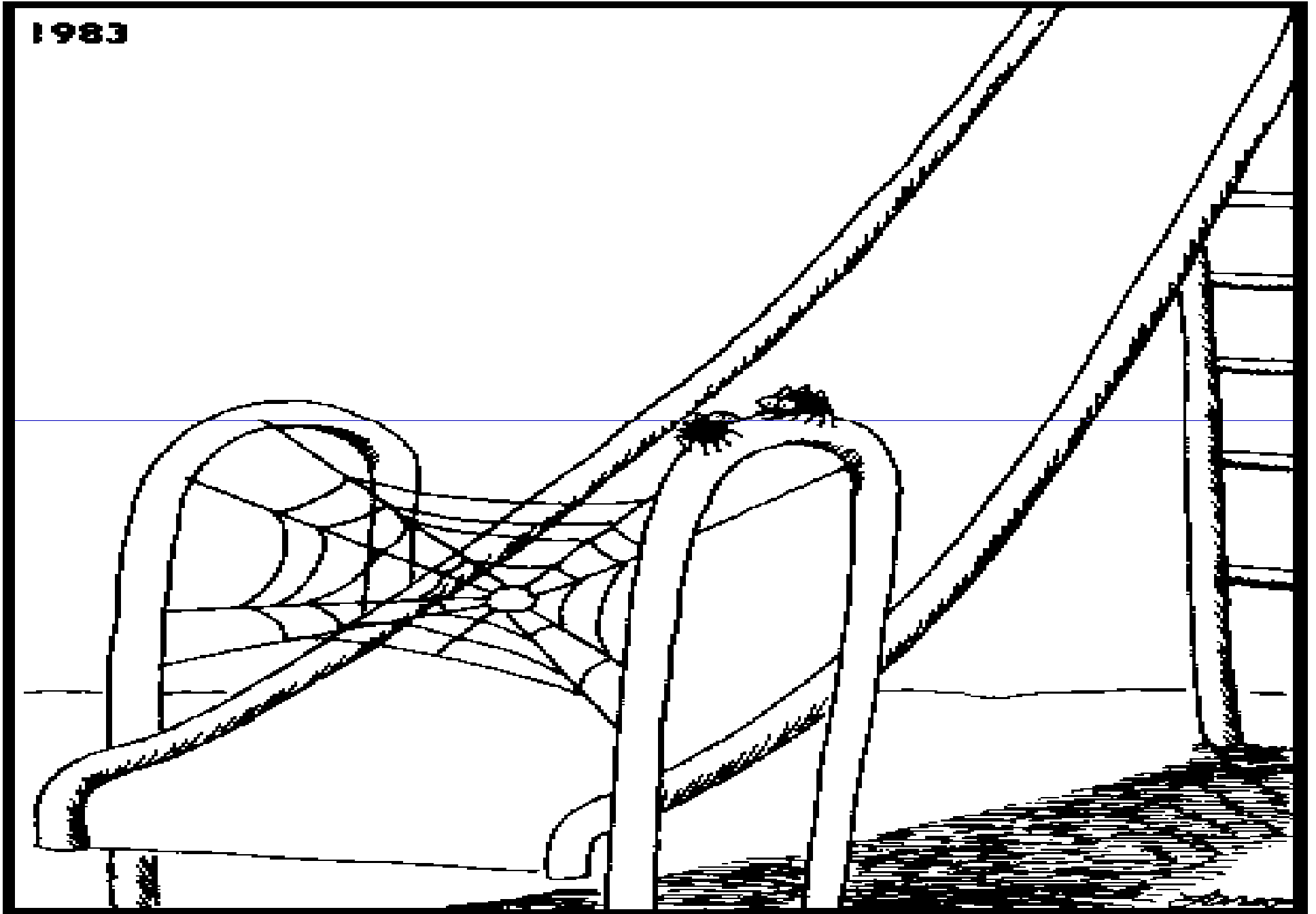


1983



“If we pull this off, we’ll eat like kings.”

# Large Cohort Studies as the “Population Laboratories” for Genetic Research

- Cohort Selection Allows Assessment of Larger Number of Risk Factors and Intermediate Factors
- Biologic repository is Collected Prior to Incident Event
- Ability to Assess Change in Risk Markers
- Reduced Bias in Outcome Assessment
  - (i.e., validation of outcomes, includes events with rapid mortality, can limit to **incident** events, etc)

**Human Genome Project**

```
graph TD; A[Human Genome Project] --> B[HapMap Project]; B --> C[Candidate Gene Evaluations]; C --> D[Genome Wide Association Studies];
```

**HapMap Project**

**Candidate Gene Evaluations**

**Genome Wide Association Studies**

# Unprecedented Evolution in Cohort Studies During the Last Decade

## Defining Genetic risk

## Data Analysis Issue

Family history  $\longrightarrow$  Smaller Scale Local analyses

Targeted SNPs ( $N \sim$  tens)  $\longrightarrow$  Smaller Scale Local analyses

Larger # of SNPs ( $N \sim$  thousands)  $\longrightarrow$  Larger distributed analyses  
with required replication

GWA Studies ( $N \sim$  millionish)  $\longrightarrow$  Huge Data Issues: being  
made available to larger  
scientific community

**Next Evolution?**  $\longrightarrow$  **Next Evolution?**

# **Existing vs. New Cohorts?**

## **Advantage of Existing Cohorts**

- **Cost Efficient**
- **Time Efficient (events available)**
- **Well Characterized Phenotypes**
- **Stored Biologic Specimens**

# **Existing vs. New Cohorts**

## **Challenges of Existing Cohorts**

- **Data usually Disease Specific: often limited to common diseases (CA, CVD, DM, etc.)**
- **Consent from pre-GWA era**
- **No input on Data Collected/Outcomes**
- **Fewer Large Cohorts of Children**
- **Harmonizing phenotypes across cohorts**

# Next Steps:

## Cohorts/Population Laboratories?

- **How can we enhance and standardize:**
  - Informed Consent/Participant Genetic Data Security
  - Harmonization of Exposures and Outcomes
  - Data Specimen Collection
    - frozen blood based specimens (DNA, Serum, etc) are inadequate for the “omics” revolution
    - what are needs of future population labs (e.g., tissue, fresh specimens, etc)?
- **How can we enhance use and publication of GWA data from existing Cohort Studies by the broader scientific community?**
- **When should new cohort studies be initiated?**
  - Should they span full age and outcome spectrum?
- **Should we embark on strategies to enhance efficiency?**
  - Initial use of case-control, medical record linkage studies, etc
  - Using existing well characterized cohorts to verify findings and better assess mediating factors?

# What should be next for cohort studies?

- **What are the priorities for the above issues?**
- **Who will lead these effort?**
  - **NHGRI?**
  - **Other?**