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A Comprehensive Survey on Lexicase Selection

Alina Geiger, Dirk Schweim, Martin Briesch, David Wittenberg, Franz Rothlauf, Dominik Sobania

Abstract—In recent years, rapid progress has been made in the field of lexicase selection with the development of many new variants, expanding its applicability across various domains. In order to consolidate this progress, we conduct a systematic survey on the current state of research on lexicase selection. Our analysis includes about 300 in-scope papers, making it a comprehensive overview of current literature in this field. We identify five major research streams, namely lexicase, ϵ -lexicase, batch-lexicase, down-sampled lexicase as well as hybrids and approximations. For each research stream, we highlight and discuss the main empirical and theoretical contributions. Further, we identify the major application domains of lexicase, ranging from program synthesis and symbolic regression to classification, Boolean logic, and even robot control. Based on our findings, we discuss current challenges and identify opportunities for future research directions in evolutionary computation and beyond.

Index Terms—Lexicase, Selection, Survey.

I. INTRODUCTION

Selection plays a crucial role in evolutionary computation [1] and many selection operators have been proposed over the years, like fitness-proportionate selection, tournament selection, and, last but not least, lexicase selection [2]. Classical methods like tournament selection operate on an aggregated fitness value, which is calculated by averaging the performance of a candidate solution across all training cases (input-output pairs). However, this creates an evaluation bottleneck where all the information about a candidate solution is compressed into a single value [3], [4]. This poses a dilemma, as problems of interest often exhibit structures that require different modular capabilities from a solution [2]. Spector [2] discusses the example of a computer program that should serve as a mathematical calculator and, therefore, must be able to perform very different operations such as addition, subtraction, and multiplication correctly. Discovering all those operations simultaneously during evolution is highly unlikely. In contrast, evolving a candidate solution that solves one operation at a time may be possible. With an evaluation bottleneck, however, such candidate solutions that excel in a particular modular part (e.g., addition) but perform poorly on all other modular parts will be filtered out by selection and their genetic material might be lost [5].

In contrast to those selection operators, lexicase selection does not operate on an aggregated fitness value but instead

evaluates candidate solutions based on their performance on individual training cases [2]. More specifically, for each selection event, all training cases are first shuffled in random order. All candidate solutions are then evaluated on a single training case, only preserving those individuals with the best performance on this case. This is repeated for the next training case until only one candidate solution remains. If more than one candidate solution remains after all training cases have been used for evaluation, one of the remaining candidates is chosen either at random or according to some other metric [6].

This approach not only increases diversity [7], [8] but also promotes the selection of *specialist* solutions that perform exceptionally well on training cases where others perform poorly while performing worse on average. These specialists would rarely be selected with methods based on aggregated fitness values. Specialists stand in contrast to *generalist* solutions, which perform well on average but do not excel on any particular training case compared to the rest of the population [9], [10]. Specialists can serve as stepping stones for the evolutionary search and may contribute genetic material that leads to candidate solutions capable of solving all parts of the problem [7], [10]–[13]. In the mathematical calculator example, lexicase selection may discover specialists for individual mathematical operations, which can guide the search towards a generalist solution that successfully performs all required operations [2].

Lexicase selection has demonstrated impressive performance across a variety of application domains [14]–[18], leading to the development of many different variations and extensions in the literature. Nevertheless, many researchers only use outdated or inconsistent versions of lexicase selection without accounting for recent advancements. As a result, there is a need for a comprehensive overview that brings together current knowledge, clarifies the differences among lexicase variants, and highlights its practical applications.

Therefore, in this work, we conduct a systematic literature review to identify, consolidate, and structure the current state of research on lexicase selection. We identify almost 300 papers published between 2012 and 2025 that either investigated lexicase selection or used it as part of their experiments. From these papers, we derive five major research streams: First, lexicase as a selection method for modular problem settings (Section III). Second, ϵ -lexicase selection for continuous-valued problems (Section IV). Next, the combination of lexicase with batching (Section V) and down-sampling techniques (Section VI). Lastly, methods that approximate lexicase selection as well as hybrid approaches (Section VII). Moreover, we find that the main application domains of lexicase selection are program synthesis and symbolic regression. Additional applications include classification, Boolean logic, robotics, and others. We also identify applications of lexicase selection

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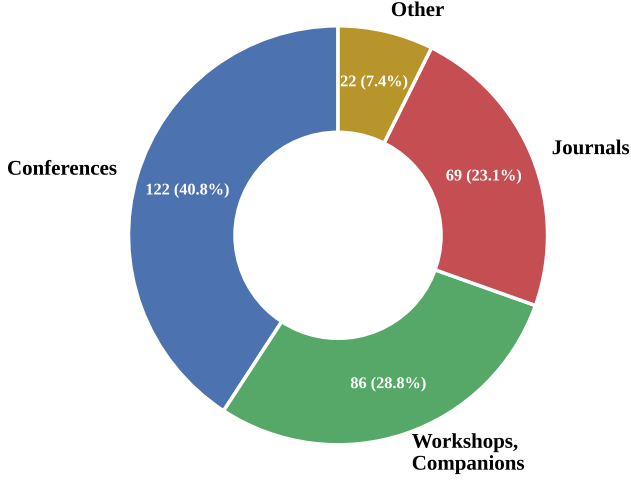


Fig. 1. Number of publications by publication type.

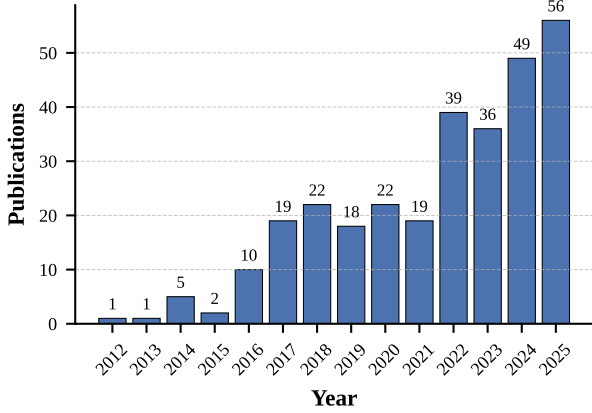


Fig. 2. Number of publications per year.

beyond the scope of evolutionary computation. Based on our findings, we discuss ongoing challenges of lexibase selection and open questions in the current body of research.

