

Computing Elo Ratings of Move Patterns in the Game of Go

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ABSTRACT

Move patterns are an essential method to incorporate domain knowledge into Go-playing programs. This paper presents a new Bayesian technique for supervised learning of such patterns from game records, based on a generalization of Elo ratings. Each sample move in the training data is considered as a victory of a team of pattern features. Elo ratings of individual pattern features are computed from these victories, and can be used in previously unseen positions to compute a probability distribution over legal moves. In this approach, several pattern features may be combined, without an exponential cost in the number of features. Despite a very small number of training games (652), this algorithm outperforms most previous pattern-learning algorithms, both in terms of mean log-evidence (-2.69), and prediction rate (34.9%). A 19×19 Monte-Carlo program improved with these patterns reached the level of the strongest classical programs.

1. INTRODUCTION

Many Go-playing programs use domain knowledge encoded into patterns. The kinds of patterns considered in this paper are heuristic move patterns. These are general rules, such as “it is bad to play in the corner of the board”, “it is good to prevent connection of two opponent strings”, “don’t fill-up your own eyes”, or “when in atari, extend”. Such knowledge may be used to prune a search tree, order moves, or improve random simulations in Monte-Carlo programs (Bouzy, 2005; Gelly *et al.*, 2006).

Move patterns may be built by hand, or generated automatically. A popular approach to automatically generate patterns is supervised learning (Araki *et al.*, 2007; Bouzy and Chaslot, 2005; Dahl, 1999; Enderton, 1991; de Groot, 2005; Marchand, 2007; Stern, Herbrich, and Graepel, 2006; Stoutamire, 1991; van der Werf *et al.*, 2003): frequent patterns are extracted and evaluated from game records of strong players. In this approach, expert knowledge is used to produce a relevant encoding of patterns and pattern features, and a machine-learning algorithm evaluates them. The advantage of automatic pattern learning over hand-made patterns is that thousands of patterns may be generated and evaluated with little effort, and little domain expertise.

This paper presents a new supervised pattern-learning algorithm, based on the Bradley-Terry model. The Bradley-Terry model is a pairwise comparison model based on the logit link function, and is the theoretical basis of the Elo rating system (Elo, 1978). The principle of Elo ratings, as applied to chess, is that each player gets a numerical strength estimation, computed from the observation of past game results. From the ratings of players, it is possible to estimate a probability distribution over the outcome of future games. The same principle can be applied to move patterns: each sample move in the training database can be considered as a victory of one pattern over the others, and can be used to compute pattern ratings. When faced with a new position, the Elo ratings of patterns can be used to compute a probability distribution over all legal moves.

1.1 Related Work

This algorithm based on the Bradley-Terry model is very similar in spirit to some recent related works, but provides significant differences and improvements.

The simplest approach to pattern learning consists in measuring the frequency of play of each pattern (Bouzy and Chaslot, 2005; de Groot, 2005). The number of times a pattern is played is divided by the number of times it is

present. This way, the strongest patterns get a higher rating because they do not stay long without being played. A major weakness of this approach is that, when a move is played, the strengths of competing patterns are not taken into consideration. In the Elo-rating analogy, this would mean estimating the strength of a player with its winning rate, regardless of the strength of opponents. By taking the strength of opponents into account, methods based on the Elo rating system can compute more accurate pattern strengths.

Stern *et al.* (2006) address the problem of taking the strength of opponents into account by using a model extremely similar to Elo ratings. With this model, they can compute high-quality probability distributions over legal moves. A weakness of their approach, however, is that they are restricted to using only a few move features, because the number of patterns to evaluate would grow exponentially with the number of features.

In order to solve the problem of combining move features, Araki *et al.* (2007) propose a method based on maximum-entropy classification. A major drawback of their approach is its very high computational cost, which forced them to learn on a restricted subset of moves, while still taking 8.75 days of computation to learn. Also, it is not clear whether their method would provide a good probability distribution over moves, because, like the frequency-based approach, it doesn't take the strength of opponent patterns into account.

