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6 trained youth soccer players aged 14 to 19 years.

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34 trained youth soccer players aged 14 to 19 years.

35

36 **Abstract**

37 Purpose: We aimed to identify distinct clusters of developmental trajectories in aerobic
38 endurance performance among highly trained youth soccer players aged 14 to 19 years,
39 moving beyond traditional group-level analyses to characterize potential heterogeneity of
40 individual developmental pathways.

41 Methods: Mixed-longitudinal data from 149 male youth soccer players from a professional
42 German academy were analyzed. Aerobic endurance was assessed using velocity at
43 individual anaerobic threshold (v_{IANT}) and velocity at 4 mmol.L⁻¹ blood lactate (v_4) from
44 incremental treadmill tests. A hybrid clustering framework combining Generalized Additive
45 Models (GAM) with model-based clustering (Mclust) was employed. Individual trajectories
46 were estimated using GAMs with factor smooth basis functions, followed by comprehensive
47 feature extraction (15 trajectory characteristics). Growth mixture models were applied across
48 3–7 clusters with multiple covariance structures, with model selection based on weighted
49 consensus scoring emphasizing BIC, silhouette coefficient, and AIC.

50 Results: Four distinct developmental trajectory clusters were identified with consistent patterns
51 across both v_{IANT} and v_4 . The overall population mean trajectory demonstrated minimal
52 systematic change across the age range. However, substantial between-player heterogeneity
53 was evident: approximately one-fifth of players showed substantial improvement followed by
54 stabilization, one-quarter exhibited progressive decline, while the majority displayed minimal
55 change patterns with either slight improvement or slight decline. Endurance performance
56 converged with increasing age across all clusters.

57 Conclusion: Aerobic endurance development in highly trained youth soccer players follows
58 distinct individual trajectories rather than uniform patterns during adolescence. These findings
59 highlight that group-level analyses mask meaningful individual differences, emphasizing the
60 importance of individualized talent evaluation and training prescription.

61

62 **Key Words:** adolescence, fitness, development, cluster

63

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68 Introduction

69 Monitoring developmental trajectories of physical performance of youth soccer players plays an
70 important role to optimally evaluate a player's current and potential future physical performance and
71 ultimately inform talent selection decisions (i.e., progression to the next age group within a
72 developmental program). Recent reviews have emphasised the need for longitudinal studies within
73 the realm of applied talent research which advances our understanding of how youth soccer players
74 naturally develop during adolescence ^{1,2}. The development of physical performance rarely follows a
75 linear pattern from early adolescence (i.e., 12 years of age) through to adulthood (i.e., 20 years of
76 age) and is influenced by a multitude of factors such as training history and biological maturation
77 that drives physiological and morphological changes ³. Without understanding these individual
78 trajectories, selection and evaluation processes risk misinterpreting temporary performance
79 differences as stable characteristics. There is therefore a pressing need to characterize the individual
80 developmental trajectories that underlie physical performance development in youth soccer players.

81 Endurance performance is one of several critical components of physical performance in soccer and
82 is associated with match running output in youth soccer players ⁴. Several longitudinal observational
83 studies have examined the development of physical performance, including endurance performance,
84 in youth soccer players, but these investigations have predominantly focused on average
85 developmental trajectories ⁵ or binary comparisons between selected vs. de-selected ⁶ or future
86 professional vs. non-professional players ⁷. While these studies provide valuable insights into
87 general developmental patterns, such broad classifications may obscure the heterogeneity of
88 individual developmental pathways that exist during adolescence. At the group level, endurance
89 performance, predominantly assessed using field-based shuttle running tests, has been shown to
90 develop throughout the entire adolescent period, with longitudinal studies demonstrating consistent
91 increases across all chronological ages and biological maturation stages ^{5,7,8}. However, despite
92 these well-established findings at the group level, empirical data regarding individual developmental
93 trajectories in youth soccer players remain limited. Moreover, laboratory-based assessments of
94 isolated aerobic capacity (e.g., lactate threshold measures) may reveal different developmental
95 patterns compared to field-based shuttle tests that encompass multiple physiological and
96 neuromuscular components. To advance talent evaluation and selection practices, there is a need
97 to move towards identifying clusters of developmental trajectories in endurance performance that
98 capture the true diversity of development in highly trained youth soccer players. This approach would
99 provide more nuanced insights into the prevalence and magnitude of endurance development during
100 adolescence.

