**Fig. S1.** Alignment of characterised AT3 acetyltransferases. Protein sequences are in the same order as Table S1 after *Salmonella* ser. Paratyphi A OafB WP\_00400612. SGNH fused acetyltransferases are indicated by a grey box. Asterisk mark residues selected for mutation from this alignment.

000400612/1-640	1 Mehlkyrpdidg <b>ir</b> aia <b>v</b> ls <b>vv</b> if <b>n</b> yfpsl
000639473/1-609	1 MIYKK FRLDING <b>LR</b> AFALIS <b>VV</b> LYEFGVPYV
U02679/1-640	1 MEHLK YRPDIDG <b>lr</b> avavls <b>v</b> vif <b>n</b> yfpsi
X87447/1-622	1 MSSIK YRPEIDGLRAIAVISVIIYELNEN.
AC99369/1-622	1 MKRTTR. YSRKYVPSIDG <b>LR</b> ALAVIAVIAVIANFS.
002245844/1-622	1 MOAVR YRPEIDG <b>LR</b> AVA <b>V</b> LS <b>VM</b> IFFILNNR
011101182/1-660	1 MNENLRSVNMSRTMSRM.NRHRM.NPRRYITGFDGIRTLAVLGVIIYHLMPA
004888877/1-615	1 MRI KWFSLVRITGLVLVLTYRFFOT
000379821/1-603	1 MOTKDFKRLE
021723064/1-605	1 MKR YVTGFNGLRTIGVLTVILLERLWPN
001220853/1-605	1 MRI KWFSLIRIIGLLLVILLYBFFOT
003687310/1-624	1 MSOAL PYRPDIDTLRAAAVLSVIVFRIEKD.
031545/1-343	1 MKRLVYMDWLRVLATIAVVTIEVSAGYV.SVLDANNAS.RW.M
010975904/1-335	1 LKT. IHGIOYLRAAAAIAVVLFRAAEKTG.
011037591/1-364	1 MNTVTGASGAWAPVQAAGARAFA.SGRSRDPRIDATKAIAILLVVFCHAKGVPH.G
	1 MITTAAITAGRVDTIAS.TVAERDWOIDVAKALAIILVALGRASGMPP.A
A75102/1-357	1 MYNKLTTSQATYLNFLRGFSAIIVLAGETLSGIP.GVI.S
8888/1-367	1 MGPSNEHSSGHRNNFDLLRLFAACOVMFSHAWNW.IHLGD.PLNG.TW
2778/1-373	1 MLDNIRAGAK
3214/1-333	1 MHKSNCFDTARLVAAMMVIVSHHYALSG.QPEP
009895914/1-394	1 MQTQTAS
004545264/1-394	1 MOTOTAS
706267/1-390	1 MHMIEINSLLLITSVILMSLLAVGLFDKISPINL.VEHGRNNOIDGMRGFLATFVLIHHAAIWNG.YLSSG.VW.EAP
W62204/1-382	1 MFEIDSLL.LITSVIILSLLAVKLFDEISPIOLVDHGRNNOIDGMRGFLAIFVLIHEAAIWHG.YLLTG.VW.KTP
000282635/1-349	1 MYNNKIDS
4131/1-367	1 MEKNSFP
004194788/1-412	1 MKSEIPVTVP.DRFDAADVLGAPOSARVADPSTLANPADRLSTHDNGFGLLRLLFATMVLWDBAFPLGG.FGADP.MW.R
001230914/1-332	1 MRKN
000170108/1-342	1 MTKV
37669/1-331	1 MOPK
70J69/1-422	1 MISPVVENTAKNKNNAAVTHLVSLTGLRGILALMVFGTHFFLAFVMAGG.SMDS.PFVQQ