The rest of the paper is structured as follows: In Section II, we present the method and results of our systematic literature review. Sections III–VII give a detailed overview of lexibase selection and its variants. Section VIII provides an overview of the application domains of lexibase selection, followed by a discussion of challenges and open questions in Section IX. Finally, Section X concludes this survey.

II. METHODOLOGY

To systematically identify the relevant literature on lexibase selection, we performed a search on Google Scholar using the keyword “lexibase”. As lexibase selection was first introduced in 2012 [2], we limited our search to papers published between 2012 and 2025. At the time of data collection, the query returned 907 publications. From these, we identified papers

that are written in English and used lexibase as part of their experiments or theoretical analysis. We excluded papers that only cite lexibase selection as part of their related work or use the term lexibase in a different context (e.g., linguistics). From the remaining papers, we included peer-reviewed articles in journals and conference proceedings, technical reports, and preprints, while excluding bachelor’s, master’s, and PhD theses.¹ In total, this resulted in 299 in-scope publications for this survey. We categorize those papers by application domain in Table III² and further discuss their application areas in Section VIII.

Figure 1 shows the number of publications by publication type. We notice that most in-scope publications on lexibase selection appear in conference proceedings (~ 41%) with *Genetic and Evolutionary Computation Conference* (45) and *European Conference on Genetic Programming* (19) being the most frequent venues. Workshop proceedings and conference companions account for approximately 29% of in-scope publications, with 55 *Genetic and Evolutionary Computation Conference Companion* and 26 *Genetic Programming Theory and Practice* publications. Around 23% of in-scope papers are published in journals such as *Genetic Programming and Evolvable Machines* and *IEEE Transactions of Evolutionary Computation*. Approximately 7% of the in-scope publications do not belong to any of the mentioned categories and are primarily preprints.

Additionally, as displayed in Figure 2, we observe a steady increase in publications over the years since the inception of lexibase selection. While initially only a few papers were published on the topic, in 2016 lexibase began to gain traction and from 2017 to 2021 around 20 papers were published every year. By 2022, this number doubled again to almost 40 papers and increased to over 50 in 2025. This trend indicates growing interest in lexibase selection and motivates a systematic review of the rapidly expanding body of literature.

Finally, we identified those papers that not only applied lexibase selection in their experiments or used it as a baseline, but also significantly contributed to the field of lexibase selection by either improving the performance of lexibase selection or contributing to the understanding of lexibase selection and its applications. We refer to these papers as “core papers” in the remainder of this survey and list them in Table I.² In total, we identified 156 core papers. Based on the core papers listed in Table I, we identified five major research streams of lexibase selection: Lexibase selection in general, ϵ -lexibase selection as a special variant of lexibase for continuous-valued problems like symbolic regression, combinations of lexibase with batching or down-sampling approaches, and approximations as well as hybridization of lexibase selection with other methods.

In the following sections, we will discuss the advancements and contributions to each research stream in detail. Although not included in the systematic review or quantitative analyses, we discuss relevant recent papers published in 2026 where appropriate.

¹For most PhD theses, the key contributions have already been published in peer-reviewed venues and are therefore already represented in this survey.

²Note that some papers appear multiple times.

TABLE I

CORE PAPERS BY DIFFERENT RESEARCH STREAMS IN CHRONOLOGICAL ORDER. WE DEFINE CORE PAPERS AS THOSE THAT ACTIVELY CONTRIBUTE TO THE UNDERSTANDING AND PROGRESS OF LEXICASE SELECTION, RATHER THAN MERELY APPLYING IT AS A SELECTION METHOD IN EC OR AS A BASELINE FOR COMPARISON.

Research Stream	References	#
Lexicase	[2], [19], [20], [21], [6], [22], [23], [7], [24], [8], [25], [12], [26], [27], [28], [29], [30], [31], [32], [33], [34], [35], [36], [37], [38], [39], [40], [41], [42], [43], [44], [45], [9], [46], [47], [48], [10], [1], [49], [50], [51], [52], [53], [54], [55], [56], [57], [58], [59], [60], [61], [62], [63], [64], [65], [66], [67], [68], [69], [70], [71], [72], [73], [74], [75], [76], [77], [78], [79], [80], [81], [82], [83], [84], [85], [86], [87], [88]	78
ϵ -Lexicase	[15], [16], [89], [27], [90], [91], [92], [93], [94], [95], [96], [97], [44], [98], [99], [100], [101], [1], [102], [103], [104], [105], [106], [107], [108], [109], [110], [56], [58], [111], [112], [113], [69], [70], [114], [115], [76], [116], [117], [86], [118], [119], [120], [121]	44
Batched Lexicase	[17], [1], [107], [122], [115], [119]	6
Down-Sampling	[123], [124], [125], [1], [126], [127], [128], [129], [130], [131], [56], [132], [133], [134], [135], [136], [137], [138], [139], [140], [141], [142], [143], [144], [119], [145]	26
Hybrids & Approximations	[22], [146], [147], [148], [95], [149], [150], [151], [152], [153], [1], [154], [155], [18], [156], [157], [56], [158], [159], [160], [161], [162], [163]	23

Algorithm 1 Lexicase selection (based on [6])

```

1:  $P' :=$  all candidate solutions  $p \in P$ 
2:  $C' :=$  all training cases  $c \in C$  in random order
3: while  $|P'| > 1$  and  $|C'| > 0$  do
4:    $c :=$  first case in  $C'$ 
5:    $\mathbf{e}_c :=$  evaluate all  $p \in P'$  on  $c$ 
6:    $P' :=$  all candidate solutions with  $e_c = e_c^*$ 
7:   remove  $c$  from  $C'$ 
8: end while
9: return random candidate solution from  $P'$ 

```

III. LEXICASE SELECTION

In contrast to other selection methods, lexicase selection does not use a single aggregated fitness value for selection. Instead, lexicase selection operates on a case-by-case basis, selecting candidate solutions that perform particularly well on individual training cases more frequently, even if their aggregated performance is worse than that of other candidate solutions [2].

