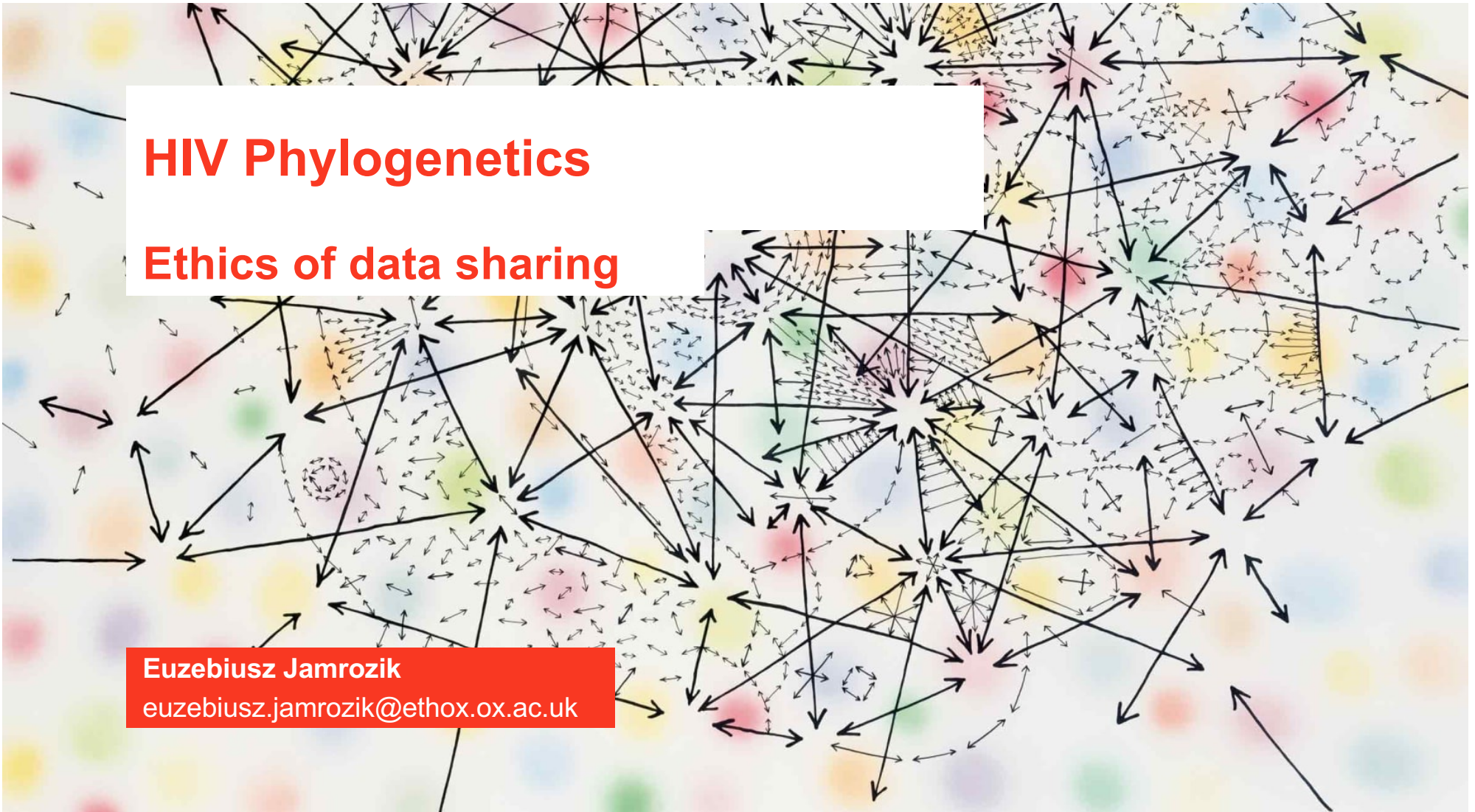


## HIV Phylogenetics

## Ethics of data sharing

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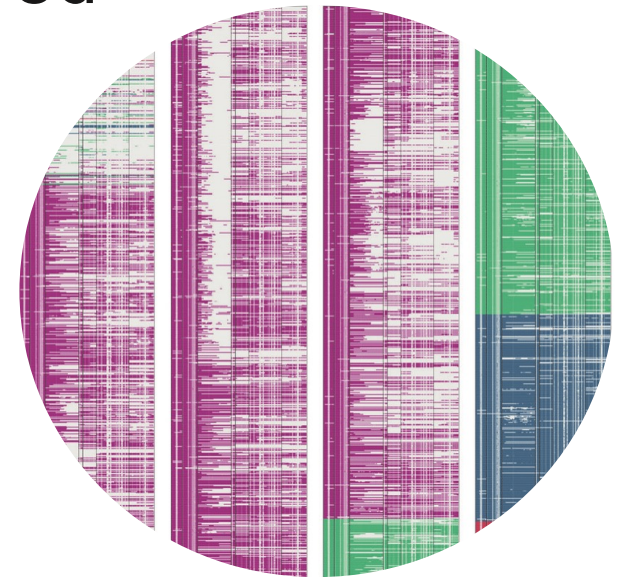


# Background

*Technical aspects* of how data are collected, linked, analysed, and shared influence

*Ethical considerations* such as:

- Potential benefits
- Potential harms
- Respect for persons
  - Privacy, stigma, etc.

