PRML studies(Ch3.4, 3.5)

- Main consideration of Model comparison
 - Compare between different model families (ex. SVM vs CNN)
 - Compare between same model families with different hyperparameter values/options (ex. Branch number in Decision Tree based models, model architecture in Neural Network models, scale of weight norm penalty in parametric models)

- Primary Model selection methods in frequentist approach:
 - 1. Criteria based: AIC, BIC, Mallows's Cp... compare models by calculating [(Train data Negative Log Likelihood) (number of parameters or model complexity penalty)] form criteria for each model. But this may yield results selecting overly simple models.
 - Holdout Method/Cross Validation: Check the model's generalizability by calculating likelihood or performance metric of separated(independent) data set different from the data which model has trained.

- In Bayesian approach, we use probabilities to represent uncertainties in the choice of models.
- Suppose we wish to compare a set of L models $\{M_i\}$ where i=1,2,...,L. Here, model refers to a probability distributions over observed data(training data) D.
- We think that data is generated from one of these models, but we are uncertain which one. And our uncertainty is expressed through a prior probability $p(M_i)$.
- First, we evaluate posterior distribution $p(M_i|D)$.
- Dropping constant term with respect to M_i ,

$$p(M_i|D) \propto p(M_i)p(D|M_i)$$

^{*}Note that $p(D|M_i)$ is called "model evidence" or sometimes "marginal likelihood"

^{*}And the ratio of model evidences $p(D|M_i)/p(D|M_i)$ is known as "Bayes factor"

• Once we know the posterior distribution $p(M_i|D)$, the predictive distribution in supervised tasks for new unobserved target variables can be represented as:

$$p(t|\underline{x},D) = \sum_{i=1}^{L} p(t|\underline{x},M_i,D)p(M_i|D)$$

- It is just a weighted average of L distributions weighted by posterior probability.
- However, its distributional form can be multi-modal.
- Simple approximation to this model averaging is to use single most probable model alone to make predictions.
 - => model selection
- Or, we can combine multiple models to make predictions.
 - => model averaging, simplest forms of model ensemble methods

• For a model governed by a set of parameters w, model evidence can be represented as:

$$p(D|M_i) = \int p(D|\underline{w}, M_i) p(\underline{w}|M_i) dw$$

Evaluating posterior distribution over parameters by Bayes formula:

$$p(\underline{w}|D, M_i) = \frac{p(D|\underline{w}, M_i)p(\underline{w}|M_i)}{p(D|M_i)}$$

• Thus, from sampling perspective, the marginal likelihood can be viewed as the probability of generating the data set *D* from a model whose parameters are sampled from prior at random.

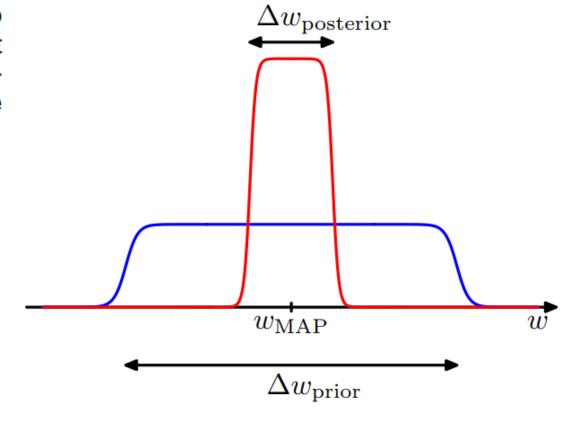
- Also, we can analyze about model evidence to gain insights more by making simple approximation to the integral over parameters.
- If we assume posterior distribution is sharply peaked around the most probable value w_{MAP} , with width $\Delta w_{posterior}$, then we can approximate the integral by the value of the integrand at its maximum times the width of the peak.
- If we further assume that prior is flat with width Δw_{prior} so that $p(w) = 1/\Delta w_{prior}$,

$$p(D) = \int p(D|w)p(w)dw \cong p(D|w_{MAP}) \frac{\Delta w_{posterior}}{\Delta w_{prior}}$$

$$\ln p(D) \cong \ln p(D|w_{MAP}) + \ln \frac{\Delta w_{posterior}}{\Delta w_{prior}}$$

 First term(log likelihood) represents the fit to data given by the most probable parameter value and second term penalizes the model according to its complexity.

Figure 3.12 We can obtain a rough approximation to the model evidence if we assume that the posterior distribution over parameters is sharply peaked around its mode $w_{\rm MAP}$.



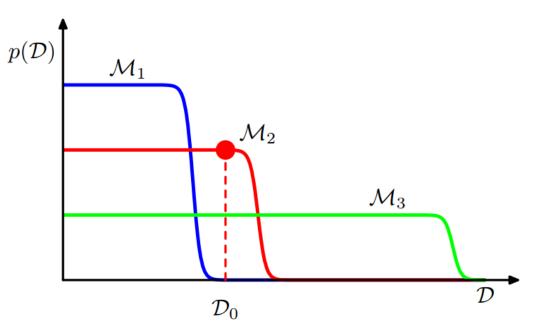
• For models which has *M* parameters, by using same procedures, we can get:

$$\ln p(D) \cong \ln p(D|\underline{w}_{MAP}) + M \ln \frac{\Delta \underline{w}_{posterior}}{\Delta \underline{w}_{prior}}$$

• If we increase the complexity of model, the first term will typically decrease, because a more complex model is better able to fit the data, where second term will increase due to the dependence on M.

