

Machine Learning

Held by Prof. D. Loiacono at Politecnico di Milano 2023/2024

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1. Disclaimers and preface

These notes were taken during AY 2023/2024 using older material, your mileage may vary. They're meant to accompany the lectures and in no way aim to substitute a professor yapping away at an iPad 30m away.

For any questions/mistakes you can reach me [here](#).

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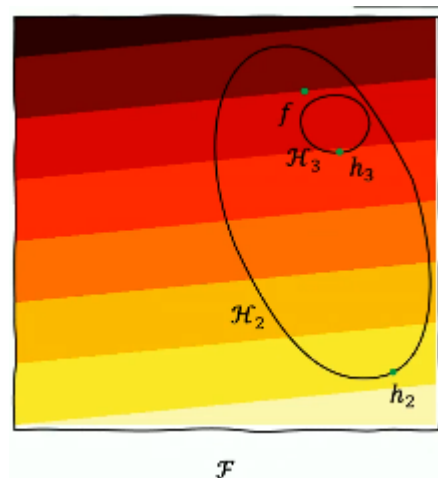
2. Overview of supervised learning

2.1. What kind of problem are we trying to solve ?

We're trying to find/approximate a function f , which is "generating" our dataset D .
The steps are:

- Define a loss function L
- Choose the hypothesis space \mathcal{H}
- Find in \mathcal{H} an approximation h of f that minimizes L .

We might be tempted to enlarge the hypothesis space, but if we do that we can have a larger risk of approximating f very poorly as we don't know if we'd be expanding \mathcal{H} in the correct direction.



2.2. The elements of a supervised learning algorithm

- The representation meaning the hypothesis space \mathcal{H} and how it's designed.
- The evaluation, so how the loss function is designed.
- The optimization algorithm, how you're looking for candidate solutions.

Some examples of representations are:

- Linear models.
- Neural networks.

Evaluation:

- Accuracy, the % of time you're correctly classifying a datapoint.
- If you have a regression problem you could use MSE.

Finally, the optimization technique really depends on your loss function:

- Gradient descent.
- Greedy search.
- Linear programming (in cases where you have some sort of constraint)

3. A supervised learning taxonomy

There are many ways you can try to classify/organize/compare, some more general paradigms other than the previous three.

3.1. Parametric vs Non-parametric

There are some learning algorithms for which the design of the hypothesis space is such that when you decide to apply a learning algorithm to a problem you have to define a set of parameters that are fixed and won't change with the data these are called **Parametric**, while non parametric models scale the number of parameters with the training set.

3.2. Frequentist vs Bayesian

No meaningful definition given by the prof here.

3.3. Direct, discriminative or generative

They're different ways to see the same problem from different perspectives, in the **direct** approach you don't really care about the probabilistic interpretation, you're just optimizing the loss function. In the **discriminative** you interpret your problem as a conditional density $p(t|x)$ you essentially try to learn the distribution and then compute the expected mean. In the **generative** tries to model the *joint* density $p(x,t)$, this allows to then infer the conditional density and generate novel samples by computing the conditional mean.

4. Linear regression

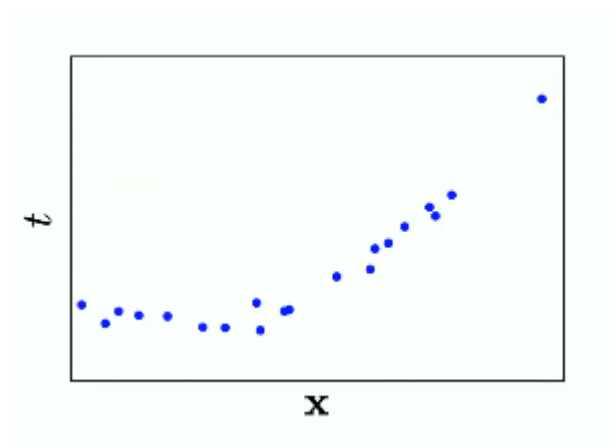
We'll focus on regression as our first type of problem, more specifically on linear models to solve.