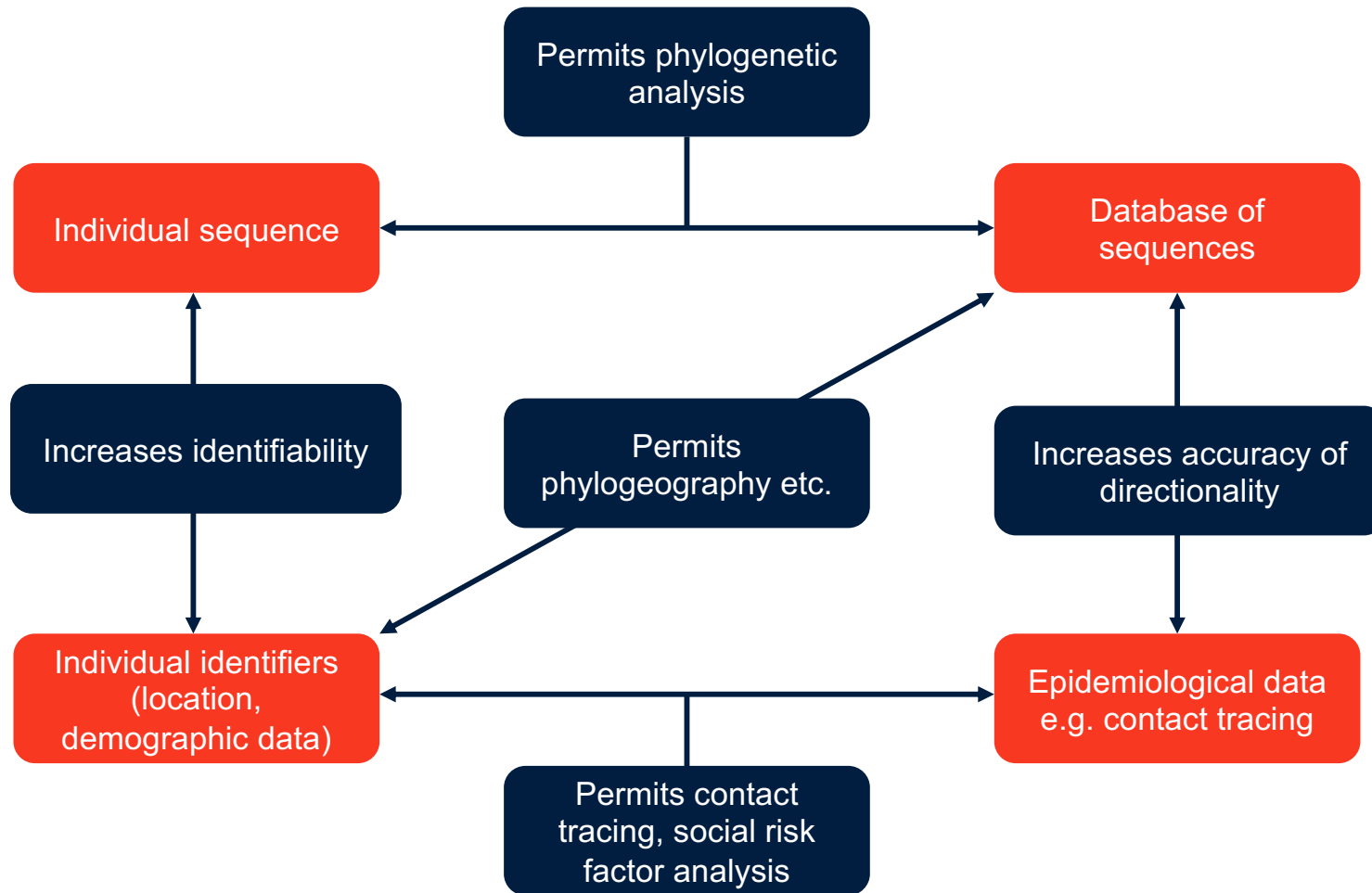


## Open science: ethical tradeoffs

- **Maximise benefits** by transparent data sharing
  - Increased validity
  - Increased (meta-)analysis
  - More (and more accurate) results
- **Minimise harms** by reducing
  - Identifiability & stigma
  - Misuse of data



# Linkage





## Minimalism: data and linkage

Scientific and ethical rationale for research data maximalism (including genomic data)

Minimalism might reduce risks by limiting:

- *Number of individuals' data* shared/linked
- *Types of data* linked
- *Level of detail* (e.g. fuzzy location data)
- *Duration of linkage* (e.g., “ephemeral phylogeny”)



Image: Celeste Boursier Mougnot

# Institutions & Governance

Relevant data held by different (global) research groups, government/public health agencies, technology companies

**Co-ordination** problems in data sharing

- Different regulations
- Need for appropriate gate-keeper

