Ensemble methods are used to form combinations of predictors that achieve a bias-variance trade-off better than the one achieved by the algorithm generating the predictors in the combination. Many stochastic gradient descent algorithms, like Pegasos, can be viewed as ensemble methods because they output a combination of all the predictors generated during the sequential run over the training set. We now look at ensemble methods that are not based on online algorithms. As usual, our focus is on binary classification.

Fix a training set \((x_1, y_1), \ldots, (x_m, y_m)\) for a binary classification problem with zero-one loss, and assume an ensemble of classifiers \(h_1, \ldots, h_T\) is available (later, we describe how to obtain the ensemble). Consider the majority classifier \(f\) defined by

\[
f(x) = \text{sgn} \left( \sum_{i=1}^{T} h_i(x) \right)
\]

Clearly, \(f\) is wrong on \(x\) if and only if at least half of the classifiers \(h_1, \ldots, h_T\) are wrong on \(x\) (assume \(T\) is odd to avoid ties). We now study the conditions under which the majority classifier achieves a small training error. Assume

\[
P(h_1(x_Z) \neq y_Z \land \cdots \land h_T(x_Z) \neq y_Z) = \prod_{i=1}^{T} P(h_i(x_Z) \neq y_Z)
\] (1)

where \(Z\) is a random variable uniformly distributed on the set \(\{1, \ldots, m\}\) of indices of training examples. In other words, each classifier is wrong independently of the others with respect to the uniform distribution over the training set. The indicator function of these events define the training error \(\hat{\ell}_S(h_i)\) of each classifier \(h_i\). Indeed,

\[
\hat{\ell}_S(h_i) = \frac{1}{m} \sum_{t=1}^{m} \mathbb{I}\{h_i(x_t) \neq y_t\} = P(h_i(x_Z) \neq y_Z)
\]

Since we use the zero-one loss, we may also assume that \(\hat{\ell}_S(h_i) \leq \frac{1}{2}\) for each \(i = 1, \ldots, T\). Indeed, if \(\hat{\ell}_S(h_i) > \frac{1}{2}\) for some \(h_i\), then the binary classifier \(-h_i\) satisfies \(\ell_S(h_i) < \frac{1}{2}\).

Now consider the majority classifier \(f\) defined by

\[
f(x) = \text{sgn} \left( \sum_{i=1}^{T} h_i(x) \right)
\]

The majority classifier is a simple example of an ensemble method, combining the predictions of
an ensemble of classifiers in order to boost the accuracy. We now bound the training error of \( f \),

\[
\hat{\ell}_S(f) = P(f(x) \neq y_Z) = P \left( \sum_{i=1}^{T} I\{h_i(x) \neq y\} > \frac{T}{2} \right) = P \left( \frac{1}{T} \sum_{i=1}^{T} I\{h_i(x) \neq y\} > \ell_{ave} + \left( \frac{1}{2} - \ell_{ave} \right) \right)
\]

where we defined

\[
\ell_{ave} = \frac{1}{T} \sum_{i=1}^{T} \hat{\ell}_S(h_i)
\]

Now introduce the Bernoulli random variables \( B_1, \ldots, B_m \) defined by \( B_i = I\{h_i(x) \neq y\} \). Note that these random variables are independent, due to our assumption (1). Also, \( E[B_i] = \hat{\ell}_S(h_i) \) and

\[
\frac{1}{T} \sum_{i=1}^{T} E[B_i] = \ell_{ave} < \frac{1}{2}
\]

A slight generalization of Chernoff-Hoeffding bounds to independent random variables with unequal expectations gives

\[
P \left( \frac{1}{T} \sum_{i=1}^{T} B_i > \ell_{ave} + \varepsilon \right) \leq e^{-2\varepsilon^2 T}
\]

By setting \( \gamma_i = \frac{1}{2} - \hat{\ell}_S(h_i) \) we get

\[
\hat{\ell}_S(f) \leq \exp \left( -2T \left( \frac{1}{2} - \ell_{ave} \right)^2 \right) = \exp \left( -2T \left( \frac{1}{T} \sum_{i=1}^{T} \gamma_i \right)^2 \right) \leq e^{-2T\gamma^2}
\]

where for the last inequality we assumed \( \gamma_i \geq \gamma > 0 \) for all \( i = 1, \ldots, T \).

