```
## 1) Project structure (recommended)
. . .
hospital-readmit/
                          # put your CSV(s) here (e.g. patients_30000.csv)
⊢ data/
  configs/
   └ config.yaml
  - src/
   data_loader.py
    preprocess.py
    - features.py
    train_models.py
    — tune_optuna.py
    stacking.py
    - serve/
      └ api.py
                        # FastAPI app
    - explainability.py # SHAP helpers
    utils.py
 - mlflow/
                         # mlflow server + backend store (optional)
 - docker/
    – Dockerfile
    docker-compose.yml
  frontend/
                          # React app scaffold (example component provided)
   └ app/
  README.md
## 2) Quick start commands (CLI)
# 1) Create venv and install
python -m venv venv
source venv/bin/activate
pip install -r requirements.txt
# 2) Train all models (defaults read configs/config.yaml)
python src/train_models.py --config configs/config.yaml
# 3) Start MLflow UI
mlflow ui --port 5000
# 4) Start API (after model saved)
uvicorn src.serve.api:app --host 0.0.0.0 --port 8000
# 5) Start frontend (frontend/app)
cd frontend/app
npm install
npm start
## 3) Minimal `configs/config.yaml` (example)
```yaml
data_path: data/patients_30000.csv
target_col: readmitted_30d
id_col: patient_id
```

```
models:
 - name: logistic_regression
 - name: random_forest
 - name: xgboost
 - name: lightgbm
 - name: catboost
 - name: simple_nn
training:
 n_splits: 5
 random_state: 42
 optuna_trials: 60
mlflow:
 tracking_uri: http://localhost:5000
 experiment_name: hospital_readmit
4) core code highlights (short excerpts)
`data_loader.py` (load & quick checks)
```python
import pandas as pd
def load_data(path):
    df = pd.read_csv(path)
    # quick sanity
    print(df.shape)
    print(df.isnull().mean().sort_values(ascending=False).head())
    return df
### `preprocess.py` (impute / encode / feature engineering)
```python
from sklearn.impute import SimpleImputer
from sklearn.preprocessing import StandardScaler
import category_encoders as ce
1. fill missings
imputer = SimpleImputer(strategy='median')
2. target encoding for high-cardinality categories
te = ce.TargetEncoder(cols=['diagnosis_code', 'hospital_id'])
3. scaling numeric features
scaler = StandardScaler()
`train_models.py` (train multiple models + CV + MLflow)
```python
import mlflow
import joblib
from sklearn.model_selection import StratifiedKFold
from sklearn.metrics import roc_auc_score, average_precision_score
# sample model dict
from xgboost import XGBClassifier
from lightgbm import LGBMClassifier
from catboost import CatBoostClassifier
from sklearn.linear_model import LogisticRegression
from sklearn.ensemble import RandomForestClassifier
models = {
```

```
'logistic': LogisticRegression(max_iter=1000),
  'rf': RandomForestClassifier(n_estimators=200, n_jobs=-1),
  'xgb': XGBClassifier(use_label_encoder=False, eval_metric='logloss',
n_{jobs=-1},
  'lgbm': LGBMClassifier(n_jobs=-1),
  'cat': CatBoostClassifier(verbose=0)
}
# training loop (sketch)
for name, model in models.items():
    scores = []
    skf = StratifiedKFold(n_splits=cfg['training']['n_splits'], shuffle=True,
random_state=cfg['training']['random_state'])
    for train_idx, val_idx in skf.split(X, y):
        model.fit(X[train_idx], y[train_idx])
        preds = model.predict_proba(X[val_idx])[:,1]
        scores.append(roc_auc_score(y[val_idx], preds))
    mean_auc = sum(scores)/len(scores)
    mlflow.log_metric(f"{name}_cv_auc", mean_auc)
    # fit on full data and save
    model.fit(X, y)
    artifact = f"models/{name}.pkl"
    joblib.dump(model, artifact)
    mlflow.log_artifact(artifact)
### `tune_optuna.py` (Optuna example for LightGBM)
```python
import optuna
from lightgbm import LGBMClassifier
def objective(trial):
 params = {
 'num_leaves': trial.suggest_int('num_leaves', 20, 200),
 'learning_rate': trial.suggest_float('learning_rate', 1e-3, 0.3,
log=True),
 'n_estimators': trial.suggest_int('n_estimators', 100, 2000)
 model = LGBMClassifier(**params)
 # cross-validate and return mean AUC
 return cv_mean_auc(model, X, y)
study = optuna.create_study(direction='maximize')
study.optimize(objective, n_trials=cfg['training']['optuna_trials'])
print(study.best_params)
`stacking.py` (simple stacking meta-learner)
```python
from sklearn.ensemble import StackingClassifier
from sklearn.linear_model import LogisticRegression
estimators = [(n, models[n]) for n in ['xgb','lgbm','cat']]
meta = LogisticRegression()
stack = StackingClassifier(estimators=estimators, final_estimator=meta, cv=5,
n_jobs=-1)
stack.fit(X,y)
joblib.dump(stack, 'models/stack.pkl')
### `serve/api.py` (FastAPI minimal)
```python
```