More specifically, Algorithm 1 (adopted from [6]) describes how lexicase selects a single parent in detail. First, the candidate pool P' is initialized by copying the whole population of candidate solutions $p \in P$ (line 1). Similarly, the case pool C' is created, containing all training cases $c \in C$ in random order (line 2). Afterwards, all candidate solutions in P' are evaluated on the first training case $c \in C'$ (line 4–5) to obtain a vector of error values for that case, denoted by $\mathbf{e}_c$.³

Only those candidate solutions with the lowest error $e_c = e_c^*$ on that training case are retained, while all other candidates (i.e., those that do not achieve the best performance on that case) are removed from P' (line 6). The first training case c is then removed from C' (line 7) and the process is repeated with the next training case in C' . If only a single candidate solution remains in P' , that candidate solution is selected and returned. Otherwise, if C' contains no more training cases, one of

³In practice, the error of each candidate solution on each training case is usually only calculated once per generation and then cached for all selection events.

TABLE II

EXAMPLE POPULATION OF FIVE CANDIDATE SOLUTIONS EVALUATED ON FOUR TRAINING CASES WITH THEIR CORRESPONDING ELITE CASES, AGGREGATED PERFORMANCE IN TERMS OF THE MAE, AND SELECTION PROBABILITY USING LEXICASE A_{lex} (BASED ON [2], [44]).

Candidates	Error per case				Elite cases	MAE	A_{lex}
	e_1	e_2	e_3	e_4			
p_1	0	2	2	3	$\{c_1, c_3\}$	1.75	0.167
p_2	0	3	2	2	$\{c_1, c_3\}$	1.75	0.25
p_3	0	0	6	3	$\{c_1, c_2\}$	2.25	0.333
p_4	0	1	3	3	$\{c_1\}$	1.75	0
p_5	2	5	5	1	$\{c_4\}$	3.25	0.25

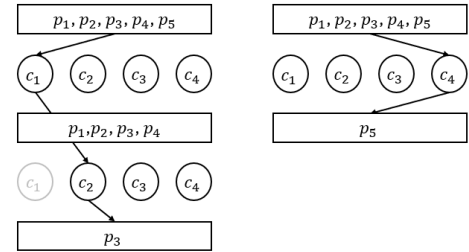


Fig. 3. Two example parent selection events with lexicase selection for the example population described in Table II (based on [44])

the remaining candidate solutions in P' is selected uniformly at random (line 9). To construct the parent population, the selection procedure is repeated using a newly shuffled ordering of C' each time until the desired number of parents has been selected.

To further illustrate lexicase selection, consider the example displayed in Table II (based on [2], [44]), which shows an example population consisting of five candidate solutions $p \in P$ and the error vectors of four training cases $c \in C$. For each candidate solution p_i , we indicate the elite cases (i.e., the cases where the individual achieves the lowest error e_c^*) and report its aggregated error in terms of the mean absolute error (MAE).

Figure 3 (based on [44]) illustrates two selection events

with the example population described in Table II. In the first selection event (left), the case c_1 is drawn first, where four candidate solutions are elite. The candidate solution p_5 is excluded from the candidate pool, because it is outperformed by the other candidates. Next, c_2 is drawn, where p_3 performs best among the remaining candidate solutions. Hence, p_3 is selected as a parent in this selection event. In the second example (right), the case c_4 is drawn first, where only p_5 is elite, so this candidate is selected immediately, even though it would never be selected based on aggregated fitness scores because it has the highest average error.

Overall, the case-by-case approach of lexicase selection alters the selection probabilities and shifts selection pressure from global average performance to excellence in specific training cases [36], effectively negating the evaluation bottleneck of other selection methods [5]. In Table II, the selection probabilities A_{lex} indicate the probability of an individual p_i being selected using lexicase selection, which can be calculated using the following recursive function taken from La Cava et al. [44]:

$$A_{\text{lex}}(p|P, C) = \begin{cases} 1 & \text{if } |P| = 1, \\ \frac{1}{|P|} & \text{if } |C| = 0, \\ \frac{1}{|C|} \sum_{k_s \in K_p} A_{\text{lex}}(p | P(q | k_s \in K_q), C \setminus \{k_s\}) & \text{otherwise.} \end{cases} \quad (1)$$

where C denotes the set of training cases, P the set of candidate solutions, and K_p the subset of cases from C for which candidate p is elite. If only one candidate remains ($|P| = 1$), this candidate is selected. If there are no training cases left ($|C| = 0$), all remaining candidates are selected with equal probability.

If multiple candidates and cases remain, the probability of selecting candidate p depends on the probability of drawing a case $k_s \in K_p$ (meaning a case where p is elite), which is $\frac{1}{|C|}$ for each case. If a case $k_s \in K_p$ is drawn, the probability of selecting candidate p further depends on the subset of candidates that are also elite on case k_s , $P(q | k_s \in K_q)$, as well as the remaining cases $C \setminus \{k_s\}$ [44]. Calculating these probabilities is NP-hard [70].

