

Study the effect of pH levels by Buffer type

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11:23 AM

Buffer Type label correspondence with actual types

0: Ace (Acetate)

1: His (Histidine)

2: Cit (Citrate)

3: Phos (Phosphate)

Additive_label Correspondence

0: Arginine, solid shape

1: Glycine, hollow shape

Acetate

Buffer_Type_label	0.0
Sugar_Salt_label	0.0
Additive_label	1.0
pH	4.5
task_ind	0.0

Histidine

Buffer_Type_label	1.0
Sugar_Salt_label	0.0
Additive_label	1.0
pH	6.5
task_ind	1.0

Cit (Citrate)

Buffer_Type_label	2.0
Sugar_Salt_label	0.0
Additive_label	0.0
pH	5.0
task_ind	1.0

Phos (Phosphate)

Buffer_Type_label	3.0
Sugar_Salt_label	0.0
Additive_label	0.0
pH	7.0
task_ind	1.0

Model accuracy affect pattern

Use original scale for embedding 3

pH level and target: linear? Model accuracy, use another model

Acetate: 3.7-5.7

Histidine: 5-7

Citrate: 3.7-5.7

Phosphate: 6.2-8.2

Citrate, Phosphate: why choose Arginine

Citrate

```
np.mean(new_train_mean[df_X_train_buffer['Additive_label']==1])  
91.14
```

```
np.mean(new_train_mean[df_X_train_buffer['Additive_label']==0])  
91.61
```

Phosphate

```
np.mean(new_train_mean[df_X_train_buffer['Additive_label']==1])  
90.82
```

```
np.mean(new_train_mean[df_X_train_buffer['Additive_label']==0])  
90.56
```

MK-6204 dataset

Ignore glycine

If excipients column has Control: didn't add any excipient

NR_CE_SDS: device

Target properties: add up to 100

NT: non-tested

Keep Sugar and Additive as the same with optimal training data solution

```
=====Ace=====  
==Among training data==  
Buffer_Type_label    0.0  
Sugar_Salt_label     0.0  
Additive_label       1.0  
pH                   4.5  
task_ind             0.0  
Name: 1, dtype: float64
```

```
==After simulation==  
Buffer_Type_label    0.0  
Sugar_Salt_label     0.0  
Additive_label       1.0  
pH                   3.8  
task_ind             0.0  
Name: 16, dtype: float64
```

```
=====His=====  
==Among training data==
```

```
====Among training data====
Buffer_Type_label    1.0
Sugar_Salt_label     0.0
Additive_label       1.0
pH                   6.5
task_ind             1.0
Name: 35, dtype: float64
```

```
==After simulation==
Buffer_Type_label    1.0
Sugar_Salt_label     0.0
Additive_label       1.0
pH                   5.0
task_ind             2.0
```

```
Name: 50, dtype: float64
=====Cit=====
====Among training data====
Buffer_Type_label    2.0
Sugar_Salt_label     0.0
Additive_label       0.0
pH                   5.0
task_ind             1.0
Name: 36, dtype: float64
```

```
==After simulation==
Buffer_Type_label    2.0
Sugar_Salt_label     0.0
Additive_label       0.0
pH                   3.7
task_ind             1.0
Name: 38, dtype: float64
```

```
=====Phos=====
====Among training data====
Buffer_Type_label    3.0
Sugar_Salt_label     0.0
Additive_label       0.0
pH                   7.0
task_ind             1.0
Name: 46, dtype: float64
```

```
==After simulation==
Buffer_Type_label    3.0
Sugar_Salt_label     0.0
Additive_label       0.0
pH                   6.2
task_ind             1.0
Name: 38, dtype: float64
```