## not certain

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## 0.1 Highest Correlated Features by Model and Scaling type

```
[]: # Standard Scaling
     dt_importance_std = dt_model.feature_importances_
     rf_importance_std = rf_model.feature_importances_
     xgb_importance_std = xgb_model.feature_importances_
[]: # Min-Max Scaling
     dt_importance_mm = dt_model.feature_importances_### STD Model Comparison Table
     results = {
         'Model': ['Linear Regression', 'Decision Tree', 'Random Forest', 'XGBoost'],
         'RMSE': [np.sqrt(mean squared error(y test 1, y pred lr)),
                 np.sqrt(mean_squared_error(y_test_1, y_pred_dt)),
                 np.sqrt(mean_squared_error(y_test_1, y_pred_rf)),
                 np.sqrt(mean_squared_error(y_test_1, y_pred_xgb))],
         'R2 Score': [r2_score(y_test_1, y_pred_lr),
                     r2_score(y_test_1, y_pred_dt),
                      r2_score(y_test_1, y_pred_rf),
                      r2_score(y_test_1, y_pred_xgb)]
     }
     results_df = pd.DataFrame(results)
     print(results_df)
     rf_importance_mm = rf_model.feature_importances_
     xgb_importance_mm = xgb_model.feature_importances_
                               RMSE R2 Score
                   Model
       Linear Regression 16.423524 -0.052112
    1
           Decision Tree 23.314402 -1.120204
    2
           Random Forest 20.237325 -0.597480
                 XGBoost 21.281860 -0.766641
    3
[]: feature_names = X_train_1.columns
[]: def plot_feature_importance(importances, feature_names, model_names, title):
        plt.figure(figsize=(15, 15))
```

```
# Create a DataFrame with feature importances
        df = pd.DataFrame(importances, index=model_names, columns=feature names)
         # Sort features by average importance across all models
        avg_importance = df.mean()
        sorted_features = avg_importance.sort_values(ascending=False).index
         # Create a custom color map from blue to cerulean
        colors = ["#0000FF", "#00BFFF"] # Blue to Cerulean
        n bins = 100
         cmap = mcolors.LinearSegmentedColormap.from list("custom", colors, N=n bins)
         # Create heatmap
         sns.heatmap(df[sorted features], annot=True, cmap=cmap, fmt='.2f')
        plt.title(title)
        plt.xlabel('Features')
        plt.ylabel('Models')
        plt.xticks(rotation=45, ha='right')
        plt.yticks(rotation=0)
        plt.tight_layout()
        plt.show()
[]: # Standard Scaling
     importances_std = [dt_importance_std, rf_importance_std, xgb_importance_std]
     model names = ['Decision Tree', 'Random Forest', 'XGBoost']
     plot_feature_importance(importances_std, feature_names, model_names, 'Feature_u
      →Importance - Standard Scaling')
     # Min-Max Scaling
     importances_mm = [dt_importance_mm, rf_importance_mm, xgb_importance_mm]
     plot feature importance(importances mm, feature names, model names, 'Feature,
      →Importance - Min-Max Scaling')
```



