```
In [1]: # Install these packages if you haven't already!
        # pip install pandas
        # pip install scikit-learn
        # pip install joblib
        # pip install numpy
        # pip install xgboost
        # pip install matplotlib
        # pip install seaborn
        # author
                      = "Siem Vonk"
        # studentID = "i6290798"
        import pandas as pd
        import numpy as np
        from sklearn.metrics import mean_squared_error, r2_score
        import joblib
        import os
        print(os.getcwd())
        os.makedirs("files", exist_ok=True)
```

d:\Coding\!MSc\Course 5 ML\MSc_ML_Assignment

Training of the models

```
In [2]: data_original = pd.read_csv("data_regression_train_df.csv")
    data = data_original.dropna(subset=['Viability']) # 0 missing rows for viability

# Missing values for each variable
    missing_values = data.isnull().sum()
    print(missing_values)

# Select variables with less than 3000 missing values (around 1/3rd of the samples)
    columns_to_keep = missing_values[missing_values < 3000].index
    data = data[columns_to_keep]

# Export columns to keep for new data (not included in training so they have to be
    columns_to_keep.to_series().to_csv("files/columns_to_keep.txt", index=False, header</pre>
```

```
Coating
                            6133
       core size nm
                            644
       hydro_size_nm
                            6804
       Surf_charge_mV
                           4779
       Surface_area_m2_g
                            8380
       shape
                            5511
       Dose_microg_mL
                            165
                             277
      Duration h
       Cell_name
                               0
      Cell_species
                               0
       cell_Organ
                              0
                             17
      Cell_morphology
                             596
       Cell_age
       cell type
                              77
                            2429
       sex
                               0
      Assay
                             195
       Test_indicator
       nanomaterial_group
                              0
       Viability
                               0
       dtype: int64
In [3]: from sklearn.model_selection import train_test_split
        # Separate features and target
        X = data.drop(columns=['Viability'])
        y = data['Viability']
        # Split the data into training and testing sets
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_sta
In [4]: # Export test set for testing new data pipeline (should return same results)
        test df = X test.copy()
        test_df['Viability'] = y_test
        test_df.to_csv("test_set.csv", index=False) # You can import this at the top to tes
In [5]: # Impute X_train and use the same fitted imputer for X_test
        from assignment_functions import assignment_impute
        imputer = assignment_impute(max_iter=10)
        X_train= imputer.fit_transform(X_train)
        X_test = imputer.transform(X_test)
        # Save the imputer for new data
        joblib.dump(imputer, 'files/imputer.pkl')
Out[5]: ['files/imputer.pkl']
In [6]: # Encode categorical variables
        from sklearn.preprocessing import OneHotEncoder
        # Encode based on training data only
        categorical_cols = X_train.select_dtypes(include='object').columns
        encoder = OneHotEncoder(sparse_output=False, handle_unknown='ignore') # Ignore any
```

NP_type

0

```
X_train_encoded = pd.DataFrame(
            encoder.fit_transform(X_train[categorical_cols]),
            columns=encoder.get_feature_names_out(categorical_cols),
            index=X_train.index
        X_test_encoded = pd.DataFrame(
            encoder.transform(X test[categorical cols]),
            columns=encoder.get_feature_names_out(categorical_cols),
            index=X_test.index
        # Find indices of numeric columns in X_train (same as X_test)
        numeric_cols = X_train.select_dtypes(include=[np.number]).columns
        # Merge encoded categorical features and the numeric features
        X_train_final = pd.concat([X_train[numeric_cols], X_train_encoded], axis=1) # axis
        X_test_final = pd.concat([X_test[numeric_cols], X_test_encoded], axis=1)
        # Save the encoder and feature order
        joblib.dump(encoder, 'files/encoder.pkl')
        joblib.dump(X_train_final.columns.tolist(), 'files/feature_order.pkl')
Out[6]: ['files/feature_order.pkl']
In [7]: # Scaling X_train and X_test
        from sklearn.preprocessing import StandardScaler
        scaler = StandardScaler() # Mean=0, std=1
        X_train_scaled = scaler.fit_transform(X_train_final)
        X_test_scaled = scaler.transform(X_test_final)
        joblib.dump(scaler, 'files/scaler.pkl') # Save the scaler for new data
        # Convert back to dataframes with column names and index preserved
        X_train_scaled_df = pd.DataFrame(X_train_scaled, columns=X_train_final.columns, ind
        X_test_scaled_df = pd.DataFrame(X_test_scaled, columns=X_test_final.columns, index=
In [8]: # Lasso (took around 3 seconds)
        from sklearn.linear_model import LassoCV
        param_grid_lasso = {
            'alpha': [0.01, 0.1, 0.5, 1, 10, 100, 1000]
        # Lasso regression with cross-validation kfold 5
        lasso_cv = LassoCV(alphas=param_grid_lasso['alpha'], cv=5, random_state=1, max_iter
        lasso_cv.fit(X_train_scaled_df, y_train)
        y_pred_lasso = lasso_cv.predict(X_test_scaled_df)
        lasso_best = lasso_cv.alpha_ # Retrieving the best alpha value which was used
        mse_lasso = mean_squared_error(y_test, y_pred_lasso)
        r2_lasso = r2_score(y_test, y_pred_lasso)
```

```
print("Lasso Regression Results")
        print(f"Mean Squared Error (MSE): {mse lasso:.4f}")
        print(f"R2 Score: {r2_lasso:.4f}")
        print(f"Best Lasso alpha: {lasso_best}")
        Betas_Lasso = pd.Series(lasso_cv.coef_, index=X_train_scaled_df.columns)
        Betas_Lasso = Betas_Lasso.loc[Betas_Lasso.abs().sort_values(ascending=False).index]
        print(f"\nLasso Coefficients")
        print(Betas_Lasso.head(5))
        # Save the model
        joblib.dump(lasso_cv, 'files/lasso.pkl')
       Lasso Regression Results
       Mean Squared Error (MSE): 783.3916
       R<sup>2</sup> Score: 0.2565
       Best Lasso alpha: 0.1
       Lasso Coefficients
                          -4.978563
       NP_type_Ag
       NP_type_CdTe
                         -4.227897
       Assay_MTS
                          4.093269
       NP_type_MWCNT -3.978024
       NP_type_Dendrimer 3.783435
       dtype: float64
Out[8]: ['files/lasso.pkl']
In [9]: # Ridge (took around 3 seconds)
        from sklearn.linear_model import RidgeCV
        param_grid_ridge = {
            'alpha': [0.01, 0.1, 0.5, 1, 10, 100, 1000]
        }
        # Ridge regression with 5-fold CV
        ridge_cv = RidgeCV(alphas=param_grid_ridge['alpha'], cv=5)
        ridge_cv.fit(X_train_scaled_df, y_train)
        y_pred_ridge = ridge_cv.predict(X_test_scaled_df)
        ridge_best = ridge_cv.alpha_
        mse_ridge = mean_squared_error(y_test, y_pred_ridge)
        r2_ridge = r2_score(y_test, y_pred_ridge)
        print("Ridge Regression Results")
        print(f"Mean Squared Error (MSE): {mse_ridge:.4f}")
        print(f"R2 Score: {r2_ridge:.4f}")
        print(f"Best Ridge alpha: {ridge_best}")
        Betas_Ridge = pd.Series(ridge_cv.coef_, index=X_train_scaled_df.columns)
        Betas_Ridge = Betas_Ridge.abs().sort_values(ascending=False)
        print(f"\nRidge Coefficients")
        print(Betas_Ridge.head(5))
```

```
# Around the same as lasso and quite low r2 so not saved
        Ridge Regression Results
        Mean Squared Error (MSE): 784.2102
        R<sup>2</sup> Score: 0.2557
        Best Ridge alpha: 1000.0
        Ridge Coefficients
        NP_type_Ag 3.704867
NP_type_Dendrimer 3.528715
        NP_type_EudragitRL 3.359812
        NP_type_CdTe
                             3.008252
        Assay_MTS
                             2.792514
        dtype: float64
In [10]: # Elastic Net (took around 2 minutes)
         from sklearn.model_selection import GridSearchCV
         from sklearn.linear_model import ElasticNet
         EN = ElasticNet(random_state=1, max_iter=5000)
         param_grid_EN = {
             'alpha': [0.01, 0.1, 0.5, 1, 10, 100],
              'll_ratio': [0.1, 0.3, 0.6, 0.9, 1.0] # Ridge (l1_ratio=0) to Lasso (=1)
         }
         # EN regression with cross-validation kfold 5 using param grid
         # GridSearchCV to find the best alpha and l1_ratio
         EN_cv = GridSearchCV(EN, param_grid_EN, cv=5, scoring='neg_mean_squared_error')
         EN_cv.fit(X_train_scaled_df, y_train)
         y_pred_EN = EN_cv.predict(X_test_scaled_df)
In [11]: # EN results
         mse_EN = mean_squared_error(y_test, y_pred_EN)
         r2_EN = r2_score(y_test, y_pred_EN)
         best_params_EN = EN_cv.best_params_
         best_alpha_EN = best_params_EN['alpha']
         best_l1_ratio_EN = best_params_EN['l1_ratio']
         print(f"Mean Squared Error (MSE): {mse_EN:.4f}")
         print(f"R2 Score = {r2_EN:.4f}")
         print(f"Best alpha: {best_alpha_EN} and l1_ratio: {best_l1_ratio_EN}")
         # Get the best ElasticNet model from GridSearchCV
         best_en_model = EN_cv.best_estimator_
         Betas_EN = pd.Series(best_en_model.coef_, index=X_train_scaled_df.columns)
         print(f"\nElastic Net Coefficients")
```