101 Therefore, the aim of the current study was to identify clusters of developmental trajectories of
102 aerobic endurance performance in highly trained youth soccer players aged 14 to 19 years.

103 **Methods**

104 **Participants**

105 The sample included aerobic endurance performance data available for N = 149 male academy
106 soccer players and n = 1380 datapoints, respectively (chronological age range: 14.0 to 19.0 years),
107 over a nine-year period (summer 2016 to summer 2025). Players belonged to the U14, U15, U16,
108 U17, and U19 age group of one professional German youth academy and can be classified as tier 3
109 athletes (highly trained/national Level) competing at the highest national level within their respective
110 age group competitions ⁹. Upon enrolment, parents/guardians signed contracts providing consent
111 confirming that data arising as a condition of regular player monitoring procedures can be used for
112 research purposes. Ethical approval was granted by the ethics committee of [blinded for peer-review]
113 (ref.no. 24-10).

114 We acknowledge that no formal sample size calculation was performed for this study. While sample
115 size calculation for clustering approaches has traditionally been challenging due to the exploratory
116 nature of these methods, recent research has provided more systematic guidance for these analyses
117 ¹⁰. Sample size requirements for Mixture Models depend on multiple factors including the degree of
118 separation between latent classes, the relative size of the smallest clusters, the number of measured
119 variables, and the specific clustering method employed ¹¹. Given the exploratory nature of our
120 functional mixture modelling approach, we therefore employed a comprehensive model diagnostic
121 process to ensure the validity and interpretability of the clustering solution, including a comparison
122 with a traditional parametric growth mixture modelling approach.

123

124 **Study design**

125 Retrospective mixed-longitudinal observational study design.

126

127 **Procedures**

128 Players performed indoor an incremental test on a Woodway treadmill (Woodway GmbH, Weil am
129 Rhein, Germany) to determine lactate-based thresholds ^{6,12}. The slope of the treadmill was set at
130 1% incline. Running velocity started at 6 kmh⁻¹ and increased by 2 kmh⁻¹ with each stage lasting 3
131 min. Between stages, a 30-second passive recovery phase was implemented allowing for capillary
132 blood collection from the earlobe. Players received instructions to perform the test until exhaustion.
133 Blood lactate concentration analysis for each stage was performed with the Biosen C-Line Sport
134 system (EKF-diagnostics GmbH, Barleben, Germany). The Ergonizer Software (K. Roecker,
135 Freiburg, Germany) was used to compute the following outcome measures: i) velocity at the
136 individual anaerobic threshold (velocity at which blood lactate concentration was 1.5 mmol.L⁻¹ above

137 the individual aerobic threshold, determined as the velocity at which blood lactate concentration
138 begins to rise above baseline levels, v_{IANT}) and ii) velocity at the absolute lactate concentration of
139 4 mmol.L^{-1} (v_4). We acknowledge that several other threshold concepts exist and aimed to describe
140 two fundamentally different ones here: an absolute threshold based on a fixed blood lactate
141 concentration (v_4) and an individual threshold accounting for the athlete's personal lactate kinetics
142 (v_{IANT})¹³.

143

144 **Statistical analyses**

145 Statistical analyses were conducted in RStudio (version 1.2.5033). A detailed description of the
146 entire clustering methodology can be found in the Supplementary File (Supplementary 1.
147 Comprehensive clustering methodology).

148 *Data preparation*

149 Player inclusion required adequate longitudinal data coverage whereby players required data
150 spanning at least 3 distinct ages, a minimum 2-year temporal span, and at least 3 observations. We
151 implemented a hybrid-based clustering framework combining Generalized Additive Models with
152 model-based clustering (GAM + Mclust), with Latent Class Mixed Models (LCMM) computed as a
153 comparative methodological approach detailed in the Supplementary File (Supplementary 2. Results
154 from the Latent Class Mixed Models (LCMM) clustering approach).

