

logo

NTDB id 1015 ACIAD RS12005 WP 004928865.1

NTDB id 1083 HON27 RS04190 WP 168726476.1

NTDB id 110277 ACAK32 RS15100 WP 000472705.1

NTDB id 1070 ABD1 RS13145 WP 000472713.1

consensus



Sequence logo for motif 1. The logo shows the conservation of amino acids at each position. The consensus sequence is: **MFKIIILLGWIGGIALMGIDFPLIMQYEKVGEALLLLAFIFYLYKRPMFVDRPFLKAVFCLLCTTSLFMVGHYHAEKALIE**. The positions are numbered 1 to 30. The most conserved positions are 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30.

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logo

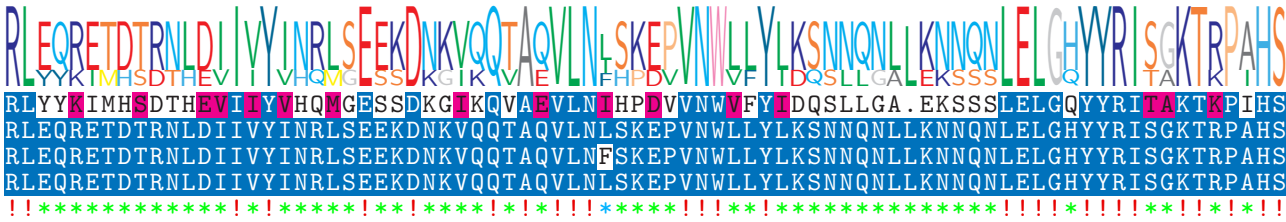
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consensus



Sequence logo for motif 2. The logo shows the conservation of amino acids at each position. The consensus sequence is: **RLEQRETDTNRNLDIIIVYNRLSEEKDNKVQQTAAQVLNLSKEPVNWLLYLKSNQNLKNNQNLLELGHYYRISGKTRPAHS**. The positions are numbered 1 to 30. The most conserved positions are 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30.

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consensus



Sequence logo for motif 3. The logo shows the conservation of amino acids at each position. The consensus sequence is: **YATPGAFDQEKWFIQRNIMSGFNVRYPLEPLSLDEIYRLGYQQHLKEQQSFSNSFRNLIEKLRLTFRQILNSSSLQQKGLI**. The positions are numbered 1 to 30. The most conserved positions are 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30.

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consensus



Sequence logo for motif 4. The logo shows the conservation of amino acids at each position. The consensus sequence is: **LALLTGDESLLSDEIQLQFQKLGISHLLAISGPHVLIFAAMVSWIFTRLTHRKPEIYLSIPQCKIVLPEFLMCVILYCA**. The positions are numbered 1 to 30. The most conserved positions are 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30.

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logo

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NTDB id 1070 ABD1 RS13145 WP 000472713.1

consensus



Sequence logo for motif 5. The logo shows the conservation of amino acids at each position. The consensus sequence is: **FVGFEIPALRTLLSYLFIQGLLLKQPIKPFILLVYSASLLLLMDPFSVLSAGFVLSYGACFILLRIYQTIQALPEQHFL**. The positions are numbered 1 to 30. The most conserved positions are 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30.

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logo

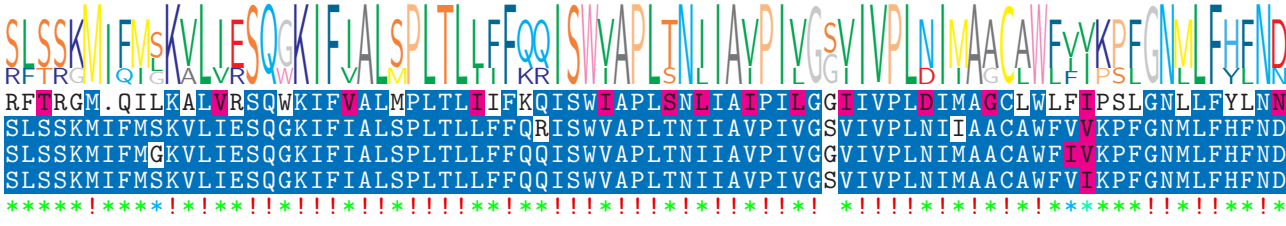
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NTDB id 110277 ACAK32 RS15100 WP 000472705.1

NTDB id 1070 ABD1 RS13145 WP 000472713.1

consensus



Sequence logo for motif 6. The logo shows the conservation of amino acids at each position. The consensus sequence is: **SLSSKMIFMSKVLIESQCKIFIALSPLTLLFFQQLSVIAPLTNIIAVPIVGSVIVPLNIIAACAWFVYKPEGNMIFHEND**. The positions are numbered 1 to 30. The most conserved positions are 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30.

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