

# Trajectory Modeling by Shape

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# Thanks

- Joint work with Brianna Heggeseth (Williams College)

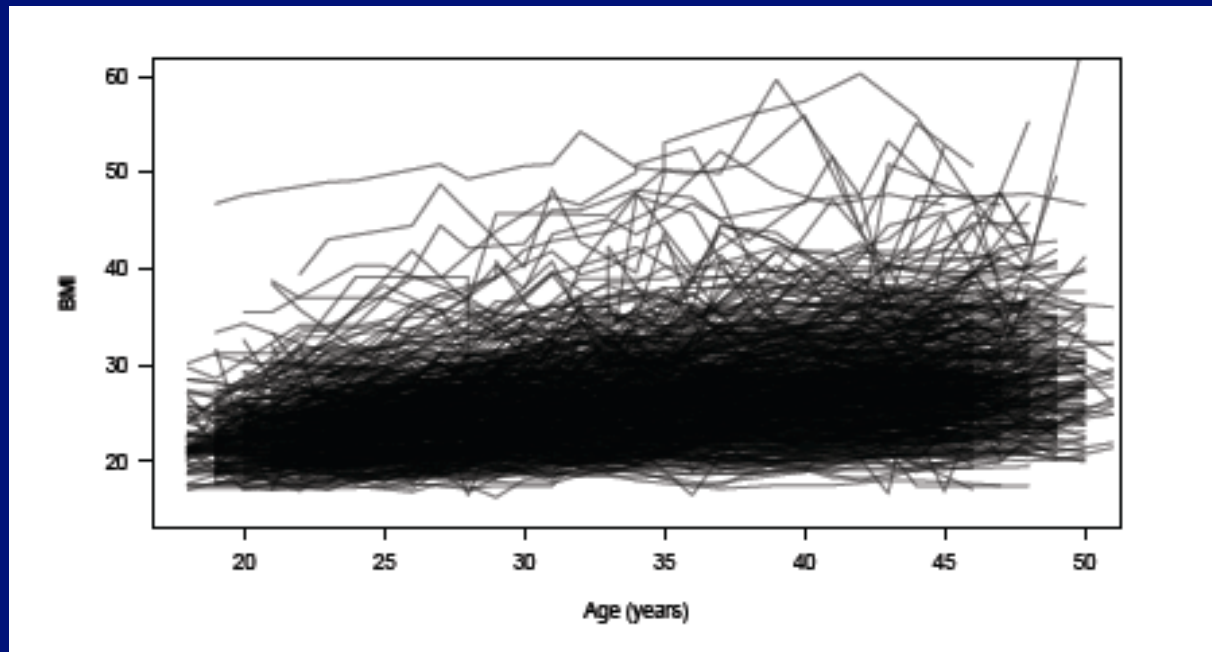
## References

- Heggeseth, BC and Jewell, NP. The impact of covariance misspecification in multivariate Gaussian mixtures on estimation and inference: an application to longitudinal modeling. *Statistics in Medicine*, 2013, 32, 2790-2803 .
- Heggeseth, BC and Jewell, NP. Vertically shifted mixture models for clustering longitudinal data by shape. Submitted for publication.

**“Understanding our world requires conceptualizing the similarities and differences between the entities that compose it”**

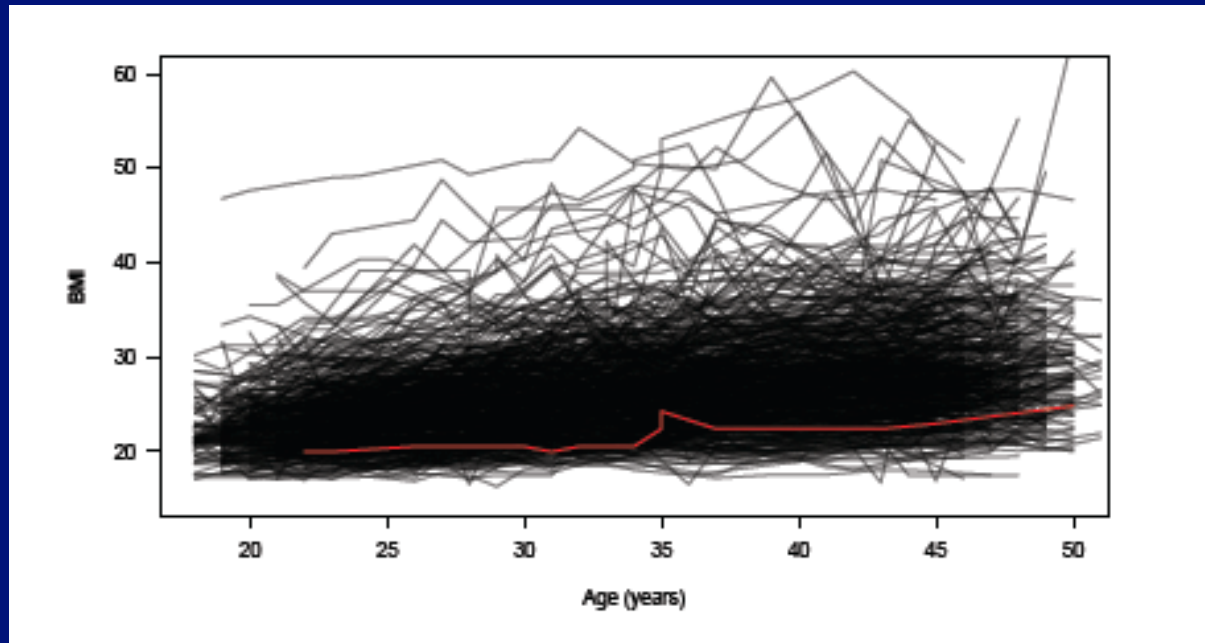
**Robert Tryon and Daniel Bailey, 1970**

# How does BMI change with age?



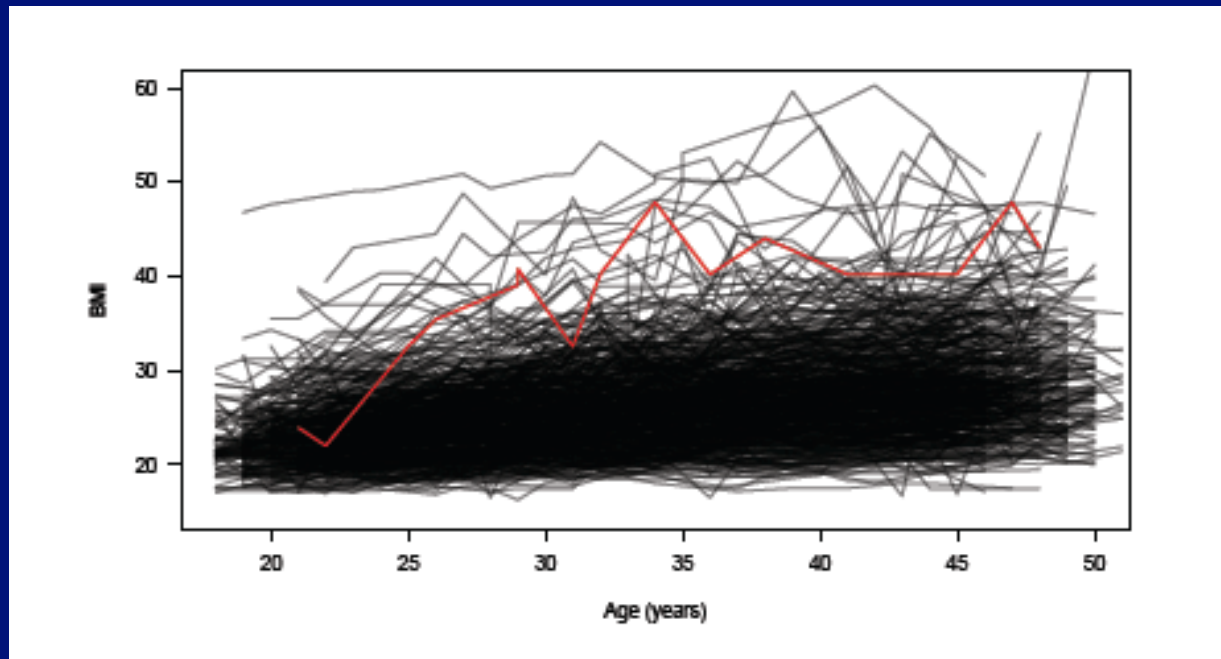
**National Longitudinal Study of Youth (NLSY) from 1979 - 2008.**

# How does BMI change with age?



**National Longitudinal Study of Youth (NLSY) from 1979 - 2008.**

# How does BMI change with age?



**National Longitudinal Study of Youth (NLSY) from 1979 - 2008.**

# Typical Longitudinal Analysis

- Use Generalized Estimating Equations (GEE) to estimate the mean outcome, and how it changes over time, adjusting for covariates
  - regression parameter estimation is consistent despite potential covariance misspecification
  - efficiency can be gained through use of a more appropriate working correlation structure
  - robust (sandwich) standard error estimators available
- But, with a heterogeneous population,
  - BMI does not change much for some people as they age
  - BMI changes considerably for some people as they age
- We don't wish to average out these separate trajectories by modeling the mean over time

# Finite Mixture Models

- **Data for  $n$  individuals:**  $\mathbf{y}_i = (\mathbf{y}_{i1}, \dots, \mathbf{y}_{im_i})$  measured at times  $t_i = (t_{i1}, \dots, t_{im_i})$
- **We assume  $K$  latent trajectories in the population that are distributed with frequencies:**  $\pi_1, \dots, \pi_K$  where  $\pi_k > 0$  and  $\sum_{k=1}^K \pi_k = 1$ .  
$$f(\mathbf{y}|\mathbf{t}, \theta) = \pi_1 \mathbf{f}(\mathbf{y}|\mathbf{t}, \beta_1, \Sigma_1) + \dots + \pi_K \mathbf{f}(\mathbf{y}|\mathbf{t}, \beta_K, \Sigma_K)$$
- **The (conditional) mixture density is**  $f(\mathbf{y}|\mathbf{t}, \beta_k, \Sigma_k)$ , **a multivariate Gaussian with mean**  $\mu_k$  **and covariance**  $\Sigma_k$ .
- **In most trajectory software, (conditional) independence is assumed as a working correlations structure:**  $(\Sigma_k = \sigma_k^2 I)$ .

$$\theta = (\pi_1, \dots, \pi_K; \beta_1, \dots, \beta_K; \Sigma_1, \dots, \Sigma_K) \quad 8$$



# Finite Mixture Models

- The mean vector  $\mu_k$  is related to the observation times as follows:

- Linear:  $(\mu_k)_j = \beta_0 + \beta_1 t_{ij}$
- Quadratic:  $(\mu_k)_j = \beta_0 + \beta_1 t_{ij} + \beta_2 t_{ij}^2$
- Splines in observation times

where the regression model (and coefficients) are assumed the same for each cluster, and  $t_{ij}$  is the  $j^{\text{th}}$  observation for the  $i^{\text{th}}$  individual where  $1 \leq j \leq m_i$

# Finite Mixture Models

- Group membership:  $\pi_k = \frac{\exp(\gamma_k z)}{\sum_{j=1}^K \exp(\gamma_j z)}$

**Z is set of same or different covariates**

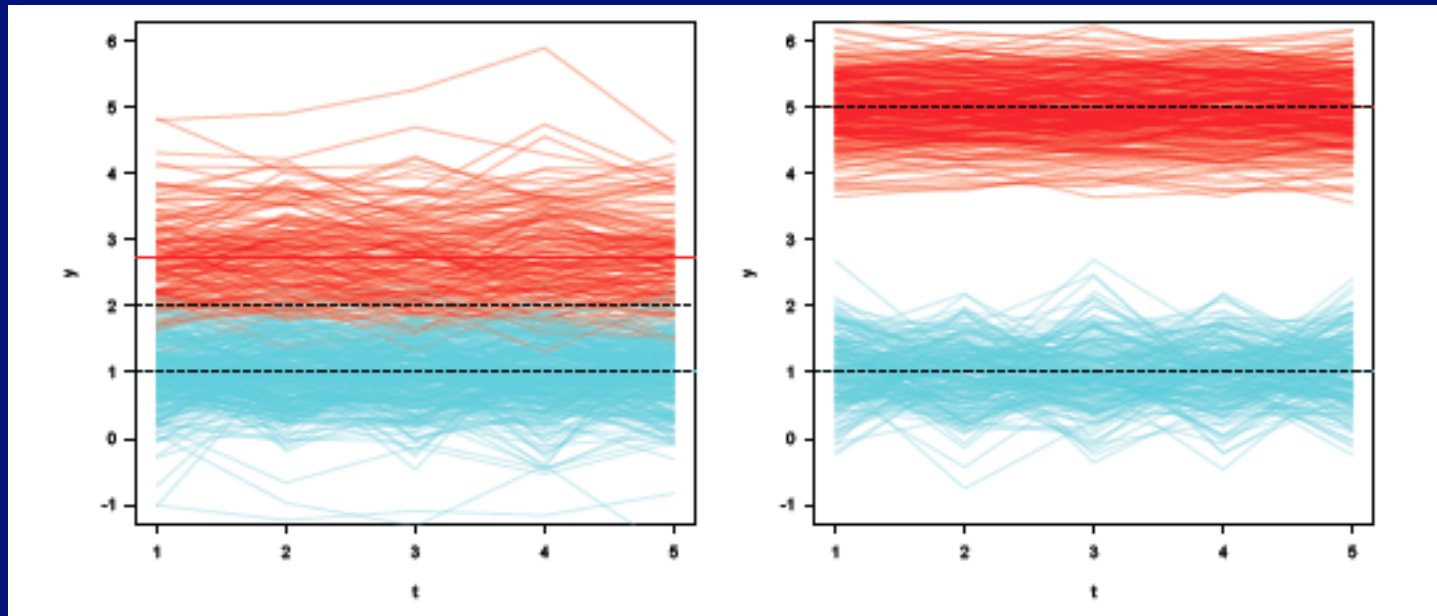
**This expands  $\theta$  to include the  $\gamma$ s also**

# Estimation for Mixture Models

- Maximum likelihood estimation for  $\theta$  via the EM algorithm
- $K$  is pre-specified; can be chosen using the BIC
- Parameter estimators are not consistent under covariance misspecification (White, 1982; Heggeseeth and Jewell, 2013).
- Robust (sandwich) standard error estimators are available.
- How bad can the bias in regression estimators be? What influences its size?