For our illustrative example (Table II), the selection probability of p_5 is 0.25, as p_5 is always selected whenever c_4 is drawn first and never otherwise. The selection probability of p_4 is 0, because it performs best on case c_1 , but so do p_1 , p_2 , and p_3 . These candidates are also elite on additional training cases, whereas p_4 is not. Consequently, p_4 is always eliminated during subsequent filtering steps. For candidate p_3 , the selection probability is 0.333, because it will be selected if case c_2 is drawn first (with probability 0.25) or if case c_1 is drawn first and subsequently c_2 (with probability 0.083). Thus, selection pressure is shifted away from aggregate performance measures such as MAE and toward case-wise elite performance. This shift in selection pressure has several important consequences.

For example, lexicase selection has a significant impact on population diversity, which is typically measured as the number of distinct error vectors in a population, commonly referred to as behavioral diversity [164]. Multiple studies show that lexicase selection increases population diversity compared to

other selection methods [7], [8]. Moreover, lexicase selection benefits from a high population diversity to effectively explore the search space, while tournament selection often gets stuck in local optima [56].

However, diversity sometimes drops significantly during runs with lexicase selection due to hyperselection events. As described in [25], hyperselection events occur when a single candidate is selected in a disproportionately large number of selection events because it is elite on a large number of training cases relative to the rest of the population. However, hyperselection events do not influence the performance of lexicase selection [25] because lexicase selection is able to re-diversify populations from a low diversity starting point or after a diversity crash, an advantage that is not observed with tournament selection [8].

While these studies suggest that this increase in diversity positively affects the performance of lexicase selection, it is not the sole reason why lexicase selection performs better [10]. Lexicase selection is also more likely to select specialists, meaning candidates that perform exceptionally well on difficult training cases despite having a comparatively high average error. In our example, candidate p_5 has a higher selection probability compared to p_1 and p_4 because it is the only one that is elite on case c_4 even though its average error is higher. In contrast, with tournament selection, the selection probability would only depend on the average error, so generalists would be preferred (in our example p_1 , p_2 , and p_4). It has been found that the selection of specialists is crucial for the problem-solving success of lexicase selection [9], [10], [36] as these specialists can be stepping stones to find a solution that solves all training cases [7], [10]–[13]. Moreover, it is assumed that lexicase selection produces higher population diversity because specialists with different behavior can coexist within a population, whereas candidates exhibiting similar behavior compete directly with each other [7], [8].

Despite all these positive effects, lexicase comes at the cost of increased runtime. For a population P and a set of training cases C , the worst-case time complexity of generating a parent population using lexicase selection is $\mathcal{O}(|P|^2|C|)$. This is worse than aggregated selection methods like tournament selection, which have a worst-case complexity of $\mathcal{O}(|P||C|)$ [44]. Fortunately, in practice, the runtime for lexicase tends to be lower than its worst-case complexity suggests, as the pool of candidate solutions is often filtered down to a single candidate solution before all training cases are used [6], [44].

Moreover, Helmuth et al. [55] showed that the expected runtime of lexicase selection depends on behavioral diversity. In highly diverse populations, the expected runtime reduces to $\mathcal{O}(|P| \cdot (|P| + |C|))$, and empirical results indicate that populations using lexicase selection are often sufficiently diverse for runtimes to be much lower than the worst-case complexity. Nonetheless, lexicase is slower than aggregated methods, especially for large populations.

The base lexicase algorithm can be adjusted in multiple ways. For example, if more than one candidate solution remains after all training cases have been considered, the selection within the remaining candidate pool does not have

Algorithm 2 Dynamic ϵ -lexicase selection (based on [44])

```

1:  $P' :=$  all candidate solutions  $p \in P$ 
2:  $C' :=$  all training cases  $c \in C$  in random order
3: while  $|P'| > 1$  and  $|C'| > 0$  do
4:    $c :=$  first case in  $C'$ 
5:    $\mathbf{e}_c :=$  evaluate all  $p \in P'$  on  $c$ 
6:   calculate  $\epsilon_c$  from  $\mathbf{e}_c$ 
7:    $P' :=$  all candidate solutions with  $e_c \leq e_c^* + \epsilon_c$ 
8:   remove  $c$  from  $C'$ 
9: end while
10: return random candidate solution from  $P'$ 

```

to be random (line 9 in Algorithm 1). Alternatively, a second objective like program size could break ties between candidate solutions by applying parsimony pressure as discussed in [157]. This favors smaller solutions, effectively reducing unnecessary code growth. However, other tie-breaking metrics could be applied here as well.

Further, Helmuth et al. [10] describe a pre-selection step for lexicase selection that groups candidate solutions with identical behavior and only initializes P' with one representative of each group. This does not change the selection probabilities, but reduces the computational time. Otherwise, if only candidates with identical behavior are left, the algorithm will continue to compare them on all remaining cases without reducing the candidate pool any further.

Additionally, the ordering of C' (line 2) does not have to be random but can instead be biased in different ways. So far, no case ordering has been shown to significantly improve performance [148]. However, ordering the training cases by their discriminatory power (so that those cases with more variation in their corresponding error vectors are more likely to appear earlier in the sequence) reduces runtime while keeping the same performance, which is especially beneficial for large data sets [156], [158]. Others describe variants of lexicase selection that only use a subset of the available training cases to initialize C' [123], [124], [138], which we discuss further in Section VI.

Likewise, instead of evaluating individuals iteratively on a single training case at a time (line 4-5), one can use a batch of training cases for each filtering event [17], [122] (see Section V). Lastly, relaxing the elitism in line 6 can be beneficial, particularly in continuous or noisy application domains [15] (see Section IV).

IV. EPSILON LEXICASE SELECTION

So far, we have only considered lexicase selection with a strictly elitist filtering approach. This works particularly well for application domains with discrete fitness values. However, in application domains with continuous (potentially noisy) fitness values per training case, such as symbolic regression, strict elitism can become a major drawback since ties in performance are very rare [15], [44]. Therefore, selecting only the best-performing candidate on each case typically results in only a single training case being used per selection event. This can lead to a significant decrease in performance [15], [44].

