# Imputing using fancyimpute

DEALING WITH MISSING DATA IN PYTHON



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# fancyimpute package

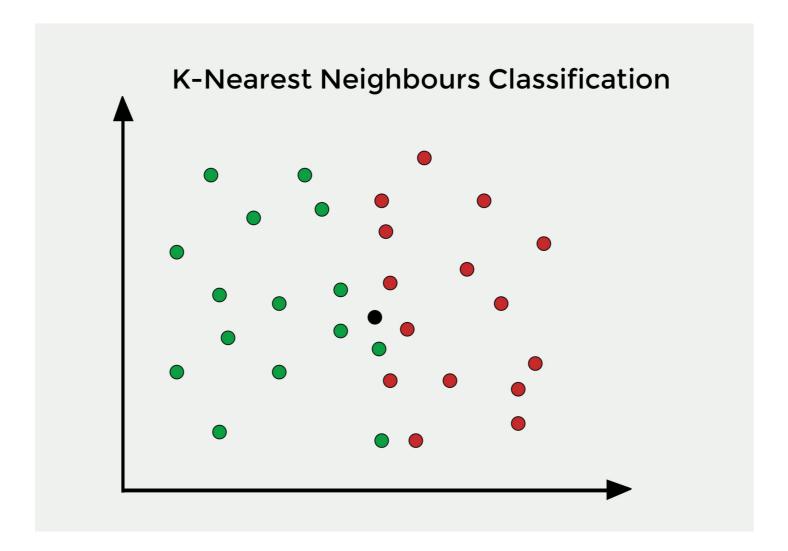
- Package contains advanced techniques
- Uses machine learning algorithms to impute missing values
- Uses other columns to predict the missing values and impute them

# Fancyimpute imputation techniques

- KNN or K-Nearest Neighbor
- MICE or Multiple Imputation by Chained Equations

# K-Nearest Neighbor Imputation

- Select K nearest or similar data points using all the non-missing features
- Take average of the selected data points to fill in the missing feature



# K-Nearest Neighbor Imputation

```
from fancyimpute import KNN
knn_imputer = KNN()
diabetes_knn = diabetes.copy(deep=True)
diabetes_knn.iloc[:, :] = knn_imputer.fit_transform(diabetes_knn)
```

# Multiple Imputations by Chained Equations (MICE)

- Perform multiple regressions over random sample of the data
- Take average of the multiple regression values
- Impute the missing feature value for the data point

# Multiple Imputations by Chained Equations(MICE)

```
from fancyimpute import IterativeImputer

MICE_imputer = IterativeImputer()

diabetes_MICE = diabetes.copy(deep=True)

diabetes_MICE.iloc[:, :] = MICE_imputer.fit_transform(diabetes_MICE)
```



# Summary

- Using Machine Learning techniques to impute missing values
- KNN finds most similar points for imputing
- MICE performs multiple regression for imputing
- MICE is a very robust model for imputation

# Let's practice!

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# Imputing categorical values

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# Complexity with categorical values

- Most categorical values are strings
- Cannot perform operations on strings
- Necessity to convert/encode strings to numeric values and impute

# Conversion techniques

#### **ONE-HOT ENCODER**

Color	Color_Red	Color_Green	Color_Blue
Red	1	0	0
Green	0	1	0
Blue	0	0	1
Red	1	0	0
Blue	0	0	1
Blue	0	0	1

#### **ORDINAL ENCODER**

Color	Value
Red	0
Green	1
Blue	2
Red	0
Blue	2
Blue	2

# Imputation techniques

- Fill with most frequent category
- Impute using statistical models like KNN

# Users profile data

```
users = pd.read_csv('userprofile.csv')
users.head()
```

ш								
ı		smoker	drink_level	dress_preference	ambience	hijos	activity	budg
ı	0	False	abstemious	informal	family	independent	student	medi
ı	1	False	abstemious	informal	family	independent	student	low
ı	2	False	social drinker	formal	family	independent	student	low
ı	3	False	abstemious	informal	family	independent	professional	medi
I	4	False	abstemious	no preference	family	independent	student	medi

# **Ordinal Encoding**

```
from sklearn.preprocessing import OrdinalEncoder
# Create Ordinal Encoder
ambience_ord_enc = OrdinalEncoder()
# Select non-null values in ambience
ambience = users['ambience']
ambience_not_null = ambience[ambience.notnull()]
reshaped_vals = ambience_not_null.values.reshape(-1, 1)
# Fncode the non-null values of ambience
encoded_vals = ambience_ord_enc.fit_transform(reshaped_vals)
# Replace the ambience column with ordinal values
users.loc[ambience.notnull(), 'ambience'] = np.squeeze(encoded_vals)
```

