

The benchmark dataset S containing 550 RNA samples, of which 176 RNA sequence samples belong to the positive subset S^+ and 374 RNA sequence samples belong to the negative subset S^- .

Positive subset S^+

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>P_H. sapiens_4
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>P_H. sapiens_6
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>P_H. sapiens_7
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>P_H. sapiens_8
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Negative subset S⁻

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