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WP_000400612/1-640	31	LPGGFVG <b>V</b> DI	FV <b>isgyl</b> itsi	ILKSAS.NK.S	FSYLDFYKRRVL	RIF.PALSI.VL	VSCL.IVGWIY.LE	Q.D.DY.KLLGKHVF.
WP_000639473/1-609	32							ST.N.EY.EALSKNAI.
SIU02679/1-640	31							Q.D.DY.KSLGKHVF.
AAX87447/1-622	30							Y.N.DF.NKLRKTIE.
CAC99369/1-622	35							H.P.EILKNLRGDAI.
WP_002245844/1-622	30							Y.E.DF.NQMRKTVE.
WP_011101182/1-660	51							Q.R.SLLTNIRATVL.
YP_004888877/1-615	26							D.H.DFVTDIGKQVA.
WP_000379821/1-603	41							'K.P.ELIIQMKRDAI.
WP_021723064/1-605	28							Q.P.NLWAGLRSNFL.
WP_001220853/1-605	26							R.Q.DYVAGIGGQIA.
WP_003687310/1-624	31							K.D.DF.FLLWKSAL.
YP_031545/1-343	43	N.LFESISRAS <b>V</b> PI		T K D				P.A.S
WP_010975904/1-335	29							F.P.NLVLTLEHVL.
WP_011037591/1-364	55							T.R.NI.GEK.
AAA86375/1-379	48	YKLFAYSFH <b>V</b> PL						JS.A.VL.NGT.
AAA75102/1-357	40							AK.A.DL.ASMPELKEWV.
P08888/1-367	48							GG.L.NV.TGIL.
Q52778/1-373	59							P.MKS
P23214/1-333	35							'S.A.E
WP_009895914/1-394	47	~						R.T.GYLGTPTGNMIDSL.
WP_004545264/1-394	47							R.T.GYLGTPKGNIIDAL.
NP_706267/1-390	77							K.S.GWRMQVS.
EFW62204/1-382	75							K.S.RWSVHVS.
WP_000282635/1-349	40							GG.E.TI.DYIYNHIY.
Q54131/1-367	39							N.V.RY.GIHTGDAF.
WP_004194788/1-412	79							Q.A.GTLHGYWNAALG
WP_001230914/1-332	40				ITYLYILQ			
WP_000170108/1-342	44							Q.K.LA.GASV
P37669/1-331	44							TSI.NM.ELALKNLL.
Q70J69/1-422	59	HVLFSRAGGTA <b>V</b> SC	THLESEFVLAYI	AKPG	DSTRSFYRRRIA	KLY.PVHLI.ST	AVAF.ILV.SV	R.Y.EV.PGMEVTL.

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WP_000400612/1-640	108	SGSFFISNFTLWS.ESG.YFDSKSY.L.KPLLHLWSIGEEQFYIIWPVVILLCF.RSK.N
WP_000639473/1-609	106	SSLLFYSNNYYAI.HSS.YFDSSSE.F.NFLLHTWSLSVEWQFYILYPLLVIIVK.KLR.F
SIU02679/1-640	108	SGAFFISNLTLWS.ESG.YFDSQSY.L.KPLLHLWSLGIEEQFYILWPLVILLCF.KSK.Y
AAX87447/1-622	108	LAIAFLSNFYLGL.TQG.YFDLSAN.E.NPVLHIWSLAVEEQYYLIFPLILILAY.KKF.R
CAC99369/1-622	113	ASFFYVSNWWFIFHNVS.YFDSFGL.P.SPLKNLWSLAIEEQFYLIWPAFLLVFL.KWV.K
WP_002245844/1-622	108	LSAVFLSNIYLGF.QQG.YFDLSAD.E.NPVLHIWSLAVEEQYYLLYPLLLIFCC.KKT.K
WP_011101182/1-660	129	TNLVYVYNWFEINHGQS.YFDRFNG.E.SPFTHLWSLSIEGQYYLFWPLVIGILM.VIF.K
YP_004888877/1-615	104	AALGFTTNYFEIA.TGG.SYENKFI.P.HLFVHTWSLAVEMHFYILWGLIAWLVA.KIS.R
WP_000379821/1-603	119	AAIFYVSNWWYISQNVD.YFNQFAI.EPLