# NHL Aging Curves using Functional Principal Component Analysis

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#### Abstract

When considering future performance in sport, age is an important feature for prediction models. On average, players tend to improve from their rookie (earliest) season, plateau, and then decline in performance until they retire from the league. In this paper we apply Functional Principal Component Analysis to the careers of players from the National Hockey League in order to construct individual aging curves. The approach is nonparametric in the sense that a parametric structure is not imposed on the aging curves. A main aspect of our work is the consideration of selection bias.

**Keywords** : Aging curves, Functional data analysis, National Hockey League, Principal component analysis, Sports analytics.

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## **1** INTRODUCTION

Many decisions made by front office staff of major sports clubs involve projecting the future performance of players. In the case of drafting younger players, teams attempt to predict a player's future performance given their abilities relative to their peers in junior leagues. However, decisions involving trades and free-agency are different from drafting decisions. In these cases, a team must forecast the performance of players who have already been entrenched in the league, and this is subject to the constraint that the number of players on team rosters is fixed. The composition of players is constantly changing as older players retire and younger players take on more prominent roles. In fantasy sports, the relationship between performance and age is also greatly discussed (Cockcroft 2023). Therefore, it is of vital interest to predict the future values of players as they age. The results of such prediction models are referred to as aging curves. Aging curves can be difficult to construct in team sports due to the fact that player performance is highly dependent on teammates and the number of minutes played.

Aging curves have been studied in many sports including: golf (Berry, Reese and Larkey 1999), football (Young and Weckman 2008), baseball (Bradbury 2009), soccer (Swartz, Arce and Parameswaran 2013), hockey (Brander, Egan and Yeung 2014), tennis (Mlakar and Tušar 2015), cricket (Saikia, Bhattacharjee and Mukherjee 2019), basketball (Wakim and Jin 2014) and snooker (McHale 2023).

The effect of aging on the body is a common issue for all athletes (Distefano and Goodpaster 2018). For many sports, players reach their peak performance before the age of 30 years, and then generally decline as they age due to decreasing athleticism and increased injury risk. The observance of this "peak" in performance has been studied, for example, by Dendir (2016) and Bradbury (2009). Bradbury (2009) demonstrates that different skills decline in baseball at different rates; baserunning, for example, declines at a much faster rate than power. Most research reaches the general consensus that age effects are dependent on position. For example, Brander, Egan and Yeung (2014) conclude that the peak age for forwards in the National Hockey League (NHL) is between ages 27-28, while the peak age for defensemen is between 28-29. Each player has a unique aging curve (due to different body composition or previous athletic history, for example). However, it seems reasonable that there should be some agreement in age curves between players.

In the literature, there are two prominent general approaches for the construction of aging curves; (1) the so-called "Delta method" and (2) regression methods. The Delta method calculates the difference between a player's performance between years, and then averages these differences over players. A feature of the Delta method is that it is not impacted by the differences in quality between players. There are variations to the Delta method; see for example, Lichtman (2009) and EvolvingWild (2017). With regression methods, the response variable of interest (player performance) is regressed against the covariate age. Regression methods may differ in the form of regression and whether additional covariates are considered in the regression model (e.g., a player effect). Early regression methods tended to take a parametric approach. For example, Fair (2008) and Bradbury (2009) imposed quadratic shapes on aging curves in baseball. Villaroel, Mora and Gonzalez-Parra (2011) consider both quadratic and cubic shapes with respect to the performance of triathletes. Although convenient, parametric approaches do not permit freedom in the shape of aging curves. For example, it has been observed in chess (Roring and Charness 2007) that performance improvement leading to peak performance may occur at a steeper incline than the decline in performance following the peak. Many of the more recent regression methods tend to take a nonparametric or semiparametric approach. For example, Turtoro (2019) uses generalized additive models (GAM) for constructing aging curves in the NHL.

Another distinguishing feature in the construction of aging curves is whether or not an approach considers selection bias. Selection bias is a systematic statistical error caused by drawing a non-random sample from a population. In sports, samples of player performances are typically non-random because only the most talented players enter leagues at very early ages. Moreover, these players are usually the same players who stay in the league the longest (except in the case of early career altering injuries). Therefore, gifted players are typically overly represented in both the left and the right tails of age distributions. Schuckers, Lopez and Macdonald (2023) demonstrate the considerable impact of selection bias using data from the NHL. Various approaches have been proposed that consider selection bias in the construction of aging curves. These all involve imputation schemes that "add" data over missing periods. For example, after a player retires, data are imputed for the years following retirement. Lichtman (2009) takes a basic approach where regression methods impute data based on best estimates. Schuckers, Lopez and Macdonald (2023) use more sophisticated imputation schemes based on the simulation of missing data. They introduce thresholds

for player performance at different ages and ensure that simulated data do not breach the threshold. Nguyen and Matthews (2023) take the simulation approach one step further by investigating the cause of missingness, and use different simulation approaches based on the different causes of missingness.

Although there is now a considerable literature on methods developed for the construction of aging curves, there are several innovations in this paper. First, we use an approach based on Functional Principal Component Analysis (FPCA). This is a nonparametric approach which permits flexibility in the shape of growth curves. Second, within the FPCA framework, we account for selection bias by imposing an intuitive constraint on the corresponding likelihood function. Third, and most importantly, we construct aging curves for individual players. This is obviously an important contribution since players age differently. The curves have a predictive component where curves can be extended beyond a player's current age. An important aspect of the FPCA model is that there is an underlying relationship between the individual aging curves, and this enables prediction. Previously, aging curves have only been constructed for the so-called "average" player.

The proposed FPCA approach is predicated on Functional Data Analysis (FDA). In FDA, we use spline basis functions to determine functional relationships. This provides a relationship between performance and age that is essentially nonparametric. In addition, to fitting separated aging curves for each player, an FDA approach can identify clusters from principal component scores, allowing us to readily compare players. FPCA aging curves have been previously considered in the context of basketball by Wakim and Jin (2014) where implementation is based on conditional expectation using the PACE method. However, the PACE approach does not account for the selection bias issue.

In Section 2, we describe the data and the player evaluation metric. In Section 3, we carry out some exploratory data analysis to investigate aging patterns. In Section 4, we outline the methods for constructing an aging curve using FPCA. Here, we briefly outline some of the underlying mathematical background. In Section 5, we present the results of our modelling. We conclude with a short discussion in Section 6.

## 2 DATA

The data in this project was scraped from Sports Reference LLC at https://www.hockeyreference.com. The website contains summary statistics (goals, points, games played, etc) for each player in the NHL during the period 1920-2022. Originally, there were approximately 50,000 rows in the dataset before adjusting for duplicate rows (due to players changing teams mid-season). This led to n = 7,393 unique players with 43,705 player-seasons worth of data. The maximum player age in the dataset is 51 years (Gordie Howe), and the minimum player age is 17 years (Wayne Gretzky). From an alternative study (Diamond 2000), the average career length in the NHL is 4.5 years.

### 2.1 Player Value

As a measure of player value, we use the point share PS statistic (Kubatko 2010). It is a measure derived from the win share metric (James and Henzler 2002) that was originally used to evaluate baseball players. Here, "points" refer to the points a team gains from winning games, and not the points a player gains from scoring a goal or assisting on a goal in hockey. Hence, the metric attempts to credit a player's contribution to their team's success. This metric was chosen because we require a composite measure of performance that adjusts for the quality of linemates. Goals and assists metrics are not composite measures since they are not reflective of contributions to team defense, and would consequently overvalue forwards.

The point share metric has been developed so that a hockey team with 100 team points (e.g., 40 wins and 10 overtime losses) will have players whose individual point shares sum to 100. Players may have a negative point share. Negative point shares indicate that a player is losing team points relative to a replacement level player. Point shares are obtained from both offensive and defensive component point shares. The offensive point share for a player during a season is based on their goals created (a weighted sum of the player's goals and assists divided by team goals and assists), adjusted by the player's minutes and adjusted for the league environment (league goals divided by league points). There are also positional adjustments for forwards and defensemen.

Over time, adjustments have been made to the point share calculation based on various

rule changes. For example, player minutes on ice have only been recorded since the 2000-2001 NHL season. Also, the number of games in the NHL season has increased over the years and the NHL previously permitted tied games, which yielded one point to both teams. Consequently the point share metric is not perfect; however it is still regarded as an advanced statistic in hockey analytics. The point share PS statistic can be obtained from https://www.hockey-reference.com. More details on it's calculation are found in Kubatko (2023).

Alternative metrics for assessing performance include those based on salary considerations (Swartz, Arce and Parameswaran 2013). Also, there are limitations on the availability of advanced match statistics for use in aging curves. For example, the NHL only started to track individual shots and plus-minus statistics in 1960-1961.

## 3 EXPLORATORY DATA ANALYSIS

In Section 5, our FPCA analyses consider forwards and defensemen separately. In this section, we provide some basic exploratory plots that motivate this distinction. It is suggested that forwards and defensemen age differently in the NHL.