# Mispecified Covariance Structure Bias and Separation of Trajectories

- Separated components lead to little bias even when you wrongly assume independence.



**Black dashed -- true means, Solid lines – estimated means**

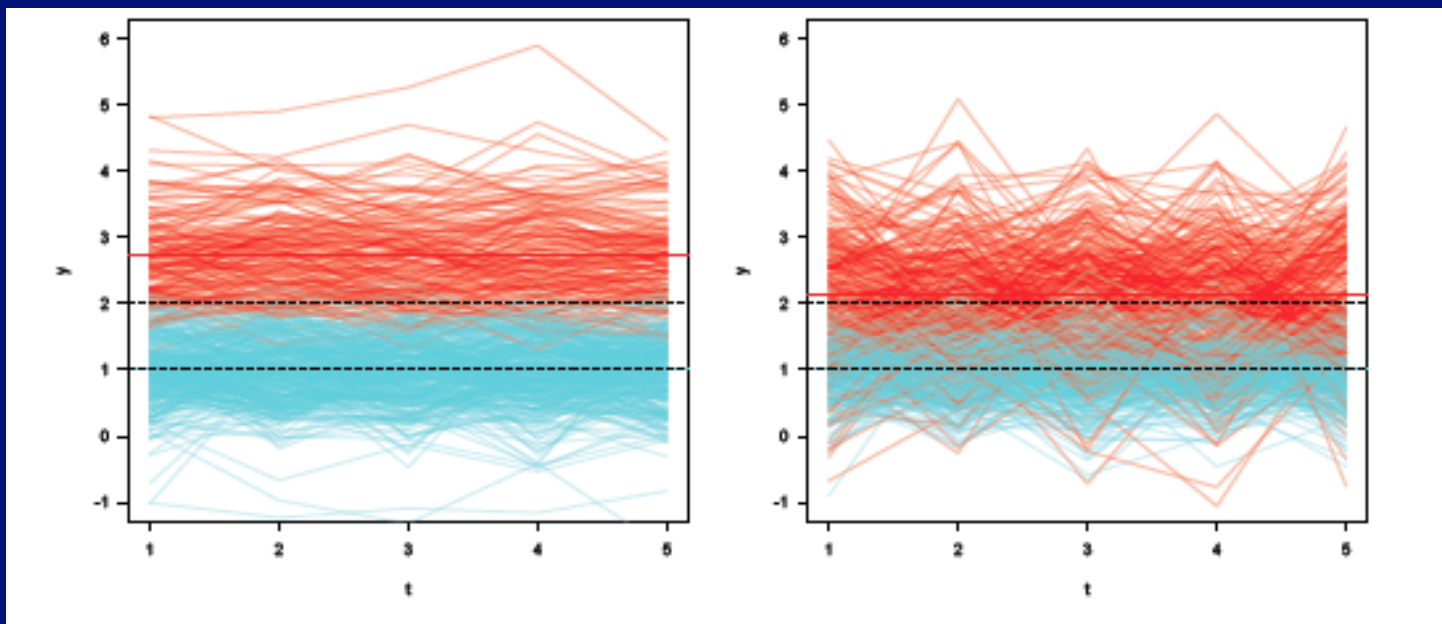
$$\hat{SE}_I(\beta_{01}) = 0.02, \hat{SE}_R(\beta_{01}) = 0.06$$

$$\hat{SE}_I(\beta_{01}) = 0.01, \hat{SE}_R(\beta_{01}) = 0.01^{12}$$

# Mispecified Covariance Structure

## Bias and Level of Dependence

- Components with little dependence lead to small bias even when you wrongly assume independence.

