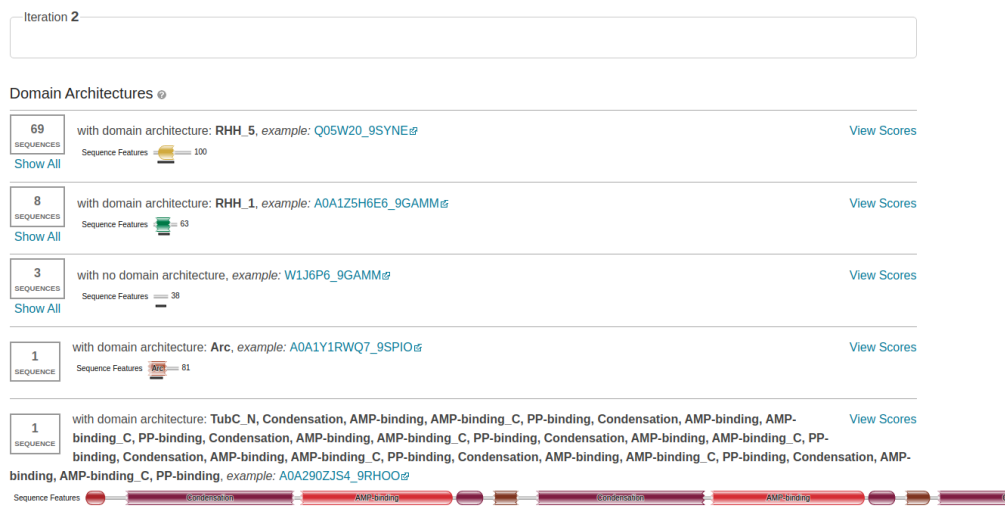


Iteration 1

- Run finished (Nothing more found or search took too long to finish)![Iteration 2](#)
 - RHH_5 - 69 Sequences [LINK](#)
 - RHH_1 - 8 Sequences [LINK](#)
 - No architecture - 3 Sequences [LINK](#)



Iteration 3

- RHH_5 - 328 Sequences [LINK](#)
- No architecture - 262 Sequences [LINK](#)
- RHH_1 - 178 Sequences [LINK](#)
- Arc - 60 Sequences [LINK](#)
- HicB - 56 Sequences [LINK](#)
- TraY - 20 Sequences [LINK](#)
- GIY-YIG RHH_5 - 4 Sequences [LINK](#)
- CopG_antitoxin - 2 Sequences [LINK](#)
- RHH_4 - 2 Sequences [LINK](#)
- HicB MobC - 2 Sequences [LINK](#)
- ParG - 2 Sequences [LINK](#)
- MobC - 2 Sequences [LINK](#)
- RepB-RCR_reg - 2 Sequences [LINK](#)
- Methyltransf_25 - 2 Sequences [LINK](#)

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