

**Devansh Pathak**

**Data Scientist – Machine Learning, Statistical Modeling, and Data Management**

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## EDUCATION

### Master of Healthcare Data Science (MS)

University of Kansas Medical Center • Kansas City, Kansas • Jul 2025

### Bachelor of Medicine, Bachelor of Surgery (MBBS)

Smt. NHL Municipal Medical College • Gujarat, India • Aug 2017

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## EXPERIENCE

### Clinical Data Assistant

Westfield Urgent Care

**Jun 2021 – May 2023**

- Improved patient wait time by 40% using time-series predictive model that identified peak congestion patterns, leading to redesign of triage workflow and increasing daily patient capacity.
- Reduced data entry errors by 25% from standardizing physician charting templates by developing an ELT pipeline and automated data validation checks to clean intake data for 100+ daily patients.
- Led a data-first resource re-allocation by analyzing clinic revenue and operational data; presented findings to stakeholders which optimized staffing schedule and improved daily clinic revenue by nearly 2x.

### USMLE/COMLEX Instructor

Institute of Medical Boards

**Oct 2020 – Dec 2021**

- Built performance dashboards and automated reporting that featured heatmaps and trend forecasting, enabling real-time feedback loops leading to an average score improvement of 80 points across 8 cohorts.
- Implemented a predictive analytics framework in Excel and Python to analyze student performance data; identified key predictors of exam success to create personalized learning pathways for over 200 students.
- Utilized a A/B testing framework to evaluate efficacy of different teaching methodologies, resulting in adoption of new curriculum that increased student pass rates by 12%.

### Volunteer Research Assistant

University of Kansas Medical Center

**Mar 2019 – Sept 2020**

- Maintained a relational database (PostgreSQL) for high-dimensional genomic data, using advanced queries to retrieve and join count matrices with phenotypic data for analysis.
- Performed statistical analysis of RNA-sequencing data in R, utilizing DESeq2 to identify differentially expressed genes and performing pathway enrichment analysis that led to 3 peer-reviewed publications.

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## PROJECTS

### Predictive Risk Modeling for Patient Safety

- Architected a serverless pipeline on AWS (S3, Lambda, RDS) to process a 6,000-patient dataset, engineering features from patient's baseline (smoking history) and automated the identification of data errors to reduce time-to-analysis.
- Developed a Random Forest Classifier (scikit-learn) to proactively identify patients at high-risk; validated model's 68% accuracy by confirming features aligned with clinical risk factors making them both accurate and interpretable.
- Deployed a live and interactive dashboard (streamlit), translating complex outputs into actionable tool, enabling clinical teams to implement targeted monitoring strategies and potentially improve patient safety outcomes.

### Ecological Analysis of Socioeconomic Health Determinants

- Engineered a nationwide spatial analysis pipeline in R to model public health outcomes, integrated data from CDC and Census Bureau APIs; corrected geographic clustering with spatial error model (spdep, spatialreg) of 3,068 counties.
- Developed an inferential model (R-squared = 0.67) that identified county-level poverty, creating a framework to identify and stratify geographic hotspots for targeted resource allocation and population health interventions.
- Validated model through comprehensive diagnostics (VIFs via car, Cook's distance) and validated coefficient stability across 4 sensitivity analysis, delivering the project in a fully reproducible environment with renv.

### Clinical Risk Stratification from Stress Test Data

- Developed a predictive framework to stratify patient risk for myocardial ischemia, leveraging a multiple linear regression model that explained 40.5% of the outcome's variance (Adjusted R-squared) in 303 patient records.
- Improved model validity by diagnosing significant heteroscedasticity (Breusch-Pagan test) and correcting its effects with standard errors (HC3) yielding valid and reliable coefficients for clinical predictors.
- Translated model insights into a clinical decision-making framework, demonstrating that ST segment slope ( $p < 0.001$ ) and age ( $p < 0.013$ ) were critical predictors, providing physicians to refine pre-test risk assessments.

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## CERTIFICATIONS

**NCA Generative AI LLM • NVIDIA (2025)**

**Machine Learning Specialization • Stanford (2025)**

**Data or Specimens Only Research • CITI (2025)**

**Biomedical Research with Children • CITI (2025)**

**IBM Professional Data Science • IBM (2024)**

**GCP for Clinical Trials (US FDA Focus) • CITI (2024)**

**CRC Foundations • CITI (2024)**

**Biomedical PI • CITI (2024)**

**Data & Safety Monitoring in Human Subjects • CITI (2024)**

**Biomedical Researchers • CITI (2024)**