Lasso seems to perform better and is a lot faster so EN is not saved

print(Betas_EN.head(5))

```
Mean Squared Error (MSE): 783.4415
        R^2 Score = 0.2564
        Best alpha: 0.1 and l1_ratio: 0.9
        Elastic Net Coefficients
        core_size_nm
                        0.680199
        Dose_microg_mL -2.820204
                      -3.118705
        Duration_h
        NP type Ag
                       -4.757026
        NP_type_Al2O3
                        0.489946
        dtype: float64
In [12]: # Random Forest default (took around 3 seconds)
         from sklearn.ensemble import RandomForestRegressor
         # Using default (MSE) criterion for regression
         rf = RandomForestRegressor(
             n estimators=100,
             random_state=123,
             n_{jobs=-1}
         rf.fit(X_train_scaled_df, y_train)
         y_pred_rf_old = rf.predict(X_test_scaled_df)
         joblib.dump(rf, 'files/random_forest_default.pkl')
         mse_rf_old = mean_squared_error(y_test, y_pred_rf_old)
         r2_rf_old = r2_score(y_test, y_pred_rf_old)
         print(f"Random Forest Regression Results")
         print(f"Mean Squared Error (MSE): {mse_rf_old:.4f}")
         print(f"R2 Score: {r2_rf_old:.4f}")
        Random Forest Regression Results
        Mean Squared Error (MSE): 471.5396
        R<sup>2</sup> Score: 0.5525
In [13]: # Random Forest parameter optimization (took around 30 seconds)
         from sklearn.model_selection import GridSearchCV
         from sklearn.ensemble import RandomForestRegressor
         # Parameter grid (keep n_estimators fixed)
         param_grid = {
             'min_samples_split': [2, 5, 10],
             'min_samples_leaf': [1, 2, 4]
         }
         # Base model
         rf_base = RandomForestRegressor(
             n_estimators=100,
             random_state=123,
             n jobs=-1
```

```
# Grid search
         grid_search = GridSearchCV(
             estimator=rf base,
             param_grid=param_grid,
             cv=5,
             scoring='r2',
             verbose=1,
             n_{jobs=-1}
         # Fit to scaled training data
         grid_search.fit(X_train_scaled_df, y_train)
         # Get the best model
         best rf = grid search.best estimator
         # Predict and evaluate
         y_pred_rf = best_rf.predict(X_test_scaled_df)
         joblib.dump(best_rf, 'files/random_forest.pkl')
         mse_rf = mean_squared_error(y_test, y_pred_rf)
         r2_rf = r2_score(y_test, y_pred_rf)
         print("\nRandom Forest Regression Results (Optimized)")
         print(f"Best Parameters: {grid_search.best_params_}")
         print(f"Mean Squared Error (MSE): {mse_rf:.4f}")
         print(f"R2 Score: {r2_rf:.4f}")
        Fitting 5 folds for each of 9 candidates, totalling 45 fits
        Random Forest Regression Results (Optimized)
        Best Parameters: {'min_samples_leaf': 1, 'min_samples_split': 5}
        Mean Squared Error (MSE): 442.7312
        R<sup>2</sup> Score: 0.5798
In [14]: # xgboost
         from xgboost import XGBRegressor
         from sklearn.metrics import mean_squared_error, r2_score
         xgb_model = XGBRegressor(n_estimators=100, learning_rate=0.1, max_depth=3, random_s
         xgb_model.fit(X_train_scaled_df, y_train)
         y_pred_xgb = xgb_model.predict(X_test_scaled_df)
         joblib.dump(xgb_model, 'files/xgb.pkl')
         mse_xgb = mean_squared_error(y_test, y_pred_xgb)
         r2_xgb = r2_score(y_test, y_pred_xgb)
         print("XGBoost Regression Results:")
         print(f"Mean Squared Error (MSE): {mse_xgb:.4f}")
         print(f"R2 Score: {r2_xgb:.4f}")
         # Worse than random forest but better than lasso and EN (R2 of 0.33)
```

```
R<sup>2</sup> Score: 0.3331
In [15]: from sklearn.svm import SVR
          # SVR model (RBF kernel is default)
          svr = SVR(C=1.0, epsilon=0.1)
          svr.fit(X_train_scaled_df, y_train)
         y_pred_svr = svr.predict(X_test_scaled_df)
          mse svr = mean_squared_error(y_test, y_pred_svr)
          r2_svr = r2_score(y_test, y_pred_svr)
          print("\nSVR Regression Results:")
          print(f"Mean Squared Error (MSE): {mse_svr:.4f}")
          print(f"R2 Score: {r2_svr:.4f}")
         # very low R2 score (0.08) and quite slow so not saved
        SVR Regression Results:
        Mean Squared Error (MSE): 972.2903
        R<sup>2</sup> Score: 0.0772
In [16]: # KNN regression (almost instant)
         from sklearn.neighbors import KNeighborsRegressor
          knn = KNeighborsRegressor(n_neighbors=5)
          knn.fit(X_train_scaled_df, y_train)
         y_pred_knn = knn.predict(X_test_scaled_df)
          mse_knn = mean_squared_error(y_test, y_pred_knn)
          r2_knn = r2_score(y_test, y_pred_knn)
          print("KNN Regression Results")
          print(f"Mean Squared Error (MSE): {mse_knn:.4f}")
          print(f"R2 Score: {r2_knn:.4f}")
          # Decent R2 score (0.39)
          joblib.dump(knn, 'files/knn.pkl')
        KNN Regression Results
        Mean Squared Error (MSE): 640.7020
        R<sup>2</sup> Score: 0.3919
Out[16]: ['files/knn.pkl']
In [17]: # KNN parameter optimization
          from sklearn.neighbors import KNeighborsRegressor
          from sklearn.model_selection import GridSearchCV
          param_grid = {'n_neighbors': range(1, 21)}
          knn_opt = KNeighborsRegressor()
          grid_search = GridSearchCV(knn_opt, param_grid, cv=5, scoring='r2')
```

