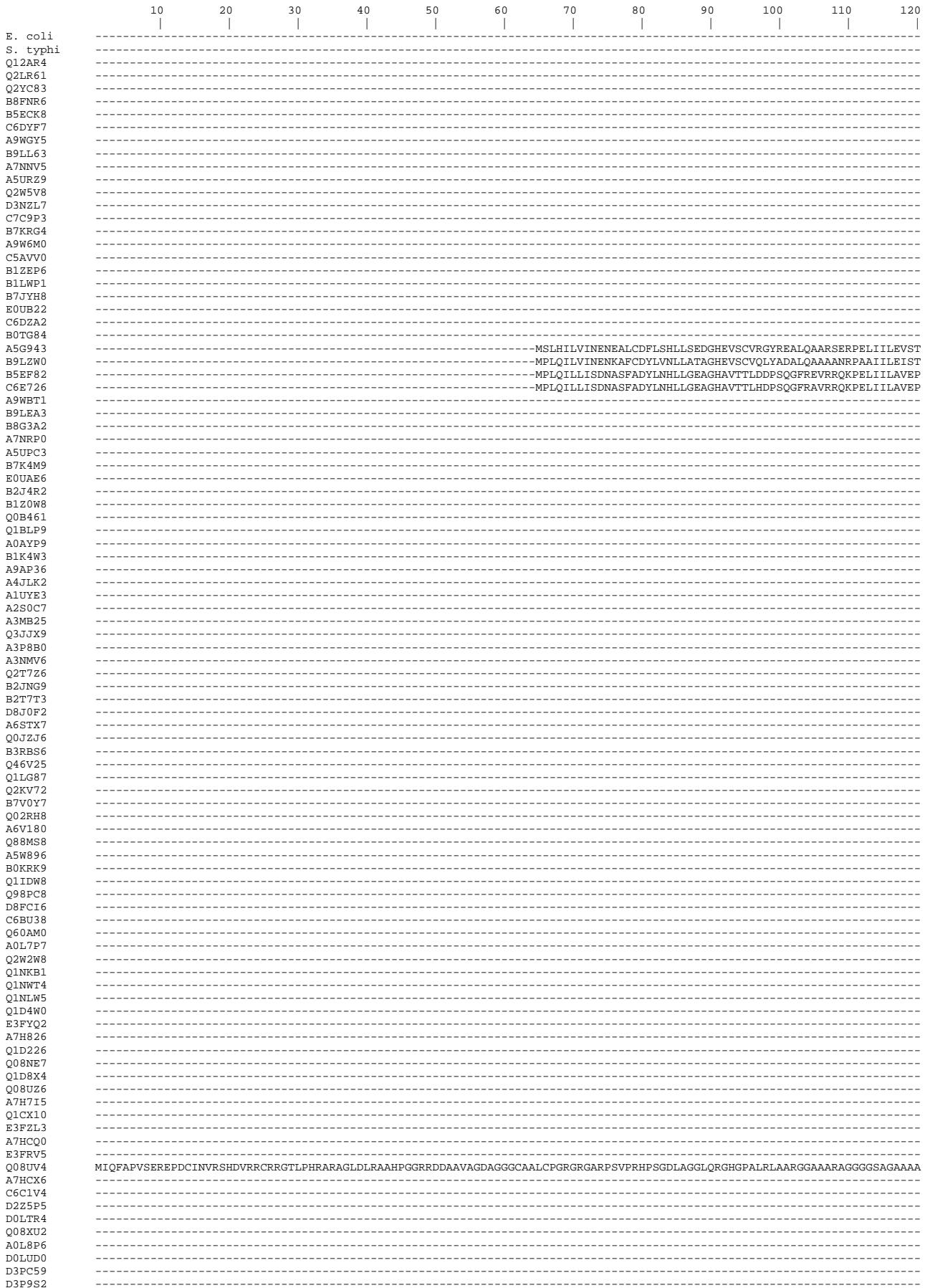


Figure S4A



	250	260	270	280	290	300	310	320	330	340	350	360
E. coli	GHYLN-----	LLESNQHS-----	GEWQAF INSLLNTLAFF-----	YEAHHFPLLADHAR-----		RRSGE-----	YRVWSAAASTGEEPYSIAMTLADTLGT-APGR-W					
S. typhi	GRYLS-----	MLEANQNS-----	AEWQAFIFNALNTLTAFF-----	YEAHHFPLLADHAR-----		RRHGE-----	YRVWSAASTGEEPYSIATTLADALGM-APGR-W					
Q12AR4	A-----	CIDWLSSAPPTRA-----	OLQVLVLAGHLTVGETTYFFF-----	DRQTLLEVLTESALPELIRARRGG-----		EQRLRRIWSAGCCSCEEPPYLAILLHEALPDFG-NWQV						
Q2LR61	P-----	FVSWLLSTKLQR-----	QIETLASHLTVGETTYFFF-----	TEASALEKYIPLPELIESRSRS-----		TRSRLRRIWSAGCATGEEPYTIAMLSRLIPDIS-KWQV						
Q2YC83	T-----	AARHLLSTSLLTRR-----	QIEIIACRCLTVGETTYFFF-----	EQKCLNLVEQSILPELIRCKCES-----		NRQLRRIWSAGCTGEEPPYSIAIILLDRLFKHKGEWNA						
B8PNR6	S-----	CIOGLLGSVPVKV-----	QLDTLVAHLTVGETTFL-----	DESHFNALQHQILPDIPEKCAKK-----		GKPRFWSSAACCTGEEPYTMAMVVDRMKPWT-AKDF						
B5ECK8	G-----	YHLWLMSPQLSLE-----	QTAKALALALTIGETYFFF-----	DPKSYRAFQOHLLPKLLAAKGS-----		DKSLKIWSAGCSSGEEPPYSIAIMLTRALADLA-DWKV						
C6DYP7	G-----	YHLWLMSPQLSQE-----	QTAKALALALTIGETYFFF-----	DPKSYRAFQOHLLPKLLAAKGS-----		DKSLKIWSAGCSSGEEPPYSIAIMLTRALADLA-DWKV						
A9WG5		SFAQLLAMVE-----	SRPSANDELIAELTIGETYFFF-----	NAAQFAALRDIIILPDLMRQRA-----		TMRYLRLRWSAGCATGEEPYSLAITLHEVLP-AWPWQV						
B9LL63		SFAQLLAMVE-----	SRPSANDELIAELTIGETYFFF-----	NAAQFAALRDIIILPDLMRQRA-----		TMRYLRLRWSAGCATGEEPYSLAITLHEVLP-AWPWQV						
A7NNV5		TLEALYAAV-----	VDDKLHTVIEGTVGETTYFFF-----	INTSQTALRDRILPDLIARRE-----		VIERSRLRWSAGCATGEEPYSLAIILHEALPD-PDAQW1						
A5UR29		SLEALYTAAV-----	AASERVLHTI ELSVTGETTYFFF-----	NAISQFAALRDIIILPDLIARRE-----		AISRLRRIWSAGCATGEEPYSLAIILRDLTLP-FDWQNI						
Q2W5V8		DCAGYLLDLRH-----	LQGQKAEMDALIAELTIGETYFFF-----	HKEQFDALRDLIPEAERNR-----		TVRRLRRIWSAGCATGEEPYSVAIMLREFGERIAGHWV						
D3NZL7		TVGGYLAALK-----	EGPGGAEYQGLINELAVGEOF-----	YIEQFDALRAVAIAPECLRRNQ-----		GSRLRRIWSAGCSIGPEAYSVEILLQQFAVQLEGWQV						
CTC9P3		DCAAYSSL-----	GAAGREWAALEAITIGETFFF-----	YAEQFAALRKTILPELIEARS-----		EIRRLRRIWSAGCTGAEPPSLAIVLVDLLGEALADWSV						
B7KRG4		DCAAYSSL-----	GAAGREWAALEAITIGETFFF-----	YAEQFAALRKTILPELIEARS-----		EIRRLRRIWSAGCTGAEPPSLAIVLVDLLGEALADWSV						
A9W6M0		DCAAYSSL-----	GAAGREWAALEAITIGETFFF-----	YAEQFAALRKTILPELIEARS-----		EIRRLRRIWSAGCTGAEPPSLAIVLVDLLGEALADWSV						
C5AVV0		DCAAYSSL-----	GAAGREWAALEAITIGETFFF-----	YAEQFAALRKTILPELIEARS-----		EIRRLRRIWSAGCTGAEPPSLAIVLVDLLGEALADWSV						
B1ZEP6		DCTAYRVLB-----	GAAGRDEWAALEARITEITGETFFF-----	YAEQFAALRKSILPELIAARS-----		EIRRLRRIWSAGCTGAEPPSLAIVLVDLLGEALADWSV						
B1LWP1		DCAAYAALLSS-----	GAEGBEAWALEARIEITGETFFF-----	YAEQFNALRATILPALLAARA-----		SERVLRLRRIWSAGCTGAEPPSLAIVLVDLLGEALADWSV						
B7JYH8		TVEDYYHLLSETVQSQEKEWQLI-----	TLTTNNESSYFFF-----	DLGQFKLRLNPKLILPELIERNQ-----		RTKILRICSAGCTGAEPPSLAIVLKLIPN-WQWNL						
EOUB22		SPQAYYYLATTNPNSDQEWWQF-----	VCLMTNKESYFFF-----	DKGQFLPKLRLNPKLILPELIERNQ-----		KSQHRLRILSAGCSIGPEPPSLAIVLRLQMPD-FAGWHI						
C6DZA2		SEWDYYLQLQDETPESAREWCVI-----	LAAILTTGESYFFF-----	DKGQFLPKLNLKHPIELIKRK-----		RQRRLRRIWSAGCSSGEEPPSLAMWVKELLPV-LDNWQI						
BOTG84		DHEAYHRLLLDKGPAGTEEWRRLI-----	DAVTGETFFF-----	DRQFQALLRERILPILLAERR-----		DKKALRRLRWSAGCTGEEAYSIAIILVDELVGA-APGWITI						
A5G943	REYYD-----	YLIHQERRQR-----	OLEKQLLKPFLTIGETYFFF-----	YHAAFAALKRLLKLD-----	ELAT-----	KKPEGEKILRRLRWSAGCSIGPEPPSMAMTIMETVPG-WQMDI						
B9LZW0	EEYYH-----	YLVLHQESR-----	OLEKQLLKPFLTIGETYFFF-----	YHAAFAALKRLLKLD-----		LAQGERLKLRRLRWSAGCTGEEPPSLAIITVMEAISD-WKHKDI						
B5EF82	RQYLA-----	YLKLHGEDR-----	HELQKLLQFLTVGETYFFF-----	YPAHFAALRERFIP-PPPV-----		DRP-----	IRIWSAGCSTGEEPPYSIAITLMEALPD-WRDRDI					
C6E726	RDYL-----	YLKLHGEDR-----	HELQKLLQFLTVGETYFFF-----	YPAHFAALRERFNP-PPPV-----		DRP-----	IRIWSAGCSTGEEPPYSIAITLMEALPD-WRDRDI					
A9WB71	ARYLA-----	DLVLDADR-----	TELQQLAEELLNHETIFF-----	NRPHM DALRKTLLP-SLHA-----		TLPPGM-PIKWIWSAGCATGEEPPYSIAITALEALGD-PLPRPV						
B9LEA3	ARYLA-----	DLVLDADR-----	TELQQLAEELLNHETIFF-----	NRPHM DALRKTLLP-SLHA-----		TLPPGM-PIKWIWSAGCATGEEPPYSIAITALEALGD-PLPRPV						
B8G3A2	DQYLA-----	DLTNRDR-----	TELQQLAEFPFLNHEIFF-----	YPAHFAALRERFNP-PPPV-----		TLPPGV-PIKWIWSAGCATGEEPPYSIAITALELTLGE-PLPRPV						
A7NRP0	ESYER-----	HISAPAGR-----	NELHLRAEMVNHNETHFFF-----	NAQMRLARETLTLF-ELHR-----		RKPGPE-PIRISIWSAGCATGEEAYSTAIVTLETGFL-ALIRPV						
A5UPC3	EAYER-----	RITAPAGR-----	DELQRILTELVNHNETHFFF-----	YPAHFAALRERFNP-PPPV-----		RKPAGE-PIRISIWSAGCATGEEAYSTAIVTLETGFS-TMIRPV						
B7K4M9		PSYKLHRSQ-----	REFQVELVEEVISETWFFF-----	YAEQFAALRKTILPELIEARS-----		NHPQEKLRLVSLPCSTGEEAYSTAIVTLETGFS-PHQ-F						
EQUA86		KSYVQFLFTESS-----	QEWELLDQIVVPEPTWFFF-----	YAEQFAALRKTILPELIEARS-----		NHPQEKLRLVSLPCSTGEEAYSTAIVTLETGFS-PHQ-F						
B2J4R2		ESYVWRLQST-----	LELEELIELLIVPETWFFF-----	YHAEFKLHKYVKT-EWLKPCKTRRSAPQVSILRLSLPCSTGEEAYSTAIVTLETGFS-SNQ-F								
B1Z0W8		ERLPSAARTP-----	VTPEALDAYWQHNLNASA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLALE-RIAA-----	T-PGRVIRVLRLSAPCSTGEEPPYSAAAMALLDAGLD-PAS-F						
Q0B461		ERLPSAARPA-----	VTPEALDAYWQQLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLALE-RIAA-----	T-PGRVIRVLRLSAPCSTGEEPPYSAAAMALLDAGLD-PAS-F						
Q1BEP9		ERVPSAARPP-----	VTPEALDAYWQQLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIAA-----	M-PGRVIRVLRLSAPCSTGEEPPYSIAITLMEALPD-WRDRDI						
AOAYP9		ERVPSAARPP-----	VTPEALDAYWQQLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIAA-----	M-PGRVIRVLRLSAPCSTGEEPPYSIAITLMEALPD-WRDRDI						
B1K4W3		ERVPAAPRPP-----	VTPEALDAYWQQLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIAA-----	M-PGRVIRVLRLSAPCSTGEEPPYSAAAMALLDAGLE-PAS-F						
A5AP36		ERLPSVSRPP-----	VTPEALDAYWQQLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATVLAQV-RAVA-----	S-PSRVRLVSLPCSTGEEPPYSAAAMALLDAGLE-PDD-F						
A4JL2K		ERLPSAARAP-----	VTTEALRAYWARLHACA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RICAC-----	A-PGRVIRVLRLSAPCSTGEEPPYSAAAMALLDAGLD-PAS-F						
A1UYE3		-GEAAQAGRPOP-----	TADEVADAYWQCLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIFPA-----	E-PARALRILSAPCSTGEEPPYSAAAMALLDAGID-PSR-F						
A2S0C7		-GEAAQAGRPOP-----	TADEVADAYWQCLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIFPA-----	E-PARALRILSAPCSTGEEPPYSAAAMALLDAGID-PSR-F						
A3MB25		-GEAAQAGRPOP-----	TADEVADAYWQCLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIFPA-----	E-PARALRILSAPCSTGEEPPYSAAAMALLDAGID-PSR-F						
Q3JJX9		-GEAAQAGRPOP-----	TADEVADAYWQCLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIFPA-----	E-PARALRILSAPCSTGEEPPYSAAAMALLDAGID-PSR-F						
A3P8B0		-GEAAQAGRPOP-----	TADEVADAYWQCLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIFPA-----	E-PARALRILSAPCSTGEEPPYSAAAMALLDAGID-PSR-F						
A3NMV6		-GEAAQAGRPOP-----	TADEVADAYWQCLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIFPA-----	E-PARALRILSAPCSTGEEPPYSAAAMALLDAGID-PSR-F						
Q2T7Z6		DGAAAPAGRPOQ-----	TDALDAYWQQLNNSA-----	DERRALIELFVVPEPTWFFF-----	DREAFATLARLAAE-RIFPA-----	Q-PARVRLRILSAPCSTGEEPPYSAAAMALLDAGLE-PAR-F						
B2JNG9		AHDTSVCVP-----	ADIDAYWQQLTTSR-----	DERQALIEALVVPETWFFF-----	DREAYVALARLANE-RLVR-----	E-PARLRLRILSAPCSTGEEPPYSAAAMALLDDGIG-ENR-F						
B2T7T3			HAEATLADYWEAQCS-----	SMVQALIETVVPETWFFF-----	DADAFKALARLAHE-RLAE-----	RGNALPLRVLRLSAPCSTGEEPYTIAMLLDAGFD-PAQ-F						
D8J0F2	AL-----	YLSLLLSA-----	AELQALIEALVVPETWFFF-----	DREAFATLARLASD-KLAA-----	Q-PAAVIRLRLSAPCSTGEEPPSLAMALLDAGLT-PGQ-F							
A6STX7	A-----	YLLALQSA-----	GELOLIELVIVPETWFFF-----	DREALISLARLARMAQ-RLAE-----	K-PDRCRVRLRLSAPCSTGEEPPYSVAMALLDAGVP-SVN-F							
Q0JZJ6		QAYWDLLHGTA-----	DEFQALIEAVVPEPTWFFF-----	HREALLALGRLAAE-RAFA-----	D-GRTRLRRLSAPCSTGEEPPYSIAMMLDAGVP-AGR-F							
B3RB56		EAYWQLLQGTA-----	DEFQALIEAVVPEPTWFFF-----	HREALLALGRLAAE-RAFA-----	D-GRTRLRRLSAPCSTGEEPPYSIAMMLDAGVP-AGR-F							
Q46V25		AAYWNLNHSQ-----	DEFQALIEAVVPEPTWFFF-----	HREALLALGRLAAE-RVFG-----	D-HARHLRVLRLSAPCSTGEEPPYSIAMMLDAGIP-AAR-F							
Q1LG87		ATYWNLLHAPV-----	DELQALIEESVVPETWFFF-----	HREALLALRFAAQ-RVFG-----	A-GEPMRLRILSAPCSTGEEPPYSIAMMLDAGIP-GDR-F							
Q2KV72		ASYWTRLLSE-----	EMEQQLIEAVVPEPTWFFF-----	HYPESFLAKALGALR-----	D-HRGLRLRILSAPCSTGEEPPYSIAMMLDAGK-PEA-F							
B7V0Y7		DEYWRMLNGLSP-----	GEQVALIEAVVPEPTWFFF-----	HYPESFTTFLARLAFE-RLPS-----	LGGGRALRILSAPCSTGEEPPYSVAMALLDAGLS-EYL-F							
