorization: Bearer <jwt>
```

Your JWT determines which tenant and resources you can access.

Inference Options

You can run predictions in two ways:

- **Single-patient inference**
 - **Best for:** point-of-care workflows and interactive apps
 - **Pattern:** submit a single `patient_history` JSON → get a `job_id` → poll job for results
 - **Endpoint:** `/v1/job/from-patient`
 - **Multiple patients:** use the **batch** workflow (`/v1/batch`) instead of a multi-patient real-time endpoint
- **Batch inference**
 - **Best for:** large populations, scheduled risk assessments, research & analytics
 - **Pattern:** upload files → finalize → (optionally) validate → start job → poll → fetch results
 - **Endpoints:** `/v1/batch`, `/v1/batch/{batch_id}/upload`, `/v1/batch/{batch_id}/finalize`,
`/v1/batch/{batch_id}/start`, plus job endpoints under `/v1/job/...`

Quick Start – Single Patient

Use this for a single patient at a time.

Request: create job from patient

```
curl -X POST https://phenom-api.om1.com/v1/job/from-patient \
-H "Authorization: Bearer <jwt>" \
-H "Content-Type: application/json" \
-d '{
    "outcome_ids": [
        "v0p3_65plus_fixed_anchor_2023_JanToDec_04012025_any_time_hospitalization_future_1mon"
    ],
    "patient_history": {
        "patient": {
            "patient_id": "abc123",
            "birth_date": "1964-05-18",
            "sex": "F",
            "zip3": "021"
        },
        "lab_results": [
            {
                "lab_date": "2025-07-03",
                "loinc": "718-7",
                "value_abnormal": "N"
            }
        ],
        "diagnoses": [
            {
                "diagnosis_date": "2020-01-01",
                "code": "I10",
                "code_type_name": "ICD10"
            }
        ],
        "procedures": [
            {
                "procedure_date": "2024-12-15",
                "code": "93000",
                "code_type_name": "CPT"
            }
        ],
        "medications": [
            {
                "medication_start": "2023-03-01",
                "code": "C09AA02",
                "code_type": "ATC"
            }
        ]
    },
    "idempotency_key": "idem-b-20251107-abc123"
}'
```

Immediate response (example):

```
{
    "job_id": "job_01J90Q2TTM",
    "status": "QUEUED"
}
```

Poll for final result

```
# Poll job until SUCCEEDED
curl https://phenom-api.om1.com/v1/job/job_01J90Q2TTM \
-H "Authorization: Bearer <jwt>"
```

Then fetch results:

```
curl "https://phenom-api.om1.com/v1/job/job_01J90Q2TTM/results?page_size=1000" \
-H "Authorization: Bearer <jwt>"
```

Example final result payload:

```
{
  "items": [
    {
      "patient_id": "abc123",
      "outcome_id": "v0p3_65plus_fixed_anchor_2023_JanToDec_04012025_any_time_hospitalization_fut",
      "probability": 0.7421,
      "prob_upper_95_percent_bound": 0.7934,
      "prob_lower_95_percent_bound": 0.6912,
      "relative_probability": 2.18,
      "rel_upper_95_percent_bound": 2.36,
      "rel_lower_95_percent_bound": 2.02,
      "bin_id": 9,
      "num_bins": 100
    }
  ],
  "next_page_token": null,
  "total_rows": 1
}
```

Quick Start – Batch

End-to-end example: upload data, run a job, fetch results.

1. Create a batch

```
curl -X POST https://phenom-api.om1.com/v1/batch \
-H "Authorization: Bearer <jwt>" \
-H "Content-Type: application/json" \
-d '{
  "name": "october_data_load",
  "upload_method": "s3_multipart"
}'

# Response (example)
{
  "batch_id": "batch_0199bb07-3b87-7e83-8842-ad9d52a3c472",
  "status": "CREATED"
}'
```

2. Upload patient data

```
# Create upload session for one CSV
curl -X POST https://phenom-api.om1.com/v1/batch/{batch_id}/upload \
-H "Authorization: Bearer <jwt>" \
```

```
-d '{  
    "object_type": "patients",  
    "filename": "patients_2025_10.csv",  
    "mode": "single"  
}'  
  
# Use the presigned URL returned in the response to upload the file.  
# Repeat for other object_types: lab_results, diagnosis, procedures, medications.
```

3. Finalize the batch

```
curl -X POST https://phenom-api.om1.com/v1/batch/{batch_id}/finalize \  
-H "Authorization: Bearer <jwt>"  
  
# Response (example)  
{  
    "batch_id": "batch_0199bb07-3b87-7e83-8842-ad9d52a3c472",  
    "status": "FINALIZED",  
    "manifest_key": "batches/batch_.../manifest.json"  
}
```

4. Start the job

```
curl -X POST https://phenom-api.om1.com/v1/batch/{batch_id}/start \  
-H "Authorization: Bearer <jwt>" \  
-H "Content-Type: application/json" \  
-d '{  
    "outcome_ids": [  
        "v0p3_65plus_fixed_anchor_2023_JanToDec_04012025_any_time_hospitalization_future_1month"  
    ]  
}'  
  
# Response (example)  
{  
    "job_id": "job_01J90Q2TTM",  
    "status": "QUEUED"  
}
```

5. Poll job status

```
curl https://phenom-api.om1.com/v1/job/{job_id} \  
-H "Authorization: Bearer <jwt>"  
  
# When complete (example)  
{  
    "job_id": "job_01J90Q2TTM",  
    "status": "SUCCEEDED",  
    "metrics": {  
        "records_processed": 1245678,  
        "patients_scored": 101234,  
        "patients_excluded": 1123  
    }  
}
```

6. Fetch results

```
# Option 1: Paginated API results
curl "https://phenom-api.om1.com/v1/job/{job_id}/results?page_size=1000" \
-H "Authorization: Bearer <jwt>"

# Option 2: Download full results (manifest with shard URLs)
curl https://phenom-api.om1.com/v1/job/{job_id}/results/download \
-H "Authorization: Bearer <jwt>"
```

Additional job endpoints are available to inspect **exclusions** and **errors**:

- `/v1/job/{job_id}/exclusions`
 - `/v1/job/{job_id}/errors`
-

Security & Compliance

- **HIPAA & BAA:** HIPAA-compliant infrastructure with BAA coverage
- **Tenant isolation:** All resources are scoped to your tenant via JWT claims
- **Encryption:** SSE-S3 at rest; TLS in transit
- **PHI handling:** PHI appears **only** in uploaded data, not in resource names or metadata