XGBoost Regression Results:

Mean Squared Error (MSE): 702.6405

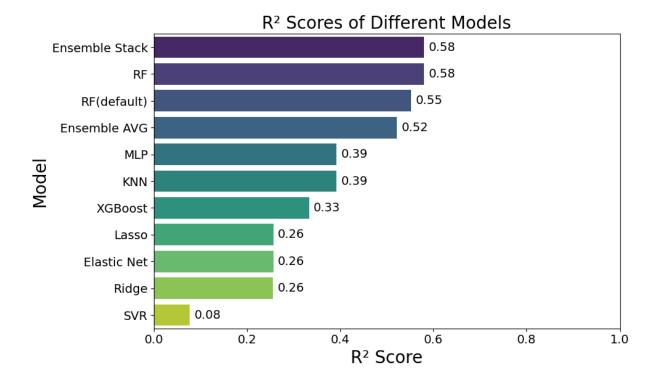
```
grid_search.fit(X_train_scaled_df, y_train)
          print(f"Best n neighbors: {grid search.best params ['n neighbors']}")
          print(f"Best R2 Score: {grid_search.best_score_:.4f}")
         # Update the model with the optimal number of neighbors
         best_knn_opt = grid_search.best_estimator_
         # n neighbors = 5 is better so not saved
        Best n_neighbors: 3
        Best R<sup>2</sup> Score: 0.3929
In [18]: # Neural Network (MLP) regression (took around 1 minute)
         from sklearn.neural network import MLPRegressor
         # Multi-layer Perceptron regressor with one hidden layer of 100 neurons, ReLU activ
         mlp = MLPRegressor(hidden_layer_sizes=(100,), max_iter=1000, random_state=123)
         mlp.fit(X_train_scaled_df, y_train)
         # Predict
         y_pred_mlp = mlp.predict(X_test_scaled_df)
In [19]: # MLP Evaluation
         mse_mlp = mean_squared_error(y_test, y_pred_mlp)
         r2_mlp = r2_score(y_test, y_pred_mlp)
          print("Neural Network Regression Results")
         print(f"Mean Squared Error (MSE): {mse mlp:.4f}")
         print(f"R2 Score: {r2_mlp:.4f}")
         # Decent R2 score (0.39)
         joblib.dump(mlp, 'files/mlp.pkl')
        Neural Network Regression Results
        Mean Squared Error (MSE): 640.0897
        R<sup>2</sup> Score: 0.3925
Out[19]: ['files/mlp.pkl']
In [20]: # Overview of all models before ensembling
         results = pd.DataFrame({
              'Model': ['Lasso', 'Ridge', 'Elastic Net', 'RF', 'XGBoost', 'SVR', 'KNN', 'MLP'
              'MSE': [mse_lasso, mse_ridge, mse_EN, mse_rf, mse_xgb, mse_svr, mse_knn, mse_ml
              'R<sup>2</sup> Score': [r2_lasso, r2_ridge, r2_EN, r2_rf, r2_xgb, r2_svr, r2_knn, r2_mlp,
          results = results.sort_values(by='R2 Score', ascending=False)
         print(results)
```

```
Model
                              MSE R<sup>2</sup> Score
                    RF 442.731225 0.579794
        8 RF (default) 471.539635 0.552452
        7
                    MLP 640.089711 0.392477
                    KNN 640.701986 0.391896
        6
                XGBoost 702.640506 0.333109
        4
        0
                 Lasso 783.391590 0.256466
        2 Elastic Net 783.441468 0.256419
        1
                  Ridge 784.210216 0.255689
        5
                    SVR 972.290282 0.077179
In [21]: # Ensemble (Averaging) of Random Forest, KNN, and MLP
         preds_rf = best_rf.predict(X_test_scaled_df)
         preds_knn = knn.predict(X_test_scaled_df)
         preds_mlp = mlp.predict(X_test_scaled_df)
         # Simple average (unweighted)
         ensemble_preds = (preds_rf + preds_knn + preds_mlp) / 3
         # Evaluation
         mse_ensemble = mean_squared_error(y_test, ensemble_preds)
         r2_ensemble = r2_score(y_test, ensemble_preds)
         print("Ensemble (Averaging) Results")
         print(f"MSE: {mse ensemble:.2f}")
         print(f"R2 Score: {r2_ensemble:.2f}")
         # R2 of 0.52 which is Lower than RF
        Ensemble (Averaging) Results
        MSE: 503.88
        R<sup>2</sup> Score: 0.52
In [22]: # Ensemble with stacking (took around 2.5 minutes)
         from sklearn.ensemble import StackingRegressor
         from sklearn.linear_model import LinearRegression
         estimators = [