# **Ordinal Encoding**

```
# Create dictionary for Ordinal encoders
ordinal_enc_dict = {}
# Loop over columns to encode
for col name in users:
    # Create ordinal encoder for the column
    ordinal_enc_dict[col_name] = OrdinalEncoder()
    # Select the nin-null values in the column
    col = users[col_name]
    col_not_null = col[col.notnull()]
    reshaped_vals = col_not_null.values.reshape(-1, 1)
    # Encode the non-null values of the column
    encoded_vals = ordinal_enc_dict[col_name].fit_transform(reshaped_vals)
```

# Imputing with KNN

```
users_KNN_imputed = users.copy(deep=True)
# Create MICE imputer
KNN_{imputer} = KNN()
users_KNN_imputed.iloc[:, :] = np.round(KNN_imputer.fit_transform(imputed))
for col in imputed:
    reshaped_col = imputed[col].values.reshape(-1, 1)
    users_KNN_imputed[col] = ordinal_enc[col].inverse_transform(reshaped_col)
```

# Summary

Steps to impute categorical values

- Convert non-missing categorical columns to ordinal values
- Impute the missing values in the ordinal DataFrame
- Convert back from ordinal values to categorical values

# Let's practice!

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# Evaluation of different imputation techniques

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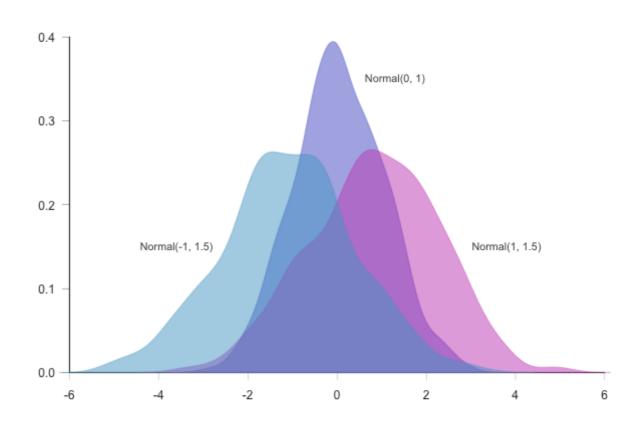
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# **Evaluation techniques**

- Imputations are used to improve model performance.
- Imputation with maximum machine learning model performance is selected.
- Density plots explain the distribution in the data.
- A very good metric to check bias in the imputations.



# Fit a linear model for statistical summary

```
import statsmodels.api as sm

diabetes_cc = diabetes.dropna(how='any')
X = sm.add_constant(diabetes_cc.iloc[:, :-1])
y = diabetes_cc['Class']
lm = sm.OLS(y, X).fit()
```

Summary:		==========				
Dep. Variable:		Class	 R-squared:		0.3	
Model:		0LS	Adj. R-square	d:	0.3	32
Method:	Least	Squares	F-statistic:		25.	30
Date:	Wed, 10 J	ul 2019	Prob (F-stati	stic):	2.65e-	31
Time:	1	5:03:19	Log-Likelihoo	d:	-177.	76
No. Observations:			AIC:		373	
Df Residuals:		383	BIC:		409	.3
Df Model:		8				
Covariance Type:	nonrobust					
=========	coef		======== t			0.975]
<hr/>						
const	-1.1027	0.144	-7.681	0.000	-1.385	-0.820
Pregnant	0.0130	0.008	1.549	0.122	-0.003	0.029
Glucose	0.0064	0.001	7.855	0.000	0.005	0.008
Diastolic_BP	5.465e-05	0.002	0.032	0.975	-0.003	0.003
	0.0017	0.003		0.506	-0.003	0.007
Serum_Insulin	-0.0001	0.000	-0.603	0.547	-0.001	0.000
BMI	0.0093	0.004		0.017	0.002	0.017
Diabetes_Pedigree		0.058	2.708	0.007	0.043	0.271
Age	0.0059	0.003	2.109	0.036	0.000	0.011



# R-squared and Coefficients

```
lm.rsquared_adj
```

0.33210

lm.params

```
const
                   -1.102677
Pregnant
                    0.012953
Glucose
                    0.006409
Diastolic_BP
                    0.000055
Skin_Fold
               0.001678
Serum_Insulin
                   -0.000123
BMI
                    0.009325
Diabetes_Pedigree
                    0.157192
                    0.005878
Age
dtype: float64
```



# Fit linear model on different imputed DataFrames

```
# Mean Imputation
X = sm.add_constant(diabetes_mean_imputed.iloc[:, :-1])
y = diabetes['Class']
lm_mean = sm.OLS(y, X).fit()
# KNN Imputation
X = sm.add_constant(diabetes_knn_imputed.iloc[:, :-1])
lm_KNN = sm.OLS(y, X).fit()
# MICE Imputation
X = sm.add_constant(diabetes_mice_imputed.iloc[:, :-1])
lm_MICE = sm.OLS(y, X).fit()
```