155 *Step 1: Individual Trajectory Estimation*

156 Generalized Additive Models were implemented using the mgcv package (version 1.9-1) with factor
157 smooth basis functions to estimate player-specific developmental trajectories for both outcome
158 measures v_{IANT} and v_4 across chronological age. Optimal basis dimensions ($k = 3-6$) were
159 determined through multi-criteria evaluation incorporating deviance explained, information criteria
160 (BIC, AIC), and effective degrees of freedom. Based on this systematic assessment, $k = 3$ was
161 selected as optimal for v_{IANT} and $k = 4$ for v_4 , representing the best balance between model fit and
162 computational stability.

163

164 *Step 2: Feature-Based Growth Mixture Modelling*

165 Individual predicted trajectories were systematically generated across age intervals from 14 to 19
166 years with 0.1-year increments. Comprehensive feature extraction yielded 15 trajectory
167 characteristics including endpoints, developmental rates, curvature indices, and variability metrics.
168 All features were standardized to ensure equal contribution to the clustering process. Growth mixture
169 models were computed using the mclust package (version 6.1.1) across 3-7 clusters and 8
170 covariance structures representing different assumptions about cluster geometry. Model selection

171 employed weighted consensus scoring emphasizing BIC (30%), silhouette coefficient (30%), and
172 AIC (25%) over traditional distance-based metrics (Calinski-Harabasz: 10%, Dunn: 2%, Davies-
173 Bouldin: 3%), reflecting the importance of both statistical model adequacy and meaningful cluster
174 distinction. We identified a 5-cluster solution with spherical, equal-volume covariance structure as
175 demonstrating optimal performance.

176 Individual predicted trajectories were systematically generated across age intervals from 14 to 19
177 years with 0.1-year increments. Comprehensive feature extraction yielded 15 trajectory
178 characteristics including performance levels, developmental rates, variability metrics, timing
179 features, and curvature indices. All features were standardized to ensure equal contribution to the
180 clustering process. Model-based clustering was performed using the mclust package (version 6.1.1)
181 across 3-7 clusters and multiple covariance structures (i.e., EVI, VEI, VVI). Model selection
182 employed weighted consensus scoring emphasizing BIC (30%), silhouette coefficient (30%), and
183 AIC (25%) over traditional distance-based metrics (Calinski-Harabasz: 10%, Dunn: 2%, Davies-
184 Bouldin: 3%). The optimal solution for both vIANT and v4 identified 4 clusters with the VEI covariance
185 structure.

186

187 *Model Diagnostics*

188 GAM model validation included residual analysis for homoscedasticity assessment, observed vs.
189 predicted value evaluation, and person-level model fit quality distribution. Mclust clustering validation
190 encompassed posterior probability analysis for assignment certainty, silhouette analysis for cluster
191 separation quality, bootstrap stability assessment using Adjusted Rand Index across 30-50
192 resampling iterations, distance-based validation metrics, entropy-based separation analysis, and
193 systematic outlier detection using Mahalanobis distances. A comprehensive overview of the
194 clustering model diagnostic results can be found in the Supplementary File (Supplementary 3.
195 Comprehensive model diagnostics for vIANT of both clustering approaches (Hybrid approach (GAM
196 + Mclust) and Latent Class Mixed Model (LCMM)) and Supplementary 4. Comprehensive model
197 diagnostics for v4 of both clustering approaches (Hybrid approach (GAM + Mclust) and Latent Class
198 Mixed Model (LCMM))).

199

200 Reference overall mean trajectory estimation

201 To estimate the overall mean trajectory of the entire sample, we fitted again a generalized additive
202 model using the mgcv package (version 1.9-1) with restricted maximum likelihood (REML).
203 Chronological age served as the independent variable and both outcome measures vIANT and v4
204 as the dependent variable. Random intercepts for player ID were modelled as ridge-penalized
205 smooths to account for repeated measures within individuals. Optimal basis functions (tp, cr, cs, ps)

206 and smoothness parameters ($k = 3$ to 6) were determined by minimising the Akaike Information
207 Criterion (AIC). Population-level predictions were obtained by excluding the random effect term,
208 providing the marginal mean trajectory with correct confidence intervals. Detailed model information
209 is provided in the Supplementary File (Supplementary 5. Comparative quality assessment across
210 both outcome measures).