References:

- *Pattern recognition and Machine learning*, Bishop

4.1. The model

We want to learn an approximation function $f(x)$:

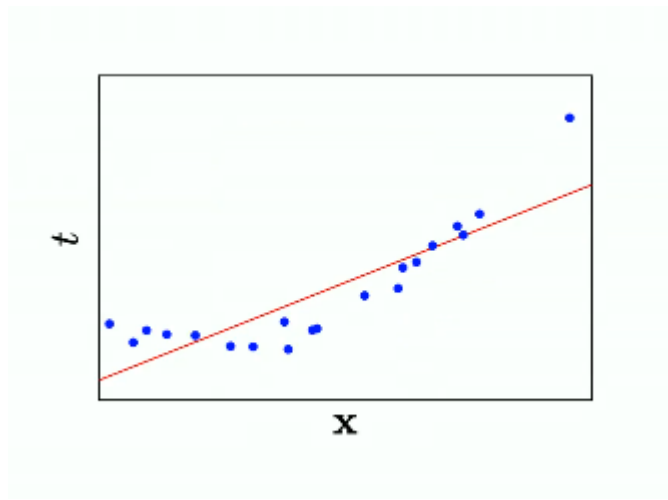


We start from a dataset

$$\mathcal{D} = \langle x, t \rangle \implies t = f(x)$$

How do we model f ? How do we evaluate our approximation ? How do we optimize our approximation ?

In linear regression we model $f(x)$ with linear functions.



A model like this clearly has room for improvement but we can very easily explain what the model is doing. Another important property is that we can solve this analytically, this is better since models like NN are hard to "debug", in this case nothing can go wrong, we have algorithms with guaranteed convergence etc...

Another thing is that these models can capture non linear interactions.

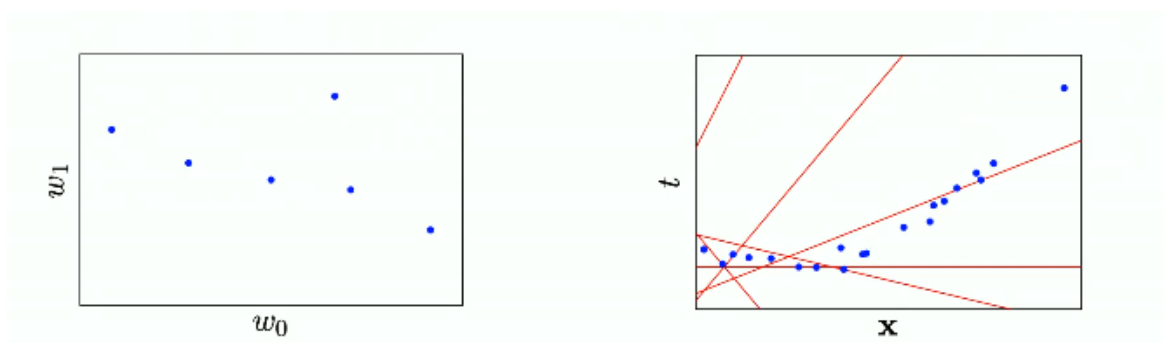
The simplest linear model can be defined as:

$$y(x, w) = w_0 + \sum_{j=1}^{D-1} w_j x_j = w^T x$$

Where:

- $x = (1, x_1, \dots, x_{D-1})$ is our input variable expressed as a vector to which we prepend 1.
- w_0 is called the bias parameter

My hypothesis space is 2D space with the two weights as dimensions.



Each point in the left image is a "solution" to our problem (a red line). We now need to *evaluate* these solutions, we need to define an error.

4.2. Loss function and optimization

A convenient error loss function is the sum of squared errors (SSE):

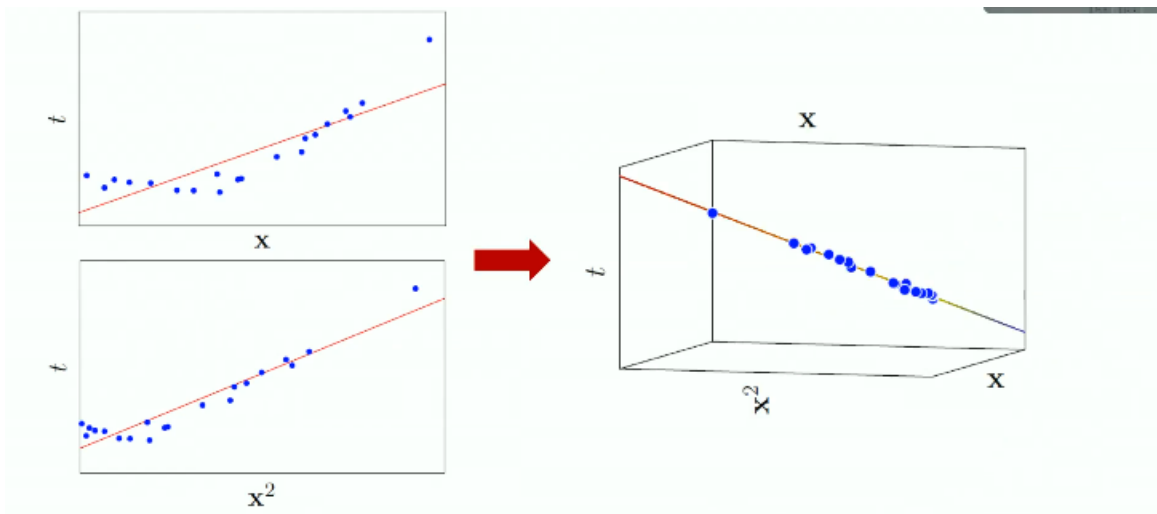
$$L(w) = \frac{1}{2} \sum_{n=1}^N (y(x_n, w) - t_n)^2$$

The reason we take the square instead of the absolute value is that ???

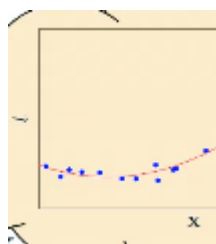
5. Linear models and basis functions

How do we capture non-linear relationships ? We can use basis functions. Instead of using x_1, x_2, \dots, x_n we can use a set of non-linear functions ϕ_i (as many as we want), my linear model will be *linear* with respect to the basis functions (we're learning the weights).

In this example i generate 2 weights for each sample, one for x and for x^2 .



If we now plot this approximation into the original space:



If we have problem specific knowledge we can pinpoint some sort of specific basis functions but generally there's no rule.

There are *families* of basis functions such as :

- Polynomial :

$$\phi_j(x) = x^j$$

- Gaussian :

$$\phi_j(x) = \exp\left(-\frac{(x - \mu_j)^2}{2\sigma^2}\right)$$

- Sigmoidal:

$$\phi_j(x) = \frac{1}{1 + \exp\left(\frac{\mu_j - x}{\sigma}\right)}$$

This can be hyper-parameterized (idk how to spell it), more on this later in the course.

6. Least squares

From now on we'll assume that our problem is solved in the feature space and not in the input space (in the basis function space in the previous case).

Prof emphasizes how important linear algebra is for this course...

$$L(w) = \frac{1}{2}RSS(w) - \frac{1}{2}(t - \phi w)^T(t - \phi w)$$

Where t is a vector with all the targets and ϕ is a matrix where on each row you have all the samples in the feature space (so on the cols you'd have features).

If you have to optimize this function with respect to a w . Let's assume it's a scalar value, how do we find the optimal value of w ?

Get the derivative and put the derivative equal to 0, right ?

Ordinary Least Squares

- For linear models, a closed-form optimization of the RSS, known as **least squares**, starting from the matrix form of the loss function:

$$L(\mathbf{w}) = \frac{1}{2}RSS(\mathbf{w}) = \frac{1}{2}(\mathbf{t} - \Phi\mathbf{w})^T(\mathbf{t} - \Phi\mathbf{w})$$

► where $\Phi = (\phi(\mathbf{x}_1), \dots, \phi(\mathbf{x}_N))^T$ and $\mathbf{t} = (t_1, \dots, t_N)^T$

- We can compute first a second derivative of $L(w)$ to find the optimal w

$$\frac{\partial L(\mathbf{w})}{\partial \mathbf{w}} = -\Phi^T(\mathbf{t} - \Phi\mathbf{w}) \qquad \frac{\partial^2 L(\mathbf{w})}{\partial \mathbf{w} \partial \mathbf{w}^T} = \Phi^T \Phi$$

$$\Rightarrow \hat{\mathbf{w}}_{OLS} = (\Phi^T \Phi)^{-1} \Phi^T \mathbf{t}$$

This is an analytical solution, the caveat is i need the matrix to not be singular cause then i'd have to invert it. It's also slow as fuck for large datasets.

You can instead apply **Gradient descent** (or SGD).

Sequential Learning

- Closed-form optimization (OLS) is not feasible with large dataset
- Instead, a **stochastic** (or **sequential**) gradient descent is possible
- **Least Mean Square** (LMS) algorithm:

$$L(\mathbf{x}) = \sum_n L(x_n)$$

$$\Rightarrow \mathbf{w}^{(n+1)} = \mathbf{w}^{(n)} - \alpha^{(n)} \nabla L(x_n)$$

$$\Rightarrow \mathbf{w}^{(n+1)} = \mathbf{w}^{(n)} - \alpha^{(n)} \left(\mathbf{w}^{(n)T} \phi(\mathbf{x}_n) - t_n \right) \phi(\mathbf{x}_n)$$

- α is called learning rate and to guarantee convergence:

$$\sum_{n=0}^{\infty} \alpha^{(n)} = +\infty \qquad \sum_{n=0}^{\infty} \alpha^{(n)^2} < +\infty$$

Prof then talks about geometric interpretation of OLS (Ordinary least squares).

7. Multiple outputs

What if my target is a multitude regression targets ?

In practice it's just like running multiple regression problems in parallel, the main optimization is using the same basis functions.

8. Regularization

What happens if i use too many features ? (p.22 to 24 of 3).

We basically add a regularizing term to the loss function that takes into account how many features we added. We can extend the loss function to take into account the complexity of our model

$$L(\mathbf{w}) = L_D(\mathbf{w}) + \lambda L_W(\mathbf{w})$$

where $L_W(\mathbf{w})$ accounts for model complexity and λ is the **regularization** coefficient and w is the vector of model coefficients (or weights).

We can design $L_W(\mathbf{w})$ in many ways, we'll look at two in particular

8.1. Ridge regression (L2 regularization)

We define $L_W(w)$ as

$$L_W(\mathbf{w}) = \frac{1}{2} \mathbf{w}^T \mathbf{w} = \frac{1}{2} \|\mathbf{w}\|_2^2$$

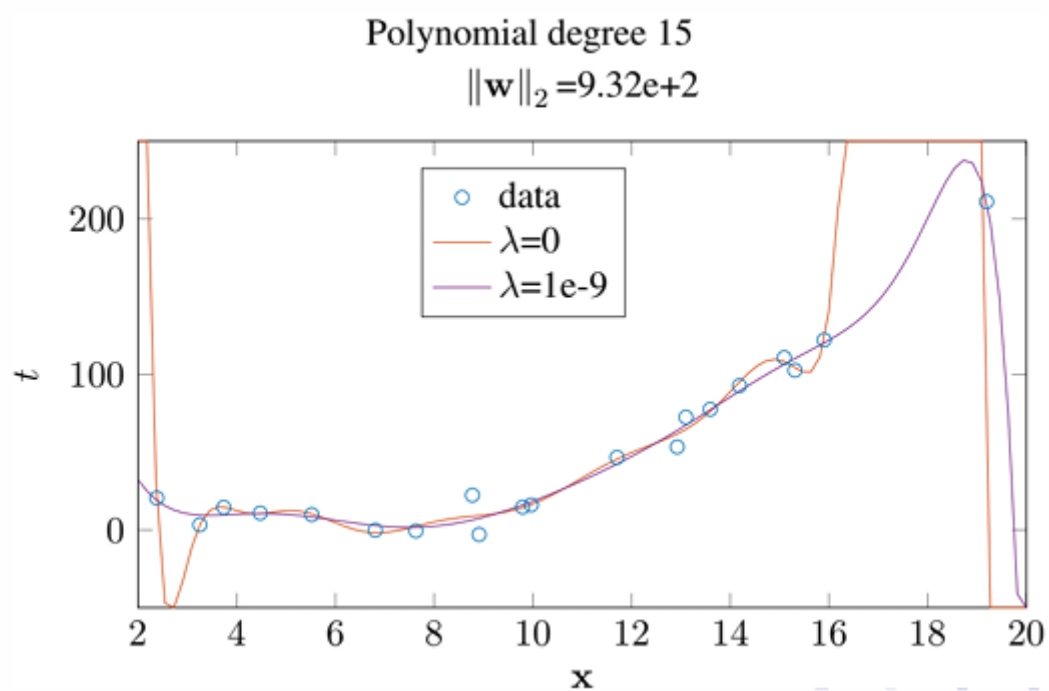
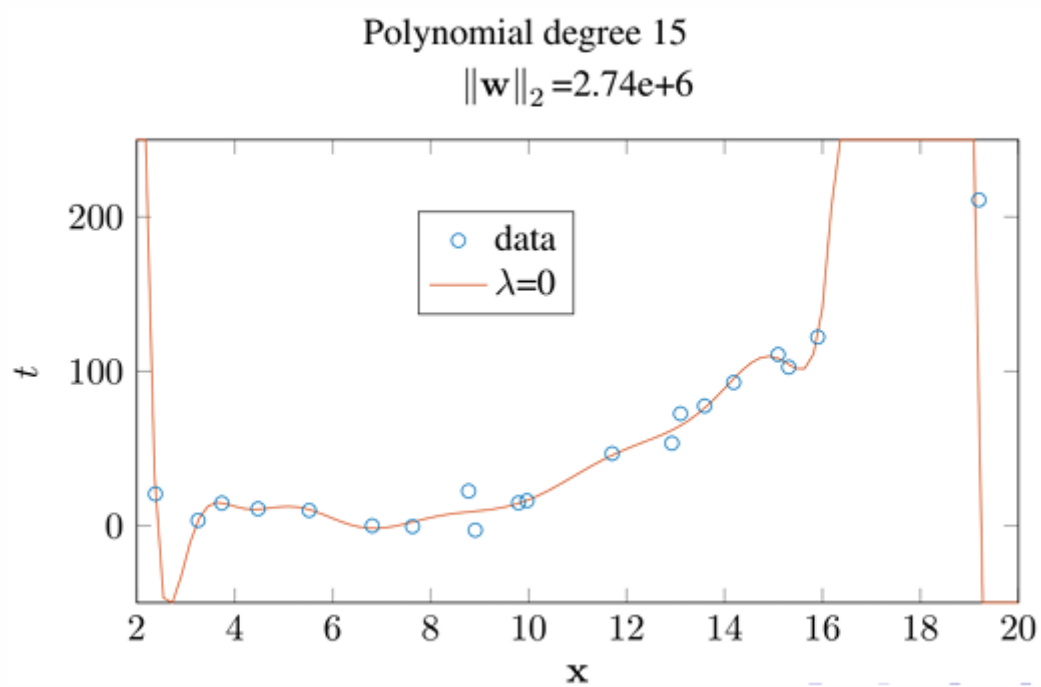
where the loss function $L(w)$ ends up being

