## Combined Scores

February 10, 2023

```
[1]: import pandas as pd
     import cobra
[2]: model = cobra.io.load_json_model('iJN1463_IPP_bypass.json')
        Load EMA-based and Opt-based targets
[3]: EMA_score = pd.read_csv('EMA-based_targets.csv', index_col=0)
     EMA_score.sort_values('EMA-based_Score', ascending=False).head()
[3]:
                  cMCS
                        EFM tool EMA-based Score
     GLCDpp
              0.671667
                             0.0
                                          0.636194
    PDH
              0.616111
                             0.0
                                          0.583573
    LDH_D
              0.500000
                             1.0
                                          0.526406
     ACACT1r
                             0.0
                                          0.497778
              0.497778
    PPS
                             1.0
              0.450000
                                          0.479047
[4]: Opt_score = pd.read_csv('Opt-based_targets.csv', index_col=0)
     Opt_score.sort_values('Opt-based_Score', ascending=False).head()
[4]:
              OptKnock
                        OptKnock_blocked_exchanges
                                                     Opt-based_Score
              0.958333
                                           0.885932
                                                            0.922132
     ACt2rpp
     CS
              0.758333
                                           0.908745
                                                            0.833539
                                           0.437262
                                                            0.697798
     HMGL
              0.958333
     BDH
              0.966667
                                           0.395437
                                                            0.681052
     TPI
              0.950000
                                           0.403042
                                                            0.676521
[5]: Combined = EMA score[['EMA-based Score']].join(Opt_score['Opt-based Score'],_
      ⇔how='outer')
     Combined.head()
[5]:
                                   Opt-based_Score
                  EMA-based_Score
     1PPDCRc
                         0.011111
                                           0.001901
     3HAACOAT100
                         0.055556
                                                NaN
     3HAACOAT120
                         0.055556
                                                NaN
                         0.055556
     3HAACOAT121
                                           0.004167
     3HAACOAT140
                              NaN
                                           0.007969
```

## 2 Filter essential reactions, transport reactions, and reactions without genes

```
[6]: gene_essentiality = pd.read_table('Gene_Essentiality.txt', index_col=0)
     gene essentiality.head()
 [6]:
               Proteins
                          RbTnSeq Data
     Genes
     PP_s0001 PP_s0001
                           spontaneous
     PP 3591
                   dpkA Non Essential
     PP_1257
                PP_1257 Non Essential
     PP 2136
                   fadB Non Essential
     PP 2008
                   fadH Non Essential
 [7]: essential_reactions = [x for x in Combined.index if ((x in model.reactions) and
                            any(gene_essentiality.loc[g.id, 'RbTnSeq Data'] ==__
       ⇔'Essential' for g in
                                model.reactions.get_by_id(x).genes if g.id in_
       ⇒gene_essentiality.index))]
      Combined.drop(essential_reactions, inplace=True)
 [8]: transporters = [x for x in Combined.index if ((x in model.reactions) and
                     (all(c in model.reactions.get_by_id(x).compartments for c in_
       all(c in model.reactions.get_by_id(x).compartments for c in_{\sqcup}
       Combined.drop(transporters, inplace=True)
 [9]: reactions_without_genes = [x for x in Combined.index if ((x in model.reactions)_
                                not model.reactions.get_by_id(x).genes)]
     Combined.drop(reactions_without_genes, inplace=True)
         Rank targets for each method and choose top priority targets
[10]: Combined_rank = Combined.rank(ascending=False)
     Combined rank['Combined rank'] = Combined rank.mean(axis=1, skipna=False).rank()
     Combined_rank.sort_values(by='Combined_rank', inplace=True)
     Top 10 targets by both EMA-based and Opt-based methods
[11]: Combined_rank_top10 = Combined_rank[(Combined_rank['EMA-based_Score'] <= 10) &__
       ⇔(Combined_rank['Opt-based_Score'] <= 10)]</pre>
      Combined rank top10
```

```
[11]: EMA-based_Score Opt-based_Score Combined_rank
HMGL 9.0 2.0 1.5
GND 5.0 6.0 1.5
BDH 9.0 3.0 3.0
```

## Top targets by EMA-based method, but not top 10 by Opt-based method

[12]: Combined\_rank.sort\_values(by='EMA-based\_Score').drop(Combined\_rank\_top10.

index)[0:5]

[12]:	EMA-based_	Score Opt-b	ased_Score	Combined_rank
LDH	H_D	1.0	NaN	NaN
ACA	ACT1r	2.0	21.0	6.0
PPS	3	3.0	91.0	28.0
KAT	Γ1	4.0	26.0	10.0
PC		6.0	44.0	16.0

## Top targets by Opt-based method, but not top 10 by EMA-based method

[13]: Combined\_rank.sort\_values(by='Opt-based\_Score').drop(Combined\_rank\_top10.

[13]:		EMA-based_Score	Opt-based_Score	Combined_rank
	CS	19.0	1.0	5.0
	TPI	21.0	4.0	7.5
	PGK	12.0	5.0	4.0
	GARFT	34.0	7.0	15.0
	ICL	57.0	8.0	18.0