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Objective

Create a machine learning model that leverages NFL football player statistics from 2017-2023 to predict players' fantasy rankings given their stats. Our model is designed to help fantasy football managers make smart decisions when selecting players for their rosters.



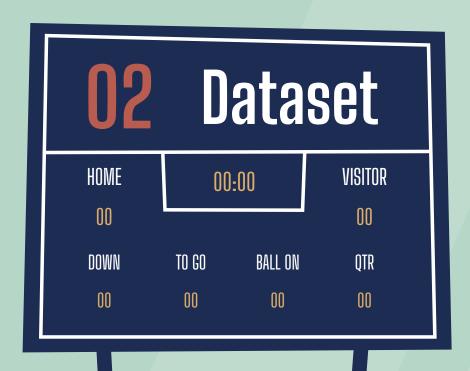
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We found <u>our dataset</u> on Kaggle:

- Football statistics by player from 2017-2023
- 27 columns (features), 3,388 rows



Our Dataset and Data Processing

Pre-processing:

- Drop columns containing player names, ids, and team
- Establish column PosRk (Position Rank) as our label
- Non-numerical column "FantPos"
 - Transform to list and onehotencoder for four unique positions (WR, RB, TE, QB)
- Scaling:
 - PowerTransformer/StandardScaler used to scale skewed numerical columns



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Step-by-step:

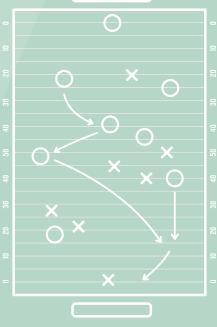
- Feature selection and pre-processing
- 2) Feature correlation analysis
- 3) Train/test split
- 4) Train multiple models
- 5) Evaluate Models
 - a) Save the best model
- 6) Test the final model

Our models: Linear Regression, Support Vector Regression, Decision Trees, Random Forest, and Gradient Boosting

Evaluate using MSE, MAE, and R-squared values

Save the best model using Pickle

One test with randomized inputs is included in the original code. Since our model is saved and can be exported, we can test it more thoroughly in our demo program.



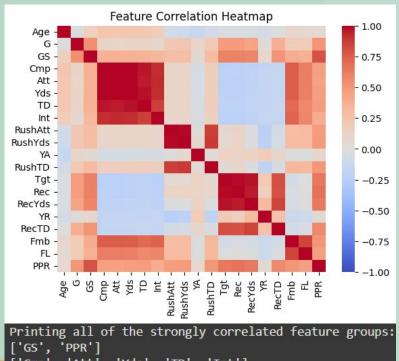
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We scaled our data this way because...



'Cmp', 'Att', 'Yds', 'TD', 'Int'] 'RushAtt', 'RushYds', 'RushTD']

'Tgt', 'Rec', 'RecYds', 'RecTD']

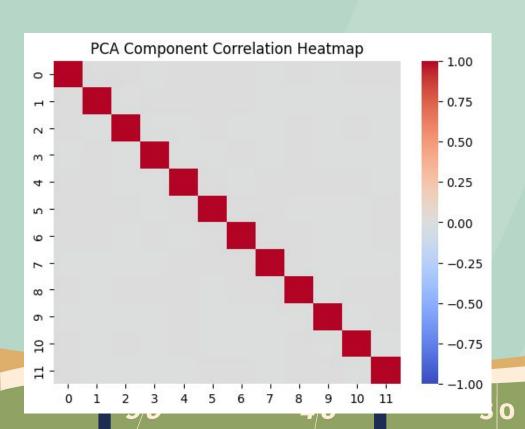
'Fmb', 'FL']

Many of these features are positively correlated, creating problems like:

- Redundant information
- Complicates model interpretability
- Can lead to overfitting

How can we fix this?

Principal Component Analysis



What does PCA do?

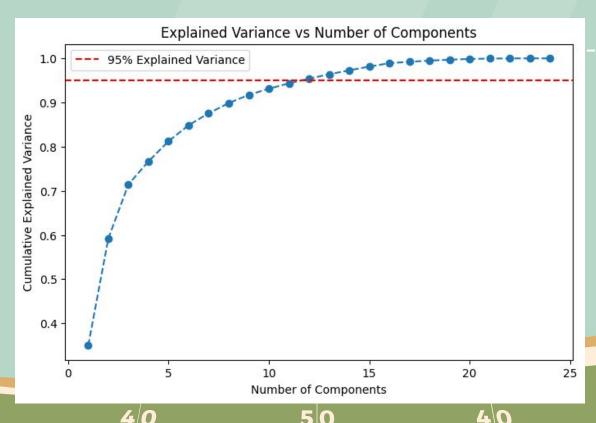
- Reduces dimensionality
- Removes multicollinearity

BUT...