A generalized Bradley-Terry model, when combined with the minorization-maximization algorithm to compute its maximum *a posteriori*, addresses all the shortcomings of previous approaches, by providing the algorithmic simplicity and efficiency of frequency-based pattern evaluation, with the power and theoretical soundness of methods based on Bayesian inference and maximum entropy.

1.2 Paper Outline

This paper is organized as follows: Section 2 explains the details of the theory of minorization-maximization and generalized Bradley-Terry models, Section 3 presents experimental results of pattern learning, and Section 4 describes how these patterns were applied to improve a Monte-Carlo program.

2. MINORIZATION-MAXIMIZATION AND GENERALIZED BRADLEY-TERRY MODELS

This section briefly explains, independently of the problem of learning patterns in the game of Go, the theory of minorization-maximization and generalized Bradley-Terry models. It is based on Hunter's paper (Hunter, 2004), where interested readers will find more generalizations of this model, with all the convergence proofs, references, and mathematical details.

2.1 Elo Ratings and the Bradley-Terry Model

The Bradley-Terry model allows to make predictions about the outcome of competitions between individuals. Its principle consists in evaluating the strength of each individual i by a positive numerical value γ_i . The stronger i , the higher γ_i . Predictions are made according to a formula that estimates the probability that i beats j :

$$P(i \text{ beats } j) = \frac{\gamma_i}{\gamma_i + \gamma_j} .$$

The Elo rating of individual i is defined by $r_i = 400 \log_{10}(\gamma_i)$, that is to say $\gamma_i = 10^{\frac{r_i}{400}}$.

2.2 Some Generalizations of the Bradley-Terry Model

The Bradley-Terry model may be generalized to handle competitions involving more than two individuals. For n players:

$$\forall i \in \{1, \dots, n\}, P(i \text{ wins}) = \frac{\gamma_i}{\gamma_1 + \gamma_2 + \dots + \gamma_n} .$$

Another interesting generalization consists in considering not only individuals, but teams. In this generalization, the γ of a team is estimated as the product of the γ 's of its members. For instance:

$$P(1-2-3 \text{ wins against } 4-2 \text{ and } 1-5-6-7) = \frac{\gamma_1 \gamma_2 \gamma_3}{\gamma_1 \gamma_2 \gamma_3 + \gamma_4 \gamma_2 + \gamma_1 \gamma_5 \gamma_6 \gamma_7} .$$

Note that the same γ may appear in more than one team. But it may not appear more than once in a team.

2.3 Relevance of Bradley-Terry Models

The choice of a Bradley-Terry model makes strong assumptions about what is being modeled, and may not be appropriate in every situation. First, a Bradley-Terry model cannot take into consideration situations where individual 1 beats individual 2 consistently, individual 2 beats individual 3 consistently, and individual 3 beats individual 1 consistently. The strengths are on a one-dimensional scale, which does not allow such cycles. Also, the generalization to teams assumes that the strength of a team is the sum (in terms of Elo ratings) of the strengths of its members. This is also a very strong assumption that may not be correct all the time.

2.4 Bayesian Inference

Bradley-Terry models, as described in the previous sections, provide a probability distribution over the outcomes of future competitions, given the strength of individuals that participate. Most of the time the exact value of parameters γ_i are unknown, and have to be estimated from the outcome of past competitions. This estimation can be done with Bayesian inference.

With $\vec{\gamma}$, the vector of parameters, and \vec{R} , past results, Bayes formula is:

$$P(\vec{\gamma}|\vec{R}) = \frac{P(\vec{R}|\vec{\gamma})P(\vec{\gamma})}{P(\vec{R})} .$$

It gives a posterior distribution over $\vec{\gamma}$, from $P(\vec{R}|\vec{\gamma})$, that is to say the Bradley-Terry model described in the previous sections, $P(\vec{\gamma})$, a prior distribution over parameters, and $P(\vec{R})$, a normalizing constant. Parameters $\vec{\gamma}$ may be estimated by finding $\vec{\gamma}^*$ that maximizes $P(\vec{\gamma}|\vec{R})$.

