

logo

NTDB id 30712 ACICU RS01365 WP 001160337.1

NTDB id 1044 HON27 RS16325 WP 206677389.1

consensus

MQFPLNSSLSHTIFRLGTYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTL LGYSLLSLVQLLCFKFIATQATRQLILFF

MQFPLNSSLSHTIFRLGTYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTL LGYSLLSLVQLLCFKFIATQATRQLILFF

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logo

NTDB id 30712 ACICU RS01365 WP 001160337.1

NTDB id 1044 HON27 RS16325 WP 206677389.1

consensus

IVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSALLAFLFFVHVG

IVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSALLAFLFFVHVG

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logo

NTDB id 30712 ACICU RS01365 WP 001160337.1

NTDB id 1044 HON27 RS16325 WP 206677389.1

consensus

IGQIAVQRFKLL EALTFHQSI ELYQLQNNRYILEQIEEGYLVLDENYDIVVSNPAACSLGIPPQFANEKYPLAKWHADLFEIL

IGQIAVQRFKLL EALTFHQSI ELYQLQNNRYILEQIEEGYLVLDENYDIVVSNPAACSLGIPPQFANEKYPLAKWHADLFEIL

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logo

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consensus

KFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLTLLILQDAQQINQQAQQLKLAALGQLSASIAHEIRNPLAAIVQANELLK

KFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLTLLILQDAQQINQQAQQLKLAALGQLSASIAHEIRNPLAAIVQANELLK

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logo

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consensus

DSDPEQQNTLRHIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLL EEDLFDVKHSIQLKISDSSLKFLFDEKQLRQVM

DSDPEQQNTLRHIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLL EEDLFDVKHSIQLKISDSSLKFLFDEKQLRQVM

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logo

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NTDB id 1044 HON27 RS16325 WP 206677389.1

consensus

INLVRNALRHNAPDSPIYITINHSQTNKIYIDVIDYGEVSKRDISQLFKPFFSTEINGTGLGLYL SHSFCEANHAKLTYVEQKQ

INLVRNALRHNAPDSPYITINHSQTNKIYIDVIDYGEVSKRDISQLFKPFFSTEINGTGLGLYL SHSFCEANHAKLTYVEQKQ

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logo

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consensus

GACFRIECPPIY

GACFRIECPPIY

!!!!!!!!!!!!!!!!

522522

 non conserved
 similar
 $\geq 50\%$ conserved