

# 140509\_49.md – AI-Powered Personalized Medicine Platform

**Theme:** AI for Industry, Responsible AI

**Mission:** Integrate genomic (DNA/RNA), clinical (EHR), and lifestyle (wearables, SDoH) data to deliver safe, explainable, privacy-preserving personalized treatment recommendations and drug discovery insights.

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## README (Problem Statement)

**Summary:** Build a platform that integrates genomic data, medical records, and lifestyle information to provide personalized treatment recommendations and drug discovery insights.

**Problem Statement:** Precision medicine requires harmonizing heterogeneous, multi-modal data and generating evidence-based recommendations while ensuring safety, fairness, and regulatory compliance. Create a system that predicts drug response, adverse reactions, and matches patients to trials, with robust privacy.

**Steps:**

- Multi-modal data integration (genomic + EHR + lifestyle)
- Drug response prediction models
- Adverse reaction prediction & DDI analysis
- Evidence-based treatment recommendation engine
- Clinical trial matching & patient stratification
- Privacy-preserving analytics & regulatory compliance

**Suggested Data:** TCGA/ICGC genomics, UK Biobank, MIMIC-III/IV (de-identified EHR), FAERS (adverse events), DrugBank/ChEMBL, wearable datasets, clinical guidelines.

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## 1) Vision, Scope, KPIs

**Vision:** A trustworthy, clinician-centered platform that accelerates precision care, reduces adverse events, and supports ethical AI in healthcare.

**Scope:**

- v1: ETL + harmonization, baseline drug-response model, ADR risk stratification, clinician UI.
- v2: trial matching, DDI engine, explainability & rationale, reporting.
- v3: federated analytics across hospitals, on-device options, RWE feedback loop.

**KPIs:**

- Drug-response AUC  $\geq 0.85$  on held-out cohorts
  - ADR model AUC  $\geq 0.90$ ; NPV  $\geq 0.95$  for high-risk flags
  - Time-to-trial-match  $\geq 70\%$
  - Clinician acceptance of recommendations  $\geq 75\%$
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## 2) Personas & User Stories

- **Oncologist/Cardiologist:** Wants interpretable, guideline-concordant recommendations.
- **Clinical Pharmacist:** Needs DDI/PGx insights (CYP variants).
- **Research Coordinator:** Wants eligible patient lists for trials.
- **Patient:** Expects privacy and understandable justifications.
- **Compliance Officer:** Requires auditability and consent management.

**Stories:**

- US'01: As a clinician, I upload VCF + EHR to get therapy ranking with genomic rationales.
  - US'05: As a pharmacist, I check DDI and PGx contraindications before prescribing.
  - US'09: As a coordinator, I get trial candidates by eligibility (biomarkers, ECOG).
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## 3) PRD (Capabilities)

1. **Data Integration & Harmonization:** FHIR/HL7 ingestion; VCF/FASTQ; wearable APIs; SDoH; normalize to OMOP CDM + patient-centric schema.
  2. **Drug Response Prediction:** Multi-task models using genomics (variants, expression), labs, phenotypes.
  3. **Adverse Reaction & DDI:** Predict ADR probability and drug-drug-gene interactions; PGx knowledge (CPIC).
  4. **Recommendation Engine:** Evidence-based ranker blending ML predictions with guidelines (NCCN/CPG), contraindications, patient prefs.
  5. **Trial Matching:** NLP parser of inclusion/exclusion; match and rank by distance to criteria; cohort builder.
  6. **Explainability & Rationale:** SHAP/PERT/attention maps; guideline citations; genomic variant evidence.
  7. **Privacy & Compliance:** Consent registry, de-ID, encryption, access controls, audit trails; federated learning.
  8. **Clinician UX:** Case timeline, what-if simulations, printable summaries.
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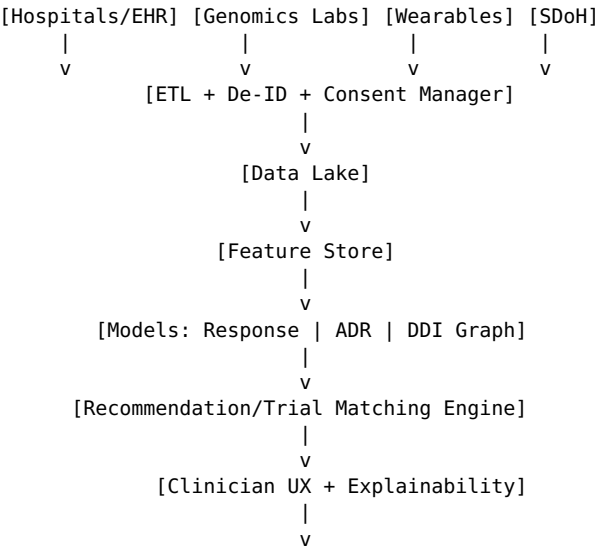
## 4) FRD (Functional Requirements)

