N-Gram Model Formulas

Word sequences

$$w_1^n = w_1...w_n$$

• Chain rule of probability

$$P(w_1^n) = P(w_1)P(w_2 \mid w_1)P(w_3 \mid w_1^2)...P(w_n \mid w_1^{n-1}) = \prod_{k=1}^{n} P(w_k \mid w_1^{k-1})$$

• Bigram approximation

$$P(w_1^n) = \prod_{k=1}^n P(w_k \mid w_{k-1})$$

• N-gram approximation

$$P(w_1^n) = \prod_{k=1}^n P(w_k \mid w_{k-N+1}^{k-1})$$

Estimating Probabilities

• N-gram conditional probabilities can be estimated from raw text based on the *relative frequency* of word sequences.

Bigram:
$$P(w_n \mid w_{n-1}) = \frac{C(w_{n-1}w_n)}{C(w_{n-1})}$$

N-gram:
$$P(w_n \mid w_{n-N+1}^{n-1}) = \frac{C(w_{n-N+1}^{n-1}w_n)}{C(w_{n-N+1}^{n-1})}$$

• To have a consistent probabilistic model, append a unique start (<s>) and end (</s>) symbol to every sentence and treat these as additional words.

Perplexity

- Measure of how well a model "fits" the test data.
- Uses the probability that the model assigns to the test corpus.
- Normalizes for the number of words in the test corpus and takes the inverse.

$$PP(W) = \sqrt[N]{\frac{1}{P(w_1 w_2 ... w_N)}}$$

• Measures the weighted average branching factor in predicting the next word (lower is better).

Laplace (Add-One) Smoothing

• "Hallucinate" additional training data in which each possible N-gram occurs exactly once and adjust estimates accordingly.

Bigram:
$$P(w_n \mid w_{n-1}) = \frac{C(w_{n-1}w_n) + 1}{C(w_{n-1}) + V}$$

N-gram:
$$P(w_n \mid w_{n-N+1}^{n-1}) = \frac{C(w_{n-N+1}^{n-1}w_n) + 1}{C(w_{n-N+1}^{n-1}) + V}$$

where V is the total number of possible (N-1)-grams (i.e. the vocabulary size for a bigram model).

• Tends to reassign too much mass to unseen events, so can be adjusted to add $0<\delta<1$ (normalized by δV instead of V).

Interpolation

• Linearly combine estimates of N-gram models of increasing order.

Interpolated Trigram Model:

$$\hat{P}(w_n \mid w_{n-2}, w_{n-1}) = \lambda_1 P(w_n \mid w_{n-2}, w_{n-1}) + \lambda_2 P(w_n \mid w_{n-1}) + \lambda_3 P(w_n)$$

Where:
$$\sum_{i} \lambda_{i} = 1$$

• Learn proper values for λ_i by training to (approximately) maximize the likelihood of an independent *development* (a.k.a. *tuning*) corpus.

Formal Definition of an HMM

- A set of N + 2 states $S = \{s_0, s_1, s_2, \dots s_N, s_F\}$
 - Distinguished start state: s_0
 - Distinguished final state: $s_{\rm F}$
- A set of M possible observations $V = \{v_1, v_2 \dots v_M\}$
- A state transition probability distribution $A = \{a_{ij}\}$ $a_{ij} = P(q_{t+1} = s_j \mid q_t = s_i) \qquad 1 \le i, j \le N \text{ and } i = 0, j = F$ $\sum_{i=1}^{N} a_{ij} + a_{iF} = 1 \quad 0 \le i \le N$
- Observation probability distribution for each state j $B = \{b_j(k)\}$ $b_j(k) = P(v_k \text{ at } t \mid q_t = s_j)$ $1 \le j \le N$ $1 \le k \le M$
- Total parameter set $\lambda = \{A,B\}$

Forward Probabilities

• Let $\alpha_t(j)$ be the probability of being in state j after seeing the first t observations (by summing over all initial paths leading to j).

$$\alpha_t(j) = P(o_1, o_2, ...o_t, q_t = s_j \mid \lambda)$$

Computing the Forward Probabilities

Initialization

$$\alpha_1(j) = a_{0j}b_j(o_1) \quad 1 \le j \le N$$

Recursion

$$\alpha_{t}(j) = \left[\sum_{i=1}^{N} \alpha_{t-1}(i) a_{ij}\right] b_{j}(o_{t}) \quad 1 \le j \le N, \quad 1 < t \le T$$

• Termination

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$$P(O \mid \lambda) = \alpha_T(s_F) = \sum_{i=1}^{N} \alpha_T(i) a_{iF}$$

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Viterbi Scores

• Recursively compute the probability of the most likely subsequence of states that accounts for the first *t* observations and ends in state *s_i*.

$$v_{t}(j) = \max_{q_{0}, q_{1}, \dots, q_{t-1}} P(q_{0}, q_{1}, \dots, q_{t-1}, o_{1}, \dots, o_{t-1}, q_{t} = s_{j} \mid \lambda)$$

- Also record "backpointers" that subsequently allow backtracing the most probable state sequence.
 - $bt_t(j)$ stores the state at time t-1 that maximizes the probability that system was in state s_j at time t (given the observed sequence).

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Computing the Viterbi Scores

Initialization

$$v_1(j) = a_{0j}b_j(o_1) \quad 1 \le j \le N$$

Recursion

$$v_t(j) = \max_{i=1}^{N} v_{t-1}(i)a_{ij}b_j(o_t) \quad 1 \le j \le N, \quad 1 < t \le T$$

Termination

$$P^* = v_T(s_F) = \max_{i=1}^{N} v_T(i)a_{iF}$$

Analogous to Forward algorithm except take max instead of sum 10

Computing the Viterbi Backpointers

Initialization

$$bt_1(j) = s_0 \quad 1 \le j \le N$$

Recursion

$$bt_{t}(j) = \underset{i=1}{\operatorname{argmax}} v_{t-1}(i)a_{ij}b_{j}(o_{t}) \quad 1 \le j \le N, \ 1 \le t \le T$$

Termination

$$q_T^* = bt_T(s_F) = \underset{i=1}{\operatorname{argmax}} v_T(i)a_{iF}$$

Final state in the most probable state sequence. Follow backpointers to initial state to construct full sequence.

Supervised Parameter Estimation

• Estimate state transition probabilities based on tag bigram and unigram statistics in the labeled data.

$$a_{ij} = \frac{C(q_t = s_i, q_{t+1} = s_j)}{C(q_t = s_i)}$$

• Estimate the observation probabilities based on tag/ word co-occurrence statistics in the labeled data.

$$b_j(k) = \frac{C(q_i = s_j, o_i = v_k)}{C(q_i = s_i)}$$

• Use appropriate smoothing if training data is sparse.

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Context Free Grammars (CFG)

- N a set of non-terminal symbols (or variables)
- Σ a set of *terminal symbols* (disjoint from N)
- R a set of *productions* or *rules* of the form $A \rightarrow \beta$, where A is a non-terminal and β is a string of symbols from $(\Sigma \cup N)^*$
- S, a designated non-terminal called the *start symbol*

Estimating Production Probabilities

- Set of production rules can be taken directly from the set of rewrites in the treebank.
- Parameters can be directly estimated from frequency counts in the treebank.

$$P(\alpha \to \beta \mid \alpha) = \frac{\text{count}(\alpha \to \beta)}{\sum_{\gamma} \text{count}(\alpha \to \gamma)} = \frac{\text{count}(\alpha \to \beta)}{\text{count}(\alpha)}$$

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