One solution to this problem is relaxing the condition for candidate solutions to be kept in P' . In particular, ϵ -lexicase selection [15] introduces a threshold value: Let e_c denote the error of a candidate solution for a given case c (lower is better) and e_c^* the best error on this case of all candidates in the candidate pool P' , then all candidate solutions with an error $e_c \leq e_c^* + \epsilon_c$ are retained in P' [15], [44]. ϵ_c is calculated for each training case as the median absolute deviation [165]

$$\epsilon_c = \text{median}(|\mathbf{e}_c - \text{median}(\mathbf{e}_c)|),$$

where $\mathbf{e}_c$ is a vector of error values of the remaining candidate solutions in P' with respect to c . The rationale of calculating ϵ_c per training case is that the selective pressure is adapted based on the variability of error values in the candidate pool P' per training case. Usually, the variance is high at the beginning of the search, and thus ϵ_c is relatively high. As the search converges, the epsilon values decrease [44].

Algorithm 2 shows the pseudocode for ϵ -lexicase selection. In comparison to standard lexicase selection, an extra line for calculating ϵ_c is added in line 6 and the condition in line 7 is relaxed using ϵ_c . This specific version, where both e_c^* and ϵ_c are determined based on the remaining candidate pool P' , is called *dynamic* ϵ -lexicase selection [44].

Alternatively, the threshold values ϵ_c and the corresponding elite errors e_c^* can also be calculated only once per generation before the filtering step using the error values of the whole population for each case, which is called *static* ϵ -lexicase selection [15], [44]. *Semi-dynamic* ϵ -lexicase selection only calculates e_c^* dynamically from the remaining candidate solutions in P' while keeping ϵ_c fixed within each generation for each case [44]. All described variations of ϵ -lexicase selection perform similarly [44].

Recently, a new method for calculating the threshold has been proposed that divides the error values for a case into two partitions so that the sum of the variances within the two groups is minimized. This has been found to be beneficial for real-world regression problems [116].

Regarding other properties of lexicase selection (such as diversity maintenance, hyperselection, and specialist selection), ϵ -lexicase selection behaves similarly to standard lexicase selection, except that the average number of cases used during selection is higher [133].

Other relaxed forms of lexicase selection for symbolic regression problems are discussed in [95]. However, ϵ -lexicase selection generally outperformed the other approaches in the reported experiments and is the most commonly used lexicase variant for symbolic regression. The concept of adding a threshold to lexicase selection has also been applied in other problem domains such as evolutionary robotics [92], [146] and learning classifier systems [17].

V. BATCHED LEXICASE VARIANTS

With lexicase selection, candidate solutions are usually evaluated on only a few training cases before a parent is selected, which can negatively affect generalization [17], [52], [122]. To solve this problem, Aenugu and Spector [17] proposed using batches of training cases instead of single training cases

Algorithm 3 Batch-lexicase selection (based on [122])

```

1:  $P' :=$  all candidate solutions  $p \in P$ 
2:  $C' :=$  all training cases  $c \in C$  in random order
3: while  $|P'| > 1$  and  $|C'| > 0$  do
4:    $B :=$  first  $b$  cases in  $C'$ 
5:    $e_B :=$  evaluate all  $p \in P'$  on  $B$ 
6:    $P' :=$  all candidate solutions with  $e_B = e_B^*$ 
7:   remove  $B$  from  $C'$ 
8: end while
9: return random candidate solution from  $P'$ 

```

to evaluate candidate solutions in each filtering step in the lexicase selection procedure as shown in Algorithm 3. In each selection event, the training cases are combined to form batches of equal size b (line 4). Then, the average performance of each candidate solution $p \in P'$ is calculated for the first batch B (line 5). Candidates that are elite on batch B proceed to the next batch (line 6). This is repeated until a single candidate remains or there are no more training cases left. Batch-lexicase selection has been found to improve generalization compared to lexicase selection in learning classifier systems [17] and for program synthesis problems [122].

This approach can be combined with a threshold depending on the problem domain. A straightforward way to incorporate a threshold is to require a candidate's average performance on a batch to exceed a manually set value before it can pass the filtering step, e.g., for classification problems [17]. Geiger et al. [136] proposed batch- ϵ -lexicase selection for symbolic regression problems, where training cases are combined into batches but the filtering step is performed using ϵ -lexicase selection. A recent study has found that batch- ϵ -lexicase selection outperforms ϵ -lexicase selection for symbolic regression problems when the search is conducted for only a short period of time [143]. The authors assume that this is because the use of batches enables the evaluation of individuals on more training cases early on, reducing the impact of case ordering [17]. However, in longer search runs, batch-free variants outperform those that use batches [143].

VI. DOWN-SAMPLING TECHNIQUES

From a computational perspective, the evaluation of candidate solutions is usually the most expensive part of evolutionary search [166]. Consequently, large training sets limit the number of candidate solutions that can be evaluated during search for a given budget [123], [124], [128]. A simple solution would be to reduce the training set beforehand in order to increase the number of generations or the population size while keeping the number of evaluations constant. However, if the training set is reduced too much, the problem space might not be adequately covered anymore [123], [124].

Therefore, down-sampling techniques can be used to sample different subsets of training cases in each generation. This not only reduces computational costs, but also improves performance and reduces overfitting and bloat [21], [167]. Using different subsets in each generation has the benefit that candidate solutions are evaluated on a large fraction of the original training set over the course of evolution [123].

