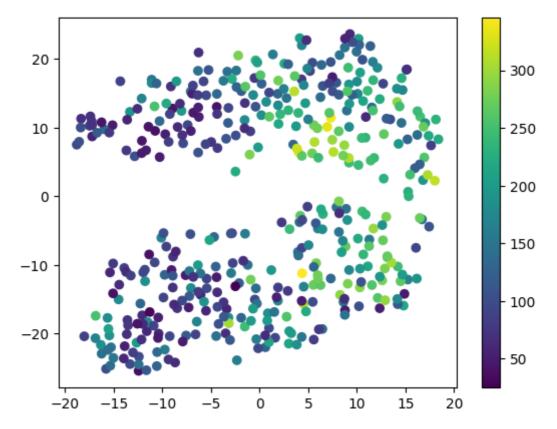
4

```
In [ ]: from IPython.display import Latex
        import numpy as np
        import matplotlib.pyplot as plt
In [ ]: #Import dataset
        import sklearn
        from sklearn import datasets
        diabetes = sklearn.datasets.load_diabetes()
In [ ]: display(Latex(r"\newpage"))
        \newpage
In [ ]: #4.a
        from sklearn.manifold import TSNE
        x = diabetes["data"]
        y = diabetes["target"]
        d = 2
        tsne = TSNE(d)
        tsne_result = tsne.fit_transform(x)
        points = plt.scatter(tsne_result[:, 0], tsne_result[:,1], c = y, cmap = 'viridis')
        plt.colorbar(points)
        c:\Users\elias\AppData\Local\Programs\Python\Python310\lib\site-packages\sklearn\m
        anifold\_t_sne.py:800: FutureWarning: The default initialization in TSNE will chan
        ge from 'random' to 'pca' in 1.2.
          warnings.warn(
        c:\Users\elias\AppData\Local\Programs\Python\Python310\lib\site-packages\sklearn\m
        anifold\_t_sne.py:810: FutureWarning: The default learning rate in TSNE will chang
        e from 200.0 to 'auto' in 1.2.
          warnings.warn(
```

Out[]: <matplotlib.colorbar.Colorbar at 0x1cff409fca0>



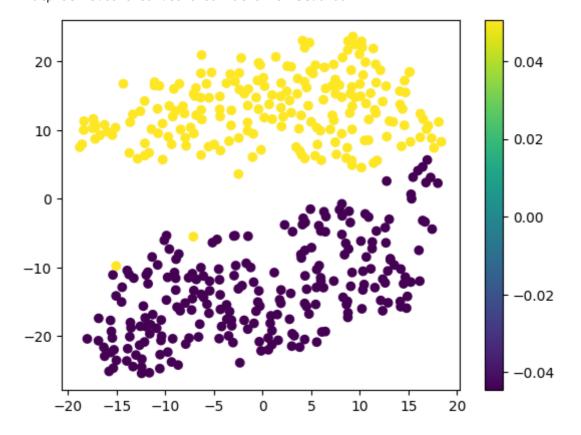
In []: display(Latex(r"\newpage"))

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4.b The feature is sex.

In []: points = plt.scatter(tsne_result[:, 0], tsne_result[:,1], c = x[:,1], cmap = 'viric
 plt.colorbar(points)

Out[]: <matplotlib.colorbar.Colorbar at 0x1cff3ed0460>



```
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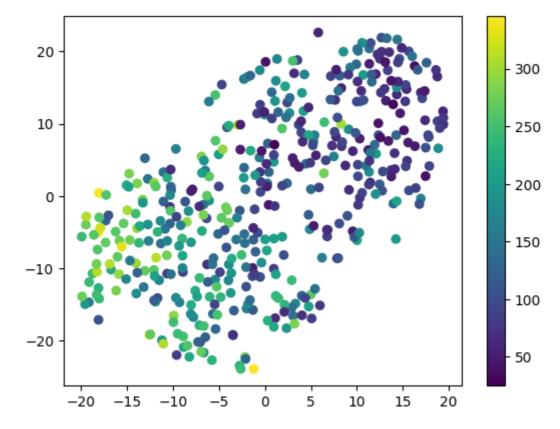
```
In []: #4.c
    new_x = np.delete(diabetes["data"], axis = 1, obj = 1)
    tsne = TSNE(d)