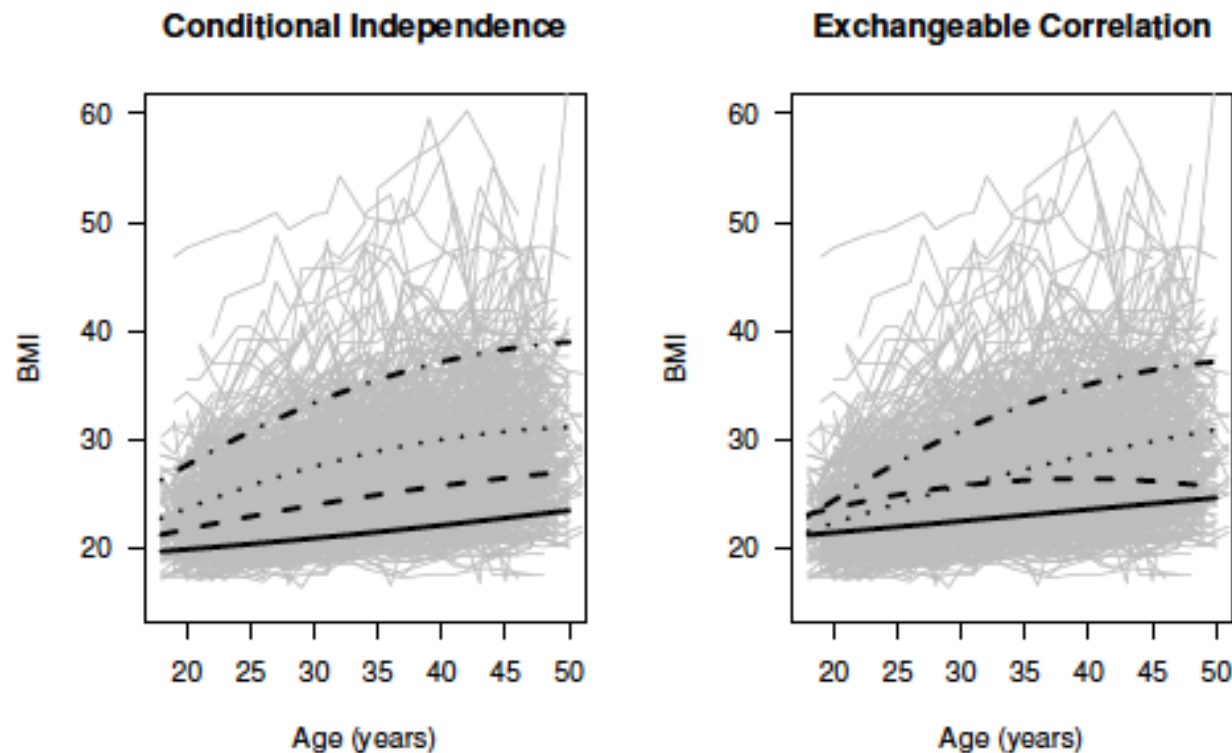


**Black dashed -- true means, Solid lines – estimated means**

$$\hat{SE}_I(\beta_{01}) = 0.02, \hat{SE}_R(\beta_{01}) = 0.06$$

$$\hat{SE}_I(\beta_{01}) = 0.03, \hat{SE}_R(\beta_{01}) = 0.04$$

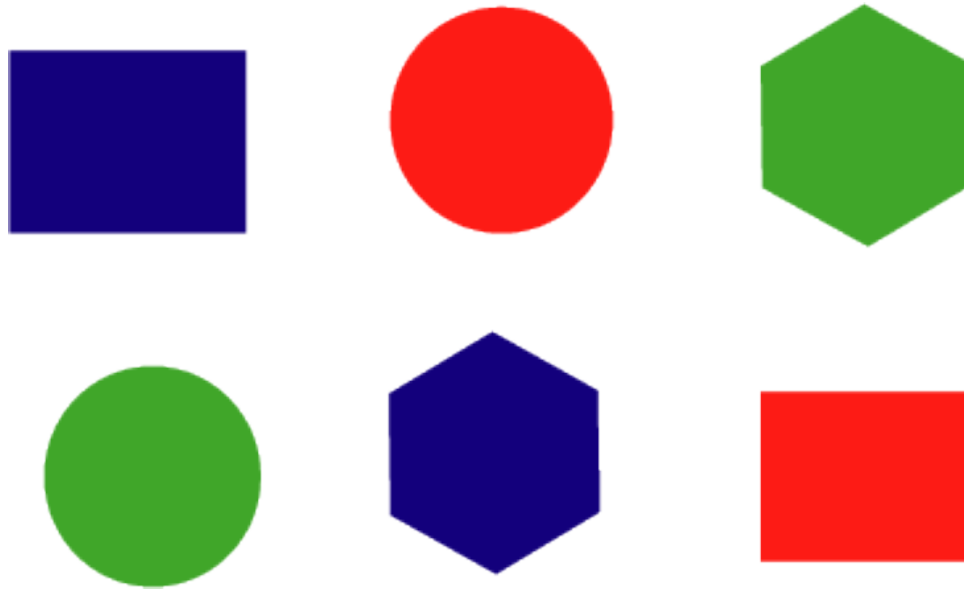
# NLSY Data Analysis



Covariance makes a difference to the trajectories

- hard to estimate bias from misspecified covariance

# How Do We Group These Blocks?

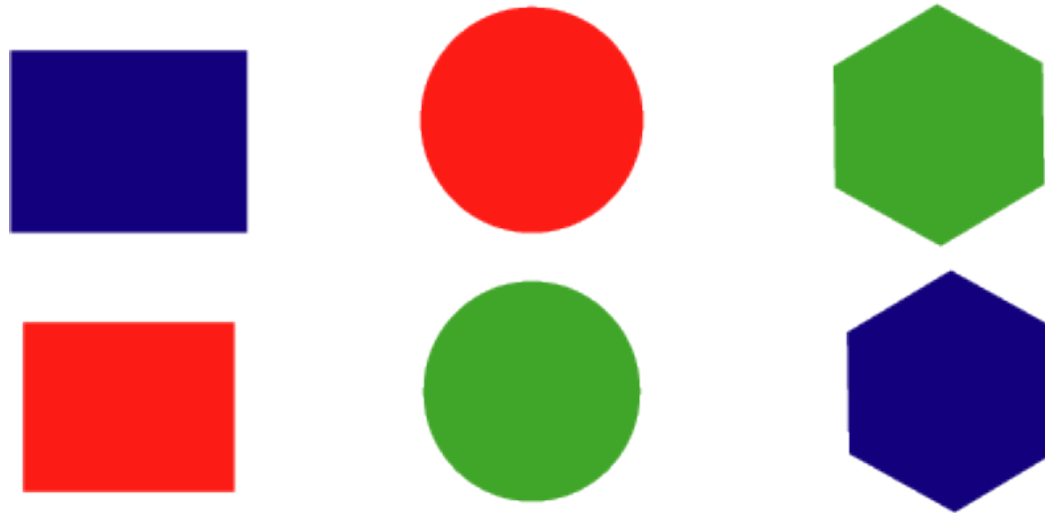


# Group by Color

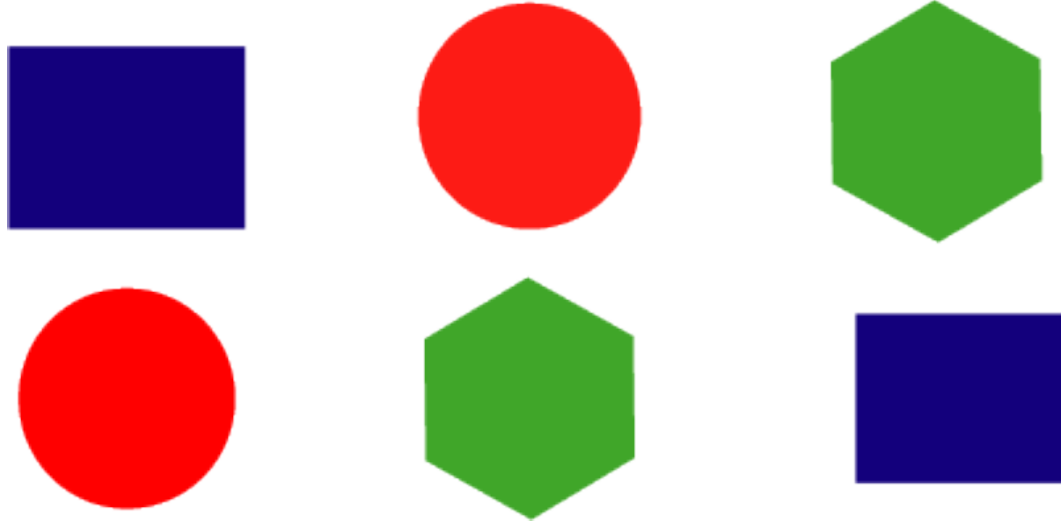




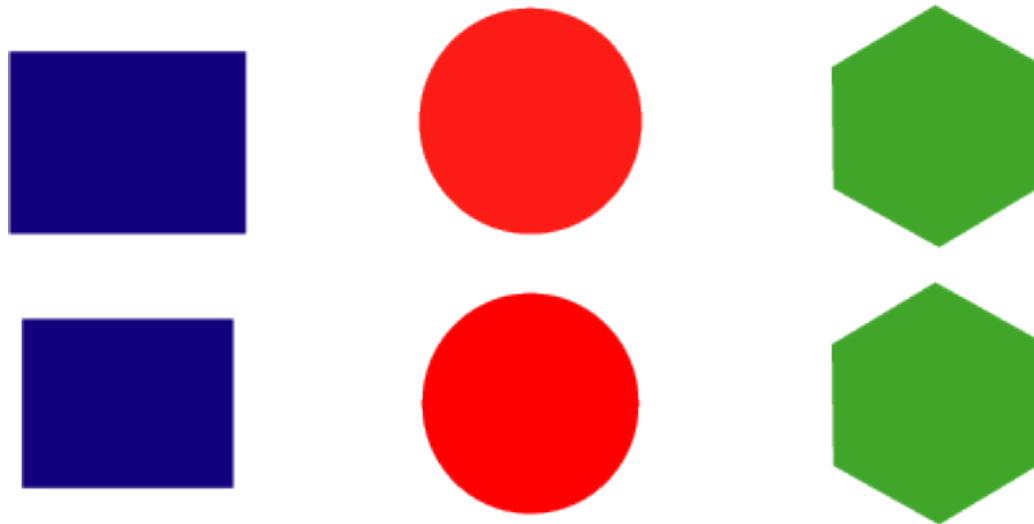
# Group by Shape



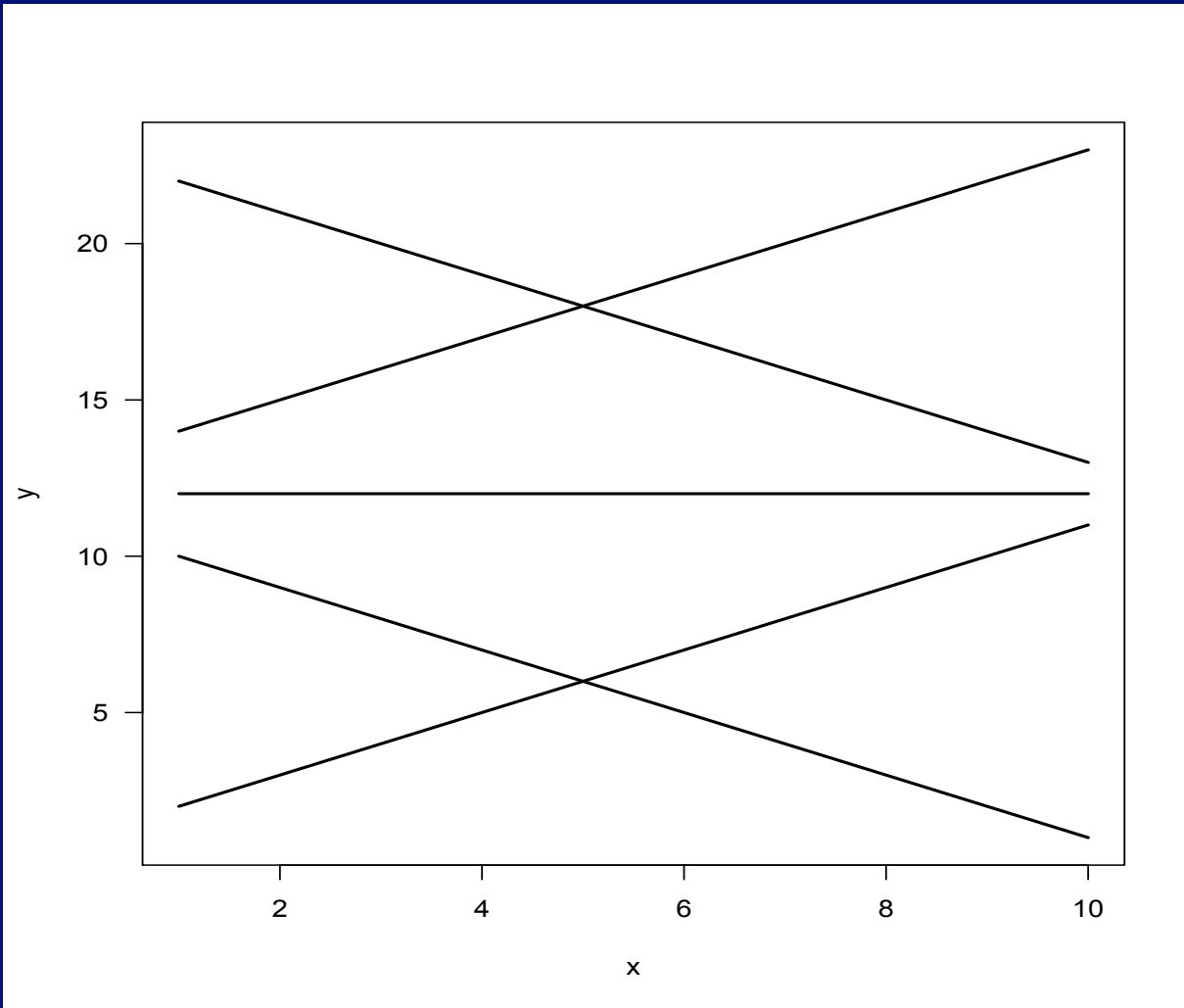
# How Do We Group These Blocks?



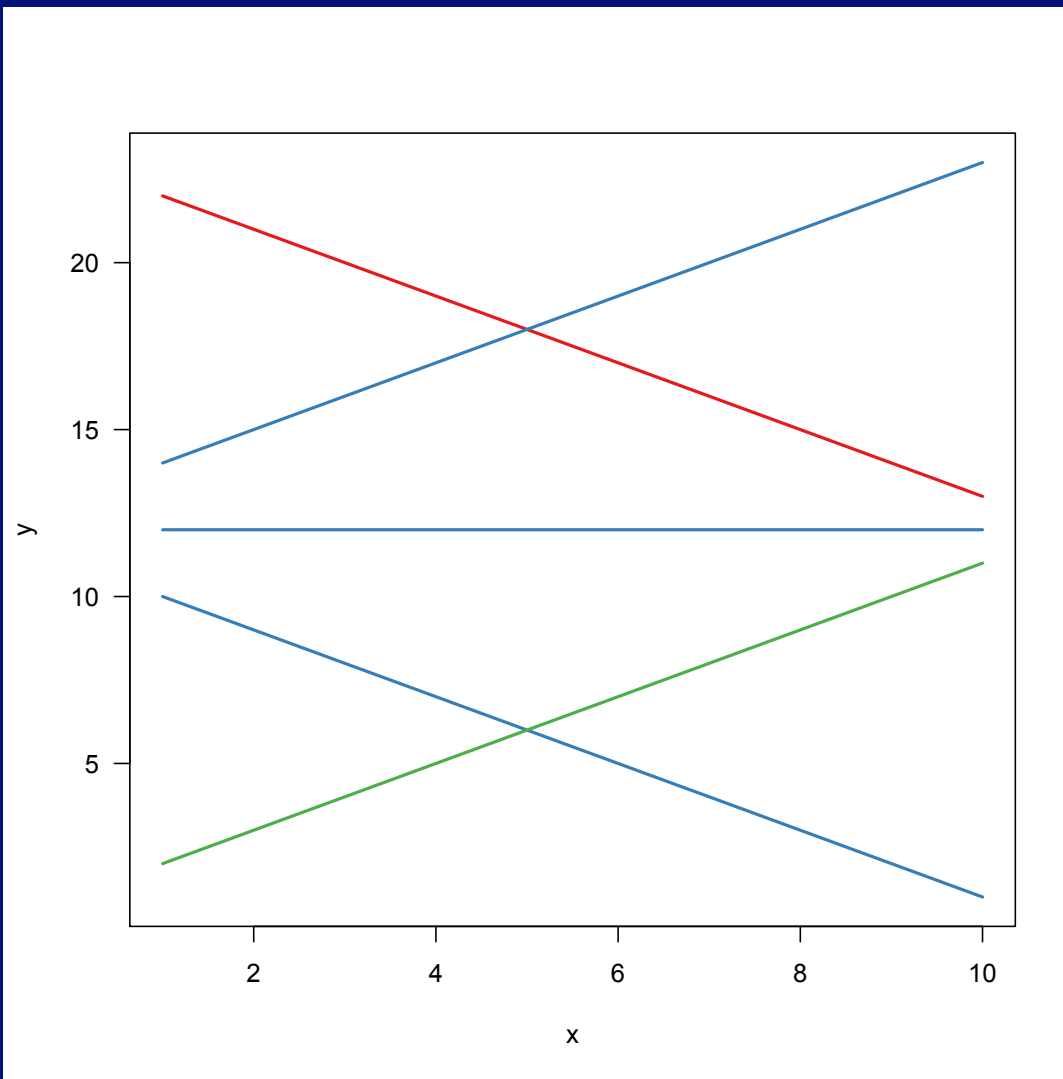
# Group by Color or Shape



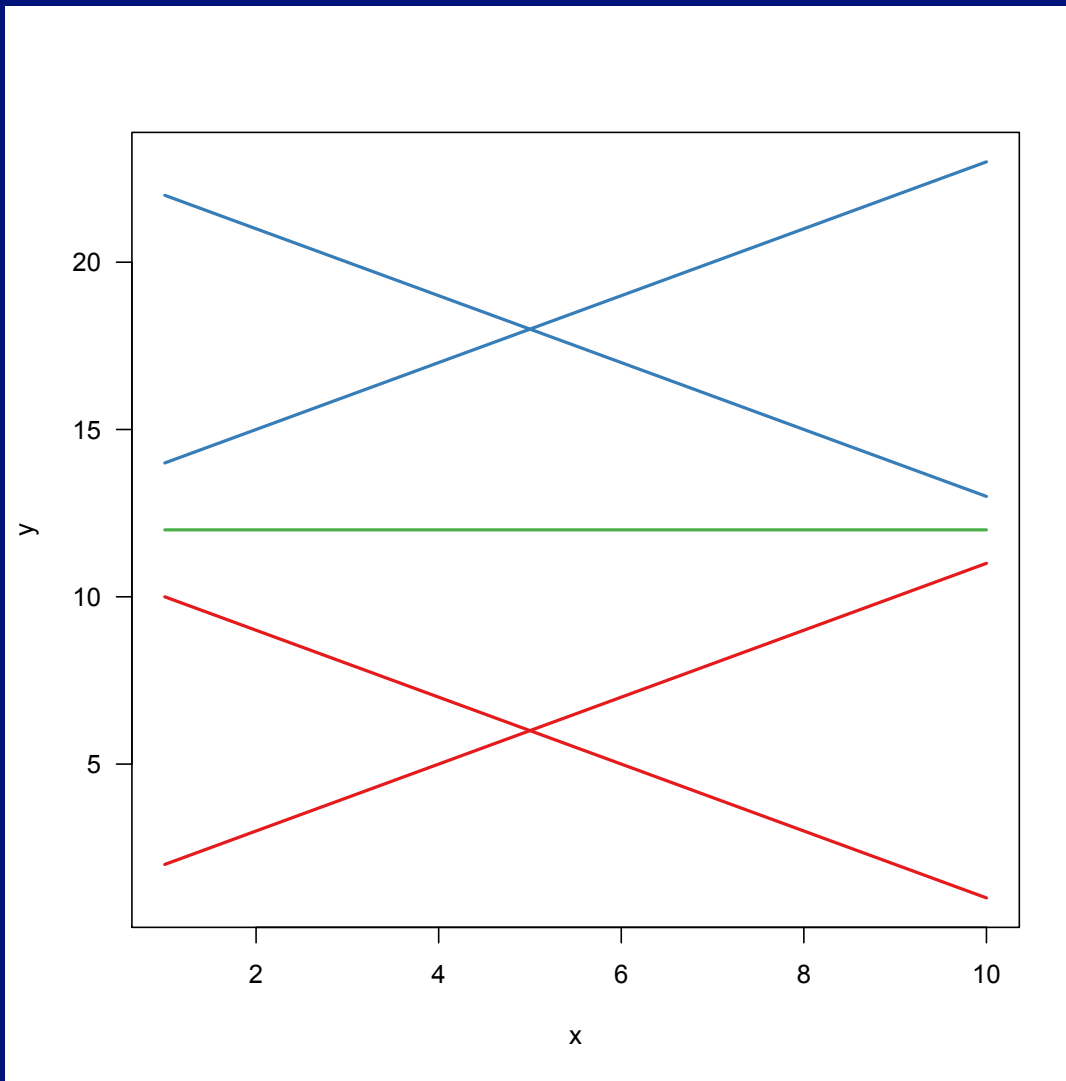
# How Do We Group These (Regression) Lines?



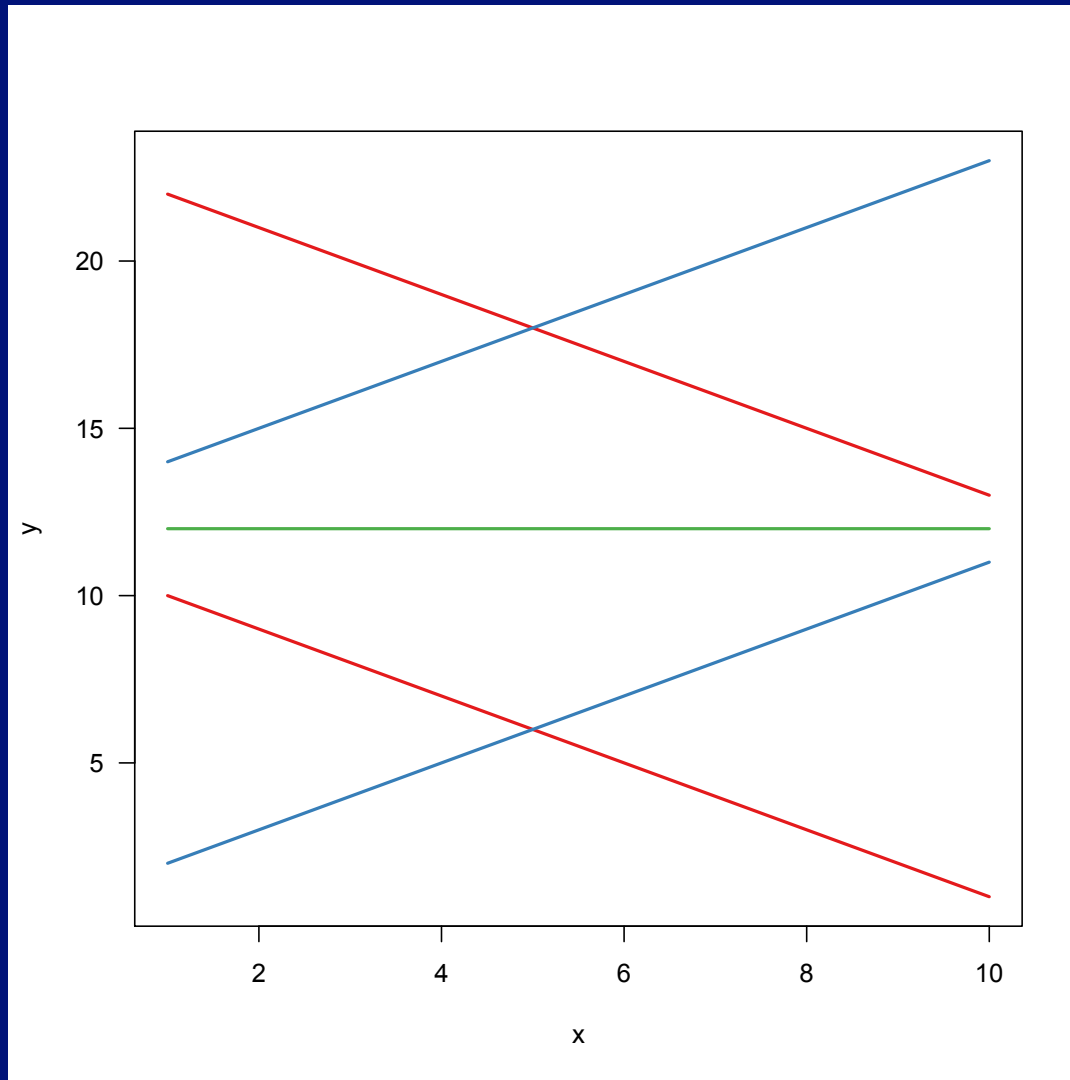
# Group by Intercept



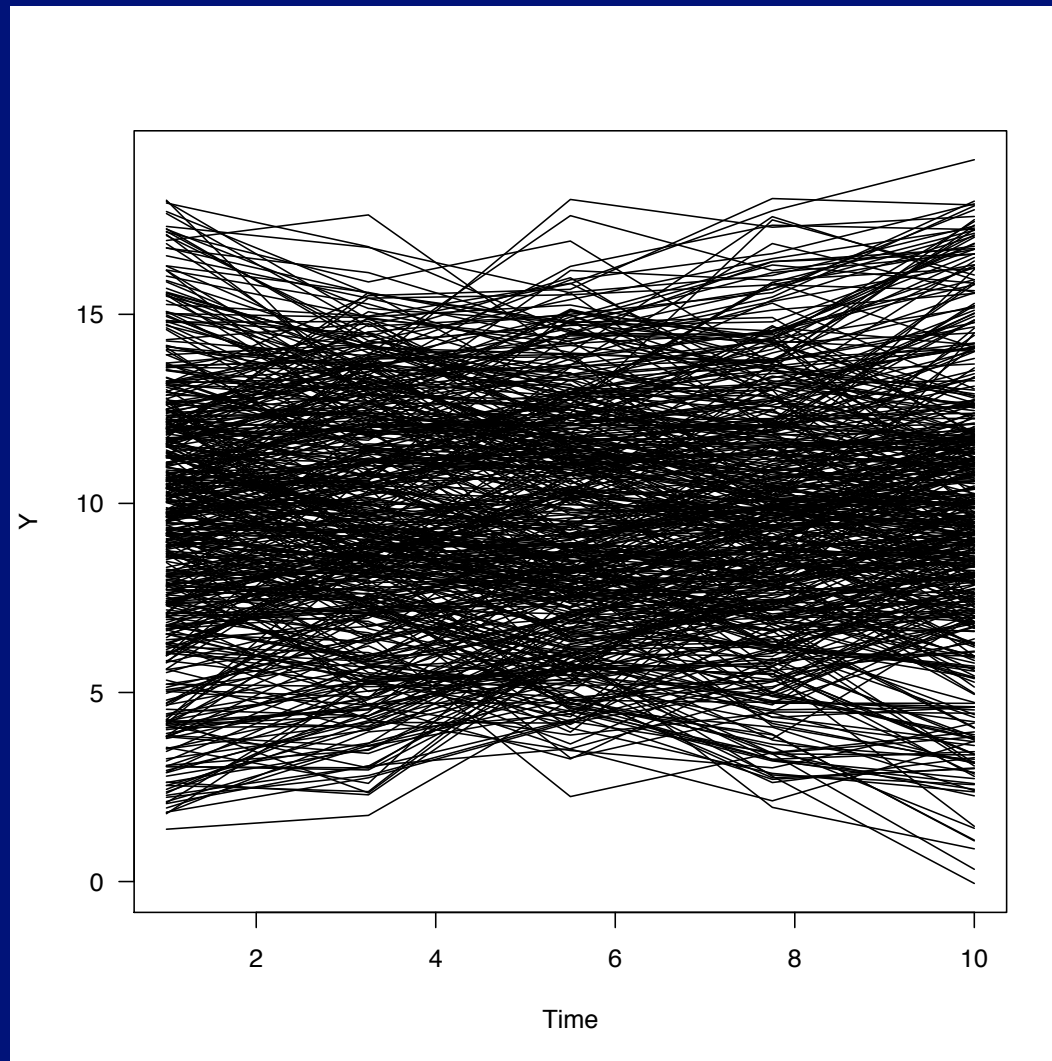
# Group by Level



# Group by Shape (Slope)



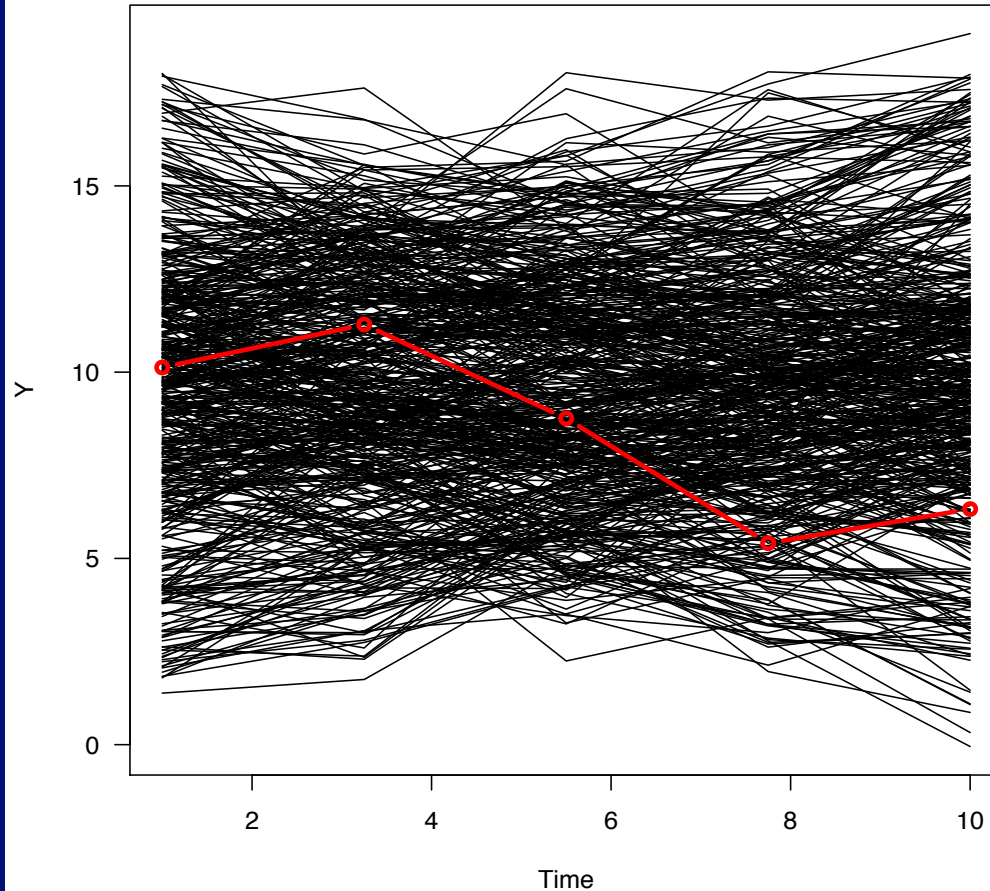
# Simulated Data



How could we group these individuals?



# Simulated Data

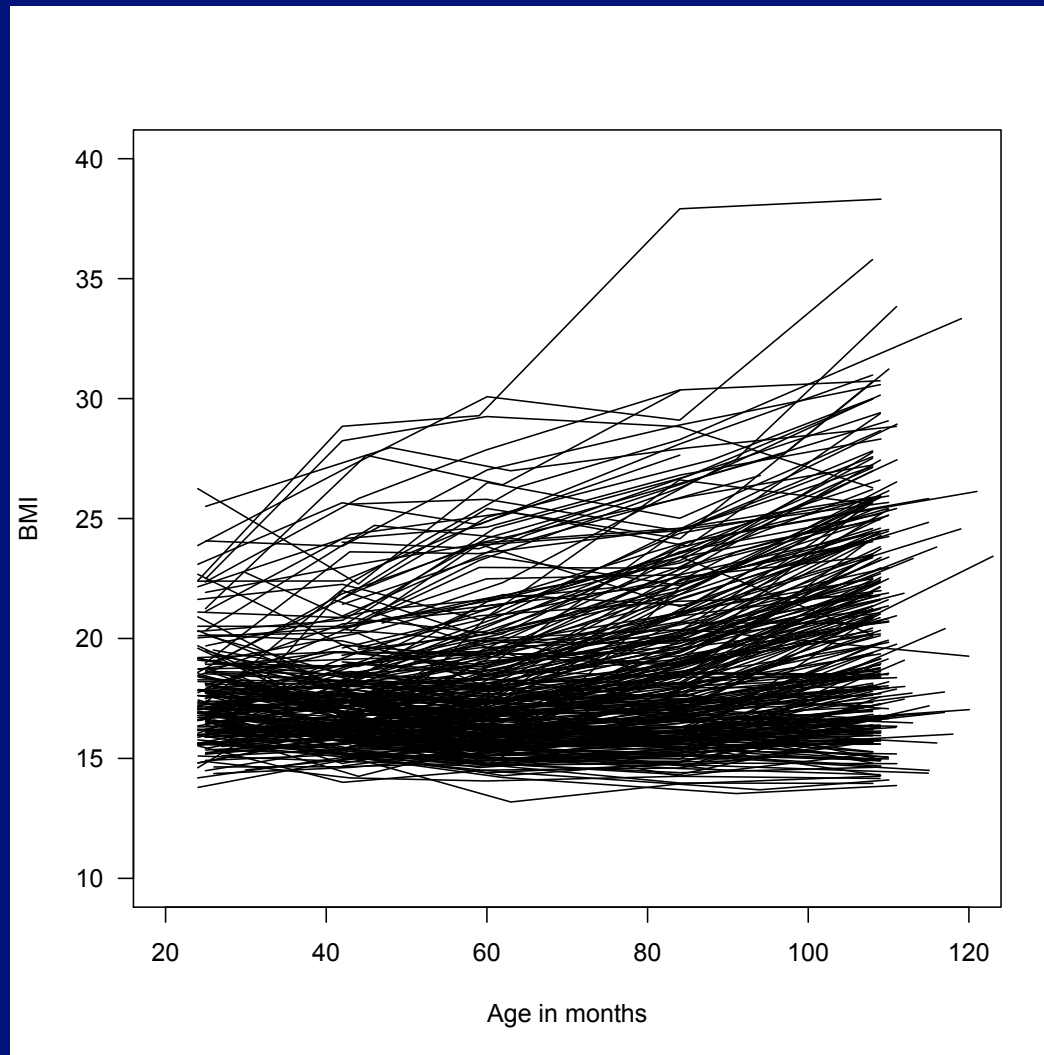


How could we group these individuals?

# Real Longitudinal Data

- **Center for the Health Assessment of Assessment of Mothers and Children of Salinas (CHAMACOS) Study**
  - **In 1999-2000, enrolled 601 pregnant women in agricultural Salinas Valley, CA.**
  - **Mostly Hispanic, agricultural workers.**
  - **Determine if exposure to pesticides and other chemicals impact children's growth patterns (BMI, neurological measures etc\_.**
- **First, focus on studying/estimating the growth patterns of children.**
- **Second, determine if early life predictors are related to the patterns**
  - **pesticide/chemical exposure in utero**
    - **ODT, PDT, PDE, BPA (bisphenol A)**

# CHAMACOS Data



How could we group these individuals?

