

Pim P(110m-0011<E) = 1 VE N=00 CONVERGENCE IN PROBABILITY  $\lim_{N \to \infty} |E[|\hat{V}_{nL} - |\hat{V}_{nL}|^2] = 0$ CONVERGENCE IN MEAN SQUARE (ASYMPTOTICALLY CONSISTENT)

§2.1
Fitting a
linear model

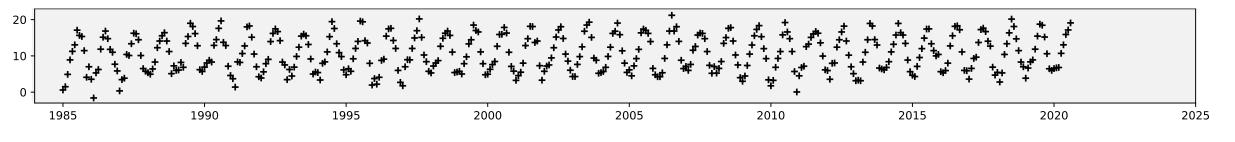
You've got to have models in your head. And you've got to array your experience – both vicarious and direct – on this latticework of models.

You may have noticed students who just try to remember and pound back what is remembered. Well, they fail in school and in life. You've got to hang experience on a latticework of models in your head.

Charlie Munger (business partner of Warren Buffet), A lesson on elementary, worldly wisdom as it relates to investment management & business.

## Monthly average temperatures in Cambridge, UK

What's a good model for this dataset?



Climate is stable? Temp(t) ~  $a + b \sin(2\pi(t + \phi)) + N(0, \sigma^2)$ 

Temperatures are increasing?

Temperatures are increasing, and the increase is accelerating?

> There are so many possible models. We want to make it easy to invent and fit new models, so we have time to explore all the possibilities.

The extremes are

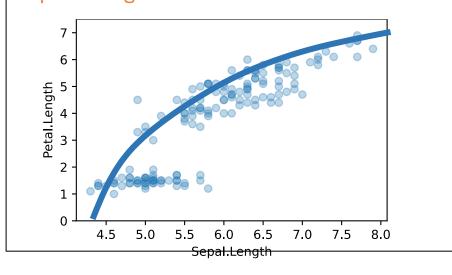
getting worse?

#### Example 2.1.1

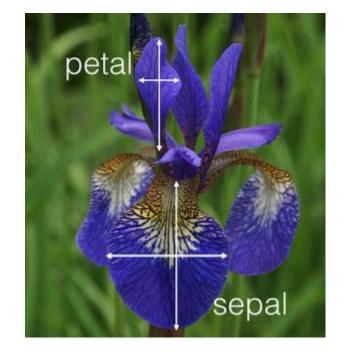
The Iris dataset has 50 records of iris measurements, from three species.

Petal. Length	Petal. Width	Sepal. Length	Sepal. Width	Species
1.0	0.2	4.6	3.6	setosa
5.0	1.9	6.3	2.5	virginica
5.8	1.6	7.2	3.0	virginica
4.2	1.2	5.7	3.0	versicolor

## How does Petal.Length depend on Sepal.Length?



Let's guess that for parameters  $\alpha$ ,  $\beta$ ,  $\gamma$ ,  $\sigma$  (to be estimated), Petal.Length  $\approx \alpha + \beta$  Sepal.Length +  $\gamma$ (Sepal.Length)<sup>2</sup>



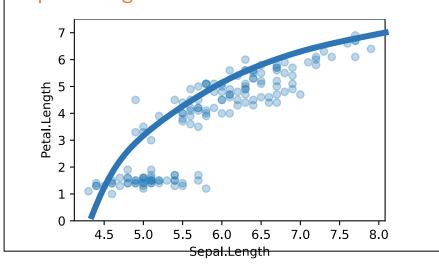
Dataset collected by Edgar Anderson and popularized by Ronald Fisher in 1936