new_result = tsne.fit_transform(new_x)

points = plt.scatter(new_result[:, 0], new_result[:,1], c = y, cmap = 'viridis')
    plt.colorbar(points)

c:\Users\elias\AppData\Local\Programs\Python\Python310\lib\site-packages\sklearn\m
    anifold\_t_sne.py:800: FutureWarning: The default initialization in TSNE will chan
    ge from 'random' to 'pca' in 1.2.
    warnings.warn(
    c:\Users\elias\AppData\Local\Programs\Python\Python310\lib\site-packages\sklearn\m
    anifold\_t_sne.py:810: FutureWarning: The default learning rate in TSNE will chang
    e from 200.0 to 'auto' in 1.2.
    warnings.warn(
```

Out[]: <matplotlib.colorbar.Colorbar at 0x1cff3f6ffd0>



We do not see any two clear clusters anymore since tsne does not differentiate on sex. In other words, most of the data points got a lot of their variance from sex meaning females and males are quite similar besides being of different genders. The fact that sex was included ended up with "pulling" the data apart and forming two clusters.

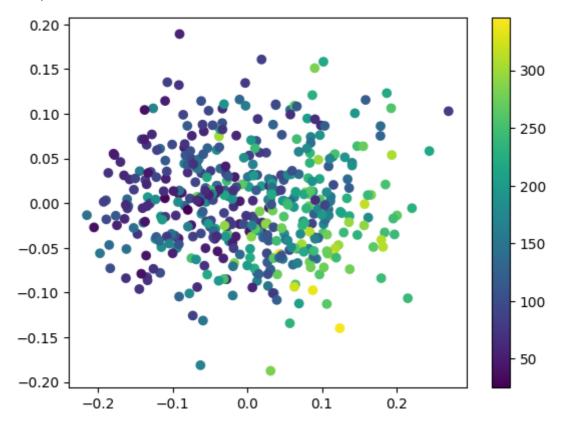
If the change did not occur and we did still see two clear clusters, it would mean that there would be another feature which was very binary in its distribution. If age was split in above 50 or below 50 this would also form two clusters.

```
In [ ]: display(Latex(r"\newpage"))
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```
In []: #4.d
    from sklearn.decomposition import PCA
    pca = PCA(d)
    pca_result = pca.fit_transform(x)
    points = plt.scatter(pca_result[:, 0], pca_result[:,1], c = y, cmap = "viridis")
    plt.colorbar(points)
```

Out[]: <matplotlib.colorbar.Colorbar at 0x1cff42b4220>



```
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```
In [ ]: #4.e
        def MSE(y_hat, y_true):
             return (1/y_hat.shape[0])*np.sum((y_hat - y_true)**2)
        X_{train} = x[:100]
        y_{train} = y[:100]
        X_{\text{test}} = x[100:]
        y_{test} = y[100:]
        X_train_mtx = np.hstack([X_train, np.ones((100,1))])
        X_test_mtx = np.hstack([X_test, np.ones((342, 1))])
        w_ols = np.linalg.inv(X_train_mtx.T @ X_train_mtx) @ X_train_mtx.T @ y_train
        y_hat = X_test_mtx@w_ols
        test_mse = MSE(y_hat, y_test)
        print(f"The test MSE is: {test_mse}")
        def c_index(y_hat, y_test):
             nr conc = 0
             nr_disc = 0
```