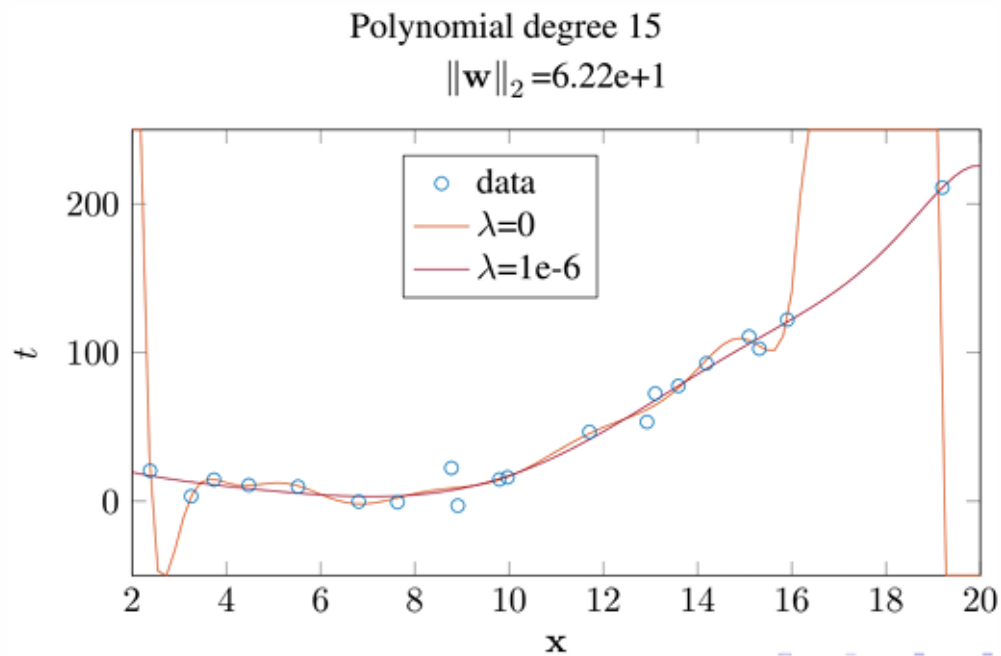
$$L(\mathbf{w}) = \frac{1}{2} \sum_{i=1}^N (t_i - \mathbf{w}^T \phi(\mathbf{x}_i))^2 + \frac{\lambda}{2} \|\mathbf{w}\|_2^2$$

the closed form (estimator) is

$$\hat{\mathbf{w}}_{ridge} = (\lambda \mathbf{I} + \Phi^T \Phi)^{-1} \Phi^T \mathbf{t}$$

