What is a probability model?

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Some notes or model comparison and selection, parameter estimation, Bayes factors, and so on.

Dear Reader & Peer, this manuscript is being peer-reviewed by you. Thank you.

> Where do probability models come from?
> To judge by the resounding silence over this
> question on the part of most statisticians,
> it seems highly embarrassing.
>
> (Dawid 1982 p. 220)

1   Motivation

The Bayesian literature on probabilistic modelling, including model comparison, hypothesis testing, and parameter estimation, often consider models for our degrees of belief having this parametric form:

$$p(D \,|\, H, I) = \int d\theta \; p(D \,|\, \theta, H, I) \, p(\theta \,|\, H, I), \qquad (1)$$

where H represents the model, I other background information and assumptions, D data, and θ a parameter with values in some manifold. This is a particular case of an exchangeable model (Bernardo et al. 2000 ch. 4). *Bayes factors* and *evidence* (Good 1985; MacKay 1992; Kass 1993; Kass et al. 1995), and model averaging (Draper 2005; Chatfield 1995; Draper 1995; Hoeting et al. 1999) are often discussed in this literature. If we have several mutually exclusive and exhaustive models $H_1, H_2, \ldots$ our degree of belief about each given some data is

$$p(H_j \,|\, D, I) = \frac{p(D \,|\, H_j, I) \, p(H_j \,|\, I)}{\sum_k p(D \,|\, H_k, I) \, p(H_k \,|\, I)}; \qquad (2)$$

this degree of belief involves the *evidence*, which is just $p(D \,|\, H_j, I)$, eq. (1). The Bayes factor between two models is just the ratio of their evidences.

The literature reports several open issues in 'model comparison' and the evaluation of evidences and Bayes factors. For example, to calculate a model's evidence we must solve the integral in eq. (1), and this requires specifying a distribution for the parameters $p(\theta \mid H, I)$. When this distribution is improper the evidence vanishes, so the ratio of two evidences becomes undetermined. Several ways of fixing this issue have been proposed in the literature (Kass et al. 1995; O'Hagan 1995; Berger et al. 1996; De Santis et al. 1997; Berger et al. 1998). It has also been pointed out if we choose just one out of several possible models, basing our choice on their evidence, and then we this model to make inferences, we can end up misrepresenting our degree of belief (Draper 2005; Chatfield 1995; Draper 1995; Hoeting et al. 1999). Our degree of belief is in fact the weighted average of those given by each model. Yet, in some concrete applications – think about the use of a neural net – we have to choose only one model. ✠ some reference justly stated that this is the domain of decision theory; can't find it

These issues suggest that we should re-examine what we're actually doing in 'model comparison' or 'model selection'. This is the purpose of the present note.

From a Bayesian point of view model comparison doesn't really exist or is superfluous, because if we are unsure about several models H_j, then our degree of belief, by the theorem of total probability, is just

$$p(D \mid I) = \sum_j p(D \mid H_j, I)\, p(H_j \mid I). \tag{3}$$

As we gather new data D' our degree of belief is updated to

$$p(D \mid D', I) = \frac{p(D, D' \mid I)}{p(D' \mid I)} \equiv \frac{\sum_j p(D, D' \mid H_j, I)\, p(H_j \mid I)}{\sum_k p(D' \mid H_k, I)\, p(H_k \mid I)}, \tag{4}$$

which can be suggestively *rearranged* as follows, multiplying and dividing by $p(D' \mid H_j, I)$:

$$p(D \mid D', I) = \sum_j \underbrace{\frac{p(D, D' \mid H_j, I)}{p(D' \mid H_j, I)}}_{=:p(D \mid D', H_j, I)} \underbrace{\frac{p(D' \mid H_j, I)\, p(H_j \mid I)}{\sum_k p(D' \mid H_k, I)\, p(H_k \mid I)}}_{=:p(H_j \mid D', I)}. \tag{5}$$

This is a weighted sum of our degrees of belief conditional on each model, the weights being our updated degrees of belief about the

model themselves. The models which have very low updated degrees of belief effectively drop out of the sum, so the probability calculus is automatically doing 'model selection' for us. From this point of view, the practice of model selection can be viewed as an approximation of the formula above. The problem with improper distributions for the parameters still remains, though: models involving such distributions assign a vanishing degree of belief to the data and therefore disappear from eqs (4) and (5). We'll discuss this issue later.

If we compare the model-averaging formula (3) with the parametric formula (1) for a specific model, we see that the two have the same structure. The second is just the continuum limit of the first. In fact many of the issues about model comparison just discussed appear also when we consider just one model of parametric form (1). Given new gathered data D', our degree of belief is updated to

$$p(D|D', H, I) = \frac{p(D, D'|H, I)}{p(D'|H, I)} \equiv \frac{\int \mathrm{d}\theta \; p(D, D'|\theta, H, I) \, p(\theta|H, I)}{\int \mathrm{d}\theta' \; p(D'|\theta', H, I) \, p(\theta'|H, I)}, \tag{6}$$

which can be suggestively *rearranged* as follows, multiplying and dividing by $p(D'|\theta, H, I)$:

$$p(D|D', H, I) =$$
$$\int \mathrm{d}\theta \; \underbrace{\frac{p(D, D'|\theta, H, I)}{p(D'|\theta, H, I)}}_{=:p(D|D',\theta,H,I)} \; \underbrace{\frac{p(D'|\theta, H, I) \, p(\theta|H, I)}{\int \mathrm{d}\theta' \, p(D'|\theta', H, I) \, p(\theta'|H, I)}}_{=:p(\theta|D',H,I)}. \tag{7}$$

This is again equivalent to the model-average case (5) but for one important difference: in the first fraction in the product above, the dependence on data D' disappear:

$$p(D|D', \theta, H, I) = p(D|\theta, H, I). \tag{8}$$

This is a typical property of parametric models: the presence of the parameters in the conditional makes all other new data in the conditional *irrelevant*. Use of an improper distribution $p(\theta|H, I)$ for the parameter usually doesn't lead to problems in this case if correctly used – that is, taking the limit only at the very end of all calculations (Jaynes 2003 ch. 15). Analogously to the model-average case, parameters with low updated density $p(\theta|D', H, I)$ give a vanishing contribution to the

continuum average (7), so the probability calculus is automatically doing a 'parameter selection' for us. From this point of view, the practice of selecting just one parameter can be viewed as an approximation of the formula above (though such approximation can be bad if the density for θ has many modes).

In some situations, however, we can't or don't want to consider all models in the model-average case (5), even when no model dominates our updated degree of belief over the others. We want just one model. Similarly, in some situations we can't or don't want to consider all parameters in the single-model case (7), even when the updated density for the parameter has no dominating peaks. We want just one parameter. These analogous situations involve *decision theory*, and therefore require us to assign gain/loss functions. But what is our decision actually about? is it really about a model or a parameter? or rather about possible data outcomes? Let's examine this question after making an important distinction between two kinds of model.

2   A tentative definition

The most general useful definition of a probability model seems to be this: given a set Y of possible data, and possibly a set X of conditional data, a *probability model* is a conjunction M of assumptions or hypotheses that allows us to assign a definite, numerical plausibility

$$p(y_1, y_2, \ldots \mid x_1, x_2, \ldots, M) \tag{9}$$

for every meaningful combination of $y_i \in Y$ and $x_i \in X$. For the moment I consider finite sets.

This definition applies in particular to exchangeable models, where X is just a set of labels for the observations, which can take values in Y; to partially exchangeable models, where X is a set of labels for the exchangeable categories; and to models used in machine learning and neural nets. Possibly it also applies to time series. This definition also include functions or maps $f \colon X \to Y$ as special cases, when the plausibility is unity for a particular y only, dependent on x: $p(y \mid x, M) = \delta[y, f(x)]$.

Let's use this definition to explore the issues in model comparison and parameter estimation previously discussed.

An important distinction can be made between two kinds of model: *learning* models and *non-learning* models, which can also be called *extremal* for reasons explained later.

A learning model M is one that yields different plausibilities about some data $(y_1, y_2, \dots) =: y$ conditional on $(x_1, x_2, \dots) =: x$ if we condition on knowledge about other data $(y'_1, y'_2, \dots) =: y'$, $(x'_1, x'_2, \dots) =: x'$:

$$p(y \mid x, y', x', M) \neq p(y \mid x, M). \tag{10}$$

A non-learning model is one for which these plausibilities are not affected:

$$p(y \mid x, y', x', M) = p(y \mid x, M). \tag{11}$$

This means that

$$p(y \mid x, M) = \prod_i p(y_i \mid x_i, M). \tag{12}$$

If we look at eq. (8) we see that the conjunction (θ, H) is defining a non-learning model. So a model like H, eq. (1), which is clearly a learning model, is given as a continuous weighted mixture of non-learning models. This is a property of most parametric models typically considered in the literature, and is a corollary of de Finetti's theorem for partially exchangeable models (de Finetti 1938; Bernardo et al. 2000 § 4.6).

In concrete applications we usually seek *non-learning* models. For example, a trained and ready-to-use neural net is a non-learning model.

Old text

When we ask about the probability of a 'model' given the data, we're asking if a given region of limit relative frequencies is more probable than another. This may not be what we want to ask, because one region as a whole can have higher probability than another region, and yet a particular frequency in the second region may be more probable than any single frequency in the first region.

I a way, our real goal is to guess the limit frequency, not guess a region in frequency space, which may be quite arbitrary and whose shape has nothing to do with our predictions.

A solution to this is to reparameterize every 'model' with a coordinate that has the same meaning across them, and then work with the union of these models, forgetting about their individualities.

