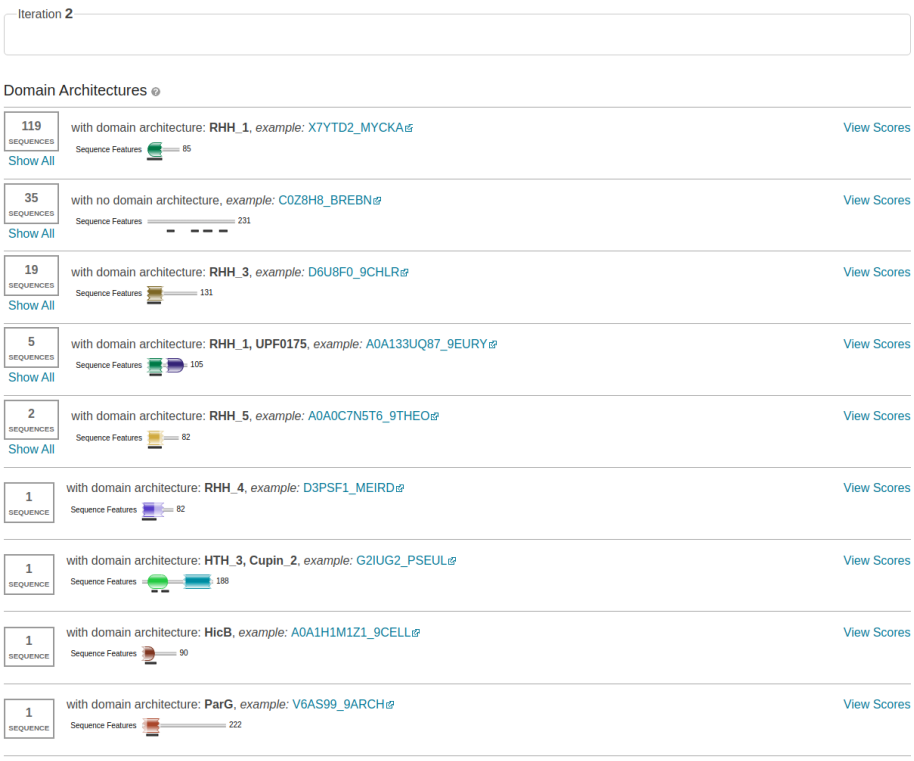
Iteration 1

- RHH_1 - 12 Sequences LINK

Iteration 2

- RHH_1 - 119 Sequences LINK
- No architecture - 35 Sequences LINK
- RHH_3 - 19 Sequences LINK
- RHH_1 UPF0175 - 5 Sequences LINK
- RHH_5 - 2 Sequences LINK



Iteration 3

- RHH_1 - 878 Sequences LINK
- No architecture - 313 Sequences LINK
- RHH_3 - 67 Sequences LINK
- HicB - 43 Sequences LINK
- RHH_5 - 16 Sequences LINK
- RHH_4 - 14 Sequences LINK
- Arc - 14 Sequences LINK
- MobC - 7 Sequences LINK
- RHH_1 UPF0175 - 7 Sequences LINK
- TraY - 6 Sequences LINK
- RHH_1 NikR_C - 5 Sequences LINK
- PSK_trans_fac - 5 Sequences LINK
- ParG - 4 Sequences LINK
- CopG_antitoxin - 3 Sequences LINK
- RHH_1 RHH_1 - 3 Sequences LINK
- ParD - 2 Sequences LINK
- RNase_H RHH_1 - 2 Sequences LINK
- RNase_H - 2 Sequences LINK
- VapB_antitoxin - 2 Sequences LINK
- HTH_26 - 2 Sequences LINK

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Iteration **3**

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Domain Architectures

878 SEQUENCES — with domain architecture: **RHH_1**, *example:* A0A1X1YG79_9MYCO — View Scores
Sequence Features 85
Show All