```
from fastapi import FastAPI
import joblib
import pandas as pd
from pydantic import BaseModel
app = FastAPI()
class Patient(BaseModel):
 patient_id: int
 age: int
 diagnosis_code: str
 # add other fields
model = joblib.load('models/stack.pkl')
@app.post('/predict')
def predict(p: Patient):
 df = pd.DataFrame([p.dict()])
 proba = model.predict_proba(df)[:,1].tolist()[0]
 return {'patient_id': p.patient_id, 'risk_30d': float(proba)}
`explainability.py` (SHAP explain endpoint helper)
```python
import shap
explainer = shap.Explainer(model.predict_proba, X_sample)
shap_values = explainer(X_patient)
# return top 10 contributing features
## 5) Frontend sketch (React) — clinician dashboard
* Show list of patients, risk score (0.0-1.0), top contributing features (SHAP)
and suggested actions (e.g., follow-up call, meds reconciliation, home visit).
* Use one main component that calls `/predict` for one patient or batch endpoint
`/predict_batch` for lists.
**Minimal example component (React)**
export default function PatientCard({patient}){
  const [risk, setRisk] = useState(null);
  useEffect(()=>{
    fetch('/api/predict', {method:'POST', body: JSON.stringify(patient)})
      .then(r=>r.json()).then(j=>setRisk(j.risk_30d))
  },[])
  return (
    <div className='card'>
      <h3>{patient.patient_id} - Risk {risk && (risk*100).toFixed(1)}%</h3>
    </div>
  )
<u>;    </u>
## 6) Best-practices & performance tips (implementation checklist)
* Use StratifiedKFold, and report AUC-ROC, PR-AUC, sensitivity at fixed
specificity thresholds.
* For imbalanced outcomes prefer PR-AUC and recall (sensitivity) for positive
```

class.

- ^r Use early stopping and regularization for boosting models.
- Try CatBoost for raw categorical-heavy EHR features (handles categories natively).
- * Use Optuna (or Ray Tune) to tune each model; then ensemble the top performers (stacking).
- Calibrate probabilities (Platt / isotonic) before showing clinical risk.
- * Log experiments and artifacts in MLflow (or other registry) with versioning.
- * Add SHAP outputs and short textual explanations next to the risk score for clinicians.
- * Implement logging + request auditing (who requested, when) for HIPAA compliance.

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- ## 7) Scaling to many datasets & multi-model runs
- * If you truly have *multiple datasets* (e.g., per hospital), treat each as a separate experiment: run the single-pipeline with different `data_path` and register model version with MLflow.
- * For training many models in parallel, use a task queue (Celery) or orchestration (Airflow / Prefect).
- * For very large runs, containerize the training job and run on a Kubernetes cluster or in a cloud batch service.

- ## 8) Monitoring and model drift
- * Track input feature distributions and output risk distribution. Alert when distribution shift crosses threshold (Population Stability Index).
- * Re-train on rolling windows (e.g., retrain monthly or when model performance drops).

- ## 9) Security, privacy & governance
- * Encrypt PHI in transit and at rest. Use secure storage for models and logs.
- * Minimize exposed PHI through the API; prefer PII-less IDs; only show explanations relevant to care.
- * Maintain audit logs and access controls.

10) Files included in this doc (copy-paste into your project)

- * `train_models.py` (full training loop)
 * `tune_optuna.py` (Optuna tuning)

- * `serve/api.py` (FastAPI server)
 * `Dockerfile` & `docker-compose.yml` (examples)
- * `README.md` with run instructions

End of document.