

Structured Data Extraction From Free-Text Melanoma Histopathology Reports Using a Locally Deployed Large Language Model

A Proof-of-Concept Study at a Tertiary NHS Melanoma Centre

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1. The Clinical Problem

The Royal College of Pathologists Melanoma structured dataset (G125, 4th ed.) mandates 20 essential fields per excision. These should exist in every first diagnosis histopathology report. However, historical report data is locked in free text and not structured for easy access within the local e-HR environment.

Clinical impact:

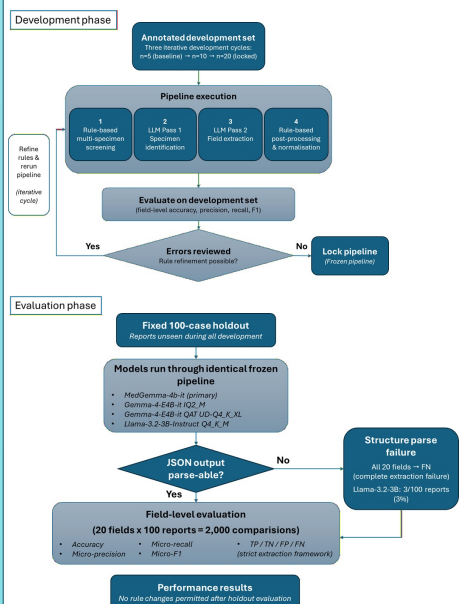
- Pathologist reporting delays due to time spent searching patient history for prognostically significant information
- MDT data extraction is manual, time-consuming, error-prone
- Staging verification and data for audit and research requires manual retrieval

2. The Approach

Feasibility challenge

- ✓ Locally deployed LLM
- ✓ No cloud APIs
- ✓ No data transfer outside the organisation
- ✓ Organisation GPU only
- ✓ DPIA reviewed, IG approved
- ✓ Incorporated within Trust-wide Dermatology Service Evaluation Framework

Pipeline Architecture



Study cohort and evaluation design

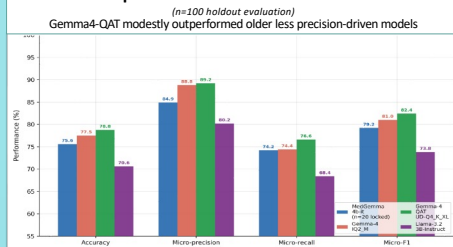
- Cutaneous malignant melanoma reports, UHS
- Pre-established frozen 100-case holdout - unseen during entire development (gold standard test set)
- 20 essential fields = 2,000 field-level comparisons
- 6 model configurations: MedGemma-4b-it (Gemma-3; 3 stages), Gemma-4-E4B-it IQ2_M and QAT, and Llama-3.2-3B
- Metrics: Accuracy • micro-F1 • Cohen's K • PPV/NPV • subtype • workload
- Single annotator gold standard • No fine-tuning

3. Results

Iterative Development – Key Findings

Development set	Accuracy	Observations
n = 5	39.5%	Baseline
n = 10	67.5%	Rule-based additions drove major gains
n = 20	75.6%	Specimen-aware context + ulceration rule addition

