


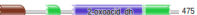



Iteration 1

- DUF2610 - 24 Sequences [LINK](#)
- DUF2610 Sel1 Sel1 Sel1 Sel1 Sel1 Sel1 Sel1 - 2 Sequences [LINK](#)

Iteration 1

Jump to the exact match for your query architecture

<div>24SEQUENCES</div> <div>Show All</div>	<div>Exact match with query architecture: DUF2610, example: A5CC80_ORITB</div> <div>Sequence Features  113</div>	<div>View Scores</div>
<div>2SEQUENCES</div> <div>Show All</div>	<div>with domain architecture: DUF2610, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, example: A0A369SKU3_9METZ</div> <div>Sequence Features  727</div>	<div>View Scores</div>
<div>1SEQUENCE</div>	<div>with no domain architecture, example: A0A1M7P688_9BACT</div> <div>Sequence Features  776</div>	<div>View Scores</div>
<div>1SEQUENCE</div>	<div>with domain architecture: Biotin_lipoyl, E3_binding, 2-oxoacid_dh, DUF2610, example: A0A2C9M548_BIOGL</div> <div>Sequence Features  475</div>	<div>View Scores</div>
<div>1SEQUENCE</div>	<div>with domain architecture: ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PS-DH, ADH_N, ADH_zinc_N_2, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PKS_DE, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PKS_DE, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PS-DH, ADH_N, ADH_zinc_N_2, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PS-DH, KR, PP-binding, example: A0A2L0F095_SORCE</div> <div>Sequence Features </div>	<div>View Scores</div>

Iteration 2

- DUF2610 - 27 Sequences [LINK](#)
- No architecture - 5 Sequences [LINK](#)
- DUF2610 Sel1 Sel1 Sel1 Sel1 Sel1 Sel1 Sel1 - 2 Sequences [LINK](#)



Iteration 3

- DUF2610 - 27 Sequences [LINK](#)
- No architecture - 23 Sequences [LINK](#)
- CHRD - 5 Sequences [LINK](#)
- DUF2610 Sel1 Sel1 Sel1 Sel1 Sel1 Sel1 Sel1 Sel1 - 2 Sequences [LINK](#)

JACKHMMER Results

Search Again

Score

Taxonomy

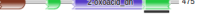
Domain

Download

Iteration 3

« previous iteration

Domain Architectures @

27 SEQUENCES Show All	with domain architecture: DUF2610 , example: A0A1M7QW60_9ACTN Sequence Features  96	View Scores
23 SEQUENCES Show All	with no domain architecture, example: A0A1M7P688_9BACT Sequence Features  776	View Scores
5 SEQUENCES Show All	with domain architecture: CHRD , example: A0A1G7WC00_9FLAO Sequence Features  645	View Scores
2 SEQUENCES Show All	with domain architecture: DUF2610, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1 , example: A0A369SKU3_9METZ Sequence Features  727	View Scores
1 SEQUENCE	with domain architecture: SLT, LysM, LysM, LysM, LysM, LysM, LysM, LysM , example: A0A1N7ELW2_9GAMM Sequence Features  907	View Scores
1 SEQUENCE	with domain architecture: Biotin_lipoyl, E3_binding, 2-oxoacid_dh, DUF2610 , example: A0A2C9M548_BIOGL Sequence Features  475	View Scores
1 SEQUENCE	with domain architecture: Sel1, Sel1, Sel1, Sel1, DUF2610 , example: A0A0B3SV27_9RHOB Sequence Features  457	View Scores
1 SEQUENCE	with domain architecture: Peptidase_C14, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, DUF2610 , example: A0A1H1JZ46_9RHIZ Sequence Features  897	View Scores
1 SEQUENCE	with domain architecture: ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PS-DH, ADH_N, ADH_zinc_N_2, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PKS_DE, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PKS_DE, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PS-DH, ADH_N, ADH_zinc_N_2, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PS-DH, KR, PP-binding , example: A0A2L0F095_SORCE Sequence Features 	View Scores
1 SEQUENCE	with domain architecture: Peptidase_C14, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, DUF2610 , example: A0A3D9Z1K3_9RHIZ Sequence Features  901	View Scores
1 SEQUENCE	with domain architecture: TPR_8 , example: I0K6Y0_9BACT Sequence Features  632	View Scores
1 SEQUENCE	with domain architecture: Peptidase_C14, TPR_2, TPR_1, TPR_12, TPR_11, DUF2610, TPR_16, TPR_8 , example: A0A4Q2UQC9_9BACT Sequence Features  1498	View Scores
1 SEQUENCE	with domain architecture: Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, DUF2610 , example: A0A1I2BU59_9BACT Sequence Features  549	View Scores
1 SEQUENCE	with domain architecture: Peptidase_C14, Sel1, Sel1, DUF2610 , example: A0A1X0T1C9_9RHIZ Sequence Features  779	View Scores
1 SEQUENCE	with domain architecture: ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PKS_DE, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PS-DH, ADH_N, ADH_zinc_N_2, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PKS_DE, KR, PP-binding, ketoacyl-synt, Ketoacyl-synt_C, KAsynt_C_assoc, Acyl_transf_1, PS-DH, KR, PP-binding, Thioesterase , example: A0A2L0ES13_SORCE Sequence Features 	View Scores
1 SEQUENCE	with domain architecture: Peptidase_C14, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, Sel1, DUF2610 , example: K9QB18_9NOSO Sequence Features  783	View Scores

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