# Cluster Analyses

- Clustering is the task of assigning a set of objects into groups so that the objects in the same group are more similar to each other than to those in other groups.
- What does it mean for objects to be more similar or more dissimilar?
  - Distance matrix
- Why do we cluster objects?

# Standard Clustering Methods

- Partition methods
  - Partition objects into  $K$  groups so that an objective function of dissimilarities is minimized or maximized.
  - Example: K-means Algorithm
- Model-based methods
  - Assume a model that includes a grouping structure and estimate parameters.
  - Example: Finite Mixture Models

# K-means algorithm

- Input: Data for  $n$  individuals in vector form. For individual  $i$ , the observed data vector is

$$\mathbf{y}_i = (y_{1i}, \dots, y_{im}).$$

- Measure of Dissimilarity: Squared Euclidean distance. The dissimilarity between the 1<sup>st</sup> and 2<sup>nd</sup> individuals is

$$d(\mathbf{y}_1 - \mathbf{y}_2) = \|\mathbf{y}_1 - \mathbf{y}_2\|^2 = (y_{11} - y_{12})^2 + \dots + (y_{im} - y_{2m})^2$$

# K-means Algorithm

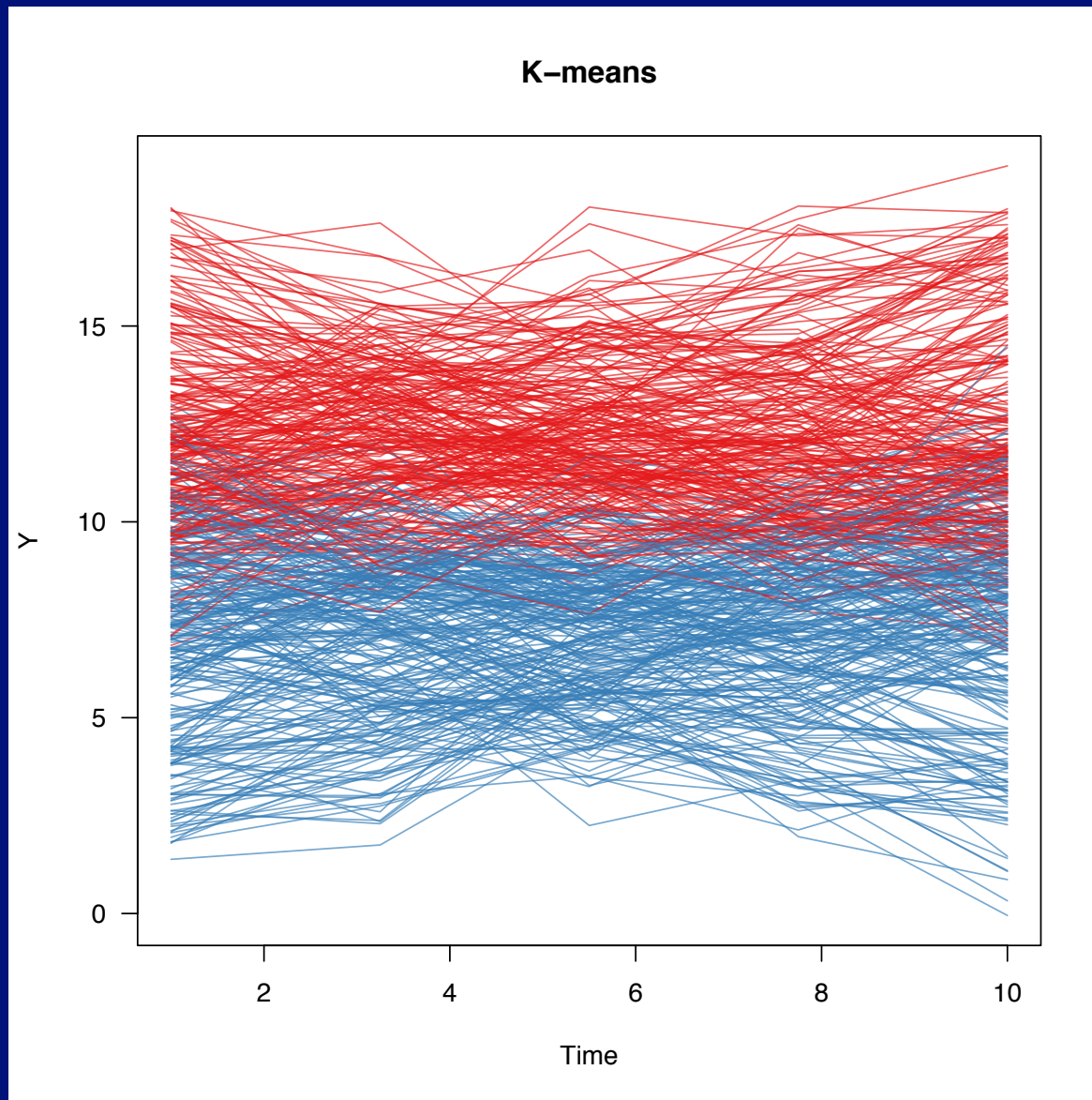
- Goal: Partition individuals into  $K$  sets  $\mathbf{C} = \{\mathbf{C}_1, \mathbf{C}_2, \dots, \mathbf{C}_K\}$  so as to minimize the within-cluster sum of squares

$$\sum_{k=1}^K \sum_{\mathbf{y}_i \in \mathbf{C}_k} \|\mathbf{y}_i - \mu_k\|^2$$

where  $\mu_k$  is the mean vector of individuals in  $\mathbf{C}_k$ .

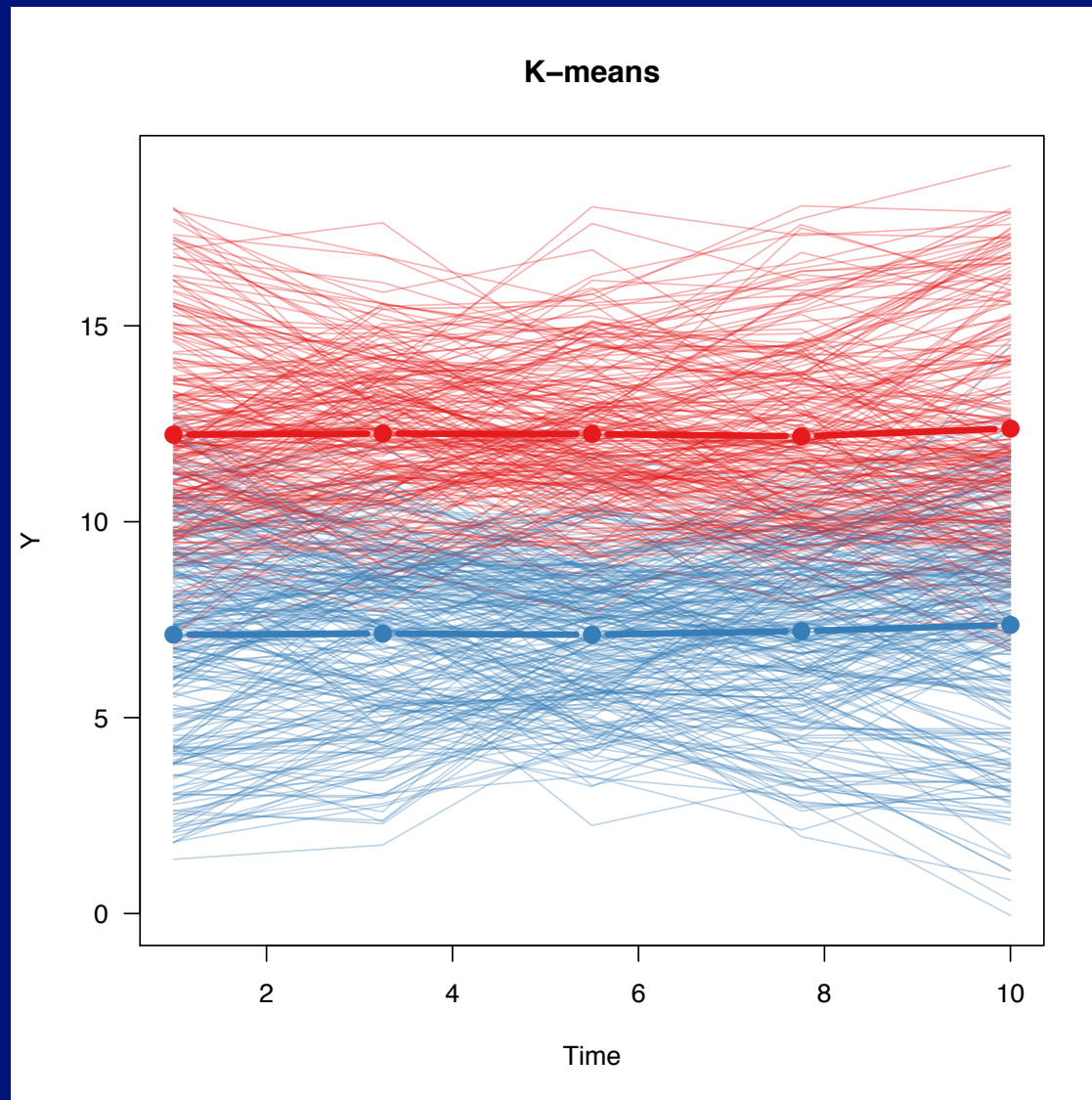
( $K$  must be known before starting  $K$ -means. There are many ways to choose  $K$  from the data that try to minimize the dissimilarity within each cluster while maximizing the dissimilarity between clusters: for example, the use of *silhouettes*.)