PCA only works on normalized data:

- applied Power Transform to skewed data
- Standard Scaling to normalize data

Principal Component Analysis



Training all the models

Many of our models require the data to be PCA transformed, while others can handle multicollinearity and lose defining information when PCA transformed.

Cannot handle multicollinearity:

Linear Regression

SVR with Linear Kernel

SVR with Polynomial Kernel

SVR with RBF Kernel

Can handle multicollinearity:

Linear Ridge Regression
Linear Lasso Regression
Decision Tree
Random Forest
Gradient Boosting



Model Evaluation Metrics (Training and Validation)

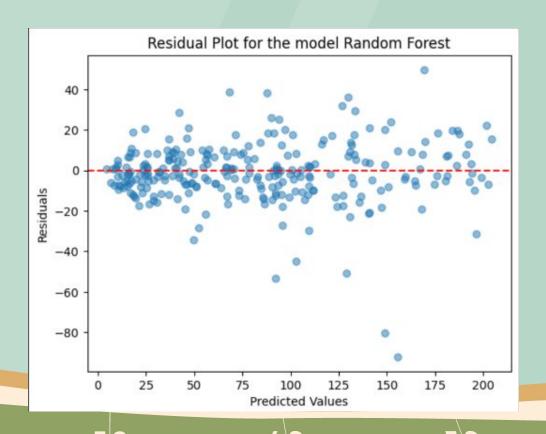
- Random Forest model achieves lowest MSE (20.91).
- High R^2 confirms strong predictive accuracy.
- Outperformed Decision Tree and Gradient Boosting models.

```
Evaluating Linear Regression ...
Average MSE on training / validation data = 447.1080153089607 / 453.76433658888766
Average MAE on training / validation data = 15.997263477422027 / 16.08469825550562
Average R2 on training / validation data = 0.8261596287655715 / 0.823160557367129
Evaluating Linear Ridge Regression ...
Average MSE on training / validation data = 447.1080242380987 / 453.76301767496153
Average MAE on training / validation data = 15.99740994876357 / 16.08481993616566
Average R2 on training / validation data = 0.8261487227207439 / 0.8231497251017691
Evaluating Linear Lasso Regression ...
Average MSE on training / validation data = 447.3359294999931 / 453.90376992892925
Average MAE on training / validation data = 16.017554778341676 / 16.103899865642205
Average R2 on training / validation data = 0.8242163622428837 / 0.8211876950990987
Evaluating SVR with linear kernal ...
Average MSE on training / validation data = 478.75933272646233 / 485.1752946718592
Average MAE on training / validation data = 15.495587395119761 / 15.617817646281605
Average R2 on training / validation data = 0.8269637227941583 / 0.8246552340304124
Evaluating SVR with polynomial kernal ...
Average MSE on training / validation data = 1045.3557389572652 / 1070.127862487328
Average MAE on training / validation data = 22.570291566711187 / 23.009611786216844
Average R2 on training / validation data = 0.21226634439387798 / 0.17617165178564015
Evaluating SVR with RBF kernal ...
Average MSE on training / validation data = 127.83003880099815 / 169.25926675073097
Average MAE on training / validation data = 5.9841686090544615 / 7.396215231375605
Average R2 on training / validation data = 0.9535084112074304 / 0.937428564685078
Evaluating Decision Tree ...
Average MSE on training / validation data = 46.24865388465027 / 51.89197961350787
Average MAE on training / validation data = 5.408853686576789 / 5.760856205163466
Average R2 on training / validation data = 0.9844434528757512 / 0.982403409215632
Evaluating Random Forest ...