This result tells us that if we manage to obtain classifiers with independent training errors in the sense of (1), then the training error of the majority vote classifier decreases exponentially with respect to \( \gamma \), which measures how better than random guessing is each classifier \( h_i \) on the training set.

**Bagging.** How can we obtain classifiers with independent training errors? A popular heuristic, known as Bagging, applies to any learning algorithm \( A \) for binary classification and to any training set \( S \). Let \( m \) be the size of \( S \). Bagging builds \( h_1, \ldots, h_T \) by drawing \( m \) examples uniformly at random with replacement from \( S \). This process is repeated \( T \) times so to obtain the resampled training sets \( S_1, \ldots, S_T \). Then \( A \) is run on each \( S_i \) setting \( h_i = A(S_i) \). The idea is that the resampling procedure helps enforce condition (1).

One may wonder how different from \( S \) any \( S_i \) can be. To find that out, we take a little detour and compute the fraction of unique data points in \( S_i \). As you see in a moment, more than one third of the points of \( S \) are missing from \( S_i \) in expectation! Let \( N \) be the number of unique points drawn,
and let $X_t$ be the indicator function of the event that $(x_t, y_t)$ is drawn. Then the probability that $(x_t, y_t)$ is not drawn is

$$P(X_t = 0) = \left(1 - \frac{1}{m}\right)^m$$

So we have

$$E[N] = \sum_{t=1}^m E[X_t] = \sum_{t=1}^m P(X_t = 1) = \sum_{t=1}^m \left(1 - \left(1 - \frac{1}{m}\right)^m\right) = m - m \left(1 - \frac{1}{m}\right)^m$$

Therefore, the fraction of unique points in $S_i$ is

$$1 - \left(1 - \frac{1}{m}\right)^m \approx 1 - \frac{1}{e} = 0.632\ldots$$

where the approximation becomes exact for $m \to \infty$.

We saw that independence of errors helps reduce the bias by driving the training error to zero. On the other hand, subsampling of the training set helps reduce the variance. If we think of the $m$ training points arranged in a $m \times d$ matrix (called the data matrix), then what bagging does is subsampling the rows of this matrix. We now briefly describe another ensemble method that increases the protection against overfitting by also subsampling the columns of the data matrix.

**Random Forest.** This ensemble method works by taking a majority vote over an ensemble $h_1, \ldots, h_T$ of tree predictors. Similarly to bagging, each tree predictor $h_i$ is obtained by running a learning algorithm over a dataset $S_i$ obtained by subsampling the rows of the full data matrix. However, the algorithm for learning tree predictors does not have direct access to $S_i$. Indeed, when considering a leaf $\ell$ for splitting, instead of being given $S_{i,\ell}$ (the set of training examples in $S_i$ that are routed to $\ell$), the algorithm has access to a version of $S_{i,\ell}$ containing a random subset of the original features (typically, $\sqrt{d}$ features are sampled from the original $d$ features). This additional sampling provides a better control on the variance at the expense of the bias. Because of its good performance on many learning tasks, Random Forest is often used as a baseline when testing a new learning algorithm.

**Boosting.** We now introduce boosting, a principled ensemble method that achieves the exponential bound (2) on the training error without requiring the demanding condition (1). Boosting is an incremental method to build classifiers of the form $\text{sgn}(f)$, where

$$f = \sum_{i=1}^T w_i h_i$$

and $w = (w_1, \ldots, w_T)$ is a vector of real coefficients. We assume $h_1, \ldots, h_T$ belong to some family $\mathcal{H}$ of base classifiers.

In practice, also in order to save computational costs, base classifiers are very simple. A typical choice for $\mathcal{H}$ is that of decision stumps. These are all classifiers of the form $h_{i,\tau} : \mathbb{R}^d \to \{-1, 1\}$ defined by $h_{i,\tau}(x) = \pm \text{sgn}(x_i - \tau)$, where $i = 1, \ldots, d$ and $\tau \in \mathbb{R}$.