We're essentially an L^2 penalty





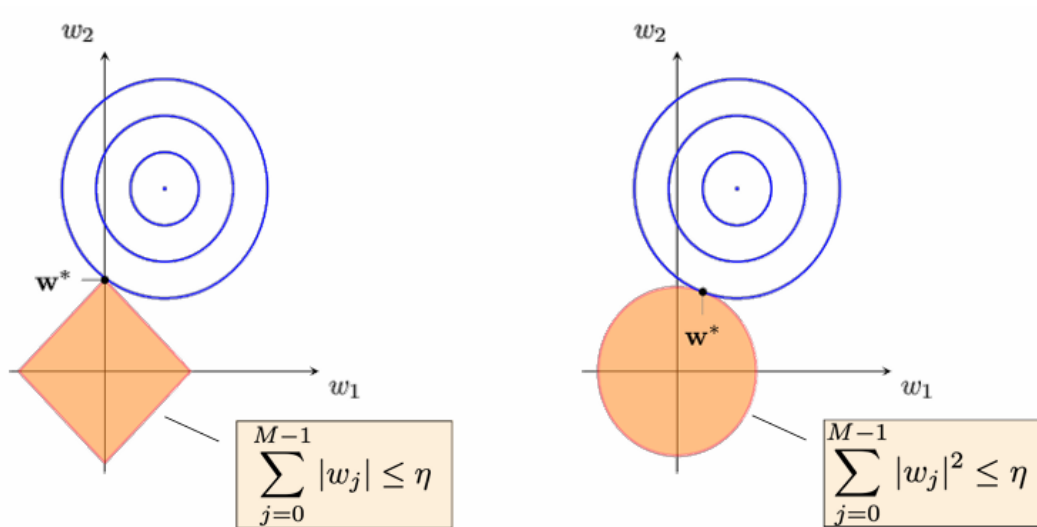
8.2. Lasso regression (L1 regularization)

Lasso stands for *least absolute shrinkage and selection operator*

$$L_W(\mathbf{w}) = \frac{1}{2} \|\mathbf{w}\|_1 = \frac{1}{2} \sum_{j=0}^{M-1} |w_j|$$

$$L(\mathbf{w}) = \frac{1}{2} \sum_{i=1}^N (t_i - \mathbf{w}^T \phi(\mathbf{x}_i))^2 + \frac{\lambda}{2} \|\mathbf{w}\|_1$$

When λ is large enough some weights might be **equal to zero**. That's because we're basically creating a constraint region in the shape of a diamond (or a cross-polytope) in the weight space. When the loss function's contours (ellipsoids) are minimized under this constraint, the solution often occurs at the corners of the diamond.



9. Least squares and Maximum likelihood

We can also approach regression in a probabilistic way by defining a probabilistic model that maps input x to outputs t using some unknown parameters w

$$y(x, w)$$

we then model the **likelihood** i.e. the probability that observed data \mathcal{D} is generated by a given set of parameters w

$$p(\mathcal{D}|w)$$

we then estimate those parameters by maximizing the likelihood that those w generated our observed sample

$$w_{ML} = \arg \max_w p(\mathcal{D}|w)$$

In the case of linear regression our model can be defined as:

$$t = y(x, w) + \epsilon = w^T \phi(x) + \epsilon$$

where $y(x, w)$ is taken to be a linear model disrupted by Gaussian noise $\epsilon \sim \mathcal{N}(0, \sigma^2)$

Given a dataset \mathcal{D} of N samples with inputs $X = \{x_1, \dots, x_N\}$ and outputs $T = \{t_1, \dots, t_N\}^T$

$$p(\mathcal{D}|w) = p(\mathbf{t}|\mathbf{X}, w, \sigma^2) = \prod_{n=1}^N \mathcal{N}(t_n | w^T \phi(\mathbf{x}_n), \sigma^2)$$

Since we're assuming the datapoints to be i.i.d. the likelihood function is the product of individual Gaussian distributions for each data points.