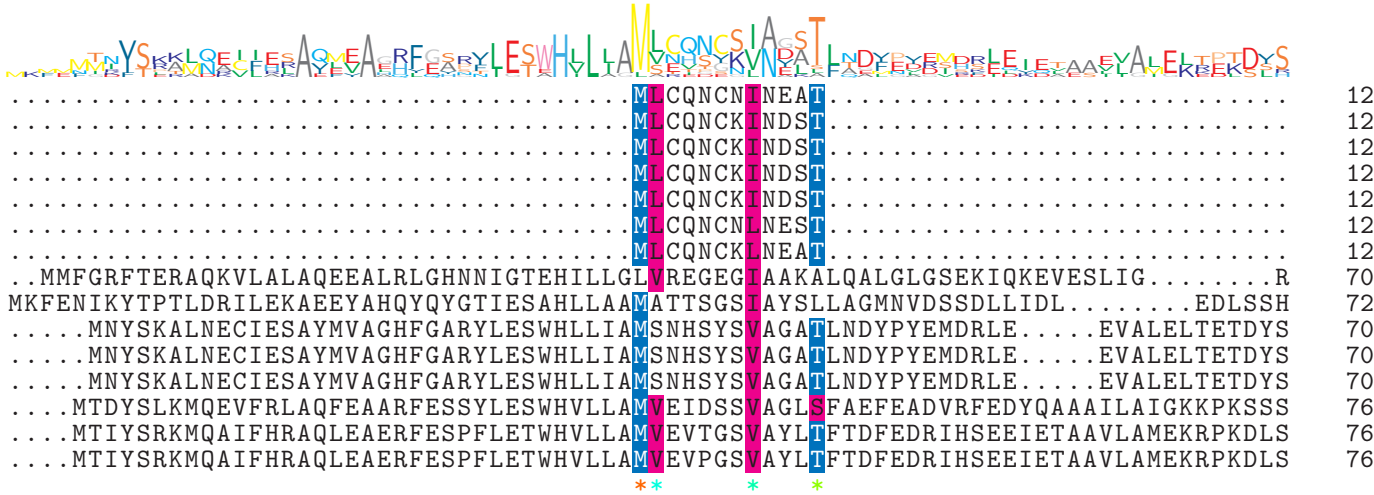


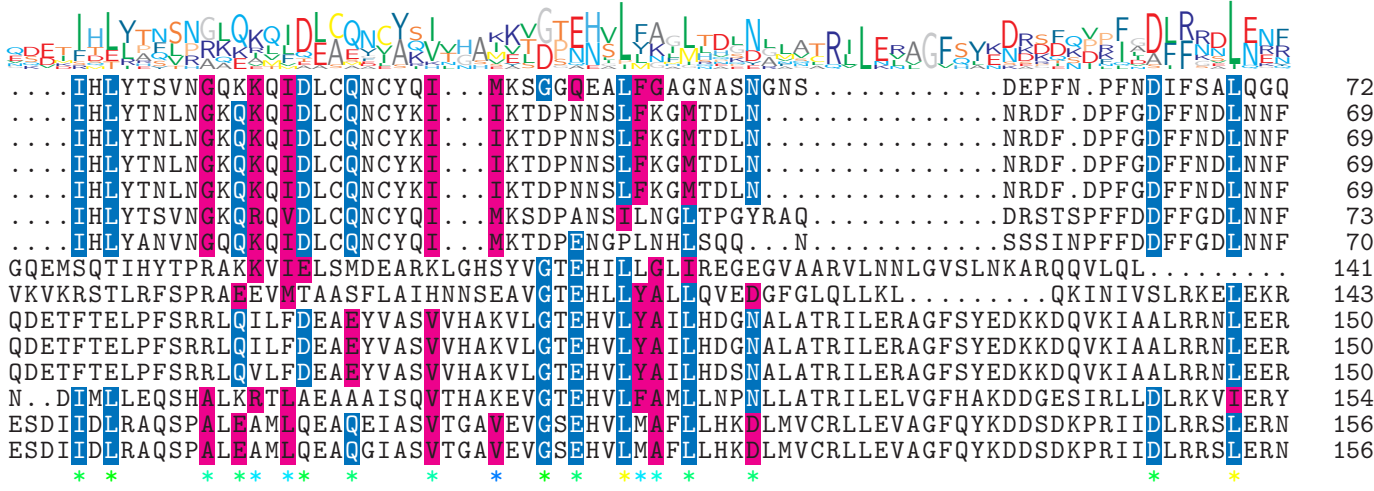
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NTDB id 85 BSU 00860 NP 387967.1
NTDB id 610 V4T04 RS10165 WP 012897346.1
NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
NTDB id 287 SP RS11210 WP 001109712.1
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NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus



logo

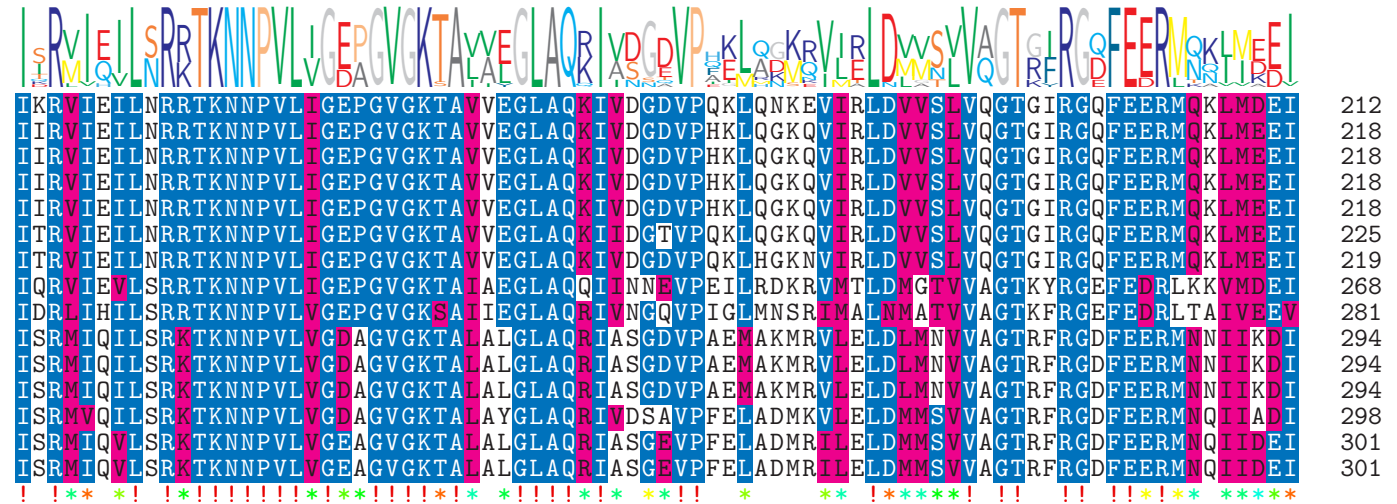
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consensus



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consensus

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logo

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NTDB id	241	SP	RS04015	WP	000882523.1	GAPPGYVGYDEAGQLTEKVRHN	607
NTDB id	1036712	ACB356	RS05745	WP	002989104.1	GAPPGYVGYDEAGQLTEQVRRN	614
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NTDB id	289	SPD	RS10700	WP	001109677.1	GAPPGYVGYDEGGELTEKVRNK	671
NTDB id	287	SP	RS11210	WP	001109712.1	GAPPGYVGYDEGGELTEKVRNK	671
NTDB id	377	SMU	RS09275	WP	002262344.1	GAPPGYVGYDEGGELTEKVRNK	674
NTDB id	297	STER	RS00545	WP	011680614.1	GAPPGYVGYDEGGELTEKVRNK	677
NTDB id	329	STU	RS10020	WP	011225298.1	GAPPGYVGYDEGGELTEKVRNK	677
consensus						!*!!!!!!!!*!*!*!!!!*!! !!!*!*!*!*!!!!!!* * !*!*?! *!! !! ! !* ***!!!!!! !	

logo

NTDB id	601	KW2	RS02615	WP	021036805.1	EASVGFGAAREG....RTKSVLGQLGDFFSPEFMNRFDGIEFSALESKENLLKIVDLMIDEVNEQICRNDIHLSTQAA	676
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NTDB id	1036712	ACB356	RS05745	WP	002989104.1	EASVGFGAAREG....RTSSVLGELSNFFSPEFMNRFDGIEFKALS	689
NTDB id	378	SMU	RS02690	WP	002263569.1	EANVGFGASREG....RTNSVLGELGNFFSPEFMNRFDGIEFQALS	683
NTDB id	85	BSU	00860	NP	387967.1	NKYVGFNVDQETQNHKDMKDKVMGELKRAFRPEFINRIDEIVFHS	747
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NTDB id	289	SPD	RS10700	WP	001109677.1	DKTVGFGAKDIRFDQENMEKRMFEELKKAYRPEFINRIDEKVVVFS	751
NTDB id	287	SP	RS11210	WP	001109712.1	DKTVGFGAKDIRFDQENMEKRMFEELKKAYRPEFINRIDEKVVVFS	751
NTDB id	377	SMU	RS09275	WP	002262344.1	DKTVGFGAKSFSQDYKAMESRIILEELKKVYRPEFINRIDEKVVVFN	754
NTDB id	297	STER	RS00545	WP	011680614.1	DKTVGFGAQTISHNHQAMQARIMEELKKSYPPEFINRIDEKVVVFS	757
NTDB id	329	STU	RS10020	WP	011225298.1	DKTVGFGAQTISHNHQAMQARIMEELKKSYPPEFINRIDEKVVVFS	757
consensus						* * !!!!!* ** *! * !!!!! ! * ! !* * * *! *!* * ** * * *	

logo



X non conserved
X similar
X ≥ 50% conserved