Average MSE on training / validation data = 2.9195638632636327 / 20.918993745387453
Average MAE on training / validation data = 1.2894917999179991 / 3.4677306273062727
Average R2 on training / validation data = 0.9990294557296885 / 0.993006565792425
Evaluating Gradient Boosting ...
Average MSE on training / validation data = 109.96074776868667 / 194.65878020481972
Average MAE on training / validation data = 7.650847135821311 / 9.771027744321193
Average R2 on training / validation data = 0.9602050545252443 / 0.927538227815664
Best model for the task is Random Forest which offers the validation MSE of 20.918993745387453
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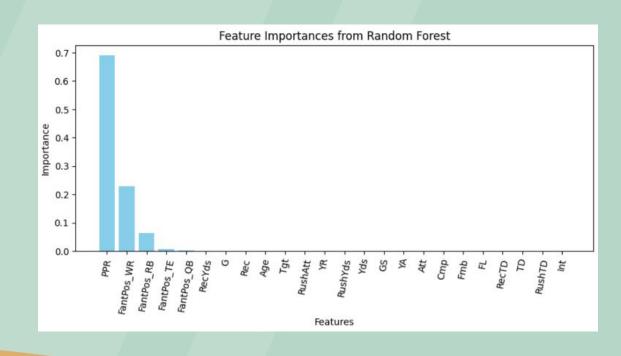
Residual Plot (Random Forest Model)

- Visualizes prediction errors (residuals) vs. actual values.
- Random scatter shows minimal bias.
- Confirms the model generalizes well across player ratings.



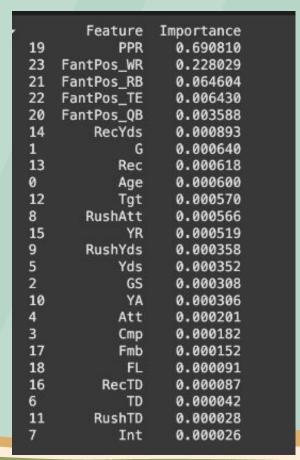
Feature Importance Bar Chart (Random Forest Model)

- Highlights top factors for predicting draftability (e.g., PPR, FantPos_WR).
- Most impactful features likely to align with real-world fantasy strategies.
- Positional data heavily influences draft decisions.



Detailed Feature Importances Table (Random Forest Model)

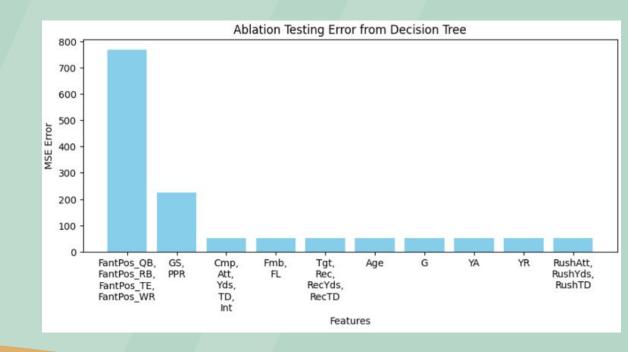
- Numerical ranking of feature importance (e.g., PPR: 0.6, FantPos_WR: 0.2).
- Adds transparency to the model's decision-making process.
- Reinforces the reliability of feature importance results.



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Ablation Testing Bar Chart (Decision Tree Model)

- Tests impact of removing features on prediction accuracy.
- Removing FantPos_-,
 PPR, and GS categories
 causes the largest error
 increase.





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Conclusions

Findings:

- PPR and positional metrics are key to draftability.
- Random Forest is the most accurate model.

Limitations:

- Temporal changes in player trends not captured.
- Lack of real-time data reduces immediate applicability.

Future Applications:

- Real-time draft tools.
- Extend to other fantasy sports.









Xavier **Worthy**

Kansas City Chiefs

ELIG WR
MANAGER Waivers
STATUS Healthy •

POSITION RANK

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AVERAGE POINTS 9.5

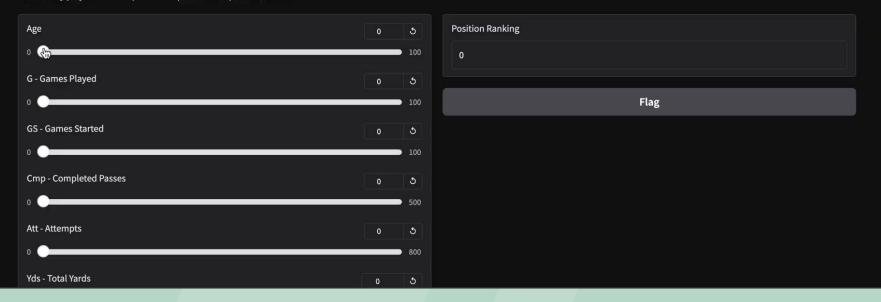
% ROSTERED **80.8** (-0.4)

https://fantasy.espn.com/football/players/add

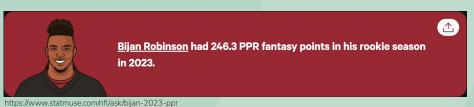
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Fantasy Football Rank Predictor

Provide key player stats and position to predict their positional rank.







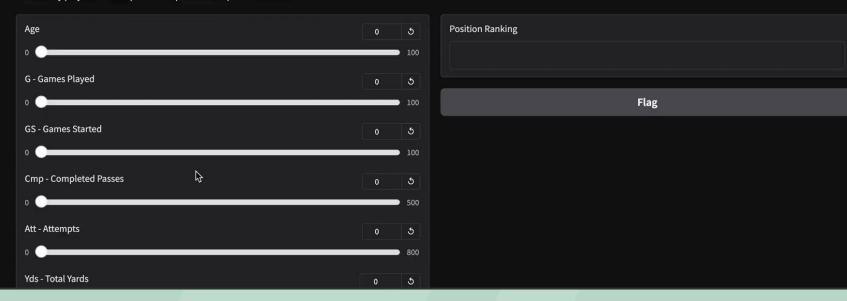


https://fantasy.espn.com/football/players/add

30 40 50 40 30

Fantasy Football Rank Predictor

Provide key player stats and position to predict their positional rank.









-Thanks for listening! TICKET SEAT 02 SEAT 20