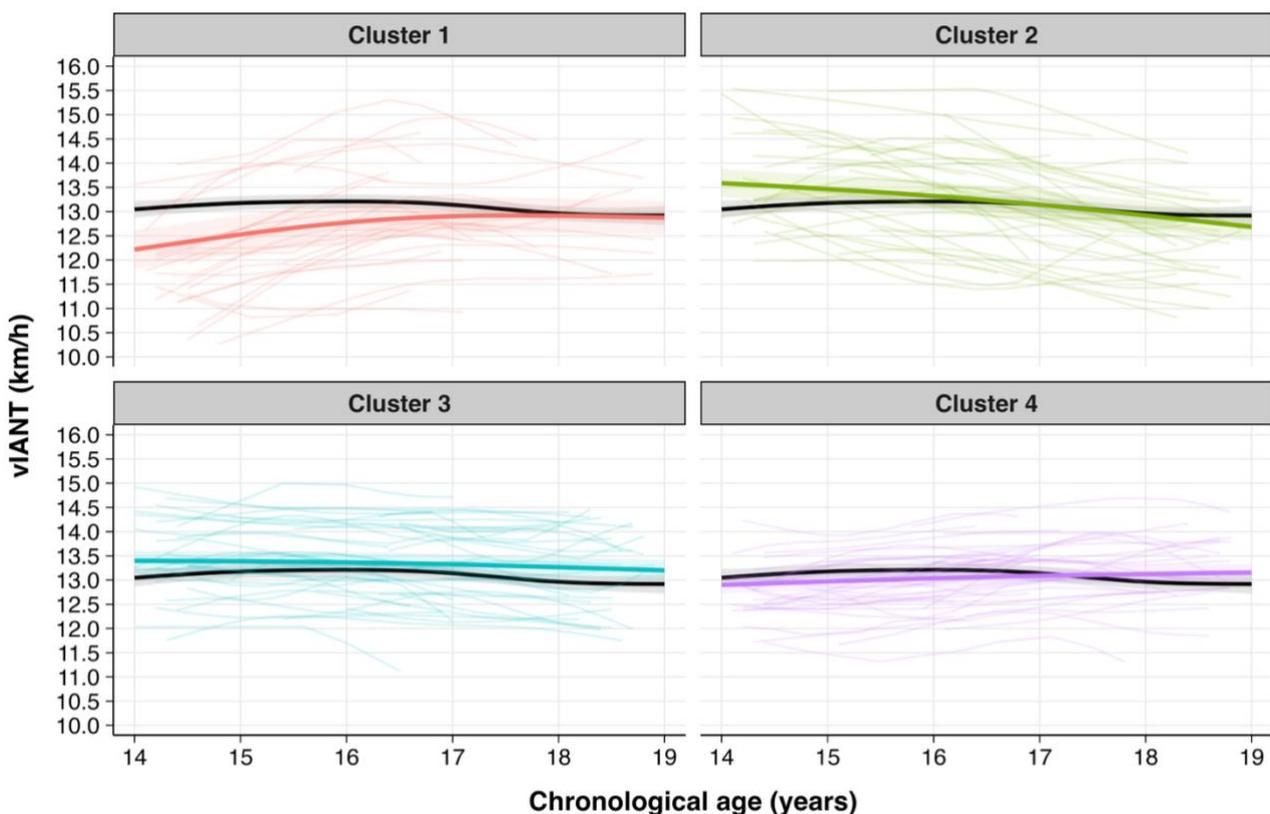
211 Results

212 Four distinct clusters of developmental trajectories were identified for both outcome measures using
213 the two-step hybrid framework combining individual GAM trajectory estimation followed by feature-
214 based clustering. Figures 1 and 2 illustrate the individual and GAM model-predicted cluster mean
215 developmental trajectories together with the mean trajectory of the entire sample as reference for
216 vIANT and v4, respectively.

