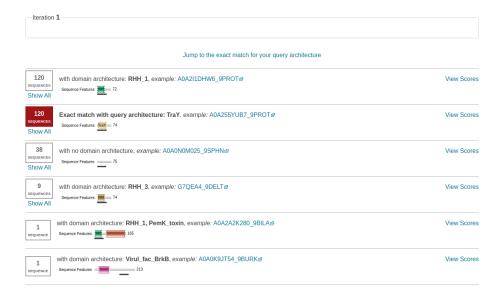
Iteration 1

- $\bullet~$ RHH_1 120 Sequences LINK
- TraY 120 Sequences LINK
- No architecture 38 Sequences LINK
- RHH_3 9 Sequences LINK



Iteration 2

- RHH_1 874 Sequences LINK
- No architecture 273 Sequences LINK
- TraY 122 Sequences LINK
- PRODH Pro_dh-DNA_bdg Pro_dh Aldedh 31 Sequences LINK
- $\bullet~$ RHH_3 21 Sequences LINK
- PSK_trans_fac 8 Sequences LINK
- RHH_5 7 Sequences LINK
- Arc 6 Sequences LINK
- HicB 6 Sequences LINK
- HTH_IclR IclR 3 Sequences LINK
- MobC 2 Sequences LINK
- RHH_1 PRODH Pro_dh-DNA_bdg Pro_dh Aldedh 2 Sequences LINK
- RHH_1 NikR_C 2 Sequences LINK

Iteration 3

- RHH_1 Next » Sequences LINK
- No architecture Last » Sequences LINK
- PRODH Pro_dh-DNA_bdg Pro_dh Aldedh 2079 Sequences LINK
- TraY 949 Sequences LINK
- HicB 292 Sequences LINK
- RHH_3 123 Sequences LINK
- RHH_5 77 Sequences LINK
- RHH_1 NikR_C 70 Sequences LINK
- Arc 69 Sequences LINK
- MobC 55 Sequences LINK
- DUF1778 34 Sequences LINK
- NIR_SIR_ferr NIR_SIR NIR_SIR_ferr NIR_SIR 25 Sequences LINK
- PSK_trans_fac 17 Sequences LINK
- ParG 15 Sequences LINK
- HicB_lk_antitox 13 Sequences LINK
- RHH_1 UPF0175 12 Sequences LINK
- YfbU 9 Sequences LINK
- RelB 8 Sequences LINK
- UPF0175 6 Sequences LINK
- CopG_antitoxin 5 Sequences LINK
- RHH_1 RHH_1 5 Sequences LINK
- DZR 5 Sequences LINK
- PRODH Pro_dh-DNA_bdg 4 Sequences LINK
- RHH_1 ParE_toxin 3 Sequences LINK
- RHH_4 3 Sequences LINK
- RHH_1 DZR 3 Sequences LINK
- ParE_toxin 3 Sequences LINK
- RHH_1 YfbU 3 Sequences LINK
- RHH_1 PRODH Pro_dh-DNA_bdg Pro_dh Aldedh 2 Sequences LINK

- NIR_SIR_ferr NIR_SIR NIR_SIR_ferr NIR_SIR ParG 2 Sequences LINK
- VapB_antitoxin 2 Sequences LINK
- HTH_IclR IclR 2 Sequences LINK
- HicB MobC 2 Sequences LINK
- RHH_1 AIRS 2 Sequences LINK
- RHH_3 NikR_C 2 Sequences LINK
- ParD 2 Sequences LINK
- PRODH Pro_dh-DNA_bdg Pro_dh 2 Sequences LINK
- Condensation AMP-binding AMP-binding_C PP-binding Condensation AMP-binding AMP-binding_C PP-binding Condensation AMP-binding Condensation AMP-binding Condensation AMP-binding_C PP-binding Condensation AMP-binding Methyltransf_12 PP-binding Condensation AMP-binding AMP-binding AMP-binding_C PP-binding Thioesterase Next » Sequences LINK
- HTH_Tnp_1 Last » Sequences LINK