# Comparing R-squared of different imputations

```
Complete Mean Imp. KNN Imp. MICE Imp.

R_squared_adj 0.332108 0.313781 0.316543 0.317679
```

# Comparing coefficients of different imputations

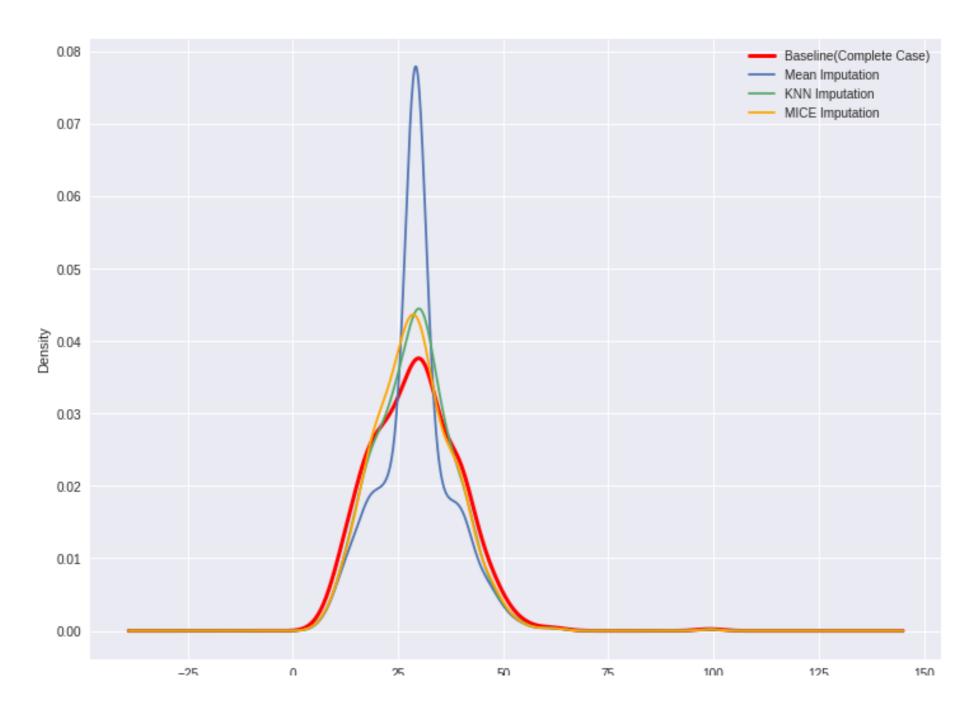
	Complete	Mean Imp.	KNN Imp.	MICE Imp.
	·	·	·	
const	-1.102677	-1.024005	-1.028035	-1.050023
Pregnant	0.012953	0.020693	0.020047	0.020295
Glucose	0.006409	0.006467	0.006614	0.006871
Diastolic_BP	0.000055	-0.001137	-0.001196	-0.001317
Skin_Fold	0.001678	0.000193	0.001626	0.000807
Serum_Insulin	-0.000123	-0.000090	-0.000147	-0.000227
BMI	0.009325	0.014376	0.013239	0.014203
Diabetes_Pedigree	0.157192	0.129282	0.128038	0.129056
Age	0.005878	0.002092	0.002046	0.002097



# Comparing density plots

```
diabetes_cc['Skin_Fold'].plot(kind='kde', c='red', linewidth=3)
diabetes_mean_imputed['Skin_Fold'].plot(kind='kde')
diabetes_knn_imputed['Skin_Fold'].plot(kind='kde')
diabetes_mice_imputed['Skin_Fold'].plot(kind='kde')
labels = ['Baseline (Complete Case)', 'Mean Imputation', 'KNN Imputation',
          'MICE Imputation']
plt.legend(labels)
plt.xlabel('Skin Fold')
```

# Comparing density plots



#### Summary

- Applying linear model from the statsmodels package
- Comparing the coefficients and standard errors
- Comparing density plots

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# Conclusion

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- Null Value operations
- Detecting missing values
- Replacing missing values
- Analyzing amount of missingness

- Types of missingness
  - MCAR
  - MAR
  - MNAR
- Correlations of missingness
  - Heatmaps
  - Dendrograms
- Visualize missingness across a variable
- Deleting missing values

- Imputation techniques
- Treating time-series data
- Graphical comparison of imputed time-series data

- Advanced imputation techniques
  - o KNN
  - MICE
- Imputing categorical data
- Evaluating and comparing the different imputations

# Congratulations!!

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