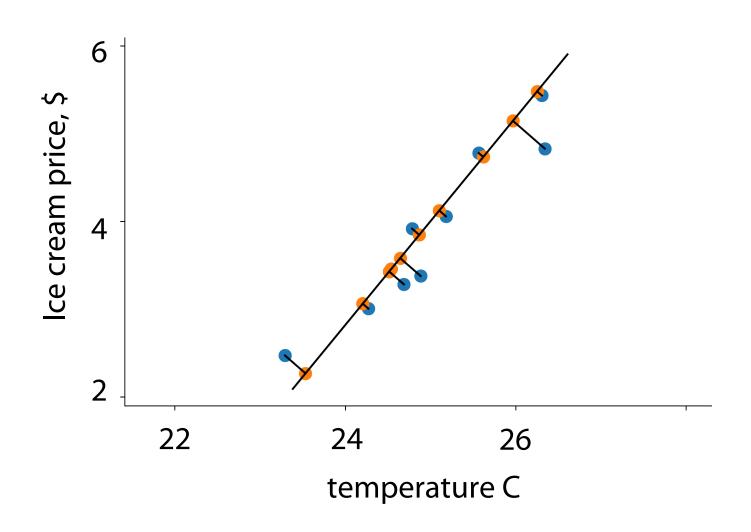
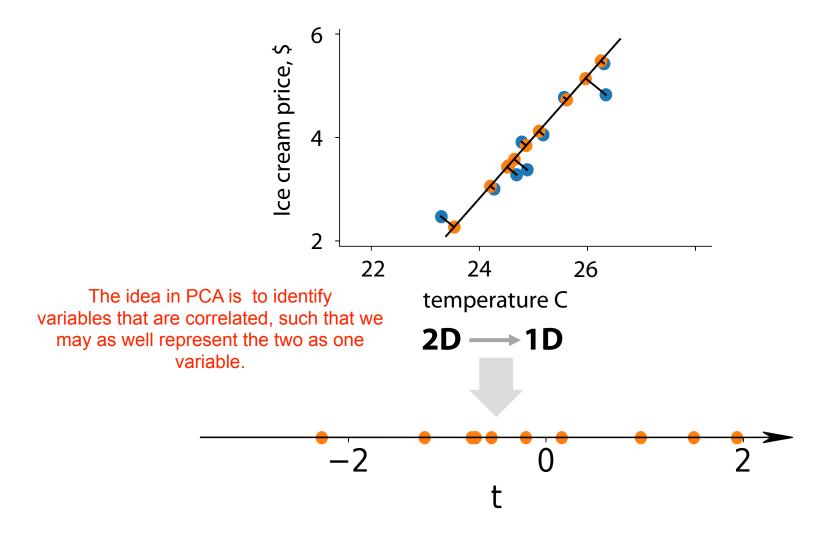
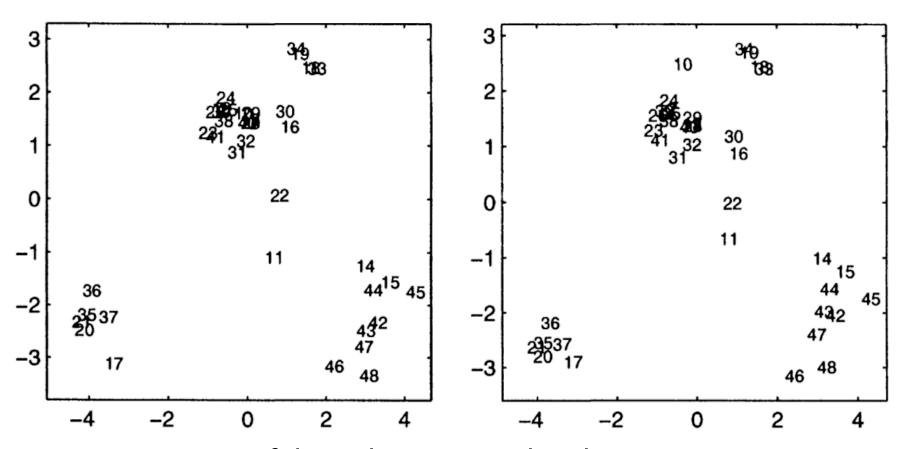
Ice Cream conspiracy









Projection of the Tobamovirus data by using PCA on the full data set and PPCA with 136 missing values

[source: Tipping, M. E., & Bishop, C. M. (1999). Probabilistic principal component analysis]

So the good thing about probabilistic PCA is that even with missing values, they are pretty much the same.

re cream price, \$ noithlos la stast.