Figure 1 is a histogram based on what is called the "participation method" (Brander, Egan and Young 2014). Here, every player-season is taken as an observation where age is the recorded variable based on player age at the end of the season. The idea is that players participate in the NHL during the seasons of their peak performance. For some, this window may be short. Therefore, the modal regions of the histogram indicate peak periods of performance. In the histograms, there are 28,871 observations corresponding to forwards and 14,834 observations corresponding to defensemen. From Figure 1, we observe that the modal age for both positions lies between 23-24 years of age. We also observe that there is greater participation longevity for defenseman than forwards.

We also provide histograms for the ages of the rookie and retirement seasons of NHL players in Figure 2 and Figure 3, respectively. From Figure 2, there is little difference between the time when forwards and defensemen begin their careers. It appears that most players begin their careers around 20-22 years of age. From Figure 3, there is some indication that defensemen end their careers slightly later than forwards. There is much



Figure 1: Histograms of the age of NHL players based on player-seasons from 1920-2022.

greater variability in the retirement age than the rookie age; most players retire before 38 years of age. Of course, "early retirement" may simply be a case of players no longer achieving the standards of NHL play.



Figure 2: Histograms of rookie age by position.



Figure 3: Histograms of retirement season age by position.

## 4 METHODS

### 4.1 Overview of FDA

FDA is a highly flexible statistical framework which is concerned with the modelling of longitudinal data. FDA is a modern approach to multivariate statistical modelling, with many applications as seen in Ramsay and Silverman (2005). A review of the current advances in the topic can be found in Wang, Chiou and Müller (2016). Recently, FDA has been used in the analysis of sports data, as seen in Guan et al. (2022) where it is used to specify conditional distributions for in-game win probabilities in rugby. In Chen and Fan (2018), FDA is used to model the score difference process in basketball. Statistical contributions to sport are highlighted in the handbook by Albert et al. (2017).

FDA is different from well-known related methods. For example, unlike time series analysis, FDA does not impose underlying assumptions regarding stationarity. Also, unlike standard multivariate statistics whose data consist of multiple measurements for each subject, FDA curves are viewed as infinite-dimensional random vectors in a functional space. There is an underlying assumption in FDA that observed samples are independent stochastic processes. FDA is particularly suited to handle sparse data.

FDA can be used to perform a number of common machine learning tasks such as classification, clustering, ANOVA, regression, principal component analysis, interpolation and extrapolation. The enhancement provided by FDA is that point estimators are replaced with functional estimators. Some benefits of FDA include the representation of observed data as smooth functions, dimensionality reduction and the ability to compute derivatives of smoothed estimators. The are numerous packages that can be used to perform FDA; the common ones being *scikit-fda* (Python) and the R package *fda* available on CRAN.

#### 4.2 Spline Regression for FDA

When performing FDA, we consider the observations from each individual arising from a random, smooth function. For example, suppose that we observe data from i = 1, ..., N players, and for the *i*th player, we observe *m* data points  $(y_{i1}, ..., y_{im})$  at time points  $t_1, ..., t_m$ . Then the core assumption in FDA is that

$$y_{ij} = X_i(t_{ij}) + \epsilon_{ij} \tag{1}$$

where the  $\epsilon_{ij}$  are independent and normally distributed with mean  $\theta$  and variance  $\sigma^2$ . Equation (1) explicitly assumes that the observations from individual *i* can be modelled by a single stochastic function  $X_i(t)$  after accounting for measurement error  $\epsilon_{ij}$ . Although we suppress additional notation, we note that the number of observed points *m* can actually vary from player to player. In order to approximate the functions  $X_i(t)$ , we introduce *Q* spline basis functions  $b_1(t), \ldots, b_Q(t)$ . Spline functions are piecewise polynomials joined at specific points, called knots. The number of knots  $\tau$  is determined by  $\tau = Q - d + 1$  where *d* is the degree of the polynomial. We write

$$X_i(t) \approx \sum_{q=1}^Q \alpha_{iq} b_q(t)$$

for sufficiently large Q where the  $\alpha_{iq}$  are the coefficients of the spline basis functions. We estimate  $\alpha_{i1}, \ldots, \alpha_{iQ}$ , by minimizing the least squares loss function

$$\sum_{j=1}^{m} \left( y_{ij} - \sum_{q=1}^{Q} \alpha_{iq} b_q(t) \right)^2 .$$

Once the  $\alpha_{ik}$  have been estimated via  $\hat{\alpha}_{ik}$ , this leads to the estimated aging curve