But does it make sense to ask whether the limit frequency distribution belongs to parametric family rather than another? It is like asking whether the limit frequency belongs to a submanifold (for example a curve) rather than another in the simplex, in the case with finite number of outcomes. The limit frequency belongs to many submanifolds at once.

If we really want to ask that question we should first choose a probability distribution in the whole limit-frequency space, a metric, and then determine the induced probability on the submanifold.

This kind of question may be useful if we are asking about several *experiments*, not just one. In this case it may make sense to ask whether their different limit frequencies belong to some common submanifold.

'Model' often seems to be mistakenly identified with the specification of likelihoods only, as if the specification of the parameter prior were not part of the model. Compare Kass et al. (1995): 'Bayes factors require priors on the parameters appearing in the models that represent the competing hypotheses' (p. 773) 'the prior distributions $\pi(\theta_k | H_k)$ on the parameters of each model must be specified' (p. 781), 'Sensitivity analysis concerns distributional forms for models $pr(\mathbf{D} | \theta_k, H_k)$ as well as priors', 'In choosing priors, just as in choosing models for data distributions, simplifications are often made' (p. 781). But see also 'the prediction rule is derived from the model H_k *(i.e., likelihood and prior)*' (p. 777, emphasis added).

The probability of hypotheses like those – concerning whole regions of limit-frequency space – cannot be computed.

The problem of 'model dimensionality' is also misplaced because we identify models with likelihoods only. In reality the dimensionality of a model is determined by the parameter prior. In fact, the very choice of likelihood can be interpreted as the choice of a particular prior from a 'non-parametric' point of view. (Compare Kass (1995), end of § 6.1.)

Also the idea of model *selection* can be dangerous, because we may be discarding the model than contains the frequency with highest likelihood.

The evidence is just an average of cross-validations (or splitting, see Kass § 6.5). Naive cross-validation is testing the wrong hypothesis.

3   Prediction and forecast

Some notation: We assume to have a possibly infinite set of observations, each of which can yield one of N outcomes, labelled by integers i. The proposition $O_i^{(a)}$ denotes that outcome i is observed at the ath observation. Such propositions also contain information about the time or place where the outcome was observed, so that from a proposition like $O_{i_2}^{(2)} \wedge O_{i_4}^{(4)}$ we can for example infer the time interval $t^{(4)} - t^{(2)}$ between observations number 4 and 2.

A statistical model is a set of assumptions M that jointly allow us to consistently assign numerical values to the probabilities

$$\mathrm{P}(O_{i_{n+1}}^{(a_{n+1})}\big|\ O_{i_1}^{(a_1)} \wedge \cdots \wedge O_{i_n}^{(a_n)} \wedge M), \tag{13}$$

for any legitimate n and any sets of observations $\{a_1, \ldots, a_{n+1}\}$ and outcomes $\{i_1, \ldots, i_{n+1}\}$; 'consistently' means that these assignments are properly related by operations like marginalization.

This definition is very general; in fact it amounts to say that a model is an assignment of the probabilities for all possible conjunctions of n outcomes, for all legitimate n.

it includes exchangeable models of various kinds, models for time series and forecasts.

Now I'd like to make a distinction between two main classes of statistical models: those that 'learn' and those what 'don't learn'.

This distinction is clear within the subclass of infinitely exchangeable models: for any such model the probability above has the form

$$\int \mathrm{p}(i_{n+1}|\,\theta, M)\,\mathrm{p}(\theta|\,i_1, \ldots, i_n, M)\,\mathrm{d}\theta, \tag{14a}$$

$$\mathrm{p}(\theta|\,i_1, \ldots, i_n, M) \propto \left[\prod_{k=1}^{n} \mathrm{p}(i_k|\,\theta, M)\right]\mathrm{p}(\theta|\,M), \tag{14b}$$

where the specific form of $\mathrm{p}(i|\,\theta, M)$ and $\mathrm{p}(\theta|\,M)$ are determined by the model. Within this subclass, models that don't learn are characterized by $\mathrm{p}(\theta|\,M) = \delta(\theta - \theta^*)$, so that the probability for an outcome does not depend on knowledge of other outcomes:

$$\mathrm{P}(O_{i_{n+1}}^{(a_{n+1})}|\; O_{i_1}^{(a_1)} \wedge \cdots \wedge O_{i_n}^{(a_n)} \wedge M) = \mathrm{P}(O_{i_{n+1}}^{(a_{n+1})}|\; M) \equiv \mathrm{p}(i_{n+1}|\,\theta^*, M). \tag{15}$$

Such a model doesn't 'learn' because it makes all knowledge about other observations irrelevant for the prediction of each observation.

Among all statistical models for a particular set of

**models (e.g. exchangeability for which accumulation of data leads to stable probabilities, and models (e.g. Markov) for which this doesn't happen.

Thanks

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('de *X*' is listed under D, 'van *X*' under V, and so on, regardless of national conventions.)

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