# Application to Simulated Data



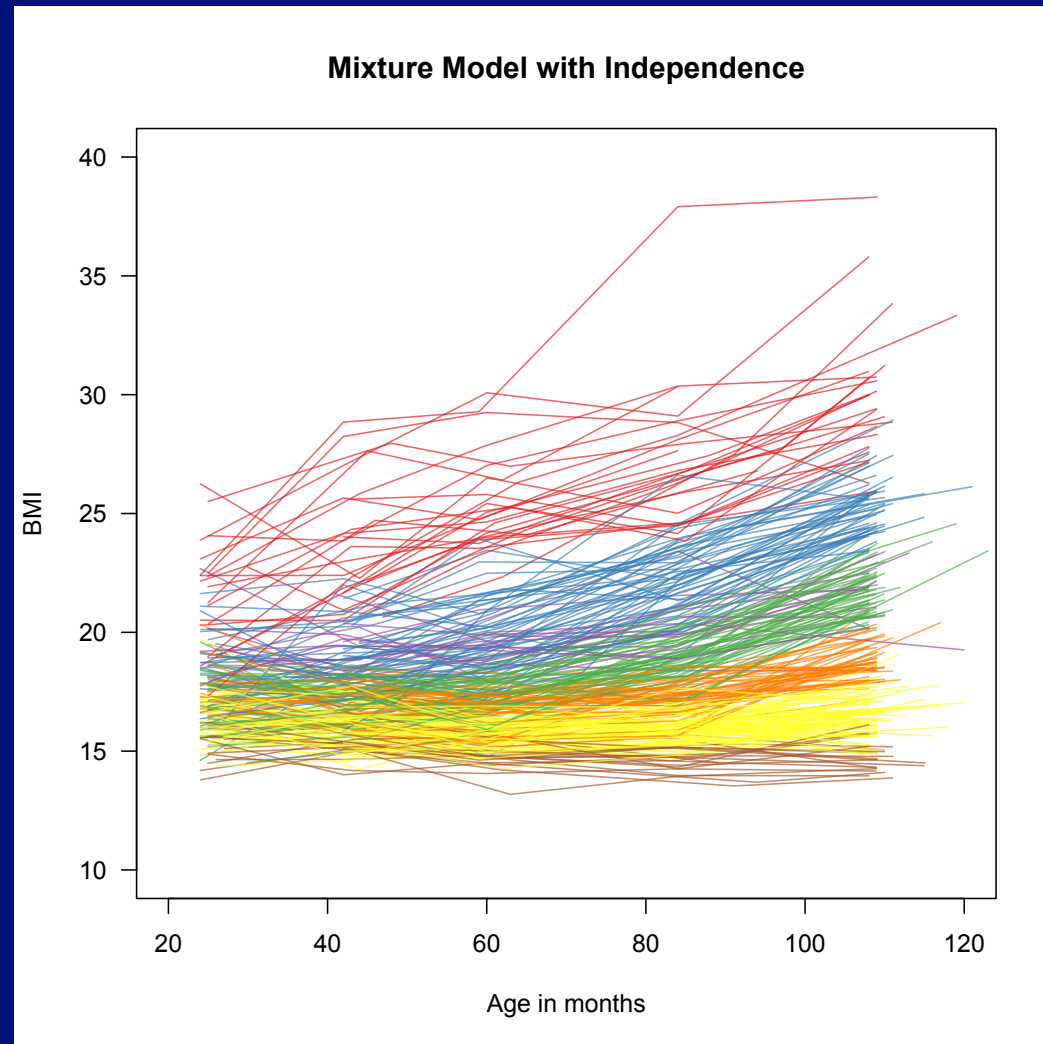


# Application to Simulated Data

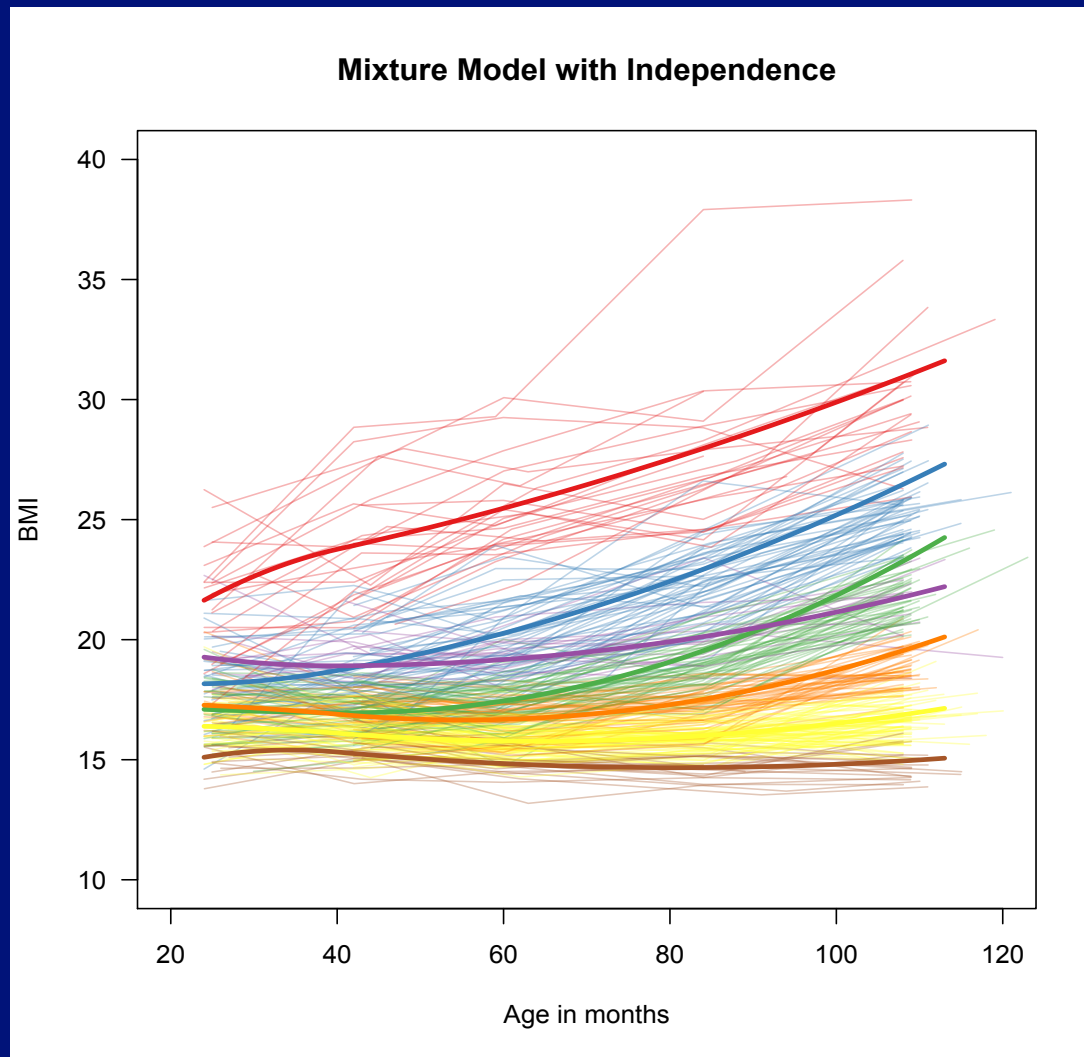


How would you describe—interpret—the group trajectories?

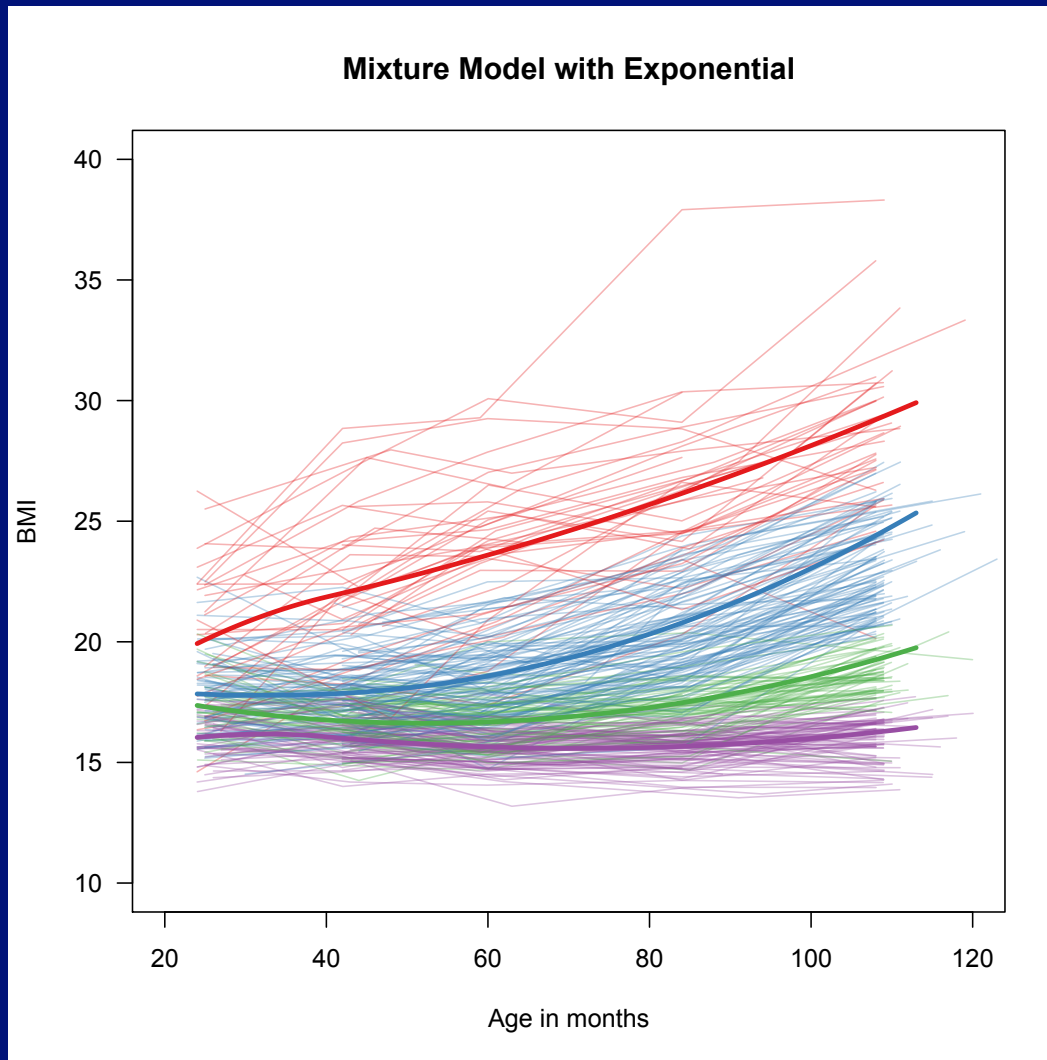
# Finite Mixture Model Applied to CHAMACOS Data



# Finite Mixture Model Applied to CHAMACOS Data



# Finite Mixture Model Applied to CHAMACOS Data



# Clustering by Shape

- Interested in *shape* not just level (which appears to dominate clustering techniques)
- Want a method that:
  - Works with irregularly sampled data
  - Includes a way to estimate the relationship between baseline risk factors and group membership
  - Groups individuals according to the outcome pattern over time ignoring the level

# Clustering by Shape Options

- Estimate slopes between neighboring observations and cluster on the “derived” observations
- Fit splines for each individual, differentiate, and cluster on coefficients of resulting derivative
- Use partition based cluster methods (like PAM) but use (i) the Pearson coefficient as a distance or dissimilarity measure

$$d_{corr}(\mathbf{x}, \mathbf{y}) = 1 - Corr(\mathbf{x}, \mathbf{y})$$

or the cosine-angle measure of dissimilarity

$$d_{cos}(\mathbf{x}, \mathbf{y}) = 1 - \frac{\sum_{j=1}^m x_j y_j}{(\sum_{j=1}^m x_j^2)(\sum_{j=1}^m y_j^2)}$$

- Vertical shifting individual trajectories

# Vertical Shifting

- For each individual, calculate

$$y_i^* = y_i - m_i^{-1} \sum_{j=1}^{m_i} y_{ij}$$

- Each individual now has mean zero and so level is removed from any resulting clustering
- Apply clustering technique to shifted data, e.g. finite mixture model

# Correlation Models for Vertical Shifted Data

- Without specifying group, suppose

$$\mathbf{y}_i^* = \lambda_i \mathbf{I}_{m_i} + \mu_i + \epsilon_i, \lambda \sim F_\lambda, \epsilon \sim N(), \Sigma)$$

where  $\mathbf{I}_{m_i}$  is an  $m_i$  length vector of 1s, and

$\mu_{ij} = \mu_k(t_{ij})$  is the  $j^{\text{th}}$  element of the vector of mean values for the  $k^{\text{th}}$  group evaluated at the observation times  $t_i$ . Thus,

$$\mathbf{y}_i^* = \mathbf{A}_i \mathbf{y}_i = \mu_i - \bar{\mu}_i + \epsilon_i - \bar{\epsilon}_i$$



# Correlation Models for Vertical Shifted Data

$$\text{Cov}(\mathbf{Y}_i^* - \mu_i) = \text{Cov}((\mathbf{A} - \mathbf{I}_{m_i})\mu_i + \mathbf{A}\epsilon)$$

## Two components of the covariance

- One induced by the averaging process
- One induced by (random) observation times

# Correlation Models for Vertical Shifted Data

## Observation Times Fixed

$$Cov(\mathbf{Y}^* - \mu) = \mathbf{A}\Sigma\mathbf{A}^T$$

suppressing the individual/group indices for simplicity ( $\Sigma$  is allowed to vary across clusters)

This covariance matrix is singular since  $\det(\mathbf{A}) = 0$

This naturally reflects the “loss” of one dimension

# Correlation Models for Vertical Shifted Data

## Observation Times Fixed

$$Cov(\mathbf{Y}^* - \mu) = \mathbf{A}\Sigma\mathbf{A}^T$$

- If  $\Sigma = \sigma^2\mathbf{I}$  (conditional independence with constant variance, then the induced covariance is exchangeable with negative correlation given by  $-1/(m-1)$  and variance decreases to  $\sigma^2(\frac{m-1}{m})$ )
- If original covariance is exchangeable with constant variance and correlation  $\rho$  then the induced covariance remains exchangeable with negative correlation and reduced variance  $\sigma^2(1-\rho)(\frac{m-1}{m})$

# Correlation Models for Vertical Shifted Data

## Observation Times Fixed

$$Cov(\mathbf{Y}^* - \mu) = \mathbf{A}\Sigma\mathbf{A}^T$$

If  $\Sigma = \sigma^2\mathbf{I}$  (conditional independence with constant variance, then the induced covariance is exchangeable with negative correlation given by  $-1/(m-1)$  and variance decreases to  $\sigma^2(\frac{m-1}{m})$ )

This induced exchangeable correlation is the lower bound for correlation in an exchangeable matrix

Thus, if “estimated”, the (true) parameter is on the boundary of the parameter space

# Correlation Models for Vertical Shifted Data

## Observation Times Random ( $\mu$ is random)

$$Cov(\mathbf{Y}^* - \mu) = m^{-2} \left( \sum_{j=1}^m Var(t_j) [\mu'(E(t_j))]^2 \right) \mathbf{1}\mathbf{1}^T + \mathbf{A}\Sigma\mathbf{A}^T$$

Sum of two non-invertible matrices, but the positive magnitude of the first matrix may counteract the negative correlations of the second.

# Correlation Models for Vertical Shifted Data

## Observation Times Random ( $\mu$ is random)

$$\text{Cov}(\mathbf{Y}^* - \mu) = m^{-2} \left( \sum_{j=1}^m \text{Var}(t_j) [\mu'(E(t_j))]^2 \right) \mathbf{1}\mathbf{1}^T + \mathbf{A}\Sigma\mathbf{A}^T$$

500 simulations of  $\mathbf{Y}_i = \mu_i + \epsilon_i$   $\epsilon_i \sim N(0, \Sigma_\rho)$

where the error covariance matrix is of exponential form with range  $\rho$

$$\mathbf{t} = \mathbf{T} + \tau \quad \tau \sim N(0, \sigma_\tau^2 \mathbf{I})$$

$$\mathbf{T} = (1, 2, \dots, 9, 10)$$

$$\mu_{ij} = \mu(t_{ij})$$

# Correlation Models for Vertical Shifted Data

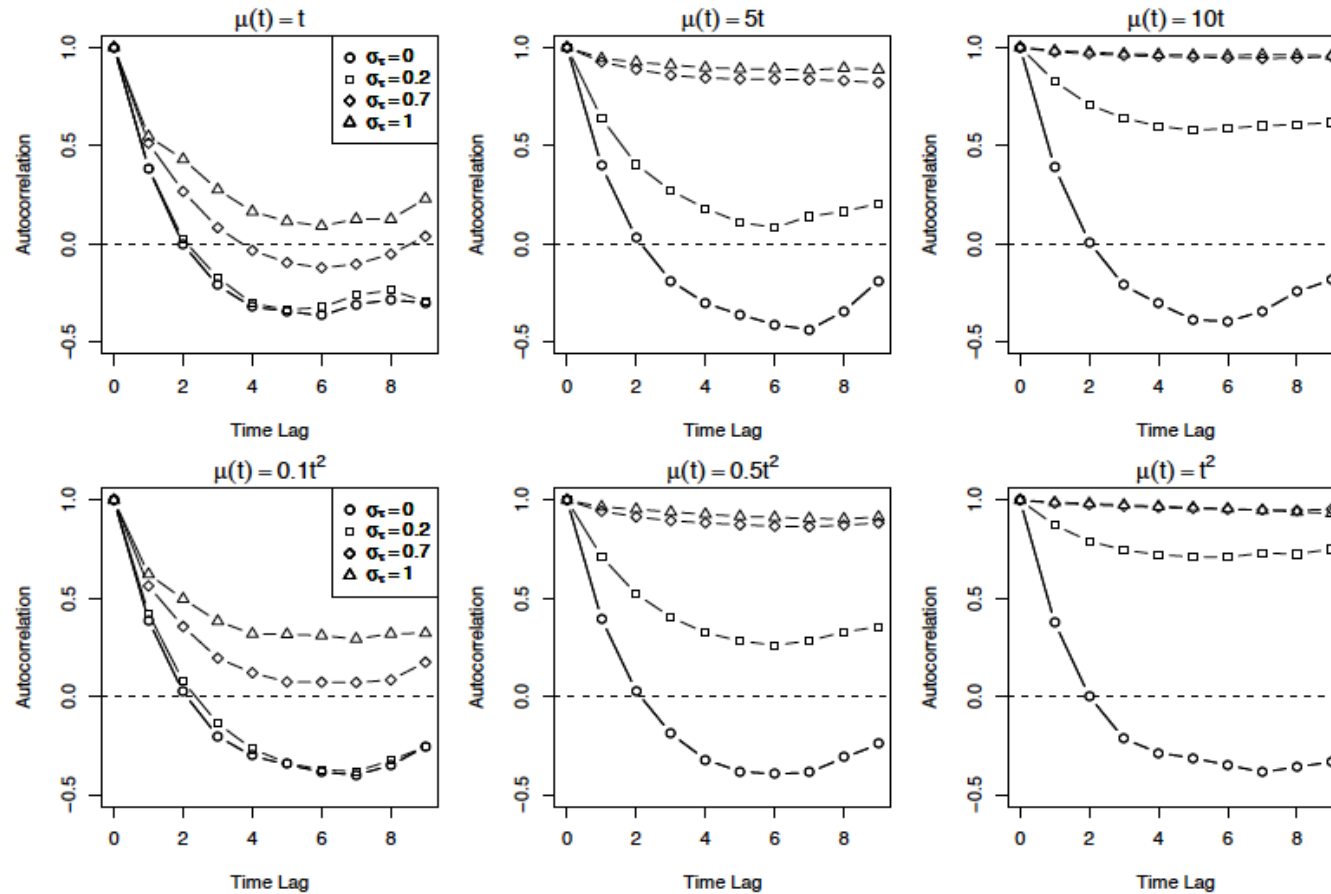
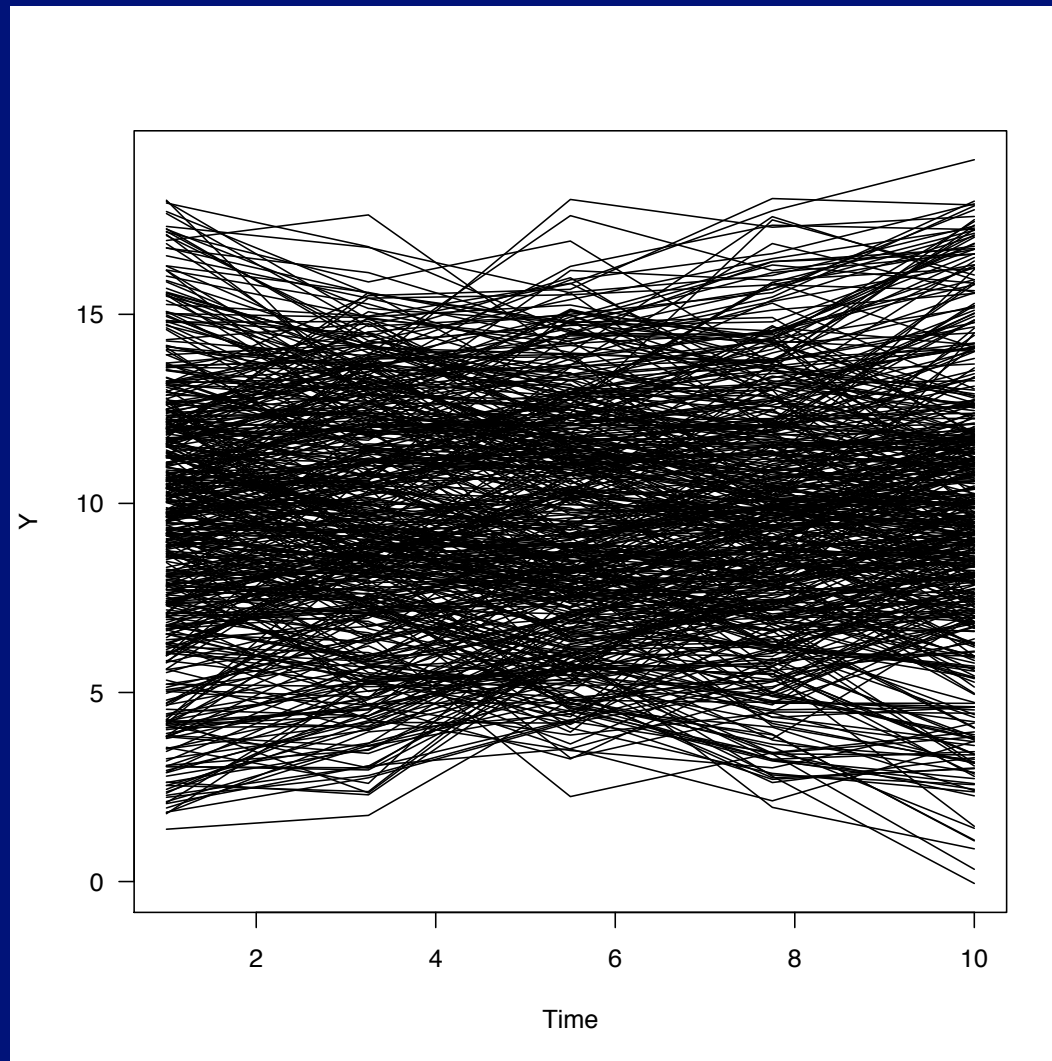


Figure 1: Estimated autocorrelation functions of the deviations from the mean from data generated with an exponential correlation error structure and random observation times under different mean functions,  $\mu(t)$ , and standard deviations of the random time perturbations,  $\sigma_\tau$ .

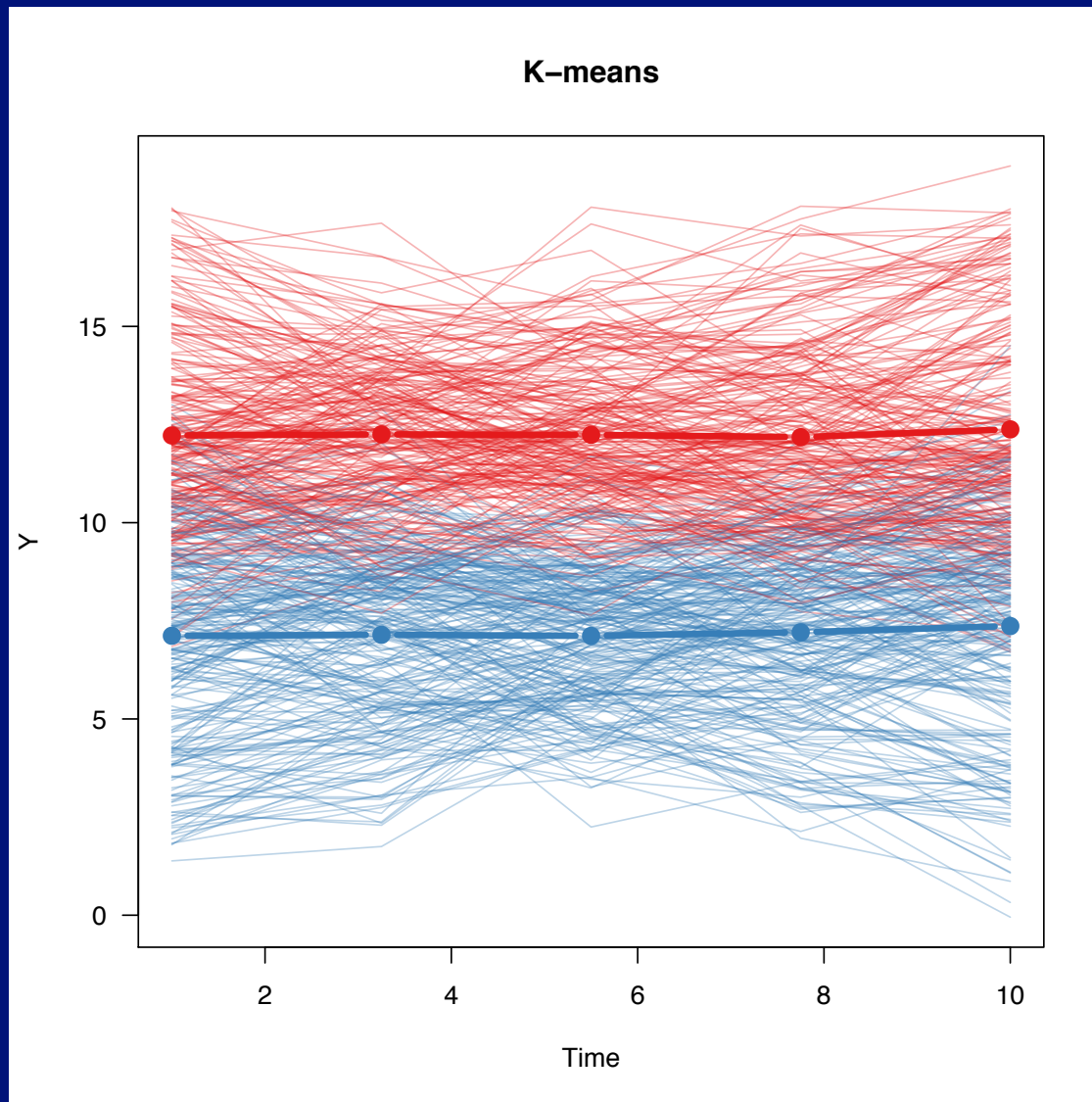
# Simulated Data



How could we group these individuals?

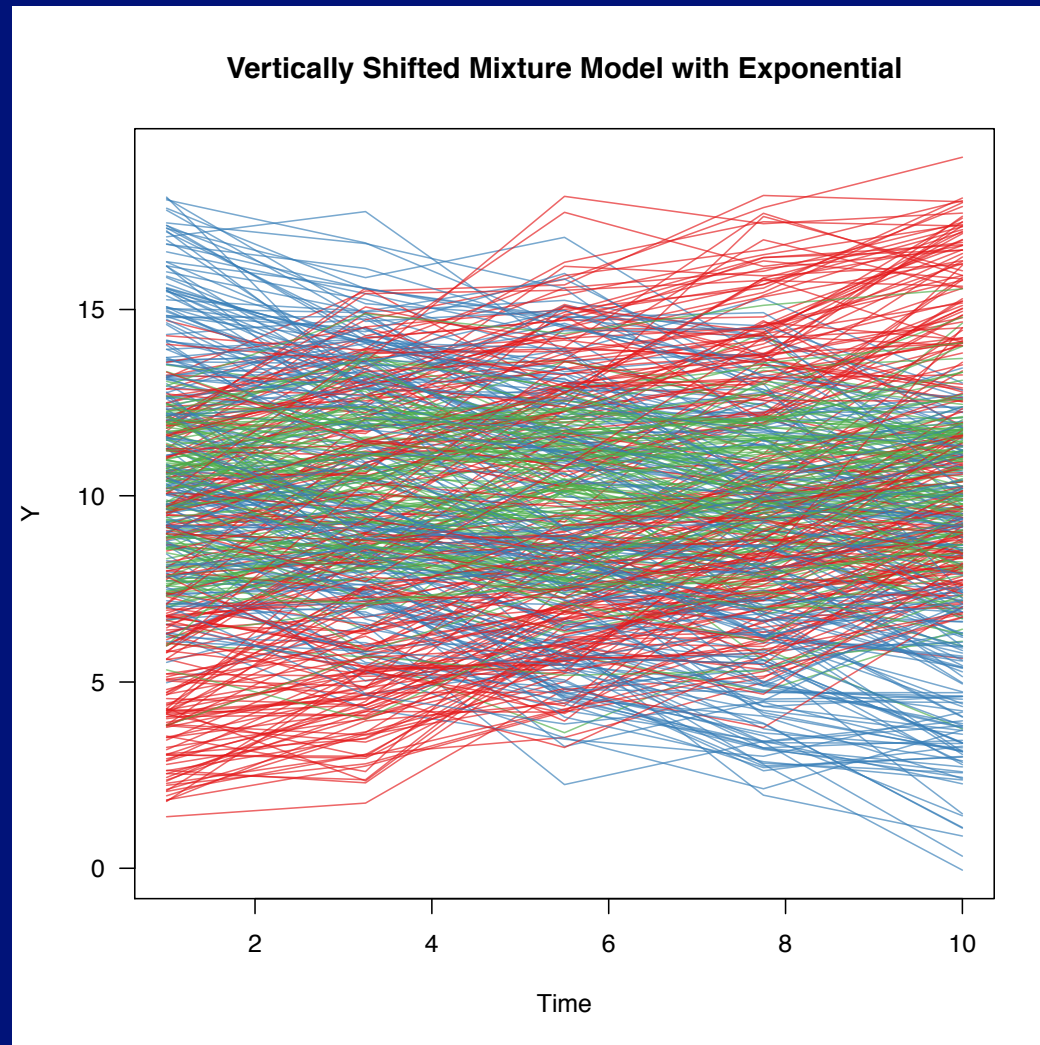


# Application to Simulated Data

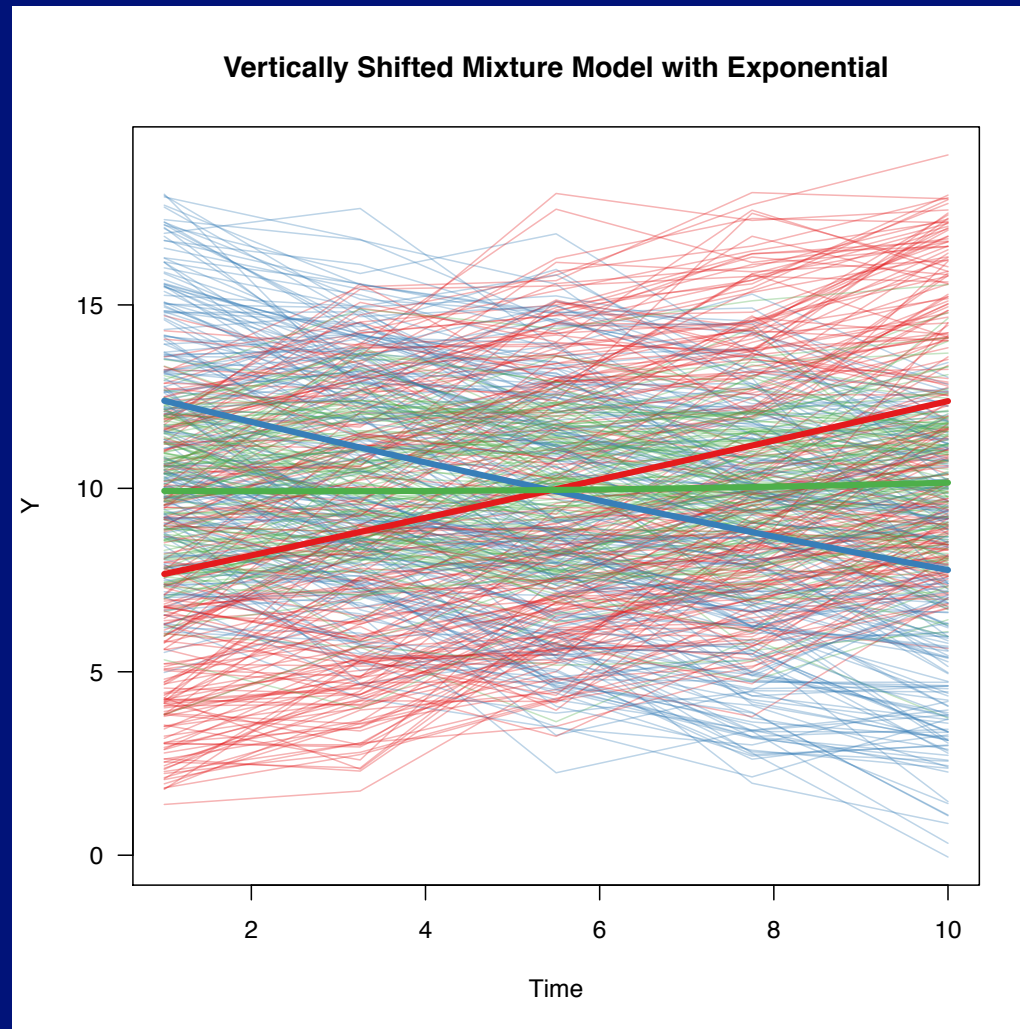


How would you describe—interpret—the group trajectories?

# Vertical Shifting Applied to Simulated Data



# Vertical Shifting Applied to Simulated Data



## 500 Simulations

$$\mu_1(t) = -1 - t$$

negative slope, low level

$$\mu_2(t) = 11 - t$$

negative slope, high level

$$\mu_3(t) = 0$$

zero slope middle level

$$\mu_4(t) = -11 + t$$

positive slope, low level

$$\mu_5(t) = 1 + t$$

positive slope, high level

**Mean functions evaluated at five equidistant points that span [1,10}  
Including ends of the interval**

## 500 Simulations

$$\mu_1(t) = -1 - t$$

negative slope, low level

$$\mu_2(t) = 11 - t$$

negative slope, high level

$$\mu_3(t) = 0$$

zero slope middle level

$$\mu_4(t) = -11 + t$$

positive slope, low level

$$\mu_5(t) = 1 + t$$

positive slope, high level

Two components to noise: random individual level perturbation  $N(0, \sigma_\lambda^2)$   
random measurement error across times  $N(0, \sigma_\epsilon^2)$

(exchangeable correlation)

# 500 Simulations

$\sigma_\epsilon$	$\sigma_\lambda$	$K = 2$	$K = 3$	$K = 4$	$K = 5$	MR
<b>Independent Gaussian Mixture</b>						
0.50	2.00	0.00	0.00	10.00	490.00	0.41
2.00	2.00	0.00	0.00	24.00	476.00	0.39
0.50	3.00	0.00	0.00	1.00	499.00	0.45
2.00	3.00	0.00	0.00	5.00	495.00	0.45
<b>K-means on Difference Quotients</b>						
0.50	2.00	0.00	500.00	0.00	0.00	0.00
2.00	2.00	483.00	17.00	0.00	0.00	0.38
0.50	3.00	0.00	500.00	0.00	0.00	0.00
2.00	3.00	483.00	17.00	0.00	0.00	0.38
<b>Correlation-based PAM</b>						
0.50	2.00	403.00	0.00	0.00	97.00	0.25
2.00	2.00	500.00	0.00	0.00	0.00	0.27
0.50	3.00	403.00	0.00	0.00	97.00	0.25
2.00	3.00	500.00	0.00	0.00	0.00	0.27
<b>Vertically Shifted Independent Mixture</b>						
0.50	2.00	0.00	499.00	0.00	1.00	0.00
2.00	2.00	0.00	498.00	2.00	0.00	0.05
0.50	3.00	0.00	499.00	0.00	1.00	0.00
2.00	3.00	0.00	498.00	2.00	0.00	0.05
<b>Vertically Shifted Exponential Mixture</b>						
0.50	2.00	0.00	500.00	0.00	0.00	0.00
2.00	2.00	0.00	499.00	1.00	0.00	0.05
0.50	3.00	0.00	500.00	0.00	0.00	0.00
2.00	3.00	0.00	499.00	1.00	0.00	0.05

Table 1: Frequency table of the number of groups chosen and average misclassification rate (MR) ( $K = 3$ ) for 500 replications of clustering methods applied to data generated under different values for  $\sigma_\epsilon$  and  $\sigma_\lambda$ .

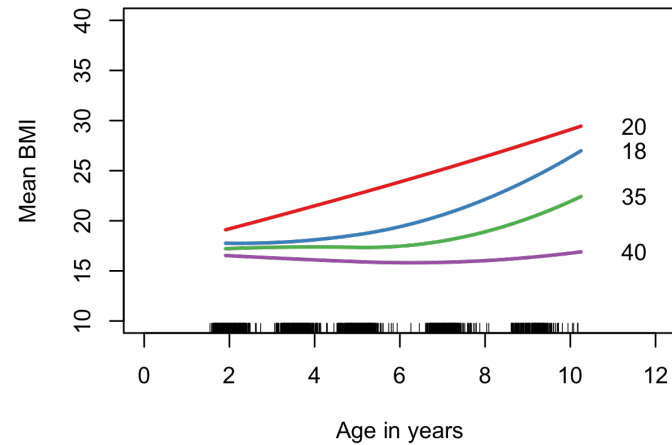
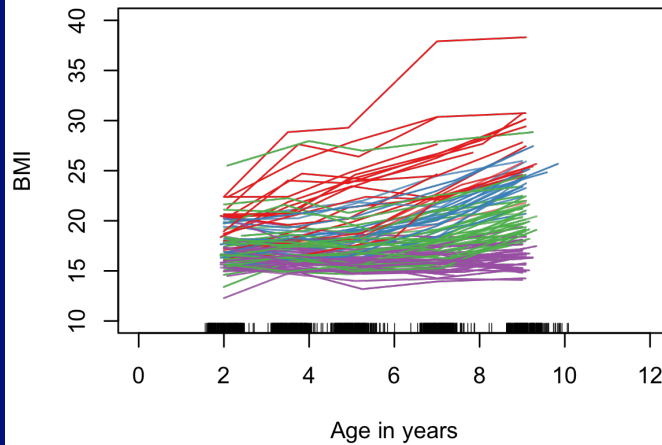
# Vertical Shifting with CHAMACOS

- **Two-part models**
  - First, use standard regression models to relate baseline predictors to BMI
  - Then, use vertically shifted shape clustering with (same or different baseline predictors for shape groups)
- **For BMI in the CHAMACOS data**
  - Works with irregularly sampled data
  - Includes a way to estimate the relationship between baseline risk factors and group membership
  - Groups individuals according to the outcome pattern over time ignoring the level

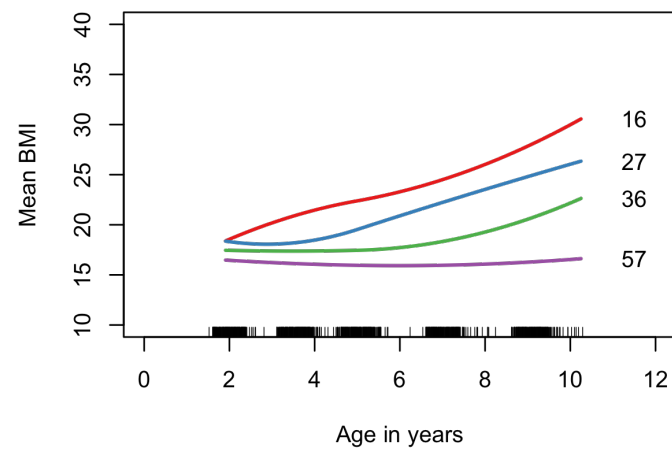
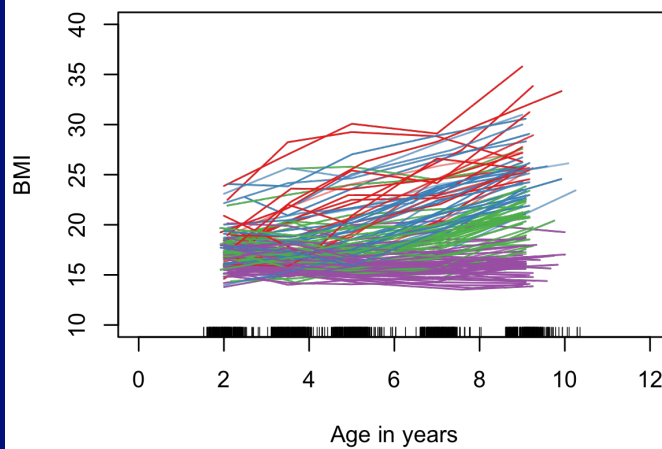


# Vertical Shifting with CHAMACOS

ODT : Boy



ODT : Girl





# Vertical Shifting with CHAMACOS

	Estimate	95% CI	$P >  z $
Red v. Purple : bmicat_preOverweight	2.494	(0.506, 12.294)	0.261
Red v. Purple : bmicat_preObese	34.068	(4.945, 234.699)	0
Red v. Purple : lng_lod2_ODT_pg	1.743	(0.343, 8.857)	0.503
Red v. Purple : yrsusa	0.97	(0.887, 1.062)	0.514
Blue v. Purple : bmicat_preOverweight	2.4	(0.408, 14.129)	0.333
Blue v. Purple : bmicat_preObese	8.636	(0.97, 76.902)	0.053
Blue v. Purple : lng_lod2_ODT_pg	6.679	(1.067, 41.812)	0.042
Blue v. Purple : yrsusa	1.048	(0.955, 1.15)	0.324
Green v. Purple : bmicat_preOverweight	1.45	(0.428, 4.917)	0.551
Green v. Purple : bmicat_preObese	15.561	(2.306, 105.031)	0.005
Green v. Purple : lng_lod2_ODT_pg	4.028	(0.598, 27.113)	0.152
Green v. Purple : yrsusa	0.953	(0.866, 1.048)	0.322

Table 1: Estimated odds ratios for Vertically Shifted Model with ODT for Boy

	Estimate	95% CI	$P >  z $
Red v. Purple : bmicat_preOverweight	6.258	(0.193, 203.281)	0.302
Red v. Purple : bmicat_preObese	64.22	(2.1, 1963.475)	0.017
Red v. Purple : lng_lod2_ODT_pg	0.358	(0.07, 1.846)	0.22
Red v. Purple : yrsusa	0.955	(0.785, 1.162)	0.646
Blue v. Purple : bmicat_preOverweight	1.785	(0.464, 6.86)	0.399
Blue v. Purple : bmicat_preObese	7.121	(1.069, 47.437)	0.042
Blue v. Purple : lng_lod2_ODT_pg	0.846	(0.276, 2.599)	0.771
Blue v. Purple : yrsusa	1.011	(0.922, 1.109)	0.815
Green v. Purple : bmicat_preOverweight	0.452	(0.15, 1.36)	0.158
Green v. Purple : bmicat_preObese	4.335	(0.998, 18.837)	0.05
Green v. Purple : lng_lod2_ODT_pg	0.637	(0.329, 1.233)	0.181
Green v. Purple : yrsusa	0.958	(0.885, 1.038)	0.295

Table 2: Estimated odds ratios for Vertically Shifted Model with ODT for Girl

# Further Thoughts

- Time-dependent covariates