             ('rf', best_rf),
             ('knn', knn),
             ('mlp', mlp),
         # Define meta-learner
         stack = StackingRegressor(
             estimators=estimators,
             final_estimator=LinearRegression(),
             cv=5,
             n_{jobs=-1}
         stack.fit(X_train_scaled_df, y_train)
```

```
Out[22]:
                                                  StackingRegressor
                            rf
                                                              knn
                                                                                            ml
                RandomForestRegressor
                                                    KNeighborsRegressor
                                                                                     MLPRegr
                                                   final_estimator
                                               LinearRegression
In [23]: # Stack ensemble predict
         stack_preds = stack.predict(X_test_scaled_df)
         mse_stack = mean_squared_error(y_test, stack_preds)
         r2_stack = r2_score(y_test, stack_preds)
         print("Stacking Regressor Results")
         print(f"MSE: {mse_stack:.4f}")
         print(f"R2 Score: {r2_stack:.4f}")
         # R2 score of 0.58 (best one so far)
         joblib.dump(stack, 'files/stack.pkl')
        Stacking Regressor Results
        MSE: 442.5417
        R<sup>2</sup> Score: 0.5800
Out[23]: ['files/stack.pkl']
In [24]: # Overview of all models (now including ensembles)
         results = pd.DataFrame({
              'Model': ['Lasso', 'Ridge', 'Elastic Net', 'RF', 'XGBoost', 'SVR', 'KNN', 'MLP'
             'MSE': [mse_lasso, mse_ridge, mse_EN, mse_rf, mse_xgb, mse_svr, mse_knn, mse_ml
              'R<sup>2</sup> Score': [r2_lasso, r2_ridge, r2_EN, r2_rf, r2_xgb, r2_svr, r2_knn, r2_mlp,
             'Saved': ['Yes', 'No', 'No', 'Yes', 'Yes', 'No', 'Yes', 'Yes', 'No', 'Yes', 'No
         })
         results = results.sort values(by='R2 Score', ascending=False)
         results.to_csv("files/model_results.csv", index=False)
         print(results)
                     Model
                                   MSE R<sup>2</sup> Score Saved
        9
            Ensemble Stack 442.541734 0.579974
                                                   Yes
        3
                        RF 442.731225 0.579794
                                                  Yes
        10
               RF(default) 471.539635 0.552452
                                                  No
              Ensemble AVG 503.878283 0.521758
        8
                                                    No
        7
                       MLP 640.089711 0.392477 Yes
                       KNN 640.701986 0.391896
        6
                                                   Yes
        4
                   XGBoost 702.640506 0.333109
                                                   Yes
                     Lasso 783.391590 0.256466
        0
                                                  Yes
        2
               Elastic Net 783.441468 0.256419
                                                    No
        1
                     Ridge 784.210216 0.255689
                                                    No
        5
                       SVR 972.290282 0.077179
                                                    Nο
```