The specific boosting algorithm we introduce is known as AdaBoost (adaptive boosting). Fix a training set $S$ with $m$ examples $(x_1, y_1), \ldots, (x_m, y_m)$, and a sequence $h_1, \ldots, h_T$ of base classifiers. We now show how to choose the coefficients $w$ so that the training error is bounded as in (2).
AdaBoost uses the convex upper bound $\mathbb{I}\{z \leq 0\} \leq e^{-z}$ on the zero-one loss function $\mathbb{I}\{f(x_t) y_t \leq 0\}$. This gives

$$\hat{\ell}_S(f) = \frac{1}{m} \sum_{t=1}^{m} \mathbb{I}\{f(x_t) y_t \leq 0\} \leq \frac{1}{m} \sum_{t=1}^{m} e^{-f(x_t) y_t} = \frac{1}{m} \sum_{t=1}^{m} e^{-\sum_{i=1}^{T} w_i h_i(x_t) y_t}$$

Note that other algorithms use different convex upper bounds on the zero-one loss. For example, SVM uses the hinge loss.

Introduce now the functions $L_1, \ldots, L_T$ defined by $L_i(t) = h_i(x_t) y_t$. Note that $L_i(t) \in \{-1, 1\}$ and $L_i(t) = 1$ if and only if $h_i(x_t) = y_t$. Recalling that $Z$ is a random variable uniformly distributed in $\{1, \ldots, m\}$, we can view each $L_i(Z)$ as a random variable and write

$$\hat{\ell}_S(f) \leq \frac{1}{m} \sum_{t=1}^{m} e^{-\sum_{i=1}^{T} w_i L_i(t)} = \mathbb{E} \left[ e^{-\sum_{i=1}^{T} w_i L_i} \right] = \mathbb{E} \left[ \prod_{i=1}^{T} e^{-w_i L_i} \right]$$

If condition (1) were true, we could write the expectation of the product as a product of expectations. In order to sidestep the condition, we change the probability space and write

$$\mathbb{E} \left[ \prod_{i=1}^{T} e^{-w_i L_i} \right] = \prod_{i=1}^{T} \mathbb{E}_i \left[ e^{-w_i L_i} \right]$$

where each expectation $\mathbb{E}_i \left[ e^{-w_i L_i} \right]$ is understood with respect to a probability $\mathbb{P}_i$ yet to be specified. Assuming (3) holds, which we verify later, we can proceed as follows

$$\hat{\ell}_S(f) \leq \prod_{i=1}^{T} \mathbb{E}_i \left[ e^{-w_i L_i} \right]$$

$$= \prod_{i=1}^{T} \left( e^{-w_i} \mathbb{P}_i(L_i = 1) + e^{w_i} \mathbb{P}_i(L_i = -1) \right)$$

$$= \prod_{i=1}^{T} \left( e^{-w_i} (1 - \varepsilon_i) + e^{w_i} \varepsilon_i \right) \tag{4}$$

where we set

$$\varepsilon_i \overset{\text{def}}{=} \mathbb{P}_i(L_i = -1) = \sum_{t=1}^{m} \mathbb{I}\{L_i(t) = -1\} \mathbb{P}_i(t)$$

Note that $\varepsilon_i$ is the error of $h_i$ with respect to the probability $\mathbb{P}_i$. Namely, $\varepsilon_i$ is the weighted training error of $h_i$ where the weights are determined by $\mathbb{P}_i$.