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 $x_i = (1,1)t_i + (25,4)$

In PCA, there is an analytical solution to reduce dimensions. It is fast. 2

But sometimes, people want to formulate PCA in probabilistic terms. See previous slide. The good thing about PPCA is that missing values can still be accounted for and results don't vary as much.

The prior for the t_i is the standard normal.

This means that the low dimensional projection will be somewhere around 0 and variance 1.

(t_i follows the standard normal pdf)

bilistic terms. The good thing and values can depend a results don't here. The points $p(t_i) = \mathcal{N}(0, I)$ to be the properties of the p

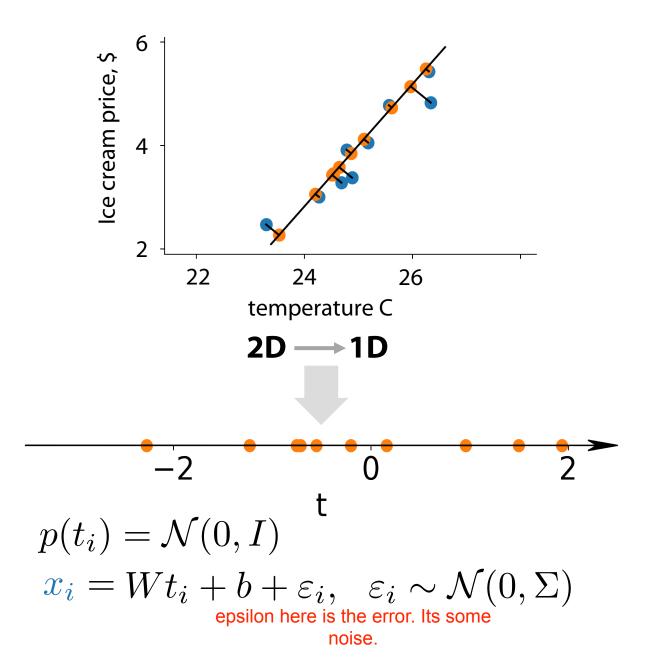
26

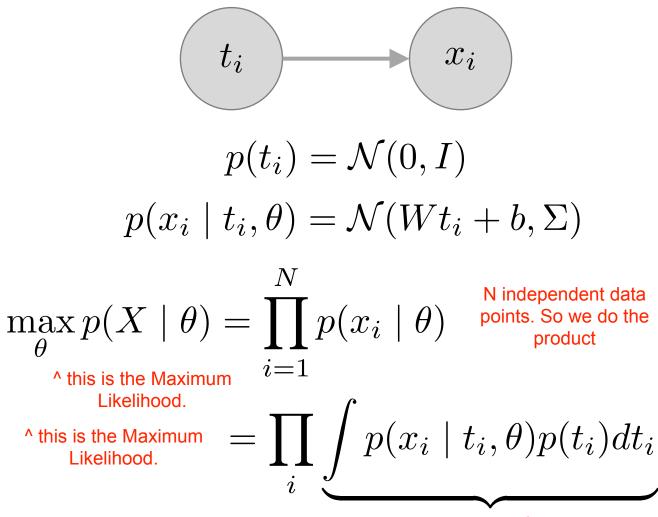
WE MAP our latent variable t i into

some two dimensional datapoint x i.

This translates to the full data.

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conjugacy, $\mathcal{N}(\mu_i, \Sigma_i)$ We marginalize out t_i here^. In general, this integral is intractable. So we use the EM algorithm to get an approximate.