$$\hat{X}_i(t) = \sum_{q=1}^Q \hat{\alpha}_{iq} b_q(t) \tag{2}$$

for the ith player. We can also compute statistics such as the mean aging curve across all players

$$\hat{\mu}(t) = \frac{\sum_{i=1}^{N} \hat{X}_i(t)}{N}$$

#### 4.3 Functional Principal Component Analysis

In our problem, assume for the time being that the number of observations m is constant across all players, i = 1, ..., N and that  $t_{ik} = t_{jk}$  for all players i, j and all ages k = 1, ..., m. FPCA begins with the estimated covariance function

$$\hat{v}(s,t) = \frac{1}{N} \sum_{i=1}^{N} \left( \hat{X}_i(t) - \hat{\mu}(t) \right) \left( \hat{X}_i(s) - \hat{\mu}(s) \right)$$

which describes the correlation of aging curves at ages s and t, where  $\hat{X}_i(t)$  is the estimated aging curve in (2) corresponding to player *i*.

Our objective is to obtain the orthonormal eigenfunctions  $\hat{\xi}_1(t), \ldots, \hat{\xi}_p(t)$  and the eigenvalues  $\rho_1, \ldots, \rho_p$  by solving the equation

$$\int \hat{v}(s,t)\xi(s) \, ds = \rho\xi(t)$$

where we introduce the dimensionality reduction  $p \leq m$ . The eigenfunctions  $\hat{\xi}_1(t), \ldots, \hat{\xi}_p(t)$  are known as functional principal components. This leads to FPC scores

$$s_{i\ell} = \int \hat{\xi}_{\ell}(t) (\hat{X}_i(t) - \hat{\mu}(t)) dt$$
(3)

for players i = 1, ..., N and the  $\ell$ th functional principal component,  $\ell = 1, ..., p$ .

It is the FPC scores  $s_{i\ell}$  in (3) that readily permit the comparison of players in lower-

dimensional settings. Once we have computed all of the FPCs terms, individual aging curves are expressed by

$$\hat{X}_{i}(t) = \hat{\mu}(t) + \sum_{\ell=1}^{p} s_{i\ell} \hat{\xi}_{\ell}(t) .$$
(4)

The formula (4) follows from the Karhunen–Loève expansion. The truncation is based on the first p principal components that explain most of the variation between curves.

#### 4.4 FPCA using imFunPCA

The conventional FPCA method assumes that missing data are missing at random (MAR). For the aging curves, we know that this is not the case due to selection bias. In Shi et al. (2021), the authors adjust for this bias using a constrained likelihood approach for missing data. For example, assuming that the data  $y_{ij}$  are normally distributed with the mean  $X_i(t_{ij}) = \mu(t_{ij}) + \sum_{\ell=1}^{p} \{s_{i\ell}\xi_\ell(t_{ij})\}$  and the variance  $\sigma^2$ , the authors show that the functional principal components can be calculated by maximizing the likelihood function

$$\prod_{i=1}^{N} \prod_{j=1}^{M} f(y_{ij})^{1-\delta_{ij}} P(y_{ij} \le c_i)^{\delta_{ij}}, \qquad (5)$$

where M is the index corresponding to maximum age,  $f(y_{ij})$  is the probability density function of  $y_{ij}$ ,  $\delta_{ij}$  is an indicator function corresponding to whether the *j*th observation for player *i* is missing where player *i* has  $m_i$  observations. Here, we allow for different numbers of observations per player. The missing data in the right tail have the natural constraint that once a player is out of the league, future values of performance are no larger than  $c_i$ ;  $c_i$ is set to be the final observation for the *i*th player. More details concerning the *imFunPCA* method are provided in Shi et al. (2021).

## 5 RESULTS

We restrict the data to include only player-years 22-34 with a minimum of 30 games played in a season. We also limit the data to players who had an NHL career lasting at least seven seasons. This provides us with 873 forwards and 438 defencemen. We randomly partitioned the data into training sets (750 forwards, 370 defensemen) and testing sets (123 forwards, 68 defensemen) such that the testing sets form approximately 15% of the data.

First, we obtained the estimated mean function  $\hat{\mu}(t)$  using the spline regression method outlined in Section 4.2 on the training data. We used Q = 6 spline basis functions of degree three, resulting in  $\tau = 4$  knots. Elijah, can you reproduce Fig 4 without normalization? The mean function is shown in Figure 4 where we observe concave shapes with peaks around 26-28 years. This period of peak performance differs from the participation peak observed in Figure 1. In keeping with the consideration of selection bias, we note that both curves in Figure 4 (forwards and defensemen) decline quickly past the age of 30 years, and that the curves are nearly indistinguishable.