omain ,	Architectures @	
874 SEQUENCES Show All	with domain architecture: RHH_1, example: Q39PP9_GEOMG@ Sequence Features 74	View Scores
273 EQUENCES	with no domain architecture, example: A0A1H3MQX7_9PROT@ Sequence Features	View Scores
122 EQUENCES	with domain architecture: TraY, example: D0L181_HALNC@ Sequence Features Inst_= 74	View Scores
31 EQUENCES	with domain architecture: PRODH, Pro_dh-DNA_bdg, Pro_dh, Aldedh, example: A0A366I5A8_9GAMMg Sequence Features	View Scores
21 EQUENCES	with domain architecture: RHH_3, example: V4J6I8_9DELT & Sequence Features 76	View Scores
8 EQUENCES	with domain architecture: PSK_trans_fac, example: A0A1I5HDY6_9PSEU@ Sequence Features 66	View Scores
7 EQUENCES	with domain architecture: RHH_5, example: A0A1Z4JFV8_LEPBY@ Sequence Features ====================================	View Scores
6 equences	with domain architecture: Arc, example: X5KJZ7_9MYCO@ Sequence Features 46	View Scores
6 EQUENCES	with domain architecture: HicB, example: GOFNL8_AMYMSig Sequence Features = 12	View Scores
3 EQUENCES	with domain architecture: HTH_IcIR, IcIR, example: Q89SX5_BRADUs6 Sequence Features	View Scores
2 EQUENCES	with domain architecture: MobC, example: D4X3K0_9BURK@ Sequence Features	View Score:
2 equences	with domain architecture: RHH_1, PRODH, Pro_dh-DNA_bdg, Pro_dh, Aldedh, example: D1NZS4_9GAMMe9 Sequence Features	View Scores
2 equences	with domain architecture: RHH_1, NikR_C, example: D2RG21_ARCPA@ Sequence Features	View Scores
1 EQUENCE	with domain architecture: YfbU, example: A0A097IJS8_9CORY@ Sequence Features	View Scores
1 EQUENCE	with domain architecture: Virul_fac_BrkB, example: A0A0K9JT54_9BURK@ Sequence Features = 213	View Scores
1 EQUENCE	with domain architecture: PRODH, Pro_dh-DNA_bdg, example: S6TZ56_PSESF@ Sequence Features	View Score
1 EQUENCE	with domain architecture: HTH_lcIR, example: A0A437M249_9PROT@ Sequence Features = 248	View Scores
1 EQUENCE	with domain architecture: HicB, MobC, example: A0A1X7BY16_9RHOBs9 Sequence Features	View Scores
1 EQUENCE	with domain architecture: UPF0175, example: A0A2B7GXD9_9EURY@ Sequence Features	View Scores
1 EQUENCE	with domain architecture: RHH_1, UPF0175, example: A0A166SQW6_9EURYØ	View Scores
1 EQUENCE	with domain architecture: DUF1778, example: A0A1V3Q8R3_9GAMMe/ Sequence Features ====================================	View Scores
1 EQUENCE	with domain architecture: RHH_1, ParE_toxin, example: A0A418UVB1_9ACTOr Sequence Features 25000000000000000000000000000000000000	View Scores
1 EQUENCE	with domain architecture: RHH_1, YfbU, example: A0A1G8MPG9_9MICC@ Sequence Features	View Scores
1 EQUENCE	with domain architecture: GIY-YIG, example: A0A2K8T3U9_9NOSO@ Sequence Features ====================================	View Scores
1 EQUENCE	with domain architecture: HNH_3, HTH_23, RHH_1, example: COQRT1_PERMH#9 Sequence Features	View Scores

with domain architecture: RHH_1, RHH_1, example: A0A125SJ70_9VIRUs

View Scores

