

logo

NTDB id 1015 ACIAD RS12005 WP 004928865.1

NTDB id 1083 HON27 RS04190 WP 168726476.1

NTDB id 110522 ACAK33 RS09185 WP 000472717.1

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.

80
80
80
80

logo

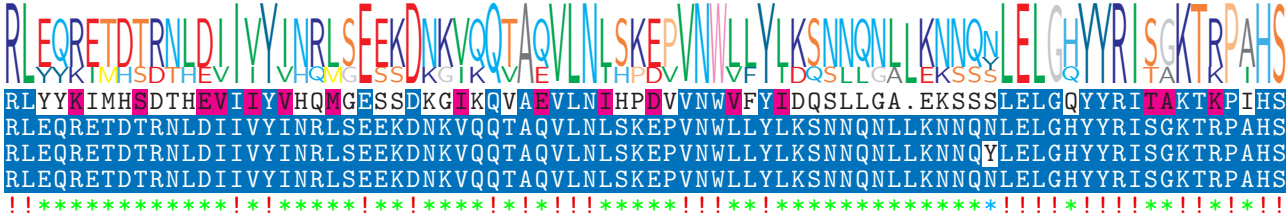
NTDB id 1015 ACIAD RS12005 WP 004928865.1

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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: **RLEQRETDRNLDIIIVYNRLSEEKDNKVQQTAAQVLNLSKEPVNWLLYLKSNQNLKNNQNLLELGHYYRISGKTRPAHS**. The positions are numbered 1 to 30.

159
160
160
160

logo

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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: **YATPGAFDQEKWFIQRNIMSGFNVRYPLEPLSLDEIYRLGYQQHLKEQQSFSSSFRNLIEKLRLTFRQILNSSSLQQKGLI**. The positions are numbered 1 to 30.

239
240
240
240

logo

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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.

319
320
320
320

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: **FVGFEIPALRTLLSYLIFIGELLKQPIKPFILLVYSASLLLLNDPFSVLSAGFVLSYGACFILIRIYQTTIAQLPEQHFL**. The positions are numbered 1 to 30.

399
400
400
400

logo

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NTDB id 110522 ACAK33 RS09185 WP 000472717.1

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: **SLSSKMIFMSKVLIESQCKIFIALSPLTLLFFQRIISWVAPLTNIIAVPIVGSVIVPLNIMAAWVFKPFGNMLFHFND**. The positions are numbered 1 to 30.

478
480
480
480