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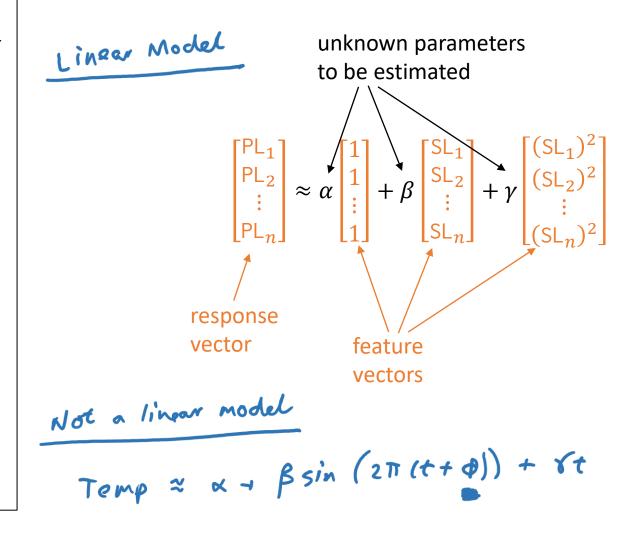
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Supervised learning Response & features are numeric Response is predicted by a linear combination of (known) feature vectors, weighted by unknown parameters. Models of this form are called *linear models* (because they're based on linear algebra).

They are flexible, and very fast to optimize.

Least squares estimation

Consider a linear model

$$y \approx \beta_1 e_1 + \dots + \beta_K e_k$$

"All models are wrong."

The vector of prediction errors is called the *residual vector*,  $\varepsilon = \gamma - (\beta_1 e_1 + \dots + \beta_K e_K)$ 

We can fit the model using *least squares estimation*. This means finding parameters  $\beta_1, \dots, \beta_K$  to minimize the mean square error

$$mse = \frac{1}{n} \sum_{i=1}^{n} \varepsilon_i^2$$

Petal.Length  $\approx \alpha + \beta$  Sepal.Length +  $\gamma$ (Sepal.Length)<sup>2</sup>

$$\begin{bmatrix} \mathsf{PL}_{1} \\ \mathsf{PL}_{2} \\ \vdots \\ \mathsf{PL}_{n} \end{bmatrix} \approx \alpha \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} + \beta \begin{bmatrix} \mathsf{SL}_{1} \\ \mathsf{SL}_{2} \\ \vdots \\ \mathsf{SL}_{n} \end{bmatrix} + \gamma \begin{bmatrix} (\mathsf{SL}_{1})^{2} \\ (\mathsf{SL}_{2})^{2} \\ \vdots \\ (\mathsf{SL}_{n})^{2} \end{bmatrix}$$

#### Fitting the model

```
2 one, SL, PL = np.ones(len(iris)), iris['Sepal.Length'], iris['Petal.Length']
```

3 model = sklearn.linear\_model.LinearRegression(fit\_intercept=False)

```
4 model.fit(np.column_stack([one, SL, SL**2]), PL)
```

```
5 (\alpha, \beta, \gamma) = model.coef_
```

#### Making predictions / getting fitted values from the model

```
6 newSL = np.linspace(4.2, 8.2, 20)
7 predPL = \alpha + \beta*newSL + \gamma*(newSL**2)
```

2

3

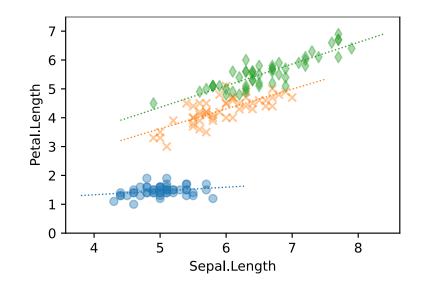
4

Petal.Length  $\approx \alpha + \beta$  Sepal.Length +  $\gamma$ (Sepal.Length)<sup>2</sup>  $\begin{array}{c|c} \Gamma L_{1} \\ PL_{2} \\ \vdots \\ PL_{n} \end{array} \approx \alpha \begin{array}{c} 1 \\ 1 \\ \vdots \\ 1 \end{array} + \beta \begin{array}{c} SL_{1} \\ SL_{2} \\ \vdots \\ SL \end{array} + \gamma \begin{array}{c} (SL_{1})^{2} \\ (SL_{2})^{2} \\ \vdots \\ (SL_{n})^{2} \end{array}$ skbarn linear model puts in the [!] feature for us by default. If we do ait want it we have to specify fit intercept = Folge. Fitting the model (cleaner code) SL, PL = iris['Sepal.Length'], iris['Petal.Length'] model = sklearn.linear\_model.LinearRegression() Making predictions / getting fitted values from the model (cleaner code) newSL = np.linspace(4.2, 8.2, 20) predPL = modul model.fit(np.column\_stack([SL, SL\*\*2]), PL) predPL = model.predict(np.column\_stack([newSL, newSL\*\*2]))