```
for i in range(y_hat.shape[0]):
        for j in range(y test.shape[0]):
            if i == j:
                continue
            else:
                y_test_i = y_test[i]
                y_test_j = y_test[j]
                y_hat_i = y_hat[i]
                y_{hat_j} = y_{hat_j}
                if y_test_i > y_test_j and y_hat_i > y_hat_j:
                    nr_conc += 1
                elif y_test_i > y_test_j and y_hat_i < y_hat_j:</pre>
                    nr_disc += 1
    return nr_conc/(nr_conc+nr_disc)
print(f"The c index is: {c_index(y_hat, y_test)}")
The test MSE is: 3430.9233826005243
```

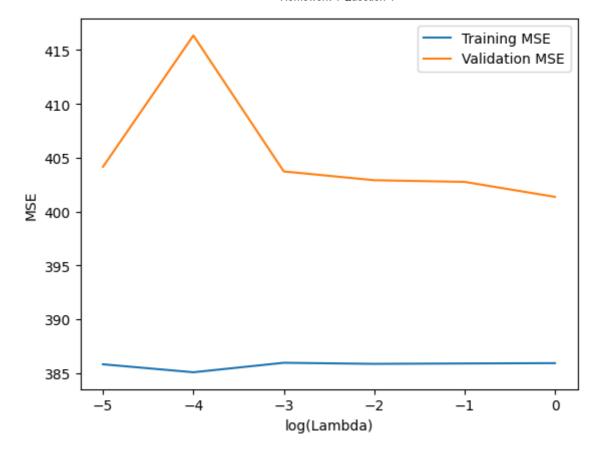
The test MSE is: 3430.9233826005243 The c index is: 0.7452930850514576

```
In [ ]: display(Latex(r"\newpage"))
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```
In [ ]: #4.f
        from sklearn.linear_model import Ridge
        def cross_validation(X, y, k, model, loss_calculator):
            n = X.shape[0]
            idx = np.random.permutation(n)
            X_{shuffled} = X[idx]
            y_shuffled = y[idx]
            size = n // k
            validation_loss = np.zeros((n,))
            training_loss = np.zeros((n,))
            for i in range(k):
                start = i*size
                end = (i+1)*size
                X test = X shuffled[start:end]
                y_test = y_shuffled[start:end]
                X_train = np.vstack([X_shuffled[:start], X_shuffled[end:]])
                y_train = np.hstack([y_shuffled[:start], y_shuffled[end:]])
                model.fit(X_train, y_train)
                training_loss[i] = loss_calculator(model.predict(X_train), y_train)
                validation_loss[i] = loss_calculator(model.predict(X_test), y_test)
            return np.mean(validation_loss), np.mean(training_loss)
        lambdas = [10**-5, 10**-4, 10**-3, 10**-2, 10**-1, 1]
        validation MSE = []
        training_MSE = []
        for 1 in lambdas:
```

```
validation_MSE_current, training_MSE_current = cross_validation(X_train, y_trai
    validation_MSE.append(validation_MSE_current)
    training MSE.append(training MSE current)
    print(f"Lambda={1} has validation loss={validation_MSE_current} and training log
   print("")
plt.plot(np.log10(lambdas), training_MSE, label='Training MSE')
plt.plot(np.log10(lambdas), validation MSE, label='Validation MSE')
plt.legend()
plt.xlabel('log(Lambda)')
plt.ylabel('MSE')
best_lambda = lambdas[np.argmin(validation_MSE)]
ridge_best_lambda = Ridge(alpha=best_lambda).fit(X_train,y_train)
ridge y hat = ridge best lambda.predict(X test)
ridge_mse = MSE(ridge_y_hat, y_test)
print(f"The best lambda was {best_lambda}.\nThis lambda gave MSE with ridge = {ridg
Lambda=1e-05 has validation loss=404.15799243503966 and training loss=385.80102276
90261
Lambda=0.0001 has validation loss=416.3534904067575 and training loss=385.06213726
177435
Lambda=0.001 has validation loss=403.72113028383114 and training loss=385.94238469
552073
Lambda=0.01 has validation loss=402.9057175061709 and training loss=385.8414729654
401
Lambda=0.1 has validation loss=402.7489796064022 and training loss=385.87803508971
024
Lambda=1 has validation loss=401.3585709192019 and training loss=385.9097279767526
The best lambda was 1.
This lambda gave MSE with ridge = 5039.062537574326.
```



In []: