

logo

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NTDB id 1115 OK783 RS00485 WP 003698035.1
NTDB id 1092 CAD91899.1 1..2286( )
NTDB id 1100 NMB RS09455 WP 002225652.1
NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1257 GC085 RS04855 WP 163621000.1
NTDB id 1284 ACE17W RS12370 WP 010946666.1
NTDB id 1009 ACIAD RS15175 WP 004923730.1
NTDB id 1052 ABD1 RS15850 WP 001017033.1
NTDB id 1039 HON27 RS01460 WP 031993779.1
NTDB id 148666 AB688 RS00800 WP 081255323.1
NTDB id 1193 PAKAF RS26665 WP 003161249.1
consensus

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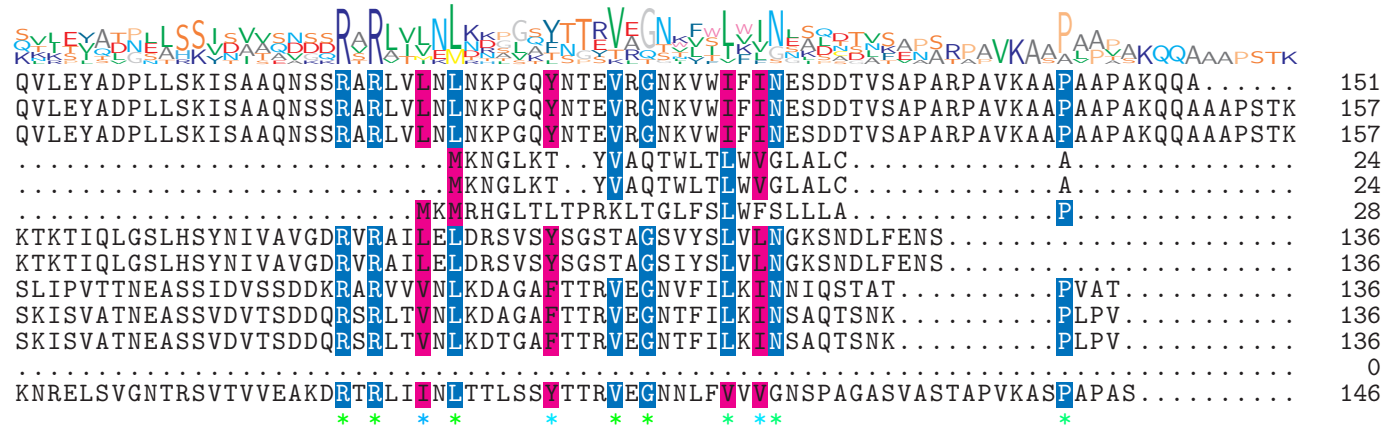


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NTDB id 1193 PAKAF RS26665 WP 003161249.1
consensus

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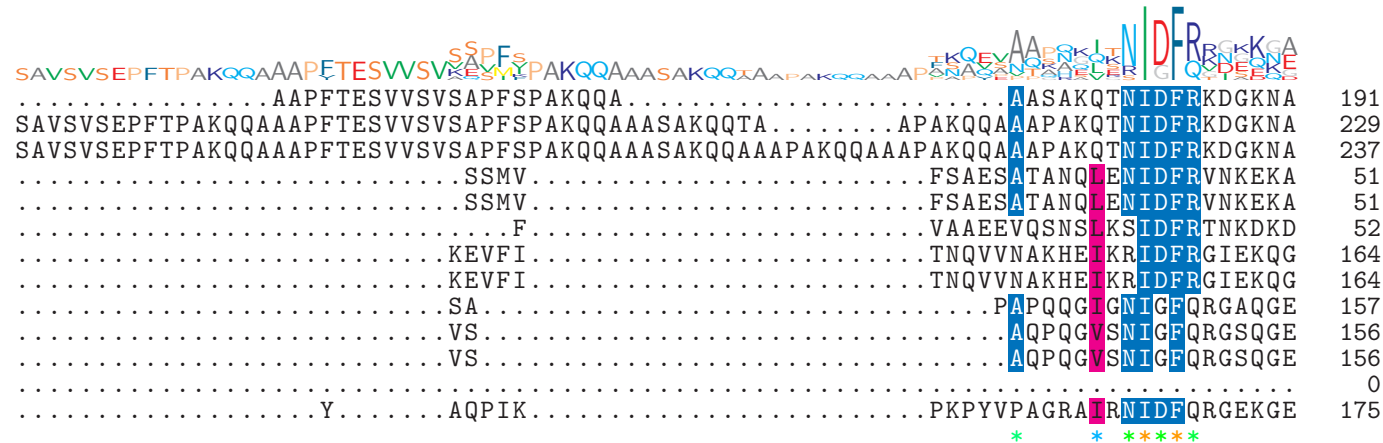


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NTDB id 1193 PAKAF RS26665 WP 003161249.1
consensus

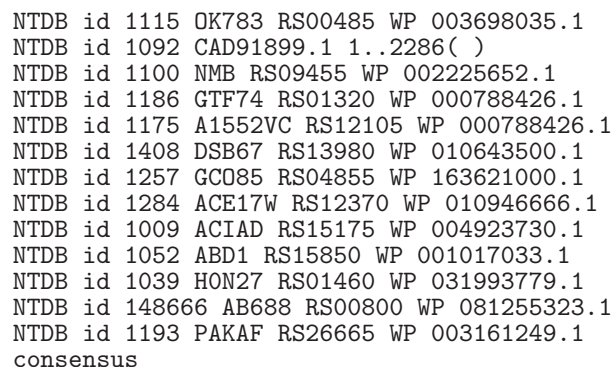
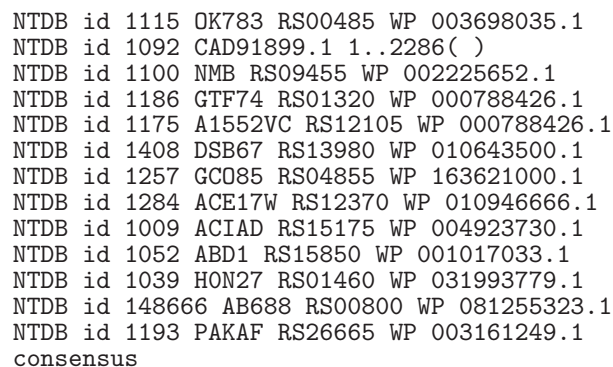
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NTDB id 1193 PAKAF RS26665 WP 003161249.1
consensus

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logo

NTDB id 1115 OK783 RS00485 WP 003698035.1	NTLVSGRGSVLIDPATNTLIVTDTRSVIEKFRKLIDELDVPAQQVMIEARIVEAADGFSRDL	487
NTDB id 1092 CAD91899.1 1..2286()	NTLVSGRGSVLIDPATNTLIVTDTRSVIEKFRKLIDELDVPAQQVMIEARIVEAADGFSRDL	525
NTDB id 1100 NMB RS09455 WP 002225652.1	NTLVSGRGSVLIDPATNTLIVTDTRSVIEKFRKLIDELDVPAQQVMIEARIVEAADGFSRDL	533
NTDB id 1186 GTF74 RS01320 WP 000788426.1	NMLSERGSISIDERTNSLLIRELPDNIIVIREIIESLDIPVKQVQIEARIVTVKEGNLEEL	338
NTDB id 1175 A1552VC RS12105 WP 000788426.1	NMLSERGSISIDERTNSLLIRELPDNIIVIREIIESLDIPVKQVQIEARIVTVKEGNLEEL	338
NTDB id 1408 DSB67 RS13980 WP 010643500.1	SMLSDRGSMTIDERTNSLLIRELPENIEVIREIIESLDIPVKQVQIEARIVTINEGNMDEL	339
NTDB id 1257 GC085 RS04855 WP 163621000.1	SLLSDRGKISVDARTNTIWIQDGTGKIEEVRELKQLDIPVKQVLIIEARIVEVTKDFSQDI	451
NTDB id 1284 ACE17W RS12370 WP 010946666.1	SLLSDRGKISVDARTNTIWIQDGTGKIEEVRELKQLDIPVKQVLIIEARIVEVTKDFSQDI	451
NTDB id 1009 ACIAD RS15175 WP 004923730.1	TNGSATVEPLGDSVGTLLSPRGITISLDPRTNTLIINDTSQKIDQIRKMIDLLDIPVKQVMIEARIVRATTSFSKEM	465
NTDB id 1052 ABD1 RS15850 WP 001017033.1	TSTSTNLNTGVDSLGNNVGSLLSPRGITITQDDRTNTLIINDTAQSIDQIRKMIDLLDVQVKQVMVEARIVRASTSFTEL	473
NTDB id 1039 HON27 RS01460 WP 031993779.1	TSTSTNLNTGVDSLGNNVGSLLSPRGITITQDDRTNTLIINDTAQSIDQIRKMIDLLDVQVKQVMVEARIVRASTSFTEL	473
NTDB id 148666 AB688 RS00800 WP 081255323.1	EPAAGRGSLSVDSRTNTLIATQSAARLAELRQLVADLDVPVRQVMIEARIVEANVDYENAL	198
NTDB id 1193 PAKAF RS26665 WP 003161249.1	GKEGGRGSITVDDRTNSTIAYQPQERLDELRRIVSQLDIPVRQVMIEARIVEANVGDKSL	467
consensus	***** !!!!!!! *!!!!!! ** ** *! **** !!!!!!!*!!!!!! * ** **	

logo

NTDB id 1115 OK783 RS00485 WP 003698035.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	539
NTDB id 1092 CAD91899.1 1..2286()	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	577
NTDB id 1100 NMB RS09455 WP 002225652.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	585
NTDB id 1186 GTF74 RS01320 WP 000788426.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	396
NTDB id 1175 A1552VC RS12105 WP 000788426.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	396
NTDB id 1408 DSB67 RS13980 WP 010643500.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	405
NTDB id 1257 GC085 RS04855 WP 163621000.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	512
NTDB id 1284 ACE17W RS12370 WP 010946666.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	512
NTDB id 1009 ACIAD RS15175 WP 004923730.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	529
NTDB id 1052 ABD1 RS15850 WP 001017033.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	541
NTDB id 1039 HON27 RS01460 WP 031993779.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	541
NTDB id 148666 AB688 RS00800 WP 081255323.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	234
NTDB id 1193 PAKAF RS26665 WP 003161249.1	GVRWGVTSKGLN...SLV...E...LW...Y...P...G...G...D...T...G...S...T...Y...P...N...Y...E...D...N...L...D...L...A...T...P...A...S...I...A...F...S...V...	536
consensus	!****! ** * * * * **** ** * **** *	

logo

NTDB id 1115 OK783 RS00485 WP 003698035.1	VRAISSGALNLELSASESLSKTKTLANPRVLTQNRKEAKIESGYEIPFTVTTASGGGNSNTNELKKAVLGLTVPNTITPD	619
NTDB id 1092 CAD91899.1 1..2286()	VRAISSGALNLELSASESLSKPKTLANPRVLTQNRKEAKIESGYEIPFTVTSIANGGSSTNTELKKAVLGLTVPNTITPD	657
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NTDB id 1186 GTF74 RS01320 WP 000788426.1	AKLGSGLTLLDLELSALQNESKAEIISPRILITTNKKPAYIEQGTEIPYLESSSSGAS...TVAFKKAVLSLKVTPQITPD	473
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NTDB id 1257 GC085 RS04855 WP 163621000.1	L..NDNILLDLELSALESEGRAELISSPRILITTNQQAIVIESGEEIPYQEATSSGAT...AVAFKKAVLSLKVTPQITPD	587
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consensus	* * *!!!!!!* * ****!!**** ** !!!!!! *!* ** ** * ****!!*!!!!!!	

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consensus

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[illegible]

logo

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NTDB id 1193 PAKAF RS26665 WP 003161249.1
consensus

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[illegible]

- X non conserved
- X similar
- X > 50% conserved