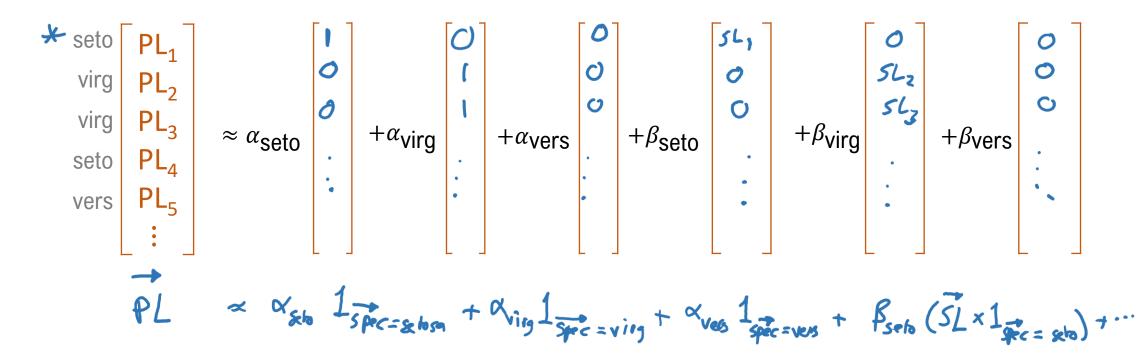
# §2.2 Feature design

How do we design features, so that linear models answer the questions we want answered?

### ONE-HOT CODING



This model hay 6 unknown parameters: <sup>CY</sup>set Xves Xving BSCF Bress Bring. So my model has 6 feature vectors.

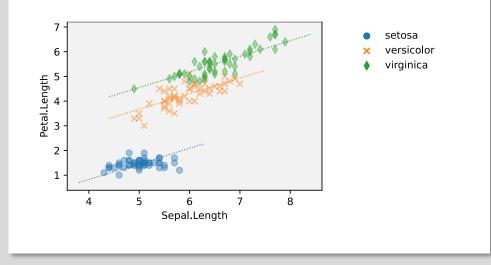


setosa versicolor

virginica

#### EXERCISE

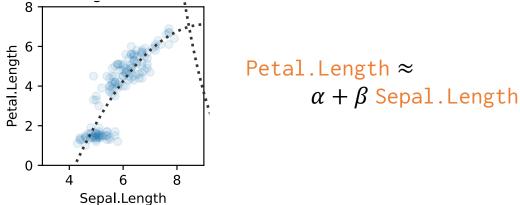
#### Fit the model with three parallel straight lines.

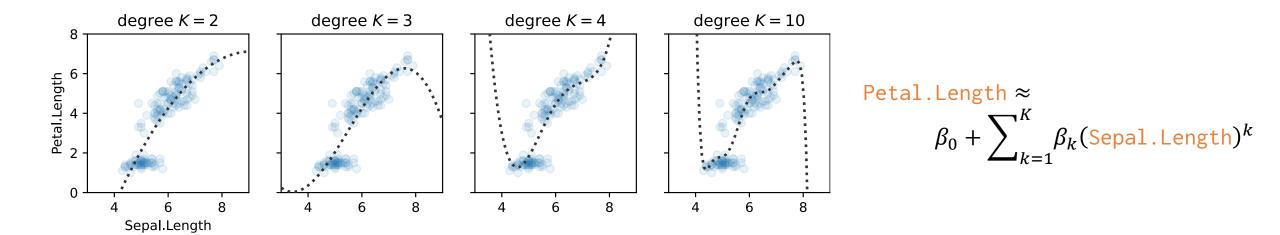


PL ~ « species + B 5L

X ast 1 spre = set + B SL

### NON-LINEAR RESPONSE





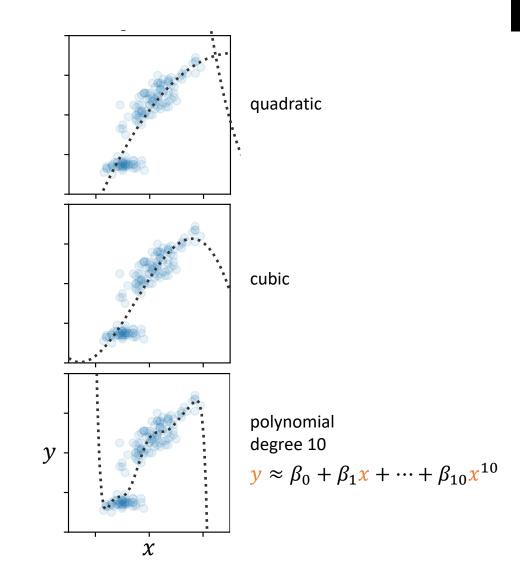
..Length ≈  $\alpha + \beta$  Sepal.Length + γ(Sepal.Length)<sup>2</sup> Q. Should we just keep adding more and more features to our model?

