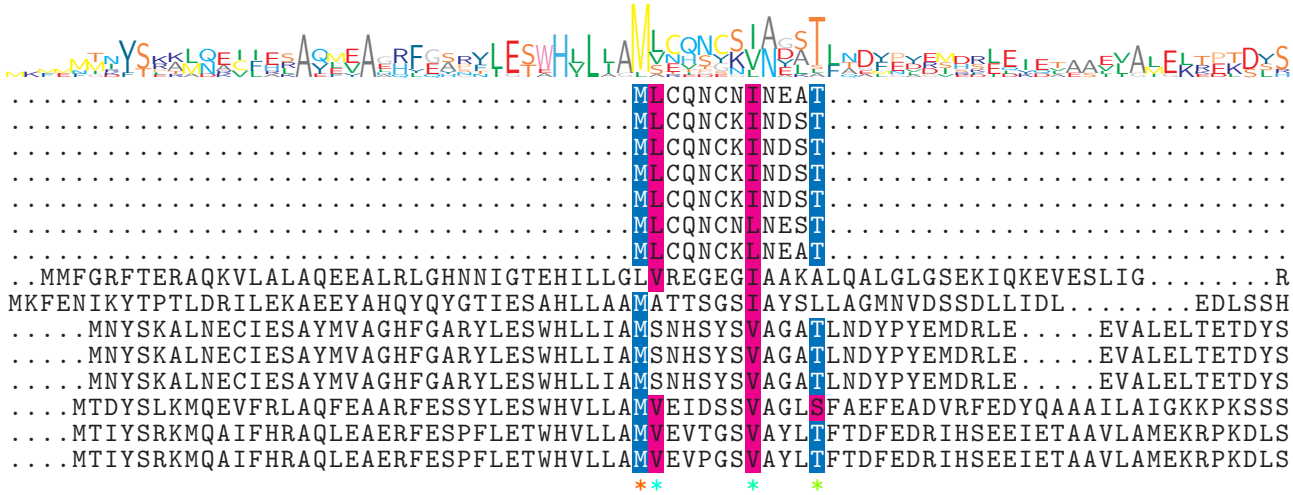


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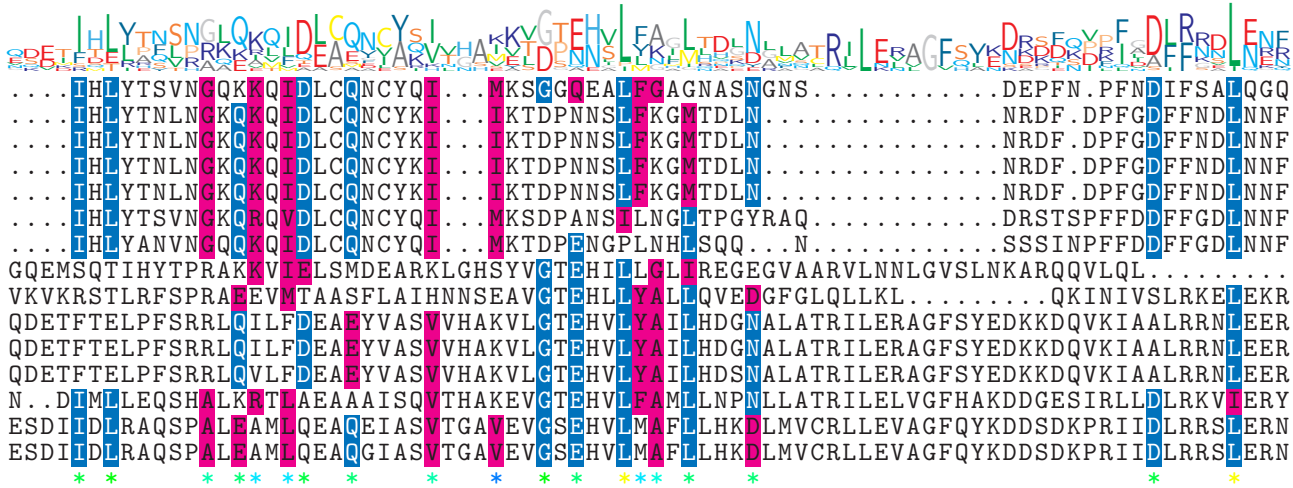
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NTDB id 287 SP RS11210 WP 001109712.1
NTDB id 377 SMU RS09275 WP 002262344.1
NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus



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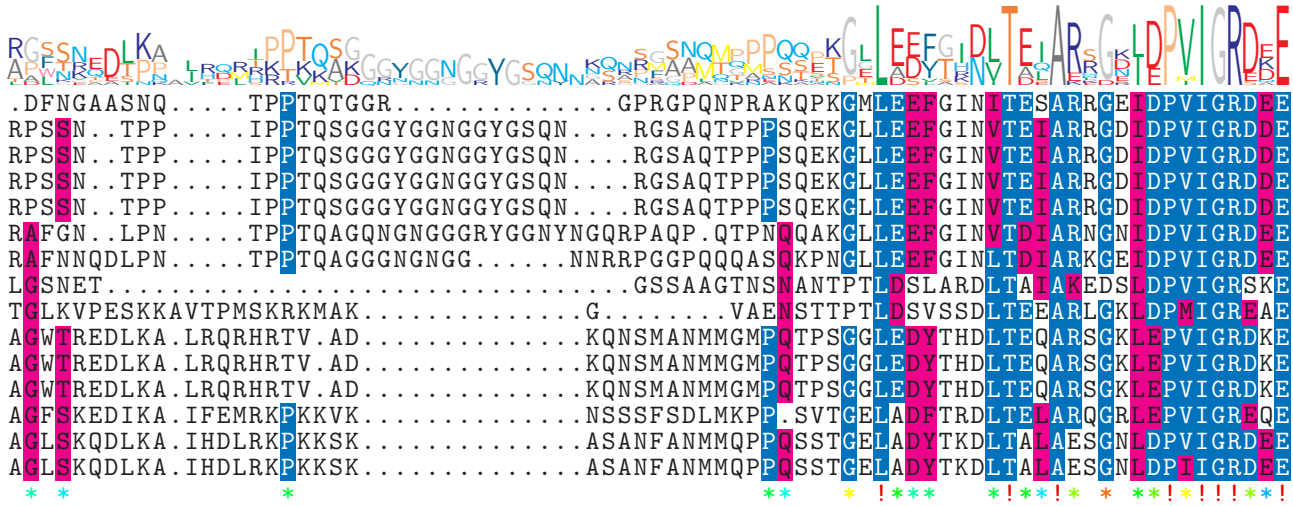
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NTDB id 329 STU RS10020 WP 011225298.1
consensus



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logo

NTDB	id	601	KW2	RS02615	WP	021036805.1
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consensus						