Q02RH8		DEYWRMLNGLSP-----	GEVQTLIEAVVPEPTWFFF-----	HYPESFTTFLARLAFE-RLPS-----	LGGGRALRILSAPCSTGEEPPYSVAMALLDAGLS-EYL-F							
A6V180		DEYWRMLNGLSP-----	GEVQALIEAVVPEPTWFFF-----	HYPESFTTFLARLAFE-RLPS-----	LGGGRALRILSAPCSTGEEPPYSVAMALLDAGLS-EYL-F							
Q88MS8		DDYWLRLQOSA-----	DEQQLALIEAVVPEPTWFFF-----	HYPESFTTFLARLAFE-RLQAK-----	LGGGRALRILSAPCSTGEEPPYSVAMALLDAGMA-PGA-F							
A5W896		DDYWLRLQOSA-----	DEQQLALIEAVVPEPTWFFF-----	HYPESFTTFLARLAFE-RLQAK-----	LGGGRALRILSAPCSTGEEPPYSVAMALLDAGMA-PGA-F							
B0KR9		DDYWLRLQOSA-----	DEQQLALIEAVVPEPTWFFF-----	HYPESFTTFLARLAFE-RLQAK-----	LGGGRALRILSAPCSTGEEPPYSVAMALLDAGMA-PGA-F							
Q1IDW8		DDYWLRLQOSA-----	EEQQLALIEAVVPEPTWFFF-----	HYPESFTTFLARLAFE-RLQAK-----	LAGTRPLRLLSAPCSTGEEPPYSIAMMLDGGVS-PAA-F							
Q98PC8		QAYREHLTGSS-----	QALQELINAVIPEPTWFFF-----	HREAFETMLARLAFE-HKRR-----	A-QSIVKLWLSAPCSTGEEAYSTVAMAMPDAGFE-AHE-F							
D8FC16	R-----	AYAEALENSD-----	DEWKRLIEMVVPETWFFF-----	HREAFYHLVRFVTD-DWLP-----	RGSGRPLSWSLPCSTGEEPPYSIAMMLDAGIQ-EDR-F							
C6BU38	S-----	YLSLVRASD-----	EELAELVEEVVPEPTWFFF-----	DNKPFELLFTETVAG-----	CSDFRVLRLSAPCSTGEEPPYSIAMMLTMLGAGLK-N---F							
Q60AM0		AYPARLAADA-----	EERARLIELDVLVPEPTWFFF-----	EPPAFLRVAESAAER-HRLQ-----	R-RSVPFHVLRLSAPCSTGEEPPSIVITALREAGMR-KGD-F							
A0L7P7		PAFLRLRLEQDR-----	EIYFELINRLTINEYFFF-----	ESEHQLQLIEHLLP-PLLNQORS-----	EPIRLVSGCWSGEEPPYSIAMALTERFGP-DQAHLF							
Q2W2W8		AGYLA RLVAEE-----	DEFQEFTVLLTINETYFFF-----	EPEQLMLLVDRLMP-RLLADESAG-----	LPITILSAGCSSGEEPPYSIAIALLEKFGE-AAARMV							
Q1NK81		EDYYRLAADA-----	GEQALVEVLTINETYFM-----	EPEYKLLLDRLIP-ELLAASGG-----	DGLKIFCAGCSCGEEPPYSLAIILLDQRYG--SAARHF							
Q1NWT4		EDYYRLAADA-----	GEQALVEVLTINETYFM-----	EPEYKLLLDRLIP-ELLAASGG-----	DGLKIFCAGCSCGEEPPYSLAIILLDQRYG--SAARHF							
Q1NLW5		EDYYRLAADA-----	GEQALVEVLTINETYFM-----	EPEYKLLLDRLIP-ELLAASGG-----	DGLKIFCAGCSCGEEPPYSLAIILLDQRYG--SAARHF							
Q1D4W0	PE-----	HYIQRLTG-----	AEELRLLPLVTVGHTFFF-----	DAKQFRALEKSVLP-DLVRSSRR-----	MRKCIWSAGCSTGEEPPYSIAMMLAELGAL-S-LEV							
E3FYQ2	PE-----	VYVQLRDLR-----	QELRSLIPLVTVGHTFFF-----	DPKQFRLERFLP-QLLAKARR-----	E-MRCVCIWSAGCATGEEAYSTVAMVMAELGAL-A-VEV							
A7H826	AR-----	AYALLRLSPAEGD-----	EELRRLIELPLVTVGKTSF-----	DERQFLRALE-ALLP-ELVARASG-----	G-GRVSIWSAGCATGEEPPYSIAMTAESLGAAP-E-EHL							
Q1D226		QYLMFLQPS-PSGV-----	TDLDEDI LAIVVNVKHTDLP-----	DEVQLAFAFRQVQLS-PLVARNR-----	RPLRVLWSAGCATGEEPVATLILLLDEAGAD---SDS							
Q08NE7		QYLLHQLS-FPSGA-----	GDLASLISVIAVHKTDLF-----	DEVQLASLFRTHVLE-PLVARRSSG-----	RPLRVLWSAGCATGEEPVATLILLLDEAGAD---SDS							
Q1D8X4		FLRLKLLVREA-----	AAVECPFIEHAVIGETYFFF-----	HPEHPLRLLTARLAQ-T-HPAPC-----	FQVWLSAGCWSGEEPPYSIAMMAEGLP-E-GRF							
Q08UZ6		FLKALLARDP-----	VVVFETIYEAVIGETYFFF-----	HPEHPLRLLTARVIAA-HGEP-----	VHIWLSAGCATGEEPPYSITMALLAAGPV-V-ENI							
A7H715		FRARLARDA-----	RCVALIEVAEVGETYFFF-----	HPEQLEVRQLVIAA-RAPRER-----	P-LRIVLSAGCATGEEPYTLMLRDRAGA-G-CRD							
Q1CX10		AYVRVLAEEP-----	SALDDLLIELTVGETYFFF-----	THEPHEFLRVSVP-L-E-LRER-----	HGPDEHTARMWLSACCSGEEPPYSIAIALLGEQWD-E-EHM							
E3FXL3		LYLGRLADSP-----	GAFDDILNELTIGETYFFF-----	THEHFDPLRQALP-B-LRRL-----	RGEDEHTVRVWLSAGCAGSCEEPPYLAMVRSPQL-AGWD-V							
A7HCQ0		AVARALLEG-----	RPLAPIEELTVGETWFFF-----	EPAQWAFIEEVLP-E-GLLRR-----	EGGVERFRAWSAGCATGEEPYTLAIVLVLDRGVR--ERV							
E3FRV5		LEAYLAKLEAGNQD-----	AELRLIAERLTVGETYFFF-----	HLAQQLQALVAEVL-RVQREG-----	RPARVLCAAGCSCGEEAYSTVAILCRENALV-EPEP-L							
Q08UV4		LEAYLAKLEAGNQD-----	AELRLIAERLTVGETYFFF-----	HLAQQLQALVAEVL-RVQREG-----	RPARVLCAAGCSCGEEAYSTVAILCRENALV-EPEP-L							
A7HCX6		LDAYVARLSEAGR-----	EIRPLAARLTVGETYFFF-----	NANDMEAFSGLVLPVERVRRGA-----	DRRLRILSAGCAGSCEEPPYLAMLVRSPQL-AGWD-V							
C6C1V4		AECLEYILSPVG-----	TDDLEQF INRLTIGETFFF-----	DTNALRVELEYILRKISGRGS-----	GLNGAARIWSSTACSTGEEPYTLAMICRRSGVS----A							
D2Z5P5	P-----	EEVFRLLRKSD-----	KAFDMISIPTHITVGETFFF-----	DRKALDAFGREIVPLRAS-----	AKRPNVWLSAGCATGEEPYTLAMIDHSILGLAGG--F							
D0L7R4		ALLMERDD-----	AEAWEMAATAMTIGETYFFF-----	DPSHFIDLRLIMRG-----	NRMRMLWSSACANGAEEPPYSMAILALEEFAAER-V							
Q08XU2												