Before computing the $\mathbb{P}_i$, we show how to pick $w_1, \ldots, w_T$ in order to minimize (4). By computing the zeros of the derivative of $e^{-w} (1 - \varepsilon_i) + e^{w} \varepsilon_i$ with respect to $w$, we find a single zero at

$$w = \frac{1}{2} \ln \frac{1 - \varepsilon_i}{\varepsilon_i}.$$ 

Note that the above expression is only defined for $0 < \varepsilon_i < 1$. As we will see, $\mathbb{P}(t) > 0$ for all $t \in \{1, \ldots, m\}$. Hence $\varepsilon_i \in \{0, 1\}$ implies that either $h_i$ or $-h_i$ has zero training error on $S$. If this
happens, than we can throw away all \( h_j \) for \( j \neq i \) and avoid using boosting altogether. Therefore, without loss of generality we may assume \( 0 < \varepsilon_i < 1 \) for all \( i = 1, \ldots, m \).

Through the study of the derivative, we find that the unique zero corresponds to a minimum of the function \( e^{-w}(1 - \varepsilon_i) + e^{w}\varepsilon_i \). Hence we set

\[
\gamma_i \overset{\text{def}}{=} \frac{1}{2} \ln \frac{1 - \varepsilon_i}{\varepsilon_i}
\]

for \( i = 1, \ldots, T \).

Substituting in (4) and simplifying, we get

\[
\hat{\ell}_S(f) \leq \prod_{i=1}^T \sqrt{4\varepsilon_i(1 - \varepsilon_i)}.
\]

Note that \( w_i = 0 \) if and only if \( \varepsilon_i = \frac{1}{2} \), meaning that the weight (according to \( \mathbb{P}_i \)) of the training points where \( h_i \) errs is exactly \( \frac{1}{2} \). Because such a \( h_i \) does not affect the value of \( f \) (since \( \varepsilon_i = \frac{1}{2} \) implies \( w_i = 0 \)) without loss of generality we may also assume that \( \varepsilon_i \neq \frac{1}{2} \).

Set \( \gamma_i \overset{\text{def}}{=} \frac{1}{2} - \varepsilon_i \) and note that \( \varepsilon_i \neq \frac{1}{2} \) implies \( \gamma_i \neq 0 \). Using the inequality \( 1 + x \leq e^x \), which holds for all \( x \in \mathbb{R} \), we get

\[
\hat{\ell}_S(f) \leq \prod_{i=1}^T \sqrt{4\varepsilon_i(1 - \varepsilon_i)} = \prod_{i=1}^T \sqrt{1 - 4\gamma_i^2} \leq \prod_{i=1}^T e^{-2\gamma_i^2} = e^{-2\sum_{i=1}^T \gamma_i^2} = e^{-2T\gamma^2}
\]

where in the last step we assumed \( |\gamma_i| > \gamma > 0 \). This is the same bound as the one we proved in (2) under the condition (1). Note, however, that the definition of \( \gamma_i \overset{\text{def}}{=} \frac{1}{2} - \varepsilon_i \) changes because \( \varepsilon_i \) now is the weighted training error of \( h_i \).

Just like (2), this bound provides a pretty strong control on the bias. Using the observation that \( \hat{\ell}_S(f) = 0 \) if and only if \( \hat{\ell}_S(f) < 1/m \), we conclude that a number

\[
T > \frac{\ln m}{2\gamma^2}
\]

of boosting rounds is sufficient to bring the training error of \( f \) down to zero.

We now move on to derive the \( \mathbb{P}_1, \ldots, \mathbb{P}_T \) satisfying condition (3). Setting \( \mathbb{P}_1 = \mathbb{P}, \mathbb{E}_1 = \mathbb{E} \), and

\[
\mathbb{P}_{i+1}(t) = \frac{\mathbb{P}_i(t)e^{-w_i L_i(t)}}{\mathbb{E}_i[e^{-w_i L_i}]} \quad \text{for } t = 1, \ldots, m \text{ and } i = 1, \ldots, T - 1 \quad (5)
\]

where

\[
\mathbb{E}_i[e^{-w_i L_i}] = \sum_{s=1}^m e^{-w_i L_i(s)} \mathbb{P}_i(s)
\]

It is easy to check that \( \mathbb{P}_1, \ldots, \mathbb{P}_T \) are indeed probability distributions on \( \{1, \ldots, m\} \). In particular, \( \mathbb{P}_i(t) > 0 \) and \( \mathbb{P}_i(1) + \cdots + \mathbb{P}_i(m) = 1 \).

For this choice of \( \mathbb{P}_i \) we can prove (3) as follows. First, we solve (5) for \( e^{-w_i L_i(t)} \) obtaining

\[
e^{-w_i L_i(t)} = \mathbb{E}_i[e^{-w_i L_i}] \frac{\mathbb{P}_{i+1}(t)}{\mathbb{P}_i(t)}
\]
Then, we write

\[
E \left[ \prod_{i=1}^{T} e^{-w_i L_i} \right] = \frac{1}{m} \sum_{t=1}^{m} \left( \prod_{i=1}^{T} E_i \left[ e^{-w_i L_i} \right] \frac{P_{t+1}(t)}{P_t(t)} \right) 
\]

\[
= \frac{1}{m} \sum_{t=1}^{m} \left( \frac{P_{t+1}(t)}{P_t(t)} \right) \left( \prod_{i=1}^{T} E_i \left[ e^{-w_i L_i} \right] \right) 
\]

\[
= \left( \sum_{t=1}^{m} \frac{P_{t+1}(t)}{P_t(t)} P_1(t) \right) \left( \prod_{i=1}^{T} E_i \left[ e^{-w_i L_i} \right] \right) 
\]

\[
= \prod_{i=1}^{T} E_i \left[ e^{-w_i L_i} \right] 
\]

concluding the proof.

These probability distributions have a simple interpretation when one studies how $P_{t+1}$ depends on $P_t$. Fix $P_t$ and suppose $\varepsilon_t < \frac{1}{2}$. Then $w_i > 0$ and each $P_{t+1}(t)$ is obtained multiplying $P_t(t)$ for the quantity $e^{-w_i L_i(t)}$, which is bigger than 1 if and only if $h_i(x_t) \neq y_t$. In other words, the weight of each training example $(x_t, y_t)$ is increased when $P_t$ is updated to $P_{t+1}$ if and only if $h_i$ errs on $(x_t, y_t)$. Intuitively, the boosting process concentrates the weight on the training examples that are misclassified by the previous classifiers. A similar argument applies to the case when $\varepsilon_t > \frac{1}{2}$.

| Input: Training set $S$ of examples $(x, y) \in \mathbb{R}^d \times \{-1, 1\}$. Learning algorithm $A$. Maximum number $T$ of boosting rounds. Initialize $P_1(t) \leftarrow 1/m$ for $t = 1, \ldots, m$. For $i = 1, \ldots, T$
| 1. Feed $A$ with $S$ weighted by $P_t$ and get $h_i$ in response
| 2. Compute $\varepsilon_i$ for $h_i$
| 3. If $\varepsilon_i \in \{0, \frac{1}{2}, 1\}$ then BREAK
| 4. Let $w_i \leftarrow \frac{1}{2} \ln \frac{1-\varepsilon_i}{\varepsilon_i}$.
| 5. Compute $P_{t+1}$ using (5).
| If for loop exited on BREAK, then deal with the special case Else output $f = \text{sgn}(w_1 h_1 + \cdots + w_T h_T)$.

We are now ready to introduce the pseudo-code for AdaBoost. The only thing we left unspecified is how the sequence $h_1, \ldots, h_T$ of classifiers is generated. In principle, we want a sequence such that the corresponding sequence $\gamma_1, \ldots, \gamma_T$ is as far away from $\frac{1}{2}$ as possible (in either direction) so to ensure an exponential decrease of the training error. It is convenient to view the boosting process as sequence of rounds between the boosting algorithm and a generic learning algorithm $A$. 

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In each round $i$, the booster $B$ gives $P_i$ to $A$ and gets $h_i$ in response. The interaction goes on for $T$ rounds, unless $A$ return $h_i$ such that $\gamma_i = 0$. In that case the boosting process stops and the booster outputs $f = \text{sgn}(w_1h_1 + \cdots + w_{i-1}h_{i-1})$. 

\[ B \xrightarrow{h_i} P_i \xrightarrow{} A \]