This optimization can be made more convenient by choosing a prior that has the same form as the Bradley-Terry model itself. That is to say, virtual results \vec{R}' will serve as a prior: $P(\vec{\gamma}) = P(\vec{R}'|\vec{\gamma})$. This way, the estimation of parameters of the model will consist in maximizing $P(\vec{R}, \vec{R}'|\vec{\gamma})$.

2.5 A Minorization-Maximization Algorithm

Minorization-Maximization is a simple algorithm to compute the maximum *a posteriori* of the Bradley-Terry model.

2.5.1 Notations.

$\gamma_1, \dots, \gamma_n$ are the strength parameters of n individuals. N results R_1, \dots, R_N of independent competitions between these individuals are known. These competitions are of the most general type, as described in Section 2.2. The probability of one competition result may be written as

$$P(R_j) = \frac{A_{ij}\gamma_i + B_{ij}}{C_{ij}\gamma_i + D_{ij}} ,$$

where A_{ij} , B_{ij} , C_{ij} , and D_{ij} are factors that do not depend on γ_i . With this notation, each $P(R_j)$ can be written in n different ways, each time as a function of one particular γ_i . For instance, the example of Section 2.2 would be

$$R_1 = 1-2-3 \text{ wins against } 4-2 \text{ and } 1-5-6-7 ,$$

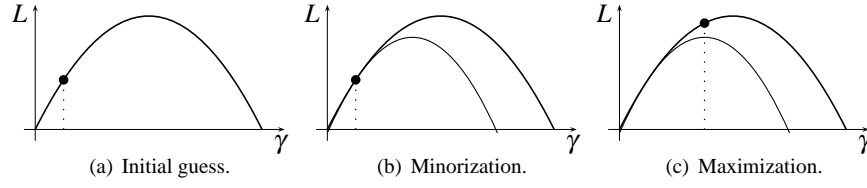


Figure 1: Minorization-maximization.

and its probability

$$P(R_1) = \frac{\gamma_2 \gamma_3 \cdot \gamma_1}{(\gamma_2 \gamma_3 + \gamma_5 \gamma_6 \gamma_7) \cdot \gamma_1 + \gamma_4 \gamma_2} ,$$

so $A_{11} = \gamma_2 \gamma_3$, $B_{11} = 0$, $C_{11} = \gamma_2 \gamma_3 + \gamma_5 \gamma_6 \gamma_7$, $D_{11} = \gamma_4 \gamma_2$. Similarly $A_{21} = \gamma_3 \gamma_1$, $A_{31} = \gamma_2 \gamma_1$, $A_{41} = 0$, etc.

E_j is defined as $E_j = C_{ij} \gamma_i + D_{ij}$, and $W_i = |\{j | A_{ij} \neq 0\}|$ is the number of wins of individual i .

The objective is to maximize:

$$L = \prod_{j=1}^N P(R_j)$$

2.5.2 Derivation of the Minorization-Maximization Formula.

(Readers who do not wish to understand all the details may safely skip to the formula)

Minorization-maximization is an iterative algorithm to maximize L . Its principle is illustrated on Figure 1. Starting from an initial guess $\vec{\gamma}^0$ for $\vec{\gamma}$, a function m is built, that *minorizes* L at $\vec{\gamma}^0$. That is to say, $m(\vec{\gamma}^0) = L(\vec{\gamma}^0)$, and, for all $\vec{\gamma}$, $m(\vec{\gamma}) \leq L(\vec{\gamma})$. The maximum $\vec{\gamma}^1$ of m is then computed. Thanks to the minorization property, $\vec{\gamma}^1$ is an improvement over $\vec{\gamma}^0$. The trick is to build m so that its maximum can be computed in closed form. This optimization algorithm is often much more efficient than traditional gradient-ascent methods.

$$L = \prod_{j=1}^N \frac{A_{ij} \gamma_i + B_{ij}}{C_{ij} \gamma_i + D_{ij}}$$

is the function to be maximized. L can be considered as a function of γ_i , and its logarithm is:

$$\log L(\gamma_i) = \sum_{j=1}^N \log(A_{ij} \gamma_i + B_{ij}) - \sum_{j=1}^N \log(C_{ij} \gamma_i + D_{ij}) .$$

Since either $B_{ij} = 0$ (ie player i is in the winning team), or $A_{ij} = 0$ (ie player i is not in the winning team), the first term can be written:

$$\sum_{j=1}^N \log(A_{ij} \gamma_i + B_{ij}) = \sum_{j, A_{ij}=0} \log(B_{ij}) + \sum_{j, B_{ij}=0} (\log A_{ij} + \log \gamma_i)$$

Terms that do not depend on γ_i can be removed, and the function to be maximized becomes:

$$f(x) = W_i \log x - \sum_{j=1}^N \log(C_{ij} x + D_{ij}) .$$

The logarithms in the right-hand part may be minorized by their tangent at $x = \gamma_i$, as shown on Figure 2. After removing the terms that do not depend on x , the minorizing function to be maximized becomes

$$m(x) = W_i \log x - \sum_{j=1}^N \frac{C_{ij} x}{C_{ij} \gamma_i + D_{ij}} .$$

Its derivative is

$$m'(x) = \frac{W_i}{x} - \sum_{j=1}^N \frac{C_{ij}}{E_j} .$$

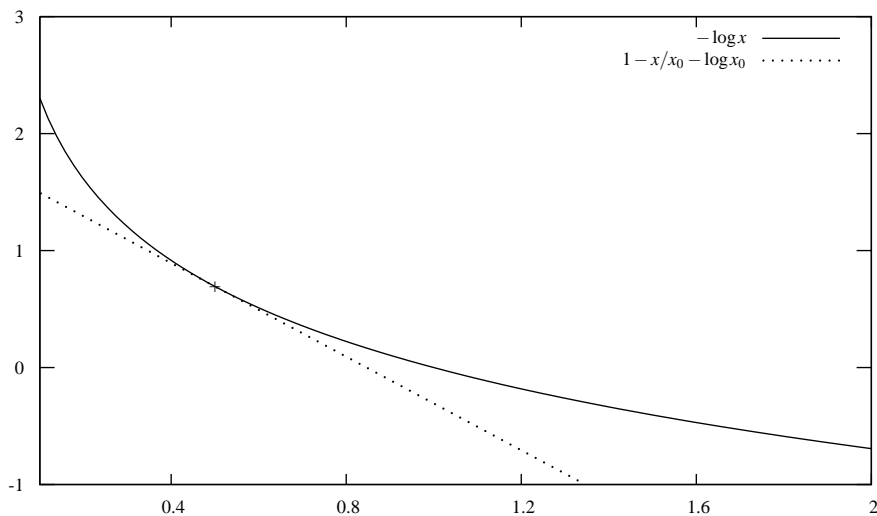


Figure 2: Minorization of $-\log x$ at $x_0 = 0.5$ by its tangent.

The maximum of $m(x)$ can be found by solving $m'(x) = 0$:

$$x = \frac{W_i}{\sum_{j=1}^N \frac{C_{ij}}{E_j}} .$$

2.5.3 Minorization-Maximization Formula.

So, minorization-maximization consists in iteratively updating one parameter γ_i according to this formula:

$$\gamma_i \leftarrow \frac{W_i}{\sum_{j=1}^N \frac{C_{ij}}{E_j}} .$$

If all the parameters are initialized to 1, and the number of participants in each competition is the same, the first iteration of minorization-maximization computes the winning frequency of each individual. So, in some way, minorization-maximization provides a Bayesian justification of frequency-based pattern evaluation. But running more than one iteration improves parameters further.