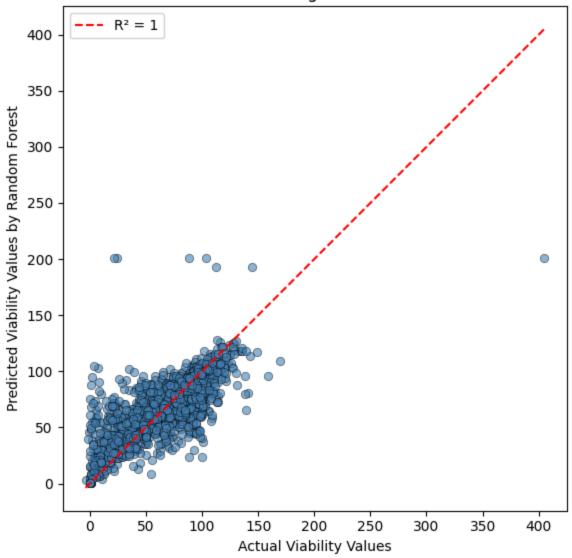
Figures

```
In [25]: ## Y_test line and y_pred line with samples on x axis
         import matplotlib.pyplot as plt
         import seaborn as sns
         import pandas as pd
         results = pd.read_csv("files/model_results.csv")
In [26]: # Bar chart of R<sup>2</sup> scores
         plt.figure(figsize=(10, 6))
         ax = sns.barplot(
             x='R2 Score',
             y='Model',
              data=results,
              hue='Model',
              palette='viridis',
              legend=False
         ax.grid(False)
         # Add R<sup>2</sup> value labels on each bar
         for container in ax.containers:
              for bar in container:
                  width = bar.get_width()
                  y_pos = bar.get_y() + bar.get_height() / 2
                  ax.text(width + 0.01, y_pos, f"{width:.2f}", va='center', ha='left', fontsi
         ax.tick_params(axis='both', labelsize=14)
         plt.title('R2 Scores of Different Models', fontsize=20)
         plt.xlabel('R2 Score', fontsize=20)
         plt.ylabel('Model', fontsize = 20)
         plt.xlim(0,1)
         plt.tight_layout()
         plt.show()
```



```
In [27]: # y_test vs y_pred_rf
         import matplotlib.pyplot as plt
         import seaborn as sns
         import numpy as np
         # Create scatter plot
         plt.figure(figsize=(8, 6))
         sns.scatterplot(x=y_test, y=y_pred_rf, color="steelblue", alpha=0.6, edgecolor='k')
         # Plot y = x line for reference
         max_val = max(max(y_test), max(y_pred_rf))
         min_val = min(min(y_test), min(y_pred_rf))
         plt.plot([min_val, max_val], [min_val, max_val], 'r--', label='R2 = 1')
         # Labels and title
         plt.xlabel("Actual Viability Values")
         plt.ylabel("Predicted Viability Values by Random Forest")
         plt.title(f"Random Forest Regression: R2 = {r2_rf:.2f} ")
         plt.legend()
         # Set 1:1 aspect ratio
         plt.gca().set_aspect('equal', adjustable='box')
         plt.tight_layout()
         plt.show()
```