	610	620	630	640	650	660	670	680	690	700	710	720
E. coli												
S. typhi												
Q12AR4					-SDVAPRTEPVTVTANGAPESG-							
Q2LR61					-YHHSSRRNSKNDAAVKGQTELGE-							
Q2YC83					--IDS-HDVHGNGDYFSV-							
B8FNR6					-NKVQGADLVDTKAESAPVKRE-							
B5ECK8					--RPLPAKAQLSALSG-							
C6DYF7					--RSLPAKAHVAALSG-							
A9WGY5					-PLQ---APLFT-NASTP-							V
B9IL63					-PLQ---APLFT-NASTP-							V
A7NNV5					-PLH---APLFD-EAIPS-							
A5UR29					-PLN---APLFD-ETAVPPP-							L
Q2W5V8					-VDRPYVVAEPPS-PPVPPP-							L
D3NZL7					-QAERAIPAAAG--PVPGA-							M
C7C9P3					-IDTPVPPPPP--VLVPA-							L
B7KRG4					-IDTPVPPPPP--VLVSA-							L
A9W6M0					-IDTPVPPPPP--VLVPA-							L
C5AVV0					-IDTPVPPPPP--VLVPA-							L
B1ZEP6					-IDTPTEPPP--VLVPA-							R
B1LWP1					-VALPGTPVDPDAPS-TGQVPG-							P
B7JYH8					-SSSEVEKITQNKCQVEKLQNQ-							L
EOUB22					-QSSVSHPAPPKTALEKLSTQ-							L
C6DZA2					-AVSLPAVTP-----LHHQ-							V
BOTG84					-APSSRGIPPAAVTQPPLPGA-							L
A5G943	N				--EQPRSKPSKT--TATVI-							KPAR
B9LZW0	E				--PAPPQRQAPAPA--KA-							KSRQ
B5EF82	Q				--PPPCAKTAPQ--AIPELPEL-							PAVPA
C6E726	Q				--PPPCVKPAPQ--AMPA-							STVPA
A9WB71					--WMP--TTTGQR-							L
B9LEA3					--WMP--TTTGQR-							L
B9G3A2					--MVA--LSVPKPSR-							PTLH
A7NRP0	H				-RSTSPPRSLP-TETRRRSPS-							VVKA
A5UPC3	R				-RTESRQPATT--AEAPQRLTB-							VATATP
B7K4M9					-ILDPPAPSS-							
EOUAE6	A				-KHKNPKPEKS--II-							
B2J4R2					--SEPLPLKLE--LLNI-							
B1Z0W8	ARYAAPTRA-				--VAPYRRAGR--TVAP-							-PAARPL-LAVVPPHTWS
Q0B461	ARYAAPTPV-				--VAPYRRAERL--TVAP-							-PAARPL-LAVVPPHTWS
Q1BLP9	ARPAAPLSA-				--AAPYPGAERF--TVAP-							-LAAPRPV-LAVAPPAWL
AOAYP9	ARPAAPLSA-				--AAPYPGAERF--TVAP-							-LAAPRPV-LAVAPPAWL
B1K4W3	ARPAAPLSA-				--AAPYPGAERF--TVAP-							-SAWSGPAPLAFAAPVAP
A9AP36	TRRATPMPA-				--AAPYRRAERF--APAP-							
A4JLK2	AGHVGGDGs				--YGRMARGGAL--GAGPTS-							-GAATASAMPLASANACASVS
A1UYE3	ASPASAQQGATRFAPSEPRQAQGDAGERPAGI	FARAPGAAEPPGEGEALARTGDAGRF			--ATAPRANAFEAAGWTAAAGAFS-							-RAPDAGRARAGLFAPASDAAR
A2S0C7	ASPASAQQGATRFAPSEPRQAQGDAGERPAGI	FARAPGAAEPPGEGEALARTGDAGRF			--ATAPRANAFEAAGWTAAAGAFS-							-RAPDAGRARAGLFAPASDAAR
A3MB25	ASPASAQQGATRFAPSEPRQAQGDAGERPAGI	FARAPGAAEPPGEGEALARTGDAGRF			--ATAPRANAFEAAGWTAAAGAFS-							-RAPDAGRARAGLFAPASDAAR
Q3JJX9	ASPASAQQGATRFAPSEPRQAQGDAGERPAGI	FARAPGAAEPPGEGEALARTGDAGRF			--ATAPRANAFETGTWTAAGAFS-							-RAPDAGRARAGLFAPASDAAR
A3P8B0	ASPASAQQGATRFAPSEPRQAQGDAGERPAGI	FARAPGAAEPPGEGEALARTGDAGRF			--ATAPRANAFETGTWTAAGAFS-							-RAPDAGRARAGLFAPAPDAAR
A3NMV6	ASPASAQQGATRFAPSEPRQAQGDAGERPAGI	FARAPGAAEPPGEGEALARTGDAGRF			--ATAPRANAFETGTWTAAGAFS-							-RAPDAGRVRAGLFAPAPDAAR
Q2T726	APSVSPPGATRGASSEPRQAQAGEAQCQAAVG	VARAPGATEPPGEEAFARARDAGGP			--AAAPRANAFDGSWTAAGAFS-							-RAPGAERARAGSFAPAIIDGV
B2JNG9	TRRAVQQAV-				--PV-----AARD--MHAP-							R
B2T7T3	TAPLATAAA				--LAMTHSQNPA--LSSP-							-HVFSAEFPFWPDVAPTP
D8J0F2	AA----VP-				--PLPPAWASLV--QAAP-							P-----A
A6STX7					--PHPTAVARKI--IAP-							
Q0JZT6	A				--ARAAPRAPHVIA--AQPP-							R
B3RBS6	R				--ARAPSYTPVHT--TQFP-							
Q46V25	P				--GEARSEQPLQIA--SAAP-							
Q1LG87	A				--SPSSHASAHT--VMPP-							Q-----P
Q2KV72					--APMPALPRRRA--PARS-							S-----
B7V0Y7	A				--PRGARPKAVID--GARP-							V-----V
Q02RH8	A				--PRGARPKAVID--GARP-							V-----V
A6V180	A				--PRGARPKPVAT--APRA-							A-----A
Q88MS8	D				--SPLAAAPAPAQT--AKRA-							F-----T
A5W896	D				--SPLAAAPAPAQP--AKRA-							F-----T
BOKRK9	D				--SAPMDVVPQAQP--VKRT-							L-----A
Q1IDW8	A				--SSAAVSLPLPV--VR-							
Q98PC8					--PVKTTAPKVS--AAIP-							
D8FC16	SG				--EKAAAWTAEK--VPVSG-							IH
C6BU38	TLLKQP				--VCPHQKCPNN--SFSGS-							G
Q60AM0					--DAAEAEPPPA--RSSRP-							
A0L7P7	LWHGLSL5				--SPSHSWRPQDE--AISWR-							G
Q2W2W8	ILP				--APHAPVPPRP--APAPA-							R
Q1NKB1	APR				--HPRPTRRAGKK--GDTIP-							V
Q1NWT4	APR				--HPRPTRRAGKK--GDTIP-							V
Q1NLW5	APR				--HPRPTRRAGKK--GDTIP-							V
Q1D4W0	P				--LRITPYPGEP--DVAARRPVPAADAFTADLRKRLPEDVPLTTRLPAVSASSVAAPGSPSVTLPALGASSSPRSVV							
E3FYQ2	R				--AKTTTPVPMSP--PVSS--SVST-----P--VPLS-----VAPAGTPPRPAPSFSLRRLRPPDAPV							
A7H826					--TPLPAPR--PIRP-							VPP
Q1D226					--ESEAPPAPL--SWNG-							
Q08NE7					--IHLAPPKRL--PASN--							L
Q1D8X4					--PRAVTPESRL--PRLGMGRNQVASP-							
Q08UZ6					--PLGSVKEAPT--PIKPLARWKQE-							
A7H715					--ERAGERVAPK--PARGRARSGSGASP-							TASR
Q1CX10					--PVPFAWSPST--GVFGTATGTSLG-							
E3FZL3					--PRSAASAPPFS--PIRAPAPATPFTP-							
A7HCQ0					--SRTASVRPTS--ARRVAPRDAAR-							
E3FRV5					--RKTGLTPVPM--AAPRP-							
Q08UV4					--RKTGLTPVPM--AAPRP-							
A7HCX6	AAAV				--PEREGPAPTP--PDART-							
C6C1V4					--FKRDDFKTSFSSAAQFDFN-							
D2Z5P5					--EEPSVPPPERYLDSAVS-							
DOLTR4					--TAPAPPPSPORTTRRPP-							
Q08XU2					--PTEPLMPAKAP--							
A0L8P6					--STSLITQSQPARPSQAALPPK-							
DOLUD0					--ASLHGALSPARIQEPKPLP-							
D3PC59					--YQDTQYSIKR--DRTKLP-							
D3P9S2					--IKNELIKKTSENINRLEN-							