When players have different strengths, C_{ij} indicates the strength of team mates of i during competition j , and E_j is the overall strength of participants. With the minorization-maximization formula, a win counts all the more as team mates are weak, and opposition is strong.

2.5.4 Batch Updates.

The minorization-maximization formula describes how to update just one γ_i . It is possible to iteratively update all the γ_i one by one, but it may be inefficient. Another possibility is to perform batch updates. A set of mutually exclusive γ_i 's may be updated in one single pass over the data. Mutually exclusive means that they cannot be members of the same team. The batch-update approach still has good convergence properties (Hunter, 2004), and offers the opportunity to re-use computations. In particular, $1/E_j$ need not be computed more than once in a batch.

3. PATTERN-LEARNING EXPERIMENTS IN THE GAME OF GO

A generalized Bradley-Terry model can be applied to supervised learning of Go patterns, by considering that each sample move is a competition, whose winner is the move in question, and losers are the other legal moves.

Each move can be considered as a “team” of features, thus allowing to combine a large number of such features without a very high cost.

3.1 Data

Learning was performed on game records played by strong players on KGS. These game records were downloaded from the web site of Kombilo (Goertz and Shubert, 2007). The training set was made of the 652 games with no handicap of January, 2006 (131,939 moves). The test set was made of the 551 games with no handicap of February, 2006 (115,832 moves). The level of play in these games may not be as high as the professional records used in previous research on pattern learning, but they have the advantage of being publicly available for free, and their level is more than high enough for the current level of Go-playing programs.

3.2 Features

The learning algorithm used 8 tactical features: pass, capture, extension, self-atari, atari, distance to border, distance to the previous move, and distance to the move before the previous move. Some of these features may take more than one value, as explained in Table 1.

The 9th feature was Monte-Carlo owner. It was computed by running 63 random games from the current position. For each point of the board, the number of final positions owned by the player to move was counted.

The 10th feature was shape patterns. Nested circles of radius 3 to 10 according to the distance defined in Table 1 are considered, similarly to Stern *et al.* (2006). 16,780 shapes were harvested from the training set, by keeping those that appear at least 625 times.

Each value that these features can take is considered as a separate “individual”, and is associated to one strength parameter γ_i . Since values within one feature are mutually exclusive, they were all updated together within one iteration of the minorization-maximization algorithm.

3.3 Prior

The prior was set by adding, for each γ_i , one virtual win, and one virtual loss, against a virtual opponent whose γ is 1. In the Elo-rating scale, this produces a symmetric probability distribution, with mean 0 and standard deviation 302.

3.4 Results

Table 1 lists the values of γ for all non-shape features.

Figure 3 plots the mean log-evidence per stage of the game, against the data of Stern, Herbrich, and Graepel (Stern *et al.*, 2006). This mean log-evidence is the mean logarithm of the probability of selecting the target move according to the Bradley-Terry model, measured over the test set. The overall mean log-evidence is -2.69, which corresponds to an average probability of 1/14.7. Uniform probability gives a mean log-evidence of -5.49, which corresponds to an average probability of 1/243.

Figure 4 is a plot of the cumulative distribution of the probability of finding the target move at a given rank, measured over the test set, and compared with other authors.