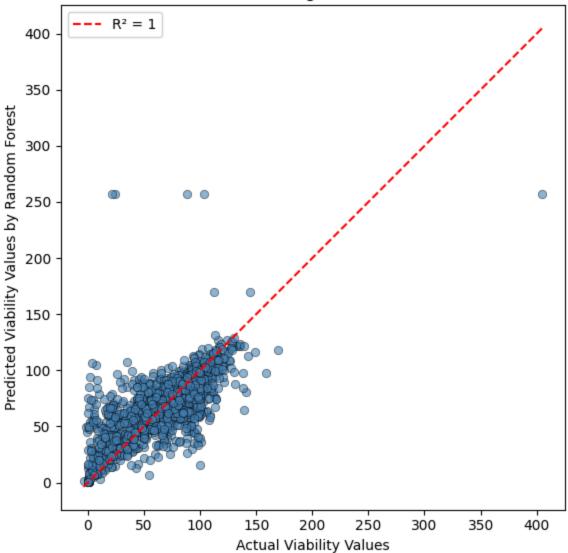
Random Forest Regression: $R^2 = 0.58$



```
In [28]:
        # The new RF seems to have less wrong predictions and removed the incorrect ones at
         # Plot the old RF (without tuning) vs y_test
         import matplotlib.pyplot as plt
         import seaborn as sns
         import numpy as np
         # Create scatter plot
         plt.figure(figsize=(8, 6))
         sns.scatterplot(x=y_test, y=y_pred_rf_old, color="steelblue", alpha=0.6, edgecolor=
         # Plot y = x line for reference
         max_val = max(max(y_test), max(y_pred_rf_old))
         min_val = min(min(y_test), min(y_pred_rf_old))
         plt.plot([min_val, max_val], [min_val, max_val], 'r--', label='R2 = 1')
         # Labels and title
         plt.xlabel("Actual Viability Values")
         plt.ylabel("Predicted Viability Values by Random Forest")
         plt.title(f"Random Forest Regression: R2 = {r2_rf_old:.2f} ")
```

```
plt.legend()
# Set 1:1 aspect ratio
plt.gca().set_aspect('equal', adjustable='box')
plt.tight_layout()
plt.show()
```

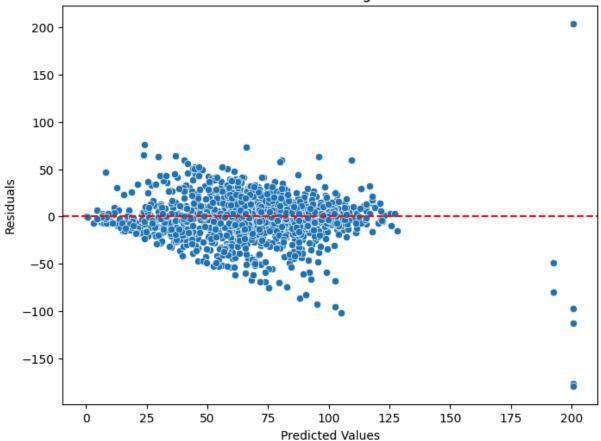
Random Forest Regression: $R^2 = 0.55$



```
In [29]: # Residuals plot for Random Forest (new model)

residuals = y_test - y_pred_rf
plt.figure(figsize=(8, 6))
sns.scatterplot(x=y_pred_rf, y=residuals)
plt.axhline(0, color='red', linestyle='--')
plt.title(f'Residuals Random Forest Regression: R2 = {r2_rf:.2f}')
plt.xlabel('Predicted Values')
plt.ylabel('Residuals')
plt.show()
```





```
In [30]: # Random Forest feature importance
import pandas as pd
import matplotlib.pyplot as plt

importances = pd.Series(best_rf.feature_importances_, index=X_train_scaled_df.colum
top_importances = importances.sort_values(ascending=False).head(10)

print("Top 10 Feature Importances:")
print(top_importances)

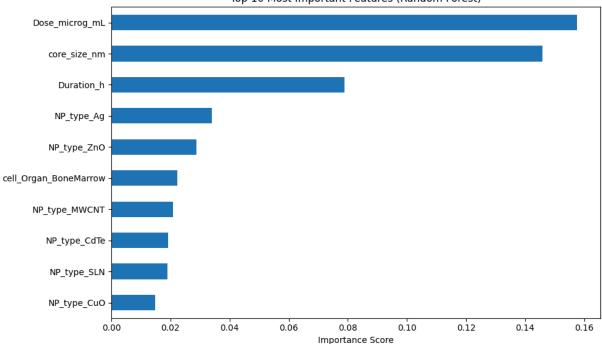
plt.figure(figsize=(10, 6))
top_importances.plot(kind='barh')
plt.gca().invert_yaxis() # Most important on top
plt.title("Top 10 Most Important Features (Random Forest)")
plt.xlabel("Importance Score")
plt.tight_layout()
plt.show()
```

```
Top 10 Feature Importances:
```

```
Dose_microg_mL
                         0.157596
                         0.145883
core_size_nm
Duration_h
                         0.078924
NP_type_Ag
                         0.033921
NP_type_Zn0
                         0.028773
cell_Organ_BoneMarrow
                        0.022332
NP_type_MWCNT
                         0.020844
NP type CdTe
                        0.019134
NP_type_SLN
                         0.018888
                         0.014804
NP_type_Cu0
```