**Tension between benefits** of local ownership of data  
**versus risks** to local communities

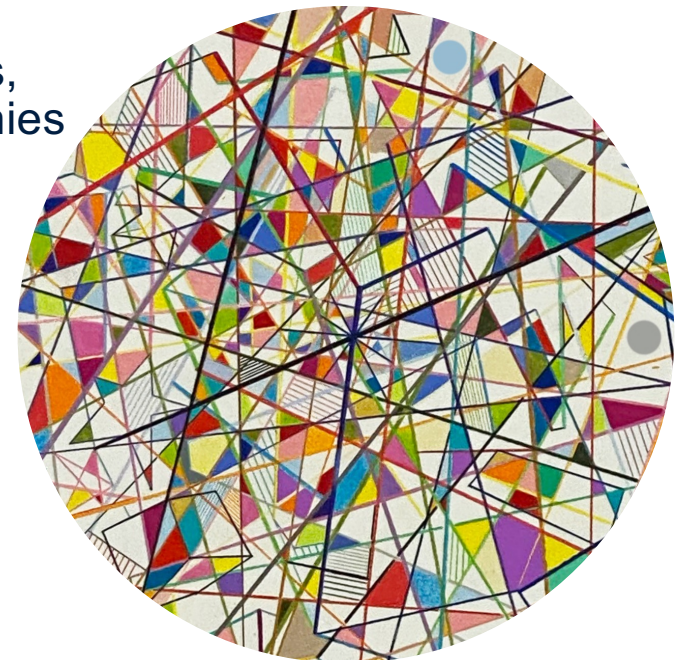


Image: Tom Friedman

# Possible technological solutions

## 1) Federated databases & analysis

- Data held by different connected institutions
- Technical barriers (computing power) & funding barriers

## 2) Trusted research environments

- Data deposited in systems held by third parties
- Need to trust centralized database security

## 3) Data packages

- Defined reasons & degree of data sharing
- Funding & staffing barriers
- Ethical tensions including post-research storage of data



Image: Chiharu Shiota

## Conclusions

Different technical approaches influence ethical features & outcomes of phylogenetic analysis

Data minimalism (including linkage minimalism) is a key consideration

Several unresolved issues for future work



# Thank you

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