3.5 Discussion

The prediction rate obtained with minorization-maximization and the Bradley-Terry model is the best among those published in academic papers. de Groot (2005) claims a 42% prediction rate, but his method for measurement is not very clear, and some recent manual testing of his program indicate that its prediction rate may be

| Feature | Level | γ | Description |
|---|-----------|----------|--|
| Pass | 1 | 0.17 | Previous move is not a pass |
| | 2 | 24.37 | Previous move is a pass |
| Capture | 1 | 30.68 | String contiguous to new string in atari |
| | 2 | 0.53 | Re-capture previous move |
| | 3 | 2.88 | Prevent connection to previous move |
| | 4 | 3.43 | String not in a ladder |
| | 5 | 0.30 | String in a ladder |
| Extension | 1 | 11.37 | New atari, not in a ladder |
| | 2 | 0.70 | New atari, in a ladder |
| Self-atari | 1 | 0.06 | |
| Atari | 1 | 1.58 | Ladder atari |
| | 2 | 10.24 | Atari when there is a ko |
| | 3 | 1.70 | Other atari |
| Distance to border | 1 | 0.89 | |
| | 2 | 1.49 | |
| | 3 | 1.75 | |
| | 4 | 1.28 | |
| Distance to previous move | 2 | 4.32 | $d(\delta x, \delta y) = \delta x + \delta y + \max(\delta x , \delta y)$ |
| | 3 | 2.84 | |
| | 4 | 2.22 | |
| | 5 | 1.58 | |
| | ... | ... | |
| | ≥ 16 | 0.33 | |
| Distance to the move before the previous move | ≥ 17 | 0.21 | |
| | 2 | 3.08 | |
| | 3 | 2.38 | |
| | 4 | 2.27 | |
| | 5 | 1.68 | |
| | ... | ... | |
| MC Owner | ≥ 16 | 0.66 | |
| | ≥ 17 | 0.70 | |
| | 1 | 0.04 | 0 – 7 |
| | 2 | 1.02 | 8 – 15 |
| | 3 | 2.41 | 16 – 23 |
| | 4 | 1.41 | 24 – 31 |
| 5 | 0.72 | 32 – 39 | |
| 6 | 0.65 | 40 – 47 | |
| 7 | 0.68 | 48 – 55 | |
| 8 | 0.13 | 56 – 63 | |

Table 1: Model parameters for non-shape features. Each feature describes a property of a candidate move in the current position. A feature might either be absent, or take one of the values indicated in the Level column. Each move in a Go position may combine several features, in which case its γ value is the product of the γ 's of those features. For instance if a move is an extension of a new string in atari ($\gamma = 11.37$), at distance two from the border ($\gamma = 1.49$), distance 3 from the previous move ($\gamma = 2.84$), distance 2 to the move before ($\gamma = 3.08$), on undecided territory with MC Owner = 4 ($\gamma = 1.41$), then its γ is $11.37 \times 1.49 \times 2.84 \times 3.08 \times 1.41$, that is to say about 209.

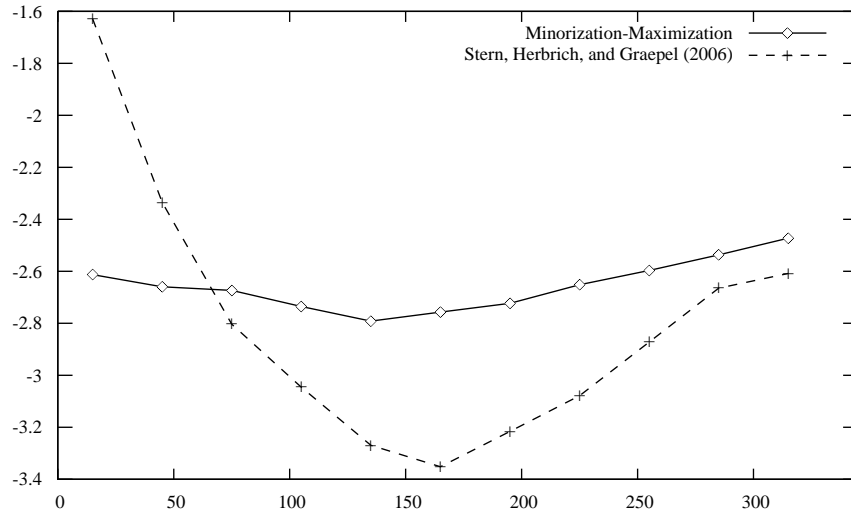


Figure 3: Mean log-evidence per stage of the game (each point is an average over an interval of 30 moves).

