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➤ 代表著作

- Associations of anomalous water temperature, salinity, and pH with change in water color of fish farming ponds. Chuang C-H, Chiu U-C, Huang C-W, and **Chang KY(*)**. Journal of the World Aquaculture Society, 54:1563–1574, 2023.
- DeepFlu: a deep learning approach for forecasting symptomatic influenza A infection based on pre-exposure gene expression. Zan A, Xie Z-R, Hsu Y-C, Chen Y-H, Lin T-H, Chang Y-S, and **Chang KY(*)**. Computer Methods and Programs in Biomedicine, 213, 106495, 2022.
- Polyphenols as Alternative Treatments of COVID-19. Wu Y, Pegan SD, Crich D, Hansen MC, Starling EB, Ellison Desrochers E, Booth C, Mullinix LN, Lou L, **Chang KY(*)**, and Xie Z-R(*). Computational and Structural Biotechnology Journal, 19:5371-5380, 2021.
- Structural and Functional Enrichment Analyses for Antimicrobial Peptides. Lo SC, Xie Z-R, and **Chang KY(*)**. International Journal of Molecular Sciences, 21(22), 8783, 2020.
- Trends in Regional Cancer Mortality in Taiwan 1992-2014. Ho YR, Ma S-P, and **Chang KY(*)**. Cancer Epidemiology, 59:185-192, 2019.

➤ 曾獲獎項

- 指導學生獲 台海大資訊工程學系專題競賽 優等 2024
 - 擴散模組在影像切割之應用
- 指導學生獲 台海大資訊工程學系專題競賽 亞軍 2023
 - 整合學習在醫學影像切割之應用
- 指導學生獲 台海大暑期學習實務體驗計畫 佳作 2023
- 指導學生獲 聯發科技物聯網開發競賽 入圍獎 2018
- 台海大院級研究優良教師 2014



From Notes to Knowledge: Accelerating Rare Disease Diagnosis with Explainable **AI** for **Phenotypic-Genomic Integration**

Bridging Clinical Narratives and Genomic Insights for Precision Medicine

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Proposal outline

- The challenge in Rare Disease Diagnosis
- Project Objectives & Innovation
- AI Framework: Key Components
- Expected Outcomes & Validation
- Alignment with Institutional Mission





Millions impacted by delayed diagnosis

Why Rare Disease Diagnosis Needs Innovation?



Critical Gaps:

Non-structured clinical data vs. genomic data
→ fragmented insights

Manual analysis → prolonged diagnosis
(weeks/months)



Current tools lack transparency → low clinician trust



Building a Transparent AI Diagnostic Framework

多源數據整合 |
Multi-Source Data
Integration

Transform free-text
EHRs → structured
phenotypes + genomic
data

致病機轉解析 |
Mechanistic Insight
Generation

Model phenotype-gene
associations using XAI

決策透明化 |
Clinician-Centric
Explainability

Traceable AI logic for
trust & actionable
decisions

How the XAI System Works



Input: EHR free text → NLP structuring → phenotype features

Integration: Merge with genomic data + biomedical databases (e.g., OMIM)

Analysis: XAI identifies candidate gene variants + ranks diagnoses

Output: Clinician-friendly report with evidence trails



Delivering Precision and Speed



AI Diagnostic Model

80% accuracy in prospective trials
(n=50)
50% faster than manual analysis



Knowledge Database

Dynamic rare disease phenotype-
genome repository



Clinical Impact

Reduced diagnostic inequality;
improved patient outcomes



Advancing Precision Medicine and Equity



- 精準醫學 | Precision Medicine: Bridging data silos for tailored care
- 診斷平等 | Diagnostic Equity: Democratizing access to advanced tools
- 創新可信度 | Trustworthy Innovation: Explainability fosters clinician adoption



Thank you for your attention!

Any Question?