	730	740	750	760	770	780	790	800	810	820	830	840
E. coli												
S. typhi												
Q12AR4	-----FPAIETSLSWEPSSA-----	-----APVAVFES-----									-LOPKEDHAVPEAPGSAL	
Q2LR61	-----WASLSTAEPPLPSDS-----	-----IPERKTEPS-----									-RMAPSVEFPSSAAMKEI	
Q2Y8C3	-----GPPSDASYKITSAS-----	-----APSPRLQGW-----									-ETPLSLESMEPQILPVAS	
B8FNR6	-----TPLPNFPARTPLTK-----	-----RPPRVERTA-----									-PPREVDRAKKMELPAD	
B5ECK8	-----PAVAAPSPFSAQL-----	-----WSEGAEPG-----									-SARKDEAAAEEAESPLE	
C6DYF7	-----PAVAVPSPLSPQL-----	-----RSEDAEPG-----									-GAREAAAAAEASPLQ	
A9WGY5	QS-----GPVVPAPP-----	-----LPASPV-----									-SLSLPPP-----RAS	
B9L63	QS-----GPVVPAPP-----	-----LPASPV-----									-SLSLPPP-----RAS	
A7NNV5	VP-----ARQMRVPLS-----	-----TTSADV-----									-ASLLLAP-----AAP	
A5UR29	SP-----ARTTPVSLPN-----	-----VSSSDV-----									-SSDLPALP-----VAL	
Q2W5V8	PL-----VSPPPVRGGL-----	-----PAKAPT-----									-SRALPKA-----LSP	
D3NZL7	P-----GPPPAAPS-----	-----PPTRPR-----									-RTAAAKPP-----AEP	
C7C9P3	PD-----LAPILVAPP-----	-----PEPAPA-----									-AAAPALPV-----PIP	
B7KRG4	PN-----LAPILVAPP-----	-----PEPAPA-----									-AAAPALPV-----PIP	
A9W6M0	PD-----LAPILVAPP-----	-----PEPAPA-----									-ATTPALPV-----PIP	
C5AVV0	PD-----LAPILVAP-----	-----EPAP-----									-AAAPALPI-----PIP	
B1ZEP6	PD-----AVPIPLAP-----	-----LPAFV-----									-AVEPAAPM-----PEP	
B1LWP1	PA-----RDPGPGASAGASS-----	-----GVSGAGL-----									-PDATVPPA-----PDP	
B7JYH8	NN-----SLLNSANKDSKK-----	-----IAIYTQDGEKNKFSYKILTTTKENIS-----									-QAIEEVNEID-----LLE	
EOUB22	NN-----TVIRPPQ-FSE-----	-----ITALPKATKPLDKIVQNLESKPQPSIPP-----T-----									-PLKSSLKDEE-----LLQ	
C6DZA2	VP-----ASARPSSAG-----	-----TKBQR-----									-LN-PAAVS-----	
BOTG84	SP-----ERQPDRIPFQP-----	-----MPAMPKR-----									-AQSAIQQAP-----LLP	
A5G943	P-----KPLLPVKTV-----	-----PPRSNAPA-----									-QSEIDL-----QLF	
B9LZW0	E-----QTAPAVNNT-----	-----APAPASP-----									-QAVDWE-----RLH	
B5EF82	Q-----AAPKAVSPP-----	-----LPAFLAP-----									-PSPEAA-----LGV	
C6E726	Q-----AAPKVVSP-----	-----PPEACAP-----									-PPPAP-----LDV	
A9WB71	P-----APALRTRAP-----	-----RPLERKV-----									-TLFTEE-----AVV	
B9LEA3	P-----APALRTRAP-----	-----RPLERKV-----									-TLFTEE-----AVV	
B9G3A2	P-----TRPTRRRPT-----	-----SPPASSD-----									-PLFTDE-----TVV	
A7NR00	S-----TPSRSKARP-----	-----VAATTAP-----									-TMEEDV-----GRV	
A5UPC3	R-----TASHGRRSP-----	-----IAPITPKP-----									-TSNSNADA-----DHL	
B7K4M9	-----PIK-----	-----PPVVSSP-----									-SNLVITV-----KKR	
EOUAE6	-----QKKIRPSN-----	-----PTELTS-----									-AQAQTH-----LKK	
B2J4R2	-----TQFQSATLQ-----	-----NPVQSLGVKN-----Y-----									-SAIDTQ-----QIK	
B1Z0W8	G-----NAASSAAALA-----	-----PAERTAFEPADRAGRSRTN-----AAT-----									-NATN-----ATN	
Q0B461	G-----NASSPAALA-----	-----PAERTAFEPLADMAGSLTAAATAAT-----									-AANAAAN-----ATT	
Q1BLP9	G-----DVRQPFFAVP-----	-----PPAVTTA-----									-ADL-----AVA	
AOAYP9	G-----DVRQPFFAVP-----	-----PPAVTTA-----									-ADL-----AVA	
B1K4W3	G-----DVRQPFFAVP-----	-----LPAVTTA-----									-ADL-----AVA	
A9AP36	-----IGGLATVH-----	-----AADVTAMADG-----									-AADL-----PAP	
A4JLK2	A-----TASGLPSSSS-----	-----PTSATRALHVPPPSG-----									-LDAARAAP-----PTAFQFRA-----PSP	
A1UYE3	A-----AHEEPFADA-----	-----AGAARAADFADPGVSADAGR-----									-ATARDAGAAPRASASLDHSFDTARPVFGTAAVAVGI-----AAP	
A2S0C7	A-----AHEEPFADA-----	-----AGAARAADFADPGVSADAGR-----									-ATARDAGAAPRASASLDHSFDTARPVFGTAAVAVGI-----AAP	
A3MB25	A-----AHEEPFADA-----	-----AGAARAADFADPGVSADAGR-----									-ATARDAGAAPRASASLDHSFDTARPVFGTAAVAVGI-----AAP	
Q3JJX9	A-----AHEEPFADA-----	-----AGAARAADFADPGVSADAGR-----									-ATARDAGAAPRASASLDHSFDTARPVFGTAAVAVGI-----AAP	
A3P8B0	A-----AHEEPFADA-----	-----AGAARAADFADPGVSADAGR-----									-ATARDAGAAPRASASLDHSFDTARPVFGTAAVAVGI-----AAP	
A3NMV6	A-----AHEEPFADA-----	-----AGAARAADFADPGGSADAGR-----									-ATARDAGAAPRASASLDHSFDTARPVFGTAAVAVGI-----AAP	
Q2T726	P-----A-AEPFADA-----	-----AGLARPAFDALPGAADAGR-----LAS-----									-MALARDAGAAPRASASLAHTFDTAWSPARGAAASGI-----AQ	
B2JNG9	S-----AAARESAK-----	-----PSLQARAF-----									-TR-AP-----LRA-----PRP	
B2T7T3	F-----RGATQFAQR-----	-----ENSATQAY-----									-THKTP-----MPAA-----TPP	
D8J0F2	P-----TPVRPDAAH-----	-----RAPARRIV-----									-ASA-----ALG	
A6STX7	P-----IQIKP-LPP-----	-----ATPVRKPF-----									-ANV-----VPP	
Q0JZJ6	-----MAAPPLR-----	-----PPASMR-----									-PVP-----PVP	
B3RBS6	-----MAAPPLR-----	-----PPASLRPL-----									-AARA-----PAP	
Q46V25	-----APAGMPP-----	-----RPVSLR-----									-VQ-----PAP	
Q1LG87	-----MIARARSAA-----	-----TAPAKH-----									-AA-----RAP	
Q2KV72	-----VAALSQAAP-----	-----RPPAFS-----									-S-----AVR	
B7V0Y7	-----AAVERASI-----	-----RPSPPP-----									-AKPR-----QRL	
Q02RH8	-----AAVERASI-----	-----RPSPPP-----									-AKPR-----QRL	
A6V180	P-----VVAERTGS-----	-----RPSPP-----									-AKPR-----QRL	
Q88MS8	-----TLPAVPYP-----	-----QPSVLP-----									-RSRR-----VLP	
A5W896	-----TLPAVPYP-----	-----QPSVLP-----									-RPLR-----VLP	
BOKRK9	-----TLPTAGYP-----	-----PPSAPLP-----									-PPLR-----VLP	
Q1IDW8	-----PLAP-----	-----PPPRVVA-----									-LPAR-----TVR	
Q98PC8	-----VRTPTPKVN-----	-----KSEPLL-----									-FRP-----TIG	
D8FC16	G-----AGSRPDENK-----	-----PKFIQKP-----									-APFK-----EVQ	
C6BU38	T-----TAAQAKAFIP-----	-----APTFQKPE-P-----									-DIGTK-----AVD	
Q60AM0	-----LAAKAVRR-----	-----APGEAKP-----									-SGP-----DLL	
A0L7B7	L-----TEIAEEDEP-----	-----LPLPLFSLDDELL-----L-----									-REETILLPEGNGEPE-----RIA	
Q2W2W8	L-----HLVPPTPPP-----	-----APP-----									-PA-----PSP	
Q1NKB1	F-----RASKGEKRD-----	-----TPAIKN-----									-RYGVSPVPGSGKVP-----SSA	
Q1NWT4	F-----RASKGEKRD-----	-----TPAIKN-----									-RYGVSPVPGSGKVP-----SSA	
Q1NLW5	F-----RASKGEKRD-----	-----TPAIKN-----									-RYGVSPVPGSGKVP-----SSA	
Q1D4W0	PGRLPAVSPHSPLPAIAARS-----	-----RVTAELPT-----	V-----								-GSVDSARPRITTEPAVATTPRP-TVEVPAWPTL-----LPP	
E3FYQ2	P-----MTP-SPRSYLEQA-----	-----ASPPSPR-----									-GSEPVVRPRPRTLEVS-ATDPRVVRRTIEVPGGAKA-----FPP	
A7H826	-----VPSAPPTV-----	-----RHVPVAP-----									-AAPRILPTPAP-----I-----P-AQOP-----LAP	
Q1D226	-----AAPPS-----	-----RKRTG-----									-ALPG-----RAP	
Q08NE7	L-----AEPPTPRPE-----	-----RRRDSGV-----									-REAAPSA-----ASP	
Q1DX4	P-----PTGWPQALDDS-----	-----RARARVPEGR-----									-LSLNEPGTSRP-----	
Q08UZ6	-----PFGTRLAAASLP-----	-----LAPLITRSP-----									-LVRNPP-----	
A7H715	T-----PIRTRTPVPGPPEAGHPEPFDSAACGRYQADR-----										-LRSAAAPAER-----	
Q1CX10	-----TTAAGLPLSGSESSSTR-----T-----	-----ALPGASAPSHPGGSAARTTTPAWSPPGAGGTSAQAEGTRAAGQPOQASAGSVLGNNTAGTGPASRSGALPRAETPTHAPSA-----AAM										
E3FZL3	-----PPLPPLP-----	-----LP-VAPP-----P-T-----PP-----									-PRPPRPPAPSVT-----ERL	
A7HCQ0	-----TLRPPPPG-----	-----HVP-----PIRS-----									-HRAAPPGP-----	
E3FRV5	-----LDVASLQAASS-----	-----VRWMSPPTPAPRMPE-----									-PFPRELG-----M	
Q08UV4	-----LDVASLQAASS-----	-----VRWMSPPTPAPRMPE-----									-PFPRELG-----M	
A7HCX6	-----AWVQAIQDASER-----	-----VAKVSAAAHAAAQPS-----									-LREALPRSGAAPREDGARV-----	
C6C1V4	-----AQLESTPADFVDLKF-----	-----EPEIPVP-----									-EVESEVPEP-----VIPR	
D2Z5P5	-----EIEPEVEA-----	-----VSDVWF-----										
DOLTR4	-----LRLYARRRS-----	-----TTLKLRT-----									-ARSORPT-----ERP	
Q08XU2	-----APRPARSG-----	-----PPRMTPP-----										
A0L8P6	-----LALSRGQTKP-----	-----ATPYVASPPAG-----									-YVAPSITNPHPP-----LVAG	
DOLUDO	-----QARVLSPRARG-----	-----VRQRQAPAAASPP-----									-ASTRASASG-----NAD	
D3PC59	-----ITDKKERYL-----	-----EIKSLELLN-----									-N-----N	
D3P9S2	-----HPFVTDIKVR-----	-----TRKISEKPE-----									-TELFKEAVN-----FMFN	