dtype: float64

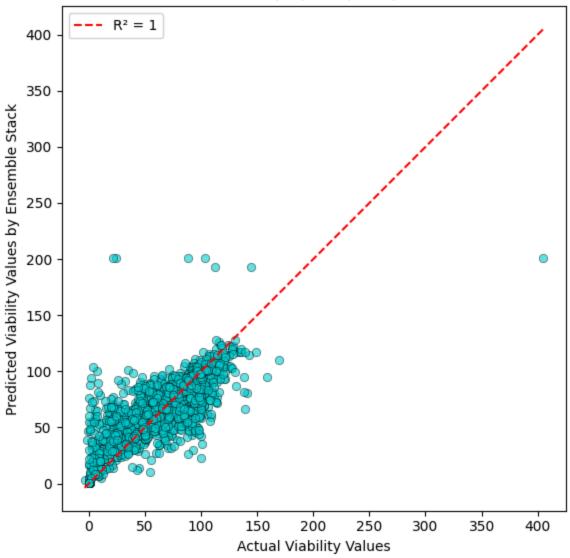




```
In [31]:
        # Plot the stack predictions vs y_test
         import matplotlib.pyplot as plt
         import seaborn as sns
         import numpy as np
         # Create scatter plot
         plt.figure(figsize=(8, 6))
         sns.scatterplot(x=y_test, y=stack_preds, color="darkturquoise", alpha=0.6, edgecolo
         # Plot y = x line for reference
         max_val = max(max(y_test), max(stack_preds))
         min_val = min(min(y_test), min(stack_preds))
         plt.plot([min_val, max_val], [min_val, max_val], 'r--', label='R2 = 1')
         # Labels and title
         plt.xlabel("Actual Viability Values")
         plt.ylabel("Predicted Viability Values by Ensemble Stack")
         plt.title(f"Ensemble Stack (RF, KNN, MLP): R² = {r2_stack:.2f} ")
         plt.legend()
         # Set 1:1 aspect ratio
         plt.gca().set_aspect('equal', adjustable='box')
```

```
plt.tight_layout()
plt.show()
```

Ensemble Stack (RF, KNN, MLP): $R^2 = 0.58$



```
In [32]: # Residuals plot for Stack

residuals = y_test - stack_preds
plt.figure(figsize=(8, 6))
sns.scatterplot(x=stack_preds, y=residuals, color="darkturquoise", alpha=0.6, edgec
plt.axhline(0, color='red', linestyle='--')
plt.title(f'Ensemble Stack (RF, KNN, MLP): R² = {r2_stack:.2f}')
plt.xlabel('Predicted Values')
plt.ylabel('Residuals')
plt.show()
```

Ensemble Stack (RF, KNN, MLP): $R^2 = 0.58$