- **ETL:** VCF to annotated variants (ClinVar, gnomAD); FHIR resources (Observation, Medication, Condition, Procedure); wearable time-series resampling.
  - **Feature Store:** patient embeddings (genotype, phenotype, labs, vitals), treatment history, outcomes; temporal windows.
  - **Models:** drug-response (per-drug head); ADR classifier; DDI graph model (drug-drug-gene).
  - **Recommender:** constrained optimizer combining efficacy, safety, guideline weights, patient prefs, costs.
  - **Trial NLP:** BERT-based criterion extraction; code eligibility functions.
  - **Cohort Ops:** filter by ICD/SNOMED, variants (e.g., EGFR L858R), lab thresholds; export to REDCap/CTMS.
  - **Explainability:** patient-specific SHAP, variant annotations, literature snippets.
  - **Reporting:** PDF/HL7 messages; audit logs of decisions.
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## 5) NFRD

- **Latency:** case query < 2 s P95; batch scoring overnight.
  - **Reliability:** 99.9% availability; graceful degradation offline.
  - **Privacy/Security:** AES-256 at rest, TLS 1.3, PHI masking, break-glass with dual approval.
  - **Compliance:** HIPAA, GDPR, 21 CFR Part 11; model change management (GxP).
  - **Fairness:** disparate impact monitoring; subgroup performance floors.
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## 6) Architecture (Logical)



## 7) HLD (Key Components)

- **ETL Pipelines:** airflow/nifi; HGVS normalization; sample QC metrics; ICD/CPT/SNOMED mapping.
  - **Feature Store:** Feast; time-aware joins; survival/longitudinal features.
  - **Drug Response:** transformer encoders for variants + clinical; survival head for time-to-progression.
  - **ADR/PGx:** model + knowledge graphs (CPIC/PharmGKB) for drug-gene rules.
  - **DDI Graph:** hetero-GNN on drug-drug-gene edges; predict interaction severity.
  - **Trial NLP:** criteria parser + boolean DSL; fuzzy tolerance ranges.
  - **Recommender:** multi-objective (efficacy, safety, QoL, cost); constraints for allergies, pregnancy, renal/hepatic function.
  - **Explainability:** local SHAP; variant evidence cards; cite guidelines.
  - **Federated Learning:** TensorFlow Federated/Flower; secure aggregation.
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## 8) LLD (Selected)

### ADR Risk Calculation:

$\text{risk\_total} = w1 * \text{model\_prob} + w2 * \text{DDI\_score} + w3 * \text{PGx\_rule} + w4 * \text{history\_flag}$

### Eligibility Function Example:

$\text{- age} \geq 18 \text{ AND ECOG in } \{0,1\} \text{ AND EGFR\_L858R} == \text{true AND creatinine\_clearance} \geq 50$

### Recommender Objective:

Maximize  $U = \hat{I}^+ * \text{Efficacy} - \hat{I}^+ * \text{ADR} - \hat{I}^+ * \text{Cost} + \hat{I}^- * \text{PreferenceMatch}$ , subject to contraindications and DDI hard constraints.

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## 9) Pseudocode (Patient Flow)

```
patient = harmonize(FHIR, VCF, wearables)
X = featurize(patient)
resp = model_response(X)
adr = model_adr(X)
ddi = gnn_ddi(patient.meds, patient.genotype)
recs = optimize(resp, adr, ddi, guidelines, prefs)
trials = match_trials(patient, protocols)
return report(recs, trials, explanations)
```

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## 10) Data & Evaluation

- **Datasets:** TCGA, UK Biobank (where licensed), MIMIC-IV, FAERS, DrugBank, CPIC, clinical guidelines.
  - **Metrics:** AUC/PR, calibration (ECE), NNT/NNH simulations, trial-matching precision/recall, time-to-decision.
  - **Validation:** temporal holdouts; clinician review panels; post-market real-world evidence.
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## 11) Security, Privacy, Governance

- Consent management; DUA enforcement; k-anonymity for exports; DP for analytics; immutable audit.
  - RBAC/ABAC; PHI tokenization; key custody/HSMs; BAA with vendors.
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## 12) Observability & Cost

- Metrics: query latency, model drift, alert rates, subgroup perf; audit lead time.
- Cost: tiered storage, GPU batch windows, quantized inference; federated to minimize data

movement.

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## 13) Roadmap

- **M1 (4w):** ETL + baseline models + clinician UI.
  - **M2 (8w):** ADR/DDI + explanations + trial matching.
  - **M3 (12w):** Federated analytics + RWE loop.
  - **M4 (16w):** Regulatory reports + on-device pilots.
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## 14) Risks & Mitigations

- **Bias & fairness:** subgroup audits, reweighting, clinician oversight.
- **Data sparsity:** transfer learning, imputation, uncertainty estimates.
- **Clinical liability:** decision support (not decision making), explainability, override workflows.
- **Privacy breaches:** strict access, DP/federation, continuous monitoring.