Figure 4: Plots of the estimated mean function for FPCA.

Figure 5 shows the estimated first functional principal component  $\hat{\xi}_1(t)$ . We notice similar trends between the two positions. We interpret the first functional principal component as highlighting performance change between a player's early career (22-28 years) and late career (28-34 years). Hence, a player *i* with a large positive FPCA score  $s_{i1}$  given by (3) corresponds to a player who performs extremely well in their early career and poorly in their late career.

Figure 6 shows the estimated second functional principal component  $\hat{\xi}_2(t)$  which appears to have an up-and-down pattern for both positions. We interpret the second functional principal component as highlighting performance change between a player's peak age and their early and late stages of their career. Hence, a player *i* with a large positive FPCA



Figure 5: The estimated first functional principal component for the forwards and the defensemen.

score  $s_{i2}$  given by (3) corresponds to a player who has an exceptional peak but is not distinguished in the early and late stages of their career.



Figure 6: The estimated second functional principal component for the forwards and the defensemen.

## 5.1 Using FPCA for Prediction

I like this section very much. I think it is important, I like the general structure of providing and discussing Table 1. However, I have confusion on several fronts. First, I am wondering

about the predictions  $X_i(t_{ij})$ . I believe that you cannot predict future seasons for a player unless you have some existing seasons for the player. The existing seasons are used to find the player-specific FPCA components. So, I am wondering about the set D below. I think we need some observed seasons, and then D might be, for example, seasons beyond 28 years of age for which  $y_{ij}$  exists. Only when  $y_{ij}$  exists, can we calculate the  $MAE_i$ . So, I think D should be described differently and carefully. Now, regarding  $MAE_i$ , I believe this is a sum of absolute errors over years where we really have data. But in Table 1, we discuss career PS differences. So I don't think the absolute value sign makes sense. Also, right now, things are not comparable for a player who is predicted for 2 years versus a player who is predicted for 5 years, since there the sum will be most likely greater for the latter. I guess I am saying that career PS differences might instead be changed to yearly PS differences.

For the prediction problem, we first estimate the first two principal components using the training data. Prediction is then based on those players included in the testing set. For the testing data, we estimated FPC scores (3)

The prediction error for the *i*th player is given by the mean absolute error

$$MAE_i = \sum_{j \in D} |y_{ij} - X_i(t_{ij})|$$

where  $j \in D$  are the observed ages,  $y_{ij}$  is the true point share performance and  $X_i(t_{ij})$  is the prediction based on FPCA. We measure average FPCA prediction error for the N players in the testing set

Test Error 
$$= \frac{1}{N} \sum_{i=1}^{N} \text{MAE}_i$$
.

Table 1 provides the test error for FPCA. The error corresponds to the difference in career point shares between the prediction and actual results. Hence a test error of 10 corresponds to a model being 10 career point shares different from the true performance of the player. For a player who plays for 10 years in the NHL, this error would correspond to a difference in one PS per year between the predicted and actual performance. To investigate the FPCA approach, we calculate the testing error from using the Delta method (Lichtman 2009) based on the same training and testing datasets. We observe that FPCA is comparable to the Delta methods in terms of forward prediction and is better in terms

of defensemen prediction.

Position	Method	Error
Forwards	FPCA	21.5
	Delta	19.2
Defencemen	FPCA	14.2
	Delta	18.3

Table 1: Prediction performance of FPCA versus the Delta method.

This investigation highlights one of the major benefits of FPCA analysis. We can approximate the FPC scores from a player's early seasons, and then use the estimated mean curve and estimated FPCA eigenfunctions to forecast the player's future performance. This may be of great value to front office staff at major sports clubs.

#### 5.2 Using FPCA for Clustering

I also like this section, but the clustering has been done using PACE. Could we do the cluster exercise using our approach according to my simple suggestion below?

Cluster analysis is an unsupervised learning technique where we attempt to form groups of subjects that share common characteristics. If the clustering is effective, then one would see significant variation between groups. We use the FPCA score estimates  $s_{ij}$  given by (3) to cluster players. We choose p = 2 principal components, and plot the pairs of points in Figure 7. We colour-code the players into two categories according to whether they were ever selected to an all-star team. From the separation in the colours, it seems that clustering was effective. (Note that Fig 7 needs to be produced as described above.)

## 6 DISCUSSION

Discuss benefits and limitations.



Figure 7: Clustering the FPC scores by position. The colour-coding refers to players' allstar status.

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