	970	980	990	1000	1010	1020	1030
E. coli	-	-	-	-	-	-	-
S. typhi	-	-	-	-	-	-	-
Q12AR4	FVLNFAFLGNLARRRGKGTGEAGKHFPANTLHLLQGQPGLLPPESDGLTATRQLAQTLASLTDP--EIAP-----						
Q2LR61	FVLAHTLGNLVRREGRKTLSSKHFHEHALLLLQNYPPEEILPESDGVTAGRLKDIIIMADLGRSNAPGA-----						
Q2YC83	LVLAHFALGSCLCTQGKSREAKRFHDNALSSLNGYSSNDILPDSGGTAGDLAHFTSIRICDSIGNKE-----						
B8FNK6	F1LAHTLGLNTMKQGKANEARRFHKNALSSLDDLDSDAVIPHSGGMTAGRLTEIVRSMILGD-----						
B5ECK8	FLLGWFAFLGNLQRQGELREAEQSFAANALRLLQRDPHEVLPPEAEGMTAGRLMQLISDIA-----						
C6DYF7	FLLAWFALGNLQRQGEPREAEQSFAANALRLLQRDPHEVLPPEAEGMTAGRLMQLISDIA-----						
A9WGY5	LIVGYIGMAHVVYQLQQPDAARRTIRSAQTTLGLADLSQVVDATGTTAAELRNY--VRLLMEKVDLR-----						
B9L63	LIVGYIGMAHVVYQLQQPDAARRTIRSAQTTLGLADLSQVVDATGTTAAELRNY--VRLLMEKVDLR-----						
A7NNV5	LYLALLAMAGILHRTGHVDEAYRAYRRLRALATLPPHPAVPSFDNATAAELQMFIQRQMEALRR-----						
A5UR29	LYLASLAMAGILQRTGHTQRLRLRSLATLQPDAPVPSLNGATAAELQMFIQRQMEALGS--TRNRMAALGS-----						
O2W5V8	LVLPHYHLGLFLARKDSTGAERSFRNAQOALLAGLAEQPVNGCEKIAVGOMREA--VAMHLKLLGG-----						
D3NZL7	FVLADYHHLALAYWRRGRVLPAQRHFRNAREALADRPGCTVTEGAGLTVQELRCM--LDLWTGEG-----						
C7CP93	FVMHAHQLGLLLARVGRGEAARLDNALRNLQGLPPETILPPEGDGVSAARLAES--ARRALAGITGRDA-----						
B7KRG4	FVMHAHQLGLLLARVGRGEAARLDNALRNLQGLPPETILPPEGDGVSAARLAES--ARRALAGITGRDA-----						
A9W6M0	FVMHAHQLGLLLARVGRGEAARLDNALRNLQGLPPETILPPEGDGVSAARLAES--ARRALAGITGRDA-----						
C5AVV0	FVMHAHQLGLLLARVGRGEAARLDNALRNLQGLPPETILPPEGDGVSAARLAES--ARRALAGITGRDA-----						
B1ZP6	FVMHAHQLGLLLARAGRGRGEAARLDNALRNLQGLSPETILPPEGDGVSAARLAES--ARRALAGITGRDA-----						
B1LWP1	FVMHAHYHLGLLILALQRPSEAVRDNALVALSQLGPDTALPEGDGSAGEIAASAAAARAALGGVGSRRS-----						
B7JYH8	SVAAYLNLANLYQQEGDQKRSEKMQMTALTILKKLPLNDTQPELGNITVQEELLTN-----						
EOUB22	SVASAYLNLANLYQQEGDQKRSEKMQMTALTILKKLPLNDTQPELGNITVQEELLTN-----						
C6DZA2	LIAPHLGEMYLSEKKYELAAKSIRSARELLKVMPGKFIALYSETTAELLRHADDLLDGGLNKEKTSDR-----						
BOTG84	HGPARIEMAHVRLILLKENDQ-----						
A5G943	FVMPHYQLSLWYARMGKDKERQRELNNNTNLIAKLGKN--STIPYSSGLSREAFIEQLKRELEMA-----						
B9LZW0	FVMAHYQLAQYLARLGRKKEERLNLNSLRIVAGLKAK--AAIPYSGGLTRELFDMDQLNKELEKAD-----						
B5EF82	FVMPRYHMGRLHLRLGRQAEAAIRENRSIRILARHDGN--DTPVFSGGLTRAVCMQMQLQNLAQVA-----						
C6E726	FVMPRYHMGRLHLRLGRQAEAAIRENRSIRILARHDGN--DTPVFSGGLTRAVCMQMQLQNLAQVA-----						
A9WB71	HPLVAFHLAECYRQRNQITEIARYRNRNARLNLNPDP--QLLEGVAQWNLIESCRWIGRLEGER-----						
B9LEA3	HPLVAFHLAECYRQRNQITEIARYRNRNARLNLNPDP--QLLEGVAQWNLIESCRWIGRLEGER-----						
B8G3A2	SPLVAFAFHLaecyRQTYDVRVADAIReYRNaeHLLHSLPPD--HLIEGVAVVWLAESCRWIRRFEGER-----						
A7NRP0	AALVSYHLALAYLARQAGRQEQMREFRSALSKLARHRS--DLIEGVEVGWLRTTCEQHLM-----						
A5UPC3	AALVSYHLAMAYRQAGKKELAAREPFRSALRKLAARYEAR--DLLEGVEVGWLRTCEQHLSLET-----						
B7K4M9	B7K4M9-----						
EOUAE6	CEEAUAE6-----						
B2J4R2	B2J4R2-----						
B1Z0W8	B1Z0W8-----						
Q0B461	HHEALTHLATLTDLVGGDAGAQWILLERARRAAG-----						
Q1BLP9	HHEALTHLATLTDLVGGDAGRAQWILLERARRSAG-----						
AOAYP9	HHEALTHLATLTDLVGGDAGRAQWILLERARRSAG-----						
B1K4W3	HHEALTHLATLTDLVGGDAGRAQWILLERARRSAG-----						
A9AP36	HHEALTHLATLTDAGGDRAGAQWILLERARRSAG-----						
A4JLK2	HHEALTHLATLTDAGGDRAGAQWILLERARRAAG-----						
A1UYE3	HYEALTHLATLTDIAGDRDGAQWLMQRARRAAQYESAA--SVTDTDGDPRGTHGTRR-----						
A2S0C7	HYEALTHLATLTDIAGDRDGAQWLMQRARRAAQYESAA--SVTDTDGDPRGTHGTRR-----						
A3MB25	HYEALTHLATLTDIAGDRDGAQWLMQRARRAAQYESAA--SVTDTDGDPRGTHGTRR-----						
Q3JJX9	HYEALTHLATLTDIAGDRDGAQWLMQRARRAAQYESAA--SVTDTDGDPRGTHGTRR-----						
A3P8B0	HYEALTHLATLTDIAGDRDGAQWLMQRARRAAQYESAA--SVTDTDGDPRGTHGTRR-----						
A3NMV6	HYEALTHLATLTDIAGDRDGAQWLMQRARRAAQYESAA--SVTDTDGDPRGTHGTRR-----						