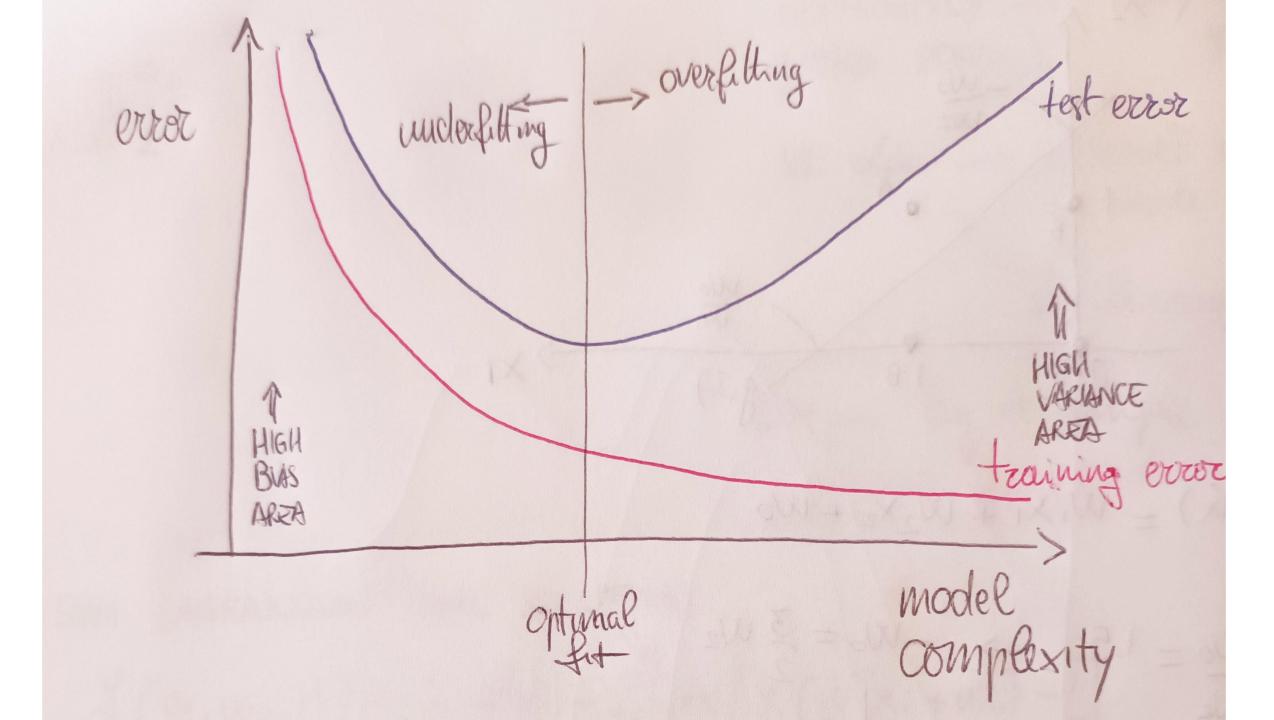
(seeing as the more features we add, the better we can fit the dataset)

A. No. If we did, we'd overfit.

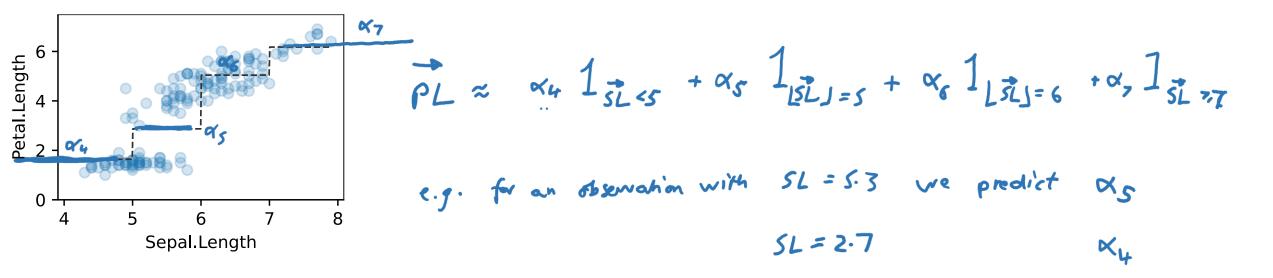
Only add in features that you (as a scientist) believe are relevant.