Algorithm 4 Random down-sampled lexicase selection (based on [123])

```

1:  $P' :=$  all candidate solutions  $p \in P$ 
2:  $S :=$  random subset of  $C$  of size  $r * |C|$ 
3:  $S' :=$  all training cases  $c \in S$  in random order
4: while  $|P'| > 1$  and  $|S'| > 0$  do
5:    $c :=$  first case in  $S'$ 
6:    $e_c :=$  evaluate all  $p \in P'$  on  $c$ 
7:    $P' :=$  all candidate solutions with  $e_c = e_c^*$ 
8:   remove  $c$  from  $S'$ 
9: end while
10: return random candidate solution from  $P'$ 

```

Hernandez et al. [123] analyzed the combination of lexicase selection with random down-sampling (RDS) [167] as shown in Algorithm 4. In each generation, a different random subset S of size $r * |C|$ (with $r < 1$) is created from the original training set C (line 2). This subset S is shuffled (line 3) and then used for parent selection with lexicase selection (line 4–10). Lexicase selection with RDS has previously been used implicitly in the domain of evolutionary robotics, where evaluations are very expensive, and therefore only a few random training cases are used for evaluation in each generation [16], [92], [97].

Furthermore, Hernandez et al. [123] proposed cohort lexicase selection, which splits the population P and the training set C into subgroups, where each candidate subgroup is evaluated only on a subgroup of C . During lexicase selection, each candidate solution competes only with solutions in the same subgroup.

Cohort sampling and RDS significantly increased the success rates of lexicase selection [123], [124]. Both reduce the number of evaluations by a factor of $D = \frac{1}{r}$ per generation. Thus, using the same number of evaluations, you can explore D times more candidate solutions (either by increasing the number of generations or by increasing the population size) [123]. As RDS and cohort lexicase selection perform similarly, while RDS is considerably simpler [123], [124], subsequent research mainly focused on analyzing RDS [125], [128]. These studies attribute the higher performance of lexicase selection with RDS to the fact that it evaluates a larger number of candidate solutions, rather than other factors, such as extending the search to more generations, changing the environment, or improving generalization. Furthermore, they found that the performance improvements are consistent across a wide range of down-sampling rates r [125], [128], as long as r is not extremely high or low [129].

In addition, RDS has been studied in combination with ϵ -lexicase selection [133] and other lexicase variants for symbolic regression problems [136], [143], where it significantly improved performance. However, RDS slightly reduced population diversity for ϵ -lexicase [133].

A downside of RDS is that it reduces the likelihood of selecting specialists, because it may exclude those training cases on which candidate solutions are specialized [124]. Consequently, selection pressure shifts toward generalists and away from specialists, which can lead to performance drops on

certain problems [56], [124]. Hernandez et al. [56] argue that RDS reduces the exploration capabilities of lexicase selection.

Moreover, RDS selects training cases with equal probability, although not all training cases are equally important. Some are redundant and provide little additional information, while others are rare and highly-informative edge cases or provide critical information about the current population. RDS may discard these critical cases for several generations, potentially harming the search [138]. To overcome this problem, Boldi et al. [138] proposed informed down-sampling (IDS).

In particular, IDS first constructs error vectors e_c for each training case by evaluating the population on all training cases. Each entry in the error vector corresponds to the performance of a single candidate solution on case c . Thus, distances between error vectors measure how differently the current population performs on distinct training cases. For example, if all candidate solutions show identical performance on two training cases, then the distance between their error vectors is zero. Consequently, larger differences in performance between candidate solutions of a population for two training cases lead to larger distances between the corresponding error vectors [138]. These distances are then used to create a diverse subset of training cases with regard to their error vectors, for example, by using the farthest-first-traversal algorithm [168]. To achieve similar computational savings as RDS, the error vectors are constructed using only a fraction of the parent population and are only updated every few generations. As a result, IDS leads to the creation of diverse and meaningful subsets, including highly informative edge cases, instead of randomly drawing training cases.

IDS combined with lexicase selection outperforms RDS for program synthesis problems [132], [138]. Consequently, IDS has been studied in combination with other selection methods as well [134], [141]–[143], [169]. For program synthesis, IDS is particularly effective for diversity-preserving selection methods such as lexicase [141]. In the domain of symbolic regression, IDS closes the performance and behavior gap between tournament and lexicase selection [142], [143]. However, it does not consistently improve performance [143]. Recent work indicates that this might be due to the tendency of IDS to over-sample outliers when applied to real-world problems [170]. Unlike edge cases, outliers deviate from the underlying pattern [171], and oversampling them potentially harms the search process and might lead to overfitting. Therefore, Geiger et al. [170] proposed ROIDS, an outlier-aware variant of IDS that excludes potential outliers by removing a small percentage of cases with the worst aggregated error value before creating subsets with IDS.

Moore and Stanton [135] proposed an alternative approach to generating more meaningful subsets by selecting training cases more frequently based on their effectiveness in filtering during the selection process. This sampling approach achieved a significant reduction in computational cost within the domain of evolutionary robotics [135], [145].

VII. HYBRIDS AND APPROXIMATIONS

As discussed in Section III, lexicase selection has improved performance in many problem domains due to characteristics

such as mitigating the evaluation bottleneck and preserving specialists. However, other selection methods exhibit strengths that complement those of lexicase selection. Consequently, several hybrid approaches have been proposed to combine the strengths of lexicase selection with those of other selection methods.

For example, novelty search [172] maintains a high population diversity by selecting parents with novel behavior. Knobely selection [149] aims to combine novelty and lexicase selection by randomly choosing either lexicase selection or novelty selection for each selection event to obtain a parent population that is both diverse and contains high-performing individuals. Similarly, novelty-lexicase selection [151] calculates the novelty scores for an individual on each training case and appends them to its list of error values. Therefore, during lexicase selection, the focus is sometimes on individuals with low errors on certain cases and sometimes on individuals exhibiting novel behavior on certain cases. Novelty-lexicase selection increased population diversity for program synthesis problems and found more solutions.

Double lexicase selection [160] reduces bloat in regression problems by performing a two-step selection process. First, multiple candidates are sampled using ϵ -lexicase selection. Then, one of them is selected as a parent based on its size with roulette wheel selection. Consequently, smaller individuals have a higher probability of being selected.