Overall performance metrics across all models

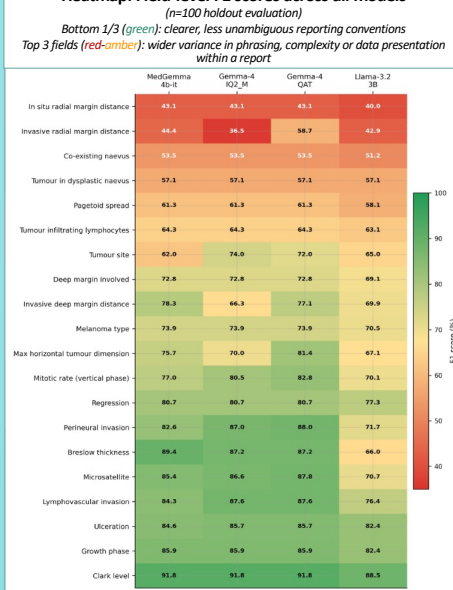


78.8%
Overall field accuracy

87.8%
Micro-precision

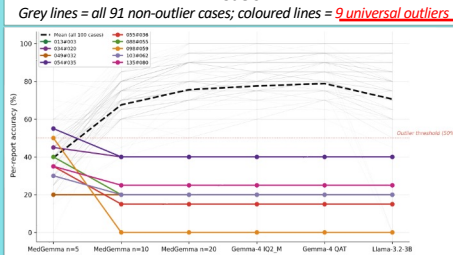
0.621
Mean Cohen's K

Heatmap: Field-level F1 scores across all models



4. Outlier Case Analysis

Accuracy trajectory of universal outlier cases across all six models



A report structure problem - Not a model problem

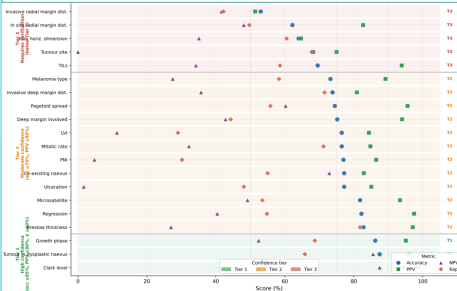
Report Structure	Failure	Cases
Multi-specimen reports	Rule-based multi-specimen screening	4
Supplementary report + amended diagnosis	Additional structure unaccounted for during development	2
External referral / non-UHS format	Model confusion and rule failure	1
Non-standard field format	Model confusion and rule failure	2

84.5%
Overall field accuracy on Gemma-4 QAT
After excluding outliers with universal failures

5. Deployment Field Confidence

Field confidence tier classification – mean metrics across four evaluation models

Tier 1 (green): Accuracy ≥85%, PPV ≥ 90%, K ≥ 60%; Tier 2 (amber): Accuracy ≥50%, PPV ≥ 80%, Tier 3 (red): Below Tier 2 thresholds



The analytical dot plot visualises the distribution of scores for each metric plotted simultaneously per field and simultaneous tier stratification

Field confidence tier classification – current deployment summary

Mean accuracy, PPV, NPV, and Cohen's K across four evaluation models (n=100 holdout)
Cell colours reflect individual cell performance

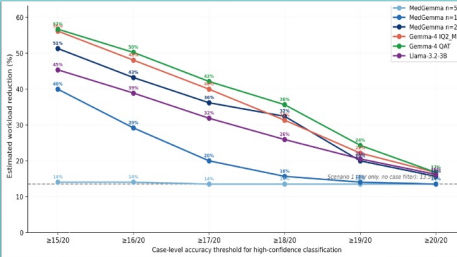
Tier 1 – High confidence				Tier 2 – Moderate confidence				Tier 3 – Requires verification			
Accuracy ≥85%, PPV ≥ 90%, K ≥ 60%				Accuracy ≥50%, PPV ≥ 80%				Below Tier 2 thresholds			
	Accuracy	PPV	K		Accuracy	PPV	K		Accuracy	PPV	K
Field	95%	95%	95%	Invasive structures	85%	85%	85%	WPA	75%	75%	75%
In situ radial margin distance	75%	75%	75%	Regression	75%	75%	75%	Melanoma type	65%	65%	65%
Invasive radial margin distance	75%	75%	75%	Microsatellite	85%	85%	85%	Max vertical dimension	65%	65%	65%
Ulceration	75%	75%	75%	Ulceration	85%	85%	85%	Ulcerated epidermal depth	65%	65%	65%
Co-existing naevus	75%	75%	75%	Regression aspects	75%	75%	75%	Invasive radial margin dist	75%	75%	75%
Tumour in dysplastic naevus	75%	75%	75%	WPA	75%	75%	75%	→ Do not use pathologically unproven measures			
Pagefold spread	75%	75%	75%	Deep margin	75%	75%	75%				
Tumour infiltrating lymphocytes	75%	75%	75%	Regression score	75%	75%	75%				
Tumour site	75%	75%	75%	Regression aspect	75%	75%	75%				
Deep margin involved	75%	75%	75%	Invasive deep margin dist	75%	75%	75%				
Invasive deep margin distance	75%	75%	75%	Melanoma type	75%	75%	75%				
Melanoma type	75%	75%	75%								

Each report field is grouped by how reliably the pipeline can extract them.

The footer of each column advises what to do with the pipeline's output for that group of fields:

- Tier 1: Trust it + occasional spot-checking
- Tier 2: Use it as a starting point + verify each value
- Tier 3: Ignore the pipeline output + manually extract

6. Estimated Workload Reduction



The more accurately the pipeline extracts a patient's report, the less manual checking a clinician needs to do.

The best model currently reduces the number of fields requiring human review by up to 42% at the ≥17/20 threshold (where risk of missing a Tier 3 field is low).

7. Next Steps & Limitations

- 4 structural failure categories identified – priority fix
- 17 of 20 fields require further development
- Tier 3 numeric fields need targeted intervention
- Consider multi-model downstream architecture
- Single annotator gold standard: inter-annotator reliability
- Workload reduction estimates are projections: time-motion study required
- Lentigo maligna subtype underrepresented in dev. cohort – requires oversampling in next dev. set

8. AI Product Development Cycle