217 For vIANS, the clusters can be characterised as: substantial improvement then stabilization (cluster
218 1, n = 26, 17%), progressive decline (cluster 2, n = 36, 24%), minimal decline (cluster 3, n = 45,
219 30%), and minimal improvement (cluster 4, n = 42, 28%) trajectories.

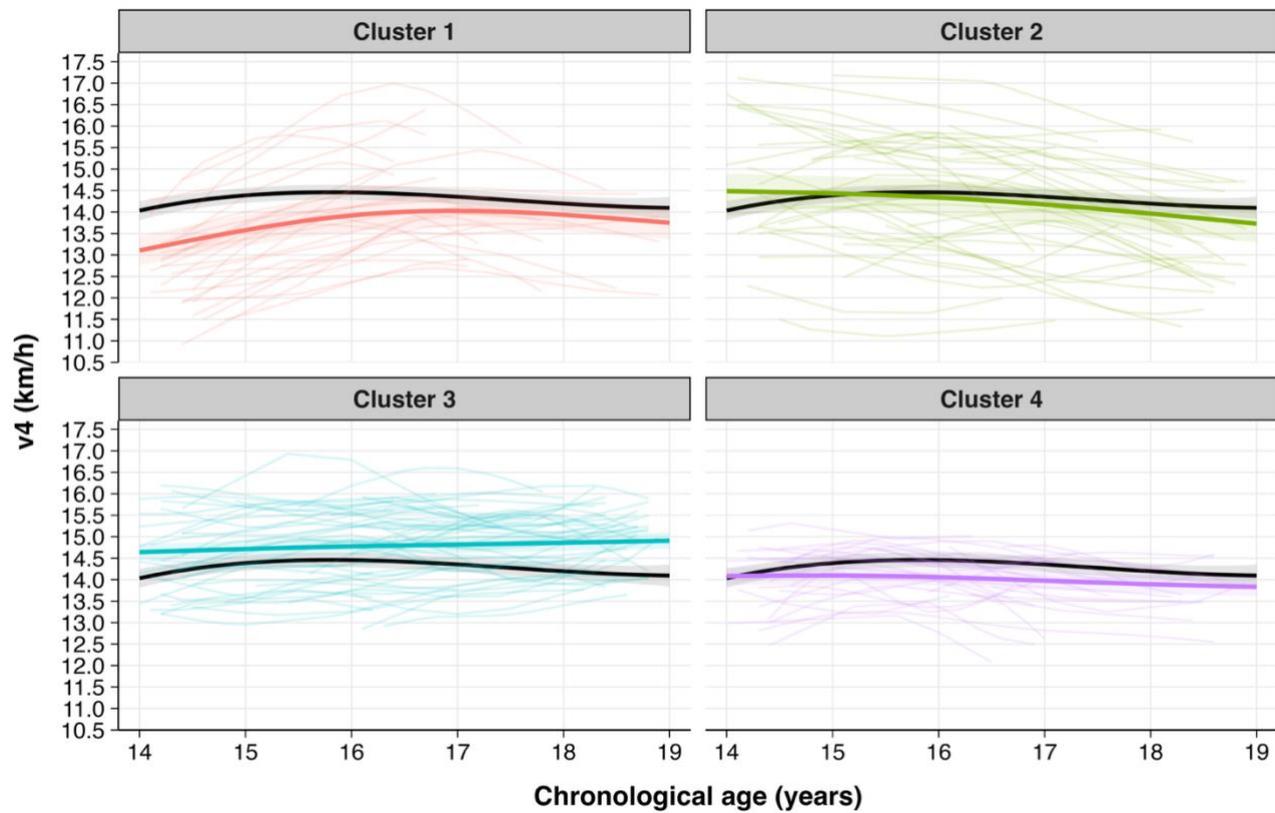
220 For v4, similar trajectory patterns emerged: substantial improvement then stabilization (cluster 1, n
221 = 26, 17%), progressive decline (cluster 2, n = 36, 24%), minimal improvement (cluster 3, n = 58,
222 39%), and minimal decline (cluster 4, n = 29, 19%) trajectories.