Or do model testing, e.g. evaluation on a holdout set. [§5–7]





NON-LINEAR RESPONSE via one-hot coding



## Memory lane





## How were listening modes calculated?

- Take a bunch of songs N<sub>s</sub> of three different music genres
- In a linear model
  - the amplitudes for each frequency would be the feature vectors;
  - the amplitude of the bass at the speaker would be the response vector
  - (dimensionality = N<sub>s</sub>)
- Calculate for each frequency the corresponding parameter

Scope of listening modes: improving audio quality

- By applying a frequency filter in the shape of the estimated distribution
  - Fostering some frequencies, reducing other frequencies



X



## Accurate listening mode





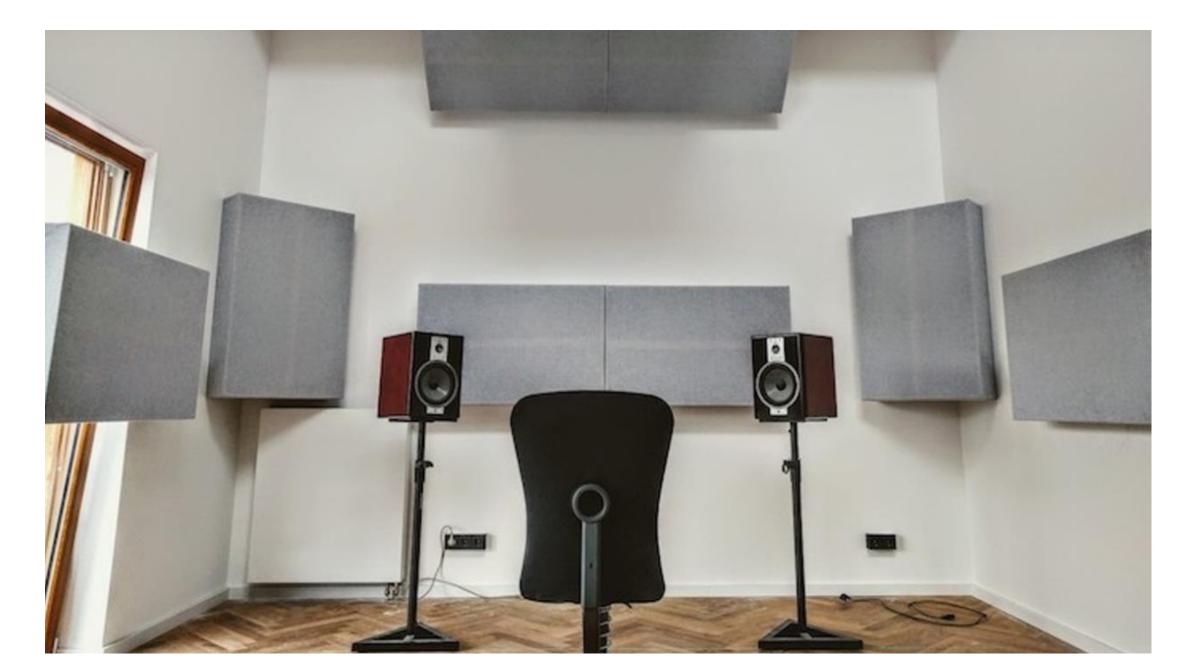
X



## Wrong listening mode



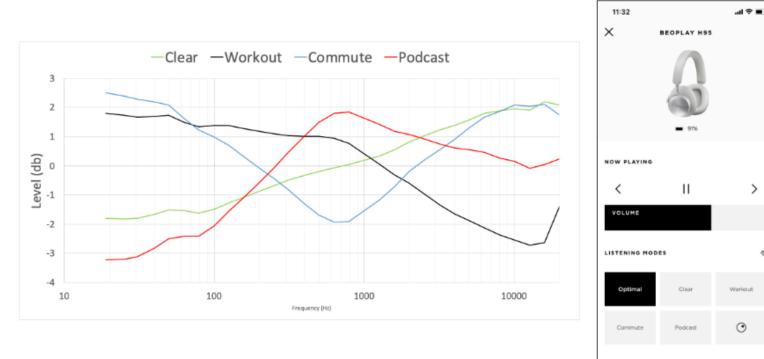
## Listening modes, today



### Listening modes, today

# **Listening Modes**

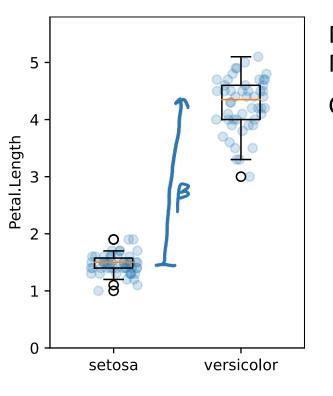
- These are the differences in frequency response relative to Optimal Mode (shown previous slide)