However, the major drawback of lexicase selection is that its high-quality solutions come at the cost of increased runtime. Therefore, recent approaches focus on reducing the runtime of lexicase selection by approximating its behavior. For example, Ding et al. [159] introduce probabilistic lexicase (plexicase) selection, which samples parents from an approximate probability distribution that estimates the probability that an individual would be selected by lexicase selection. The approximation is necessary because computing the exact probabilities is an NP-hard problem [70]. The approximation method is based on the observation that an individual’s selection probability depends on whether it ranks among the best on a given case and on the performance of the other individuals on that same case (see Section III). The advantage of plexicase selection is that once the probability distribution is calculated, parents can be sampled from the distribution directly instead of performing multiple selection events. This significantly reduces the runtime compared to standard lexicase selection.

DALex [162] approximates lexicase selection by assigning random weights to training cases and multiplying an individual’s error vector by these weights before producing an aggregated error. Because the weights are assigned randomly for each selection event, the intuition behind lexicase selection is preserved, as more information from an individual’s error vector is retained than with traditional aggregated fitness measures. The standard deviation of the weight distribution can be adjusted to easily approximate relaxed variants of lexicase selection. The performance of DALex is comparable to standard lexicase selection across several domains while being significantly faster.

TABLE III
REFERENCES BY DIFFERENT APPLICATION DOMAINS IN CHRONOLOGICAL ORDER

Application Domain	References	#
Program Synthesis	[2], [19], [20], [6], [173], [22], [14], [23], [7], [24], [8], [25], [12], [26], [28], [31], [174], [175], [13], [176], [177], [178], [179], [180], [148], [42], [32], [95], [181], [35], [182], [36], [37], [38], [39], [183], [40], [184], [41], [149], [123], [151], [43], [185], [9], [186], [187], [124], [47], [125], [188], [48], [1], [10], [189], [190], [191], [192], [52], [51], [193], [50], [194], [122], [128], [155], [195], [196], [129], [130], [131], [156], [55], [197], [198], [158], [199], [132], [159], [200], [113], [134], [201], [202], [203], [162], [138], [204], [139], [137], [140], [141], [205], [206], [207], [208], [209], [210], [85], [81], [211], [212], [144], [213], [214], [215]	106
Symbolic Regression	[21], [216], [15], [89], [217], [27], [30], [218], [177], [219], [147], [94], [95], [35], [96], [150], [44], [220], [221], [222], [223], [98], [152], [99], [100], [101], [224], [189], [225], [103], [104], [106], [107], [108], [226], [227], [228], [229], [230], [197], [231], [232], [233], [234], [235], [112], [64], [159], [133], [134], [236], [237], [160], [114], [162], [136], [116], [238], [117], [207], [239], [240], [241], [242], [243], [244], [245], [246], [247], [248], [249], [250], [251], [142], [143], [120], [118], [252], [253], [254], [255], [256], [119], [257], [258], [259], [260], [261], [262], [263], [264], [265], [266], [267], [268], [269], [270], [271], [272], [273], [274]	101
Classification	[275], [29], [30], [90], [177], [17], [46], [153], [276], [107], [49], [154], [157], [230], [62], [60], [277], [278], [231], [18], [156], [158], [112], [279], [66], [161], [162], [74], [78], [115], [163], [280], [281], [87], [121], [282], [283]	37
Boolean Logic/ Hardware Circuit Design	[22], [45], [284], [192], [285], [157], [286], [231], [287], [288], [63], [161], [78], [80], [82]	15
Robot Control	[146], [16], [91], [92], [97], [102], [289], [126], [127], [110], [135], [290], [73], [145]	14
Other/ Not Applicable	[93], [33], [34], [44], [105], [291], [109], [292], [53], [293], [61], [54], [56], [57], [58], [59], [294], [111], [67], [65], [295], [68], [69], [296], [297], [70], [72], [75], [71], [77], [298], [299], [300], [76], [79], [301], [302], [303], [304], [305], [84], [306], [86], [307], [308], [309], [310], [311], [83], [88], [312]	51

VIII. APPLICATION DOMAINS

Early applications of lexicase selection were mainly in the domains of program synthesis [6] and symbolic regression [15]. From 2016 onward, there was a noticeable expansion in both the number of papers and the variety of applications. We identified five major application domains and categorized all in-scope publications accordingly in Table III. Program synthesis and symbolic regression are still the most prominent categories with 106 and 101 publications, respectively, followed by classification with 37. Boolean logic and hardware circuit design, along with robot control, represent more niche application areas, with 15 and 14 publications, respectively. This diversification has intensified over the past two years: nearly half of the papers grouped in the category “others” were published in 2024 and 2025.

For program synthesis, lexicase selection significantly increased problem-solving success [6], [14], potentially due to its ability to preserve diversity [7], [8] and to maintain specialists [9], [10], [36]. In the domain of symbolic regression, ϵ -lexicase selection showed superior performance compared to traditional selection methods [15], [44]. In learning classifier systems, a batched lexicase variant has been found to be more robust to data errors, thereby improving the generalization in classification problems [17]. Furthermore, lexicase selection has been successfully applied to Boolean logic problems using genetic programming [22] and Boolean constraint satisfaction problems using genetic algorithms, where it found more valid solutions than traditional selection methods [45]. This is likely due to the ability of lexicase to preserve a diverse set of partial solutions that each satisfy specific constraints, rather than relying on an aggregated fitness score [45]. In evolutionary robotics, lexicase selection with down-sampling has been

shown to outperform specialized strategies [16], possibly due to its effectiveness in preserving high population diversity [92].

Other lexicase applications are combinatorial optimization [79], [311], generative art [65], [76], [86], and dynamic job scheduling [67], [295].