223



224

225 **Figure 1.** Distinct trajectory clusters for the outcome measure *velocity at the individual anaerobic*
226 *threshold (vIANT)* derived from the two-step hybrid approach combining individual GAM trajectory
227 estimation followed by feature-based clustering. Each cluster shows individual trajectories and GAM
228 model-predicted cluster means with model-based 95% confidence intervals. The black line
229 represents the mean trajectory of the entire sample as reference with 95% confidence intervals.



230

231 **Figure 2.** Distinct trajectory clusters for the outcome measure *velocity at lactate concentration of 4*
232 *mmol.L⁻¹ (v4)* derived from the two-step hybrid approach combining individual GAM trajectory
233 estimation followed by feature-based clustering, showing individual trajectories and GAM model-
234 predicted cluster means with model-based 95% confidence intervals. The black line represents the
235 mean trajectory of the entire sample as reference with 95% confidence intervals.

236 Discussion

237 In this study, we identified distinct clusters of developmental trajectories of aerobic endurance
238 performance using mixed-longitudinal data from highly trained youth soccer players aged 14 to 19
239 years. The overall mean trajectory demonstrated minimal systematic change across the age range,
240 with aerobic endurance performance remaining relatively stable between 14 and 19 years. However,
241 this average pattern masks considerable individual variation. Across both outcome measures vIANT
242 and v4, approximately one-fifth of players demonstrated substantial improvement followed by
243 stabilisation, one-quarter showed progressive decline, while the majority exhibited minimal change
244 patterns with either slight improvement or slight decline. These findings provide exploratory evidence
245 for the heterogeneity of individual developmental trajectories and challenge the assumption of
246 consistent improvement of aerobic endurance performance during adolescence.

247 The heterogeneous trajectory patterns likely reflect the complex interplay of training history,
248 accumulated training load, and individual factors during adolescence ³. We observed similar cluster
249 patterns between vIANT and v4 which can be explained by their strong correlation (repeated
250 measures correlation = 0.85, 95% CI: 0.83-0.86), highlighting that both outcome measures represent
251 the same construct of aerobic endurance performance. We observed four distinct trajectory patterns
252 which may be explained by multiple developmental factors during adolescence, including
253 asynchronous development of aerobic and anaerobic endurance ³, varying training histories and
254 accumulated loads, and systematic shifts in training emphasis toward anaerobic power and
255 neuromuscular qualities as players approach maturity. For example, cluster 1 ("substantial
256 improvement then stabilisation") likely reflects predominant aerobic development until 16-17 years,
257 followed by physiological transition toward anaerobic development ¹⁴. Conversely, cluster 2
258 ("progressive decline") challenges previous observations of continuous development of endurance
259 performance during adolescence ³, potentially indicating prioritisation shifts toward anaerobic and
260 neuromuscular adaptations at the expense of aerobic endurance ¹⁵.

261 Despite the stable overall mean trajectory, substantial between-player heterogeneity highlights that
262 group-level analyses mask meaningful individual differences in aerobic endurance development.
263 The convergence of performance levels with age (i.e., narrowing the spread of cluster means)
264 indicates that the accumulation of systematic training reduces initial disparities. Players with superior
265 aerobic endurance may accept slight declines while potentially focusing on other physical qualities
266 (e.g., speed and power), whereas players with poorer aerobic endurance performance improve
267 toward sufficient levels. These findings underscore the importance of focusing on individual
268 developmental trajectories rather than average development curves for talent evaluation and training
269 prescription. Previous research has identified conflicting evidence for potential critical periods in
270 aerobic endurance development (12-14 vs 14-16 years) ¹⁶, and recent longitudinal studies

271 emphasise individual variation in both developmental timing and trainability of aerobic fitness ¹⁴,
272 supporting our findings of distinct developmental trajectories rather than universal critical periods.

