

The dataset from *S. cerevisiae* composed of 91 true D modification site sequences (positive dataset) and 93 false D modification site sequences (negative dataset)

Positive dataset *S. cerevisiae*

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>P_S. cerevisiae_1
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>P_S. cerevisiae_2
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>P_S. cerevisiae_3
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>P_S. cerevisiae_4
UGGCCGAGCGGUCUAAGGCGCCU
>P_S. cerevisiae_5
GGCCGAGUGGUUAAGGCGAAAGA
>P_S. cerevisiae_6
GGUUGGCCGAGUGGUCUAAGGCG
>P_S. cerevisiae_7
GCCGAGUGGUCUAAGGCGGCAGA
>P_S. cerevisiae_8
UCUUGGCCCAGUUGGUUAAGGCA
>P_S. cerevisiae_9
GCCCAGUUGGUUAAGGCACCGUG
>P_S. cerevisiae_10
UAACGCGGGGAUCAGCGGUUCGA
>P_S. cerevisiae_11
UGGUUUAGUGGUAAAAUCCAACG
>P_S. cerevisiae_12
ACUUGGCCGAGUGGUUAAGGCGA
>P_S. cerevisiae_13
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>P_S. cerevisiae_14
UAAUGUGGAGAUCAUCGGUUCAA
>P_S. cerevisiae_15
CACGGCGAAGAUCGAGUUCGA
>P_S. cerevisiae_16
GGUCCAGUGGUUCAAGACGUCGC
>P_S. cerevisiae_17
GCGUGGCGUAAUGGCAACGCGUC
>P_S. cerevisiae_18
UAAUCAGAAGAUUAUGGGUUCGA
>P_S. cerevisiae_19
UGAUAGUUUAAUGGUCAGAAUGG
>P_S. cerevisiae_20
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>P_S. cerevisiae_21
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>P_S. cerevisiae_22
AGUGUAAACGGCUAUCACAUCACG
>P_S. cerevisiae_23
CCAUGGCCAAGUUGGUUAAGGCG
>P_S. cerevisiae_24
GCCAAGUUGGUUAAGGCGUGCGA
>P_S. cerevisiae_25
UAAUCGCAAGAUCGUGAGUUCAA
>P_S. cerevisiae_26
CCGUGGCGCAGUGGAAGCGCGCA
>P_S. cerevisiae_27
UAACCCUGAUGUCCUCGGAUCGA
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>P_S. cerevisiae_28
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>P_S. cerevisiae_31
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>P_S. cerevisiae_33
AUACCGAAAUGUCAGGGGUUCGA
>P_S. cerevisiae_34
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>P_S. cerevisiae_36
CAAUGGUCCAGUGGUUCAAGACG
>P_S. cerevisiae_37
CUAUAGUGUAGUGGUUAUCACUU
>P_S. cerevisiae_38
GAACCAGAAGAUUCCAGGUUCAA
>P_S. cerevisiae_39
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>P_S. cerevisiae_91
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Negative dataset_*S. cerevisiae*

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>N_S. cerevisiae_2
GCGCGCUCCCUUAGCAUGGGAGA
>N_S. cerevisiae_3
CUCCCUUAGCAUGGGAGAGGUCU
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UGGUAGCGCGCUUCCCUUGCAAG
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CGGCGUCUGGCUACGAACCAGAA
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