- Clear Mode is a upward sloped filter centered at 1kHz boosting treble above and cutting below
- Workout Mode does a similar EQ but sloped downwards
- Commute Mode boosts bass and treble
- Podcast Mode cuts bass below 250 Hz and boosts above that



Note: these curves are based on acoustical measurements of the left channel

SURROUNDINGS

### COMPARING GROUPS



Measurements for condition A:  $a = [a_1, a_2, ..., a_m]$ (ond Measurements for condition  $B: \mathbf{b} = [\mathbf{b}_1, \mathbf{b}_2, \dots, \mathbf{b}_n]$ A Can we use a linear model to compare A and B? A Α  $x \approx x_A 1_{cond} = A + x_B 1_{cond} = B$ ß В Or B  $\vec{x} = x + \beta \cdot \frac{1}{cons} = \beta$ . For a person of type A,  $x \approx \alpha$ B, x = x + B B measures the difference between the two groups.

X

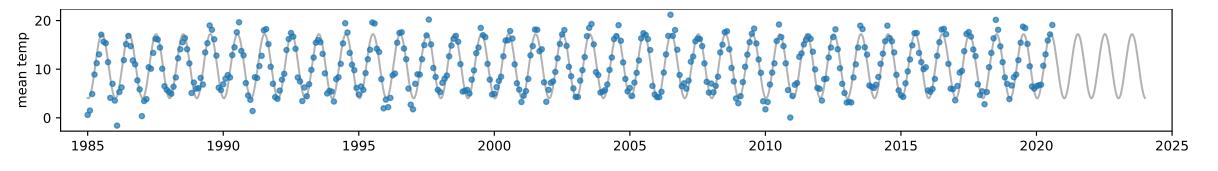
а,

d m

Ь,

P<sup>2</sup>

### PERIODIC PATTERN

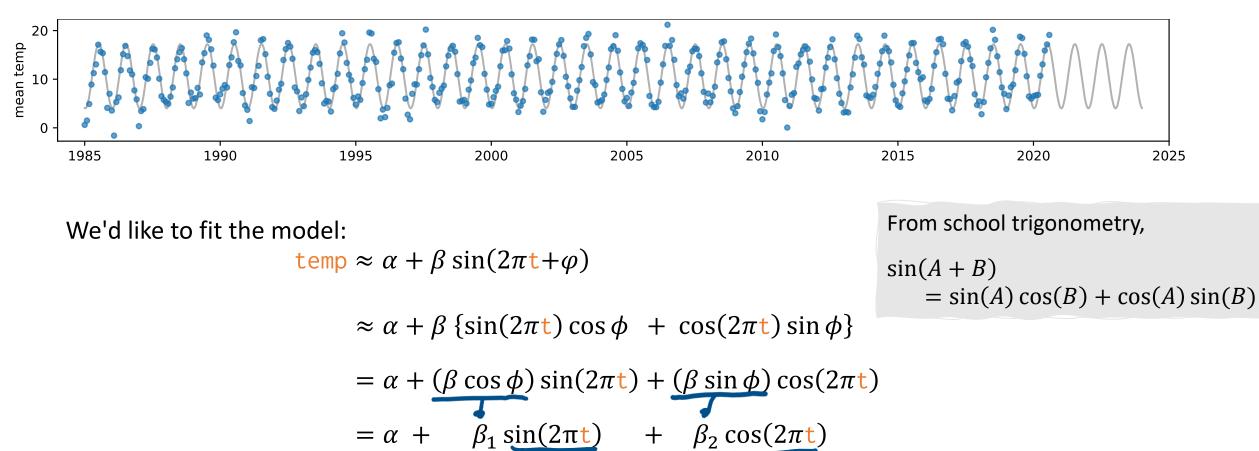


We'd like to fit the model:

 $\mathsf{temp} \approx \alpha + \beta \sin(2\pi \mathsf{t} + \varphi)$ 

It looks like we can't use sklearn.LinearRegression. That's only for linear models, e.g. temp  $\approx \alpha + \beta e + \gamma f$ 

### PERIODIC PATTERN



+

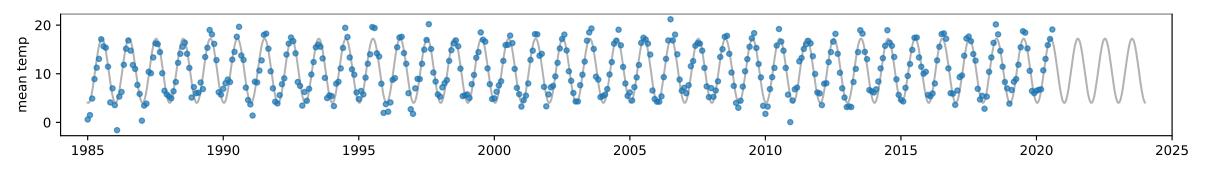
 $\beta_2$ 

 $\beta_1 e$ 

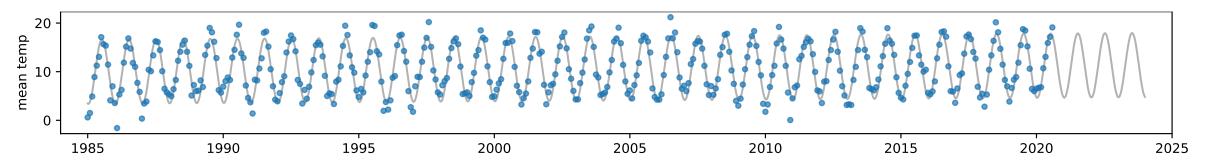
 $= \alpha +$ 

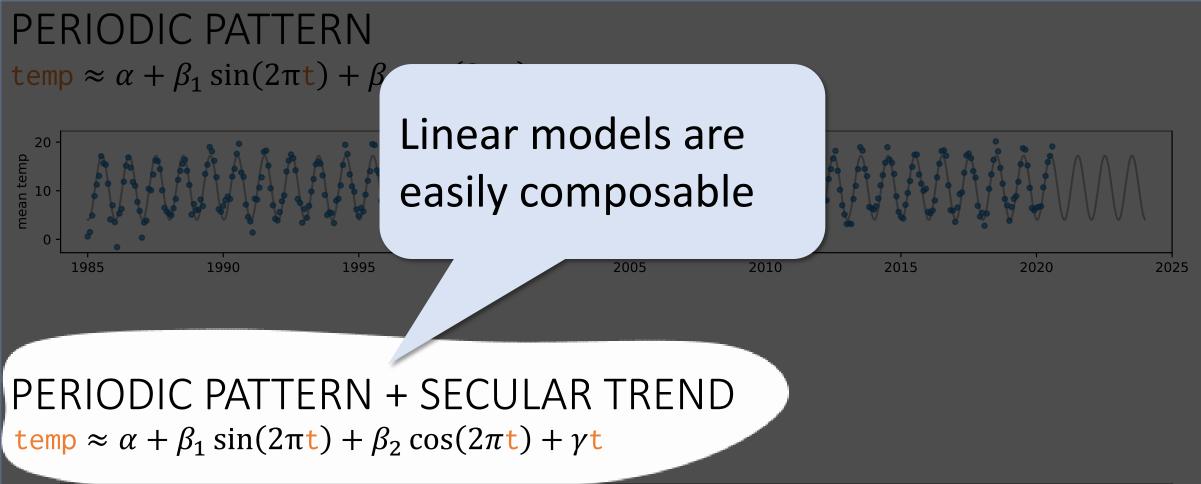
a linear model with feature vectors 1,  $sin(2\pi t)$ ,  $cos(2\pi t)$ 