273 Our two-step hybrid framework advances beyond traditional trajectory approaches ⁵ by capturing
274 individual heterogeneity that group-level classifications ^{6,7} typically miss. The GAM modelling
275 approach accounts for non-linear developmental patterns, addressing limitations of studies focusing
276 solely on group-level trends ⁸. Our findings contrast with previous literature reporting consistent
277 increases in endurance performance throughout adolescence ^{5,7,8}. This discrepancy may be
278 attributed to test-specificity: our laboratory-based aerobic threshold measures (vIANT, v4) isolate
279 aerobic capacity, whereas commonly used field-based tests such as the Yo-Yo Intermittent Recovery
280 Test assess a broader construct encompassing aerobic, anaerobic, and neuromuscular qualities
281 ^{13,17}. This methodological difference may explain why we observed minimal average development in
282 pure aerobic capacity whilst studies that included endurance performance tests encompassing
283 multidimensional constructs demonstrated improvements throughout adolescence. This aligns with
284 calls for longitudinal talent research ^{1,2} and offers practitioners a more nuanced understanding of
285 individual developmental trajectories.

286

287 *Limitations*

288 Despite the novelty and practical applicability of our trajectory clustering approach for understanding
289 aerobic endurance development, several limitations should be acknowledged. First, we did not
290 assess biological maturation status, though previous research suggests minimal associations
291 between skeletal maturation and aerobic performance in youth soccer players ¹⁸. Second, our
292 findings are derived from a single professional academy, which may limit generalizability to other
293 contexts and playing levels. Third, the relatively small sample sizes within some clusters require
294 cautious interpretation of cluster-specific patterns. Fourth, our analysis was restricted to
295 chronological ages 14 to 19 years, not encompassing the entire adolescent period and adulthood.
296 Finally, we focused exclusively on aerobic endurance without considering other physical
297 performance qualities (e.g., strength, speed, power), which prevents holistic player profiling that
298 might further explain the observed developmental trajectories. Future research should address these
299 limitations through multi-site studies incorporating early adolescence and adulthood, larger samples,
300 and comprehensive physical performance assessments.

301

302 **Practical Applications**

303 Our findings have immediate practical implications for talent development programs in youth soccer.
304 To translate these findings into practice, we developed an interactive web-based application

305 (<https://rufludwig.shinyapps.io/endurance-clustering/>) that allows practitioners to input longitudinal
306 aerobic endurance performance data (minimum 3 data points of a player) to receive cluster
307 membership probabilities with trajectory visualisations. Understanding a player's individual
308 developmental trajectory enables practitioners to make better informed evaluations of current and
309 potential future aerobic endurance performance, ultimately supporting talent selection decisions.

310

311 **Conclusion**

312 This study demonstrates that aerobic endurance development in highly-trained youth soccer players
313 follow distinct individual trajectories rather than uniform patterns during adolescence. While the
314 overall mean trajectory remained relatively stable between 14 and 19 years, substantial between-
315 player heterogeneity was evident, with approximately one-fifth demonstrating substantial
316 improvement, one-quarter showing progressive decline, and the majority exhibiting minimal change
317 patterns. These results offer nuanced insights for more individualized talent evaluation and selection
318 practices within highly-trained youth soccer players.

319

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323

324 **Disclosure statement**

325 The authors declare no conflict of interest involved with the present study.

326

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- 382

383 **Figure Legend**

384 **Figure 1.** Distinct trajectory clusters for the outcome measure *velocity at the individual anaerobic*
385 *threshold (vIANS)* derived from the two-step hybrid framework combining individual GAM trajectory
386 estimation followed by feature-based clustering. Each cluster shows individual trajectories and GAM
387 model-predicted cluster means with model-based 95% confidence intervals.

388 **Figure 2.** Distinct trajectory clusters for the outcome measure *velocity at lactate concentration of 4*
389 *mmol.L⁻¹ (v4)* derived from the two-step hybrid framework combining individual GAM trajectory
390 estimation followed by feature-based clustering, showing individual trajectories and GAM model-
391 predicted cluster means with model-based 95% confidence intervals.