313 SEQUENCES — with no domain architecture, *example:* A0A0X8D3S3_9BACL — View Scores
Sequence Features 206
Show All

67 SEQUENCES — with domain architecture: **RHH_3**, *example:* D6TXJ0_9CHLR — View Scores
Sequence Features 64
Show All

43 SEQUENCES — with domain architecture: **HicB**, *example:* A0A1H1M1Z1_9CELL — View Scores
Sequence Features 90
Show All

16 SEQUENCES — with domain architecture: **RHH_5**, *example:* A0A0C7N5T6_9THEO — View Scores
Sequence Features 82
Show All

14 SEQUENCES — with domain architecture: **RHH_4**, *example:* D3PSF1_MEIRD — View Scores
Sequence Features 82
Show All

14 SEQUENCES — with domain architecture: **Arc**, *example:* A0A372BWT9_9GAMM — View Scores
Sequence Features Arc 77
Show All

7 SEQUENCES — with domain architecture: **MobC**, *example:* A0A1Y4CF89_9BACT — View Scores
Sequence Features 102
Show All

7 SEQUENCES — with domain architecture: **RHH_1, UPF0175**, *example:* A0A133UQ87_9EURY — View Scores
Sequence Features 105
Show All

6 SEQUENCES — with domain architecture: **TraY**, *example:* A0A2L0W7A2_9RHIZ — View Scores
Sequence Features 75
Show All

5 SEQUENCES — with domain architecture: **RHH_1, NikR_C**, *example:* D2RG21_ARCPA — View Scores
Sequence Features 126
Show All

5 SEQUENCES — with domain architecture: **PSK_trans_fac**, *example:* A0A0F0GJP8_LENAE — View Scores
Sequence Features 68
Show All

4 SEQUENCES — with domain architecture: **ParG**, *example:* A0A073CA55_PLAAG — View Scores
Sequence Features ParG 81
Show All

3 SEQUENCES — with domain architecture: **CopG_antitoxin**, *example:* A0A3S8U4Y7_9RHOB — View Scores
Sequence Features 84
Show All

3 SEQUENCES — with domain architecture: **RHH_1, RHH_1**, *example:* D5IEY9_9VIRU — View Scores
Sequence Features 140
Show All

2 SEQUENCES — with domain architecture: **ParD**, *example:* A0A1H9QU33_9PROT — View Scores
Sequence Features ParD 83
Show All

2 SEQUENCES — with domain architecture: **RNase_H, RHH_1**, *example:* B0RID9_CLAMS — View Scores
Sequence Features RNase_H 251
Show All

2 SEQUENCES — with domain architecture: **RNase_H**, *example:* A0A4Q9GTZ5_9MICO — View Scores
Sequence Features RNase_H 225
Show All

2 SEQUENCES — with domain architecture: **VapB_antitoxin**, *example:* A8UV44_9AQUI — View Scores
Sequence Features 72
Show All

2 SEQUENCES — with domain architecture: **HTH_26**, *example:* A0A4P6JYT6_9CHLR — View Scores
Sequence Features 73
Show All

1 SEQUENCE — with domain architecture: **AMP-binding, AMP-binding_C, Methyltransf_12, PP-binding, Condensation, AMP-binding, AMP-binding_C, PP-binding, Condensation, AMP-binding, AMP-binding_C, Methyltransf_12, PP-binding, Condensation, AMP-binding, PP-binding, Condensation, AMP-binding, AMP-binding_C, PP-binding, Condensation, AMP-binding, AMP-binding_C, PP-binding, Condensation**, *example:* A0A1H4WEW4_9ACTN — View Scores
Sequence Features AMP-binding Condensation AMP-binding Condens

1 SEQUENCE — with domain architecture: **HTH_3, Cupin_2**, *example:* G2IUG2_PSEUL — View Scores
Sequence Features 188

1 SEQUENCE — with domain architecture: **LysM, LysM, LysM, LysM, LysM, Peptidase_M23**, *example:* W0DH41_9AQUI — View Scores
Sequence Features 458

1 — with domain architecture: **HicB, MobC**, *example:* A0A1X7BY16_9RHOB — View Scores