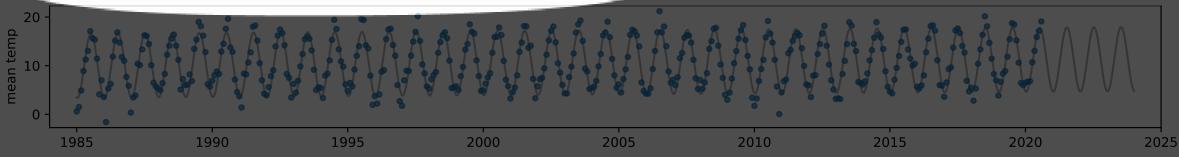
### PERIODIC PATTERN temp $\approx \alpha + \beta_1 \sin(2\pi t) + \beta_2 \cos(2\pi t)$



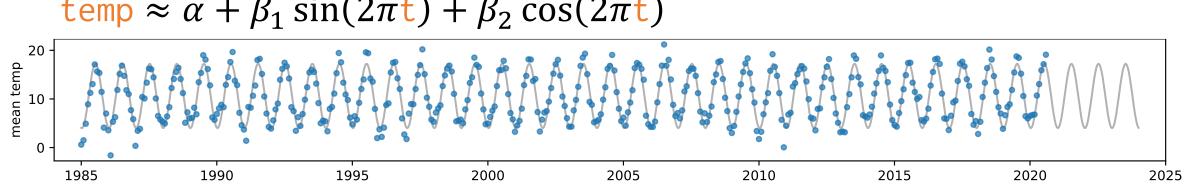
### PERIODIC PATTERN + SECULAR TREND temp $\approx \alpha + \beta_1 \sin(2\pi t) + \beta_2 \cos(2\pi t) + \gamma t$







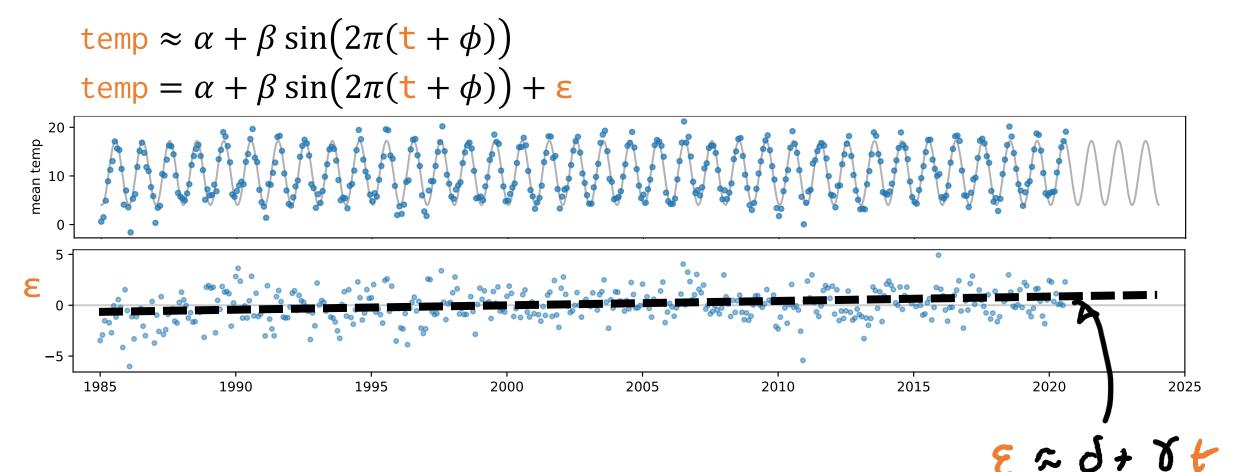
#### With our periodic model ...



 $temp \approx \alpha + \beta_1 \sin(2\pi t) + \beta_2 \cos(2\pi t)$ 

... how do we *discover* we should add a secular term  $+\gamma t$ ?

If we hadn't thought to include climate change in our temperature model ...



This suggests a revised model ...

 $\mathsf{temp} = \alpha' + \beta' \sin(2\pi(\mathsf{t} + \phi)) + \gamma \mathsf{t} + \varepsilon$ 

# Diagnosing a linear model

After fitting a model

- model.fit(..., y)
- 1. Compute the residuals
  - ε = y model.predict()
- Plot ε every way we can think of.
   If there's a systematic pattern, add a feature that describes it.