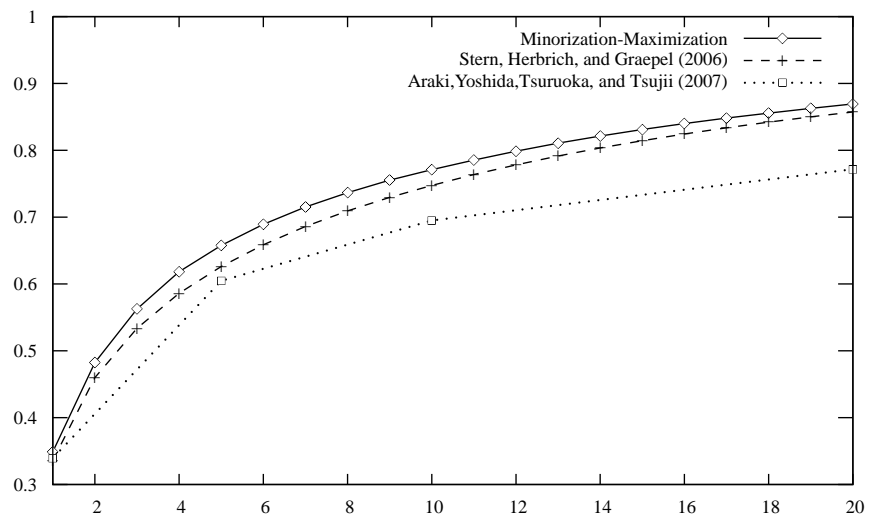


Figure 4: Cumulative distribution: probability of finding the target move within the n best estimated moves.

less (Marchand, 2007). All those prediction rates were measured on different test sets, which, although they are of similar nature, reduces the significance of the comparison.

Despite the similarity of the cumulative distributions, the mean log-evidence per stage of the game has a very different shape from that of Stern, Herbrich, and Graepel. Their algorithm provides much better predictions in the beginning of the game, and much worse in the middle. It is worth noting also that their learning experiments used many more games (181,000 instead of 652) and shape patterns (12,000,000 instead of 16,780). So they tend to learn standard opening sequences by rote, whereas our algorithm learns more general rules.

Minorization-maximization took about one hour of CPU time and 600 Mb of RAM to complete. So, to try to improve prediction further, it would be possible to use more games, and more shape patterns. Most of the computation time was taken by running the Monte-Carlo simulations. In order to learn over many more games, the slow features could be trained afterward, over a small set of games.

Although minorization-maximization is rather efficient, it is still more computationally intensive than the incremental algorithm of Stern, Herbrich, and Graepel, when applied to the same amount of data. Their incremental approach requires more approximations, and does not compute an exact maximum *a posteriori*, but makes it possible to work with a huge number of sample moves. The idea introduced in this paper, of considering moves as teams of features, could be easily transposed into such an incremental algorithm, by ranking individuals from team results with the TrueSkillTM rating system (Herbrich, Minka, and Graepel, 2006).

4. USAGE OF PATTERNS IN A MONTE-CARLO PROGRAM

Despite the clever features of this pattern-learning system, selecting the move with the highest probability still produces a terribly weak Go player. It plays some good-looking moves, but also makes huge blunders because it really does not “understand” the position. Nevertheless, the domain knowledge contained in patterns is very precious to improve a Monte-Carlo program, by providing a good probability distribution for random games, and by helping to shape the search tree. This section briefly describes how patterns are used in CRAZY STONE (Coulom, 2006).