Q2T726	HYEALTHLATLTDIAGDRDGAQWLMQRARRAAQYESAA--SVTDTDGDPRGTHGTRR-----						
B2JNG9	HYEALTHLATLTDIGGRDAGQAQNLVQRAQRSAAQ---TDASADEAQRSGFHASRR-----						
B2T7T3	HAEALAHLAALTLLELEGDRDAARLIMLQRAQRQGQQR-----						
D8J0F2	HQAALLQLAALLAQAGDASQGQMRQRAERMAKPGEGA--HG-----						
A6STX7	HYEALTHLAALLAAQGDTAGAQLMQRAQRVAASATS---QEQPH-----						
Q0JZT6	HQESLYHLLAALLDTEGDHAGAMHLQRAQRHTRKH-----						
B3RBS6	HQESLYHLLAALLDTEGDHAGAMHLQRAQRHTRKH-----						
Q46V25	HQEALHHLAALLDTEGDHAGARLQRQRAQRHTRTA-----						
Q1LGB7	HBEALYHLLAALLDASGAAQAHRLRERAQRHARMHSHG-----						
Q2KV72	HRQAMLQLAALLHAEKGQEASERLSARARRLGP-----						
B7V0Y7	HAEALAHLLAALLAARGDHAGARRLQRAARGVNKDG-----						
Q02RH8	HAEALAHLLAALLAARGDHAGARRLQRAARGVNKDG-----						
A6V180	HAEALAQLAALLAASGDHAGARRLQRAARGVNKDG-----						
Q8MS8	HPEALVHLAALLAAQGDLAGARRLQRAARAGRESER-----						
A5W896	HPEALVHLAALLAAQGDLAGARRLQRAARAGRESER-----						
BOKRK9	HPEALVHLAALLAAQGDLAGARRLQRAARAGRESER-----						
Q1IDW8	HPETLVHLAALLLASQGQVAGARRLQRAARAGRESER-----						
Q98PC8	HREALVHLAALLLRQGDHTGAEALAGLGRVQKRSGS-----						
D8FC16	HVEALNHMAMFMDLHRGNKAGAERLKRQRIIGMCGVVER--RSTENQSSGRYK-----						
C6BU38	HMESLVLHLLAALYERDEERGDEERHFRNRSAAAEGE-----						
Q6AM00	HNESLHLHLLAALYERDEERGDEERHFRNRSAAAEGE-----						
A0L7P7	HWLHYHFLAELYIAIBQSSLQREYRVLRLQDLPRTV-----EGTLLALLALAVPKQISFLCQQLAKLAVHS-----						
Q2W2W8	CWPAPHYLALYELRSADSPDLRSRSYRVALQOLSCQPEP--DGLLALPLGLPVA--EVLFLCERQAGTARSQA-----						
Q1NKB1	CWPAPHYLALYELRSADSPDLRSRSYRVALQOLSCQPEP--EARRRSFFPLSCNPNAHFVNICRHKLARL-----						
Q1NWT4	CWPAPHYLALYELRSADSPDLRSRSYRVALQOLSCQPEP--EARRRSFFPLSCNPNAHFVNICRHKLARL-----						
Q1NLW5	CWPAPHYLALYELRSADSPDLRSRSYRVALQOLSCQPEP--EARRRSFFPLSCNPNAHFVNICRHKLARL-----						
Q1D4W0	LAIGHYLLAQVHERTQDHEDAARRSRSYRVALQOLSCQPEP--RPLAGHYPEMPDSADAISRAARYALAALEEQPLR-----						
E3FYQ2	LAIGHYLLAQVHERTQDHEDAARRSRSYRVALQOLSCQPEP--RPLAGHYPEMPDSADAISRAARYALAALEEQPLR-----						
A7H826	LAALHYWLGRCREAQRDPERALARVNALTAHWRPEGKRAQFLGYPDVPEDGAFAVRAQYALAL-----						
Q1D226	HAAARYLLGILLEQCRRTAATAYRRLAQLAESGRS--RPVAFPLNPDRLRVACAHAAERLEFSIQKR-----						
Q0NE87	LAARYLLGMLMELREFEPEAGYRRLRSLEEGKA--RATPFPLNPARLQVACARVERMEVGKSR-----						
Q1DX4	LALGHATLVLTYLTYLERDEEDARQNALRDLGDDDEHPLRGVETMTAGGLRQALAPVQEAGWLGVR-----						
Q08UZ6	LALGHATLVLTYLTYLERDEEDARQNALRDLGDDDEHPLRGVETMTAGGLRQALAPVQEAGWLGVR-----						
A7H715	LAMAHAAVLVPLYARVGRDEEARRNNALEAVEGLDDTAPLGRVVEPITVGALRSLALGAVHRPQAGAGRSE-----						
Q1CX10	LVAWAHLILGRVLRRLRGDTAGALRAFRAEEELCAALPPDTAVPLSEGERASLARVHNTERVLETMTREER-----						
E3FLZ3	LAVAHLALGHLVRRRDDLALGALRAFRAEEELCAALPPDTAVPLSEGERASLARVHNTERVLETMTREER-----						
A7HCQ0	QPVHFHTGCLLRLRGDTAGASRLRTAASLAQRMPAAPVRLSHGMTAGALVEGARFLHSRLAGDGPSP-----						
E3FRV5	FALGHHLRLGILARRAQEATPARVALRILATLHLAHEQPLHLLTFGGGGFRHGLMVCQCELRACAEVP-----						
Q08UV4	FALGHHLRLGILARRAQEATPARVALRILATLHLAHEQPLHLLTFGGGGFRHGLMVCQCELRACAEVP-----						
A7HCX6	FAMPRHLHLLAKRGDDEAARRRLSRLVLLAGEEGSRILMLGGTTRDALLDVCAELNACGGIP-----						
C6C1V4	FIMAFLHLLGNLNIYEQNDSNGALRHFRISMQLQDQDPVPCSDNVTAGRLMEMVRVQQQLVQ-----						
D2Z5P5	FVMAHFALLGTAQSNDRDRARHIRNVEELLQMADEAVPVYEGEATVGDLLAALKL-----						
DOLTR4	LAFAHVVAATSFQAODERAQARRNLARVILASMPASDMVRGAQITAAELHACRORALLGERAPSAPSAGAKS-----						
Q08XU2	YLPGLLELALLERNRNGARAAAPLIMRAVHARAALKPPDQLVEGPECLPARFYKASADAYLNQGAEB-----						
A0L8P6	HMMSHFHMGMFLKIQGQRRAEAGERLRRRAELENRHAQS--QTLLDPVAGLTLIDVRAQLAGYLERDRVPL-----						
DOLUDO	SWLAPYQLAMCLEARGELEKEAEEGYRHALAVIDAGGGPFPSSRAQALAHLATTAEEVCRQRVCKRAHGN-----						
D3PC59	NFVYNLFLAFALVSKNDLKNAYKFNNMALTLVEKEESS----FESLVDREVAKLILVKVLYN-----						
D3P9S2	SSASWFYGYGLLKLNLNEKMEAKFALMFIDNNVDN-SLFYISELSGEGLKNIINYQLS-----						