• Therefore, the optimal model complexity, as determined by the maximum evidence will be given by a trade-off between these two competing terms.

Figure 3.13 Schematic illustration of the distribution of data sets for $p(\mathcal{D})$ three models of different complexity, in which \mathcal{M}_1 is the simplest and \mathcal{M}_3 is the most complex. Note that the distributions are normalized. In this example, for the particular observed data set \mathcal{D}_0 , the model \mathcal{M}_2 with intermediate complexity has the largest evidence.



- Simple models tends to have little variability and complex model have large variability in data generation.
- In the figures, for a specific data set D_0 , simple models might not generate D_0 .
- By contrast, complex models might generate D_0 , but probabilities might be relatively low.

- Implicit Bayesian model comparison framework is the assumption that true data generating distribution is contained within the set of models under consideration.
- Provided that, we can show Bayesian model comparison framework will on average favor the correct model.
- Consider two models M_1 and M_2 in which M_1 correspond to truth.
- For a given finite data set, it is possible for the Bayes factor to be larger for the incorrect model.
- However, if we average the Bayes factor over the true distribution of data sets, we obtain the expected Bayes factor in the form:

$$\int p(D|M_1) \ln \frac{p(D|M_1)}{p(D|M_2)} dD$$

$$\int p(D|M_1) \ln \frac{p(D|M_1)}{p(D|M_2)} dD$$

- This quantity is an example of "Kullback-Leibler Divergence"
- And it satisfies the mathematical property of always being positive, unless the two distributions are equal in which case it is zero.
- Thus, the Bayes factor will always favor the correct model on average.

- In conclusion, we have seen Bayesian framework avoids over-fitting and allows models to be compared with the training data alone.
- However, like any other approach to pattern recognition, Bayesian approach needs to make assumption about the form of the model.
- If the assumptions are invalid, then the result can be misleading.

• Therefore, In practical application, it will be wise to keep aside an independent test set of data on which to evaluate the overall performance of the final system.

- In fully Bayesian treatment, we would introduce prior distributions over the hyperparameters and make predictions by marginalizing with respect to these hyperparameters as well as parameters.
- However, although we can integrate over parameters or hyperparameters individually, the complete marginalization over all these variables is analytically intractable.
- By setting hyperparameters to specific values determined by maximizing the "marginal likelihood function" obtained by first integrating over the parameters, we can make approximation of the predictive distribution.

^{*}This framework is known in the statistics literature as "Empirical Bayes" or "Generalized maximum likelihood".

^{*}In Machine Learning literature, it is also called "Evidence Approximation"

• If we introduce hyperpriors $p(\underline{a})$ over hyperparameters \underline{a} , the predictive distribution $p(t^*|x^*,D)$ is obtained by marginalizing over parameters \underline{w} and hyperparameters \underline{a} . (* marks new data)

$$p(t^*|\underline{x^*},D) = \iint p(t^*|\underline{x^*},\underline{w})p(\underline{w}|D,\underline{a})p(\underline{a}|D)d\underline{w}d\underline{a}$$

Note that there might be some dependencies in some hyperparameters. For example, in linear basis models, noise precision β can be introduced and $p(t^|\underline{x}^*,\underline{w})$ can be replaced with $p(t^*|\underline{x}^*,\underline{w},\beta)$.

• From the Bayes theorem, since model evidence is posterior distribution for hyperparameter a and it is given by:

$$p(\underline{a}|D) \propto p(D|\underline{a})p(\underline{a})$$

• If the posterior distribution $p(\underline{a}|D)$ is sharply peaked around values \hat{a} , then the predictive distribution is obtained by marginalizing over \underline{w} in which \underline{a} are fixed to the values \hat{a} , so that

$$p(t^*|\underline{x^*},D) \cong p(t^*|\underline{x^*},D,\underline{\hat{a}}) = \int p(t^*|\underline{w},D) p(\underline{w}|D,\underline{\hat{a}}) d\underline{w}$$

- If the prior is relatively flat, then in the evidence framework the values of \hat{a} are obtained by maximizing the marginal likelihood function $p(D|\underline{a})$.
- This will allow as to determine values for hyperparameters from training data alone, without recourse to Holdout methods or Cross-Validation.
- Note that there are two approaches that we can take to the maximization of the log evidence, which is:
 - 1. Evaluate evidence function analytically and then set its derivative equal to zero to obtain re-estimation equations for hyperparameters.
 - 2. Use a technique called Expectation Maximization algorithm.

^{*}These two approaches converge to the same solution.

- However, there might exists a practical alternative to the evidence framework called Laplace approximation.
- Laplace approximation is based on the local Gaussian approximation centered on the mode of the posterior distribution.
- However, the integrand as a function of parameters typically has a strong skewed mode so that the Laplace approximation fails to capture the bulk of probability mass, leading to poorer results than those obtained by maximizing the evidence.

Reference

• Figures and contents are from Pattern Recognition and Machine Learning, Bishop