4.1 Random Simulations

The pattern system described in this paper produces a probability distribution over legal moves, so it is a perfect candidate for random move selection in Monte-Carlo simulations. Monte-Carlo simulations have to be very fast, so the full set of features that was described before is much too slow. Only light-weight features are kept in the learning system: 3x3 shapes, extension (without ladder knowledge), capture (without ladder knowledge), self-atari, and contiguity to the previous move. Contiguity to the previous move is a very strong feature ($\gamma = 11$), and tends to produce sequences of contiguous moves like in MOGO (Gelly *et al.*, 2006).

4.2 Progressive Widening of the Monte-Carlo Search Tree

CRAZY STONE also uses patterns to prune the search tree. This is performed at a much slower rate, so the full power of complex features can be used. When a node in the Monte-Carlo search tree is created, it is searched for a while without any pruning, selecting the move according to the policy of random simulations. As soon as a number of simulations is equal to the number of points of the board, this node is promoted to internal node, and pruning is applied. Pruning consists in searching only the n best moves according to patterns, with n growing like the logarithm of the number of random simulations. More precisely, the n th move is added when t_{n-1} simulations have been run, with $t_0 = 0$ and $t_{n+1} = t_n + 40 \times 1.4^n$. On 19×19 , thanks to the distance-to-the-previous-move feature, progressive widening tends to produce a local search, like in MOGO (Gelly *et al.*, 2006).

Progressive widening was independently invented by Chaslot *et al.* (2007) under the name of “progressive un-pruning”. It is also similar in spirit to Cazenave’s idea of iterative widening (2001).

| Pat. | P.W. | Size | Minutes/game | GNU Level | Komi | Games | Win ratio |
|------|------|---------|--------------|-----------|------|-------|-----------|
| - | - | 9 × 9 | 1.5 | 10 | 6.5 | 170 | 38.2% |
| x | - | 9 × 9 | 1.5 | 10 | 6.5 | 170 | 68.2% |
| x | x | 9 × 9 | 1.5 | 10 | 6.5 | 170 | 90.6% |
| - | - | 19 × 19 | 32 | 8 | 6.5 | 192 | 0.0% |
| x | - | 19 × 19 | 32 | 8 | 6.5 | 192 | 0.0% |
| x | x | 19 × 19 | 32 | 8 | 6.5 | 192 | 37.5% |
| x | x | 19 × 19 | 128 | 8 | 6.5 | 192 | 57.1% |

Table 2: Match results. P.W. = progressive widening. Pat. = patterns in simulations.

4.3 Performance against GNU Go

Table 2 summarizes CRAZY STONE's performance against GNU Go 3.6, on an AMD Opteron at 2.2 GHz, running on one CPU. CRAZY STONE ran, per second, from the empty position, 15,500 simulations on 9 × 9, and 3,700 on 19 × 19.

Results indicate that using patterns in simulations and progressive widening both bring significant improvements to the playing strength on 9 × 9. On 19 × 19 the contribution of progressive widening to the playing strength is huge, and playing strength scales with computational power.

5. CONCLUSION

The research presented in this paper demonstrates that a generalized Bradley-Terry model is a very powerful technique for pattern learning in the game of Go. It is simple and efficient, can combine several features, and produces a probability distribution over legal moves. It is an ideal tool to incorporate domain knowledge into Monte-Carlo tree search.

Experiment results clearly indicate that significant progress can be made by learning shapes over a larger amount of training games, and improving features. In particular, the principle of Monte-Carlo features is very powerful, and could be exploited more, as Bouzy did with history and territory heuristics (Bouzy, 2006).

Also, the validity of the model could be tested and improved. First, using all the moves of one game as sample data breaks the hypothesis of independence between samples, since consecutive positions are very similar. Sampling one or two positions per game might be better. Also, the linearity hypothesis of the generalized Bradley-Terry model, according to which the strength of a team is the sum of the strengths of its members, is likely to be wrong. Estimating the strength of some frequent feature pairs separately might improve predictions.

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