Figure S4B

Sequence alignment of E. coli and S. typhi proteins Q98BP3, Q2W002, D5CN06, and Q0F1Z8 across various regions of the genome. The alignment shows conservation of amino acids across different species and strains. Regions are numbered from 10 to 800.

Regions:

- 10-100
- 110-200
- 210-300
- 310-400
- 410-500
- 510-600
- 610-700
- 710-800

Species and Strains:

- E. coli
- S. typhi
- Q98BP3
- Q2W002
- D5CN06
- Q0F1Z8

Conservation:

- Primer Consensus (Prim. cons.)
- Protein Consensus (Protein cons.)

810 | 820 | 830 | 840 | 850 | 860 | 870 | 880 | 890 | 900

E. coli
S. typhi
Q98BP3
Q2W002
D5CN06
Q0F1Z8

RLRQLFANYETADFGMGQVDHKVDLQNLPFAADESYEFFICSHVLEHVESDDRAIRELYRITRTGGCGILVAPIIVGLERTVEDPSVKDAAGRWRLYGQDD

910 | 920 | 930 | 940 | 950

E. coli
S. typhi
Q98BP3
Q2W002
D5CN06
Q0F1Z8

HVRLYAHDDYVNKIRSHGFVAELGEEYFGEEIFHSLGLHTSILYVVSK