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logo

NTDB	id	601	KW2	RS02615	WP	021036805.1
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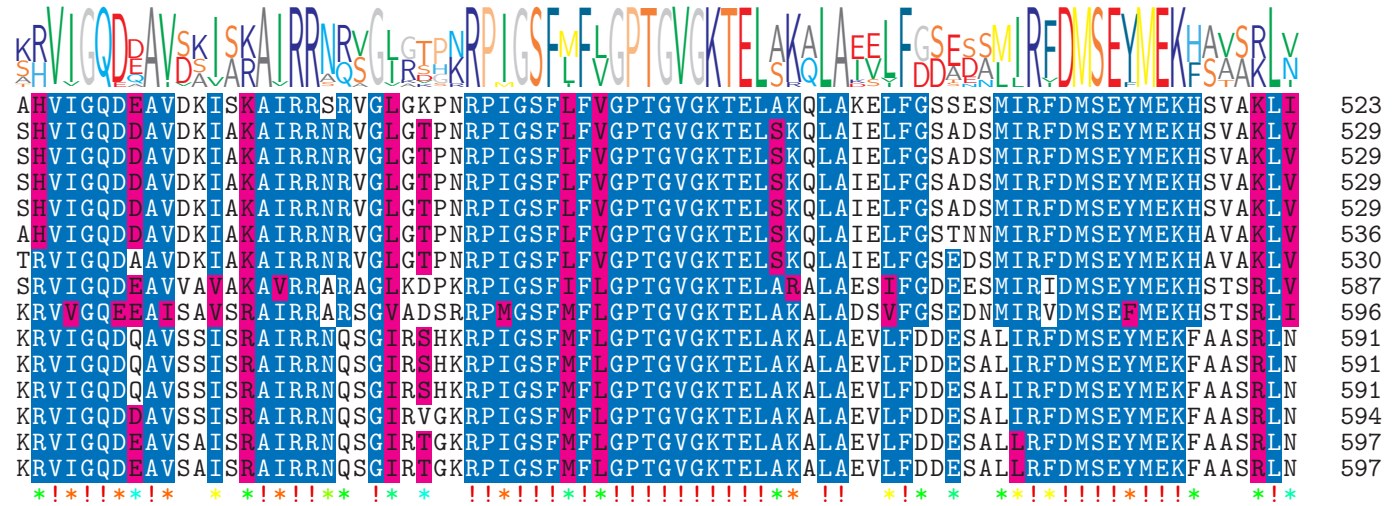
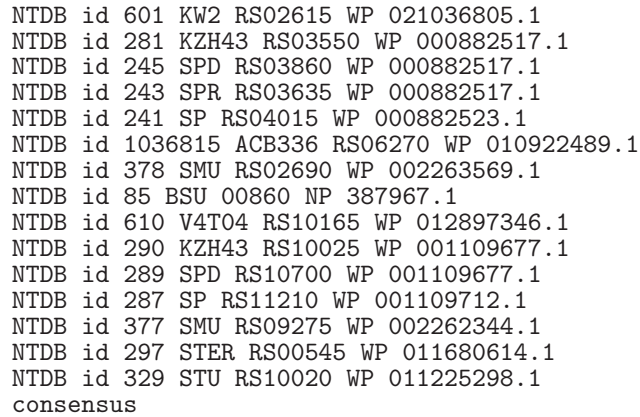


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NTDB id 601 KW2 RS02615 WP 021036805.1
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consensus

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logo

NTDB id	601	KW2	RS02615	WP	021036805.1	GAPPGYVGYDEAGQLTEKVRNKPYSLLLDEVEKAHPDVMMFLQVLDGRLTDGGRVTSFKDSLIIIMTSNAGTGK..V	601
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NTDB id	243	SPR	RS03635	WP	000882517.1	GAPPGYVGYDEAGQLTEKVRHN	607
NTDB id	241	SP	RS04015	WP	000882523.1	GAPPGYVGYDEAGQLTEKVRHN	607
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NTDB id	297	STER	RS00545	WP	011680614.1	GAPPGYVGYDEGGELTEKVRNK	677
NTDB id	329	STU	RS10020	WP	011225298.1	GAPPGYVGYDEGGELTEKVRNK	677
consensus						!*!!!!*!*!*!*!!!!*!! !!!*!*!*!*!!!!!!* * !!!*!!!!*!! !!!*!*!!!!!!*	

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NTDB id	601	KW2	RS02615	WP	021036805.1	EASVGFGAAREG....RTKSVLGQLGDFFSPEFMNRFDGIEFSALESKENLLKIVDLMIDEVNEQIGRNDIHLSTVQAA	676
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NTDB id	1036815	ACB336	RS06270	WP	010922489.1	EASVGFGAAREG....RTSSVLGELSNFFSPEFMNRFDGIEFKALS	689
NTDB id	378	SMU	RS02690	WP	002263569.1	EANVGFGASREG....RTNSVLGELGNFFSPEFMNRFDGIEFQALS	683
NTDB id	85	BSU	00860	NP	387967.1	NKYVGFNVDQETQNHKDMKDKVMGELKRAFRPEFINRIDEIVFHS	747
NTDB id	610	V4T04	RS10165	WP	012897346.1	DKTVGFGAKNITADYSAMKSRILEELKRHYRPEFLNRIDENIVFHS	756
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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