What was the rationale behind selecting Position Ranking as the target value for prediction, rather than Overall Ranking?

We chose to select Position Ranking for our label instead of the Overall Ranking because in fantasy football, scoring differs depending on the specific position of the player, whether it's QB, RB, WR, TE. Some positions will get more points than others for the same statistic so we decided that delivering the user a position rank might be more beneficial and accurate.

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What factors led to the decision to use regression models for prediction rather than other machine learning algorithms?

We used regression analysis because it provides more detailed and precise predictions among players. Given our skewed data, where many stats are clustered on the lower end, regression allows us to rank players with similar stats more effectively.

Could you provide a detailed description of the mechanics behind Power Transformation in data preprocessing?

See the following slide...

Power Transformation

At its basic description, Power Transformation changes skewed data to fit a normal distribution (Gaussian). And there are two methods:

- **Box-Cox**: Applicable only to datasets with positive values (excluding 0)
- Yeo-Johnson: Applicable to datasets with positive and negative values

For our project, we used **Yeo-Johnson** transformation as certain stats aren't applied to all player positions. For instance, quarterbacks don't have stats in receptions.

How does Yeo-Johnson transformation work technically?

For all the positive values, the Yeo-Johnson transformation applies a formula similar to Box-Cox, where

$$\psi(\lambda, y) = \begin{cases} ((y+1)^{\lambda} - 1)/\lambda & \text{if } \lambda \neq 0, y \geq 0\\ \log(y+1) & \text{if } \lambda = 0, y \geq 0\\ -[(-y+1)^{2-\lambda} - 1)]/(2-\lambda) & \text{if } \lambda \neq 2, y < 0\\ -\log(-y+1) & \text{if } \lambda = 2, y < 0 \end{cases}$$

y is the original data value and λ is a parameter that determines the specific equation to be applied. This λ parameter is chosen to maximize the likelihood of achieving a Gaussian distribution, generally iterating through possible values until the highest likelihood is achieved.





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References and Project Link

Link to the GitHub repository:

https://github.com/aneeshharwalkar/FantasyFootballDraftabilityModel

Link to the Kaggle dataset:

"Fantasy Football Data 2017 - 2023" by Gary Bolduc

https://www.kaggle.com/datasets/gbolduc/fantasy-football-data-2017-2023?select=fantasy_merged_7_17.csv

References:

Yeo, In-Kwon, and Richard A. Johnson. "A new family of power transformations to improve normality or symmetry." *Biometrika*, vol. 87, no. 4, 1 Dec. 2000, pp. 954–959, https://doi.org/10.1093/biomet/87.4.954.