Furthermore, lexicase selection has been analyzed in areas related to machine learning. These include evolving artificial neural networks [275], [290], optimizing deep neural networks [18], [156], [158], clustering [291], [292], [294], computer vision [113], feature representation [96], feature construction [64], [66], [160], [232]–[234], [237], [244], [247], [256], [260], [262], [263], [272], feature selection [46], [89], [90], [264], [279], and even evolving machine learning pipelines [112].

Several papers address multi-objective optimization [34], [72], [75], [77], [93], [111], [297], [300], [310], as lexicase selection can be interpreted as a multi-objective strategy. La Cava et al. [44] found that lexicase selection only selects individuals located on the boundaries of the Pareto front with respect to their performance on training cases (which can be viewed as different objectives). Thus, lexicase selection performs well for multi-objective problems, while also being significantly faster than traditional approaches [93]. Shahbandegan and Dolson [72] also observed this behavior for problems with many contradictory objectives.

This overview shows that lexicase selection is beneficial in a variety of problem domains, particularly when dealing with modular problem structures, where it is important to preserve specialists for behavioral niches [2]. For discrete problems, such as in program synthesis or Boolean logic, the standard form of lexicase selection is well-suited [6]. In contrast, for continuous domains such as symbolic regression, ϵ -lexicase

selection is generally recommended [15]. If overfitting is a concern, batched variants of lexicase selection offer improved generalization [122]. Additionally, down-sampling strategies are often beneficial [123], [138], especially if evaluations are very expensive, as is often the case in robot control tasks [145]. If computational efficiency is very important, approximations may be preferable [159], [162]. Lastly, for specific requirements such as bloat control or novelty promotion, hybrid approaches exist that combine the advantages of lexicase selection with those of more traditional selection methods [149], [151], [160].

IX. CHALLENGES, OPEN QUESTIONS, AND BEYOND

Our literature review demonstrates that lexicase selection has proven successful across many application domains largely because it maintains high population diversity [7], [8] and effectively promotes and preserves specialists [9], [10], [36], which can serve as stepping stones during search [11]–[13].

Besides its many advantages, the original formulation of lexicase selection suffers from high computational costs [44]. This challenge has been approached by several approximation strategies. In particular, DALex [162] significantly reduces runtime and even enables GPU-based implementations of lexicase selection. In addition, down-sampling techniques [123], [138] reduce the number of evaluations per generation, thus saving runtime while maintaining or even improving performance [142].

Furthermore, it has been observed that lexicase selection sometimes generalizes poorly to unseen test cases [52], [125]. Evaluating solutions on batches of training cases rather than single cases mitigates this shortcoming and has been shown to improve generalization [17], [122]. In symbolic regression, down-sampling techniques have also helped to improve generalization [142].

This demonstrates that many of the original limitations of lexicase selection have been effectively mitigated, making it a compelling option for a wide range of problems. Despite these advancements, lexicase selection has not yet become the default selection method in genetic programming.

We identified four key challenges that, if addressed, could significantly advance the field and promote the broader adoption of lexicase selection in the community.

- 1) We strongly recommend integrating lexicase selection and especially new lexicase variants into genetic programming frameworks to make them more accessible to researchers and practitioners.
- 2) Many new variants introduce additional parameters that require costly hyperparameter tuning and consequently raise the barrier to adoption for practitioners. Introducing adaptive parameters, for example, a dynamic adjustment of parameters based on convergence speed, would reduce the need for manual tuning and make new methods more accessible.
- 3) Despite its empirical success, lexicase selection variants lack a strong theoretical foundation. Only a few theoretical studies have been published so far [44], [69]–[71], leaving our understanding of lexicase variants

incomplete. A deeper theoretical grounding would not only improve our understanding of their behavior but also enable a more systematic development of new variants.

- 4) The growing variety of lexicase variants is currently difficult to compare. Most studies evaluate new methods on a single problem domain and a limited set of benchmarks, making it difficult to assess their relative strengths and weaknesses. To remedy this, we advocate for a unified evaluation framework that includes all relevant lexicase variants and benchmark problems for domains such as symbolic regression and classification from SRBench [103] and PMLB [313], program synthesis tasks from PSB1 [14] and PSB2 [194], as well as problems from other domains. Such a framework would allow researchers to compare methodological advances rather than implementation-specific differences across domains.

Addressing these challenges would not only deepen our understanding of lexicase selection, but also accelerate its adoption as a standard selection method in genetic programming.

Beyond genetic programming, the properties of lexicase selection are also interesting when using large language models as variation operators in evolutionary search [314]–[318]. As the output diversity of large language models is often limited [206], [319], evolutionary search converges too early and tends to generate mediocre rather than rare but high-quality outputs [320]. This poses a problem because large language models need to be used in an open-ended manner to constantly produce new and interesting outputs to make novel discoveries [321]. This diversity problem has been successfully approached using Quality-Diversity methods [320], [322]. Nevertheless, lexicase selection remains underexplored in this domain, despite its potential to serve as a general diversity-preserving optimization algorithm, making it a promising direction for future work.

X. CONCLUSIONS

Lexicase selection led to significant performance improvements across a variety of application domains due to its ability to preserve diversity and select specialists. This success inspired the development of variants that address original limitations of lexicase, thereby expanding its applicability.

Despite these advances, we observed that many researchers use outdated versions of lexicase or continue to rely on traditional selection methods. To address this problem, we provided the first comprehensive survey of the field to consolidate current research.

We identified five major research streams that emerged in recent years and discussed their contributions. Further, we identified the most relevant application domains, ranging from program synthesis and symbolic regression to classification, Boolean logic, and robot control. Based on these findings, we identified critical challenges for future work such as the need for a stronger theoretical foundation, a unified evaluation framework for developing new methods, and the integration of lexicase variants in existing frameworks to accelerate the

adoption of lexibase selection in genetic programming. Lastly, we identified several promising research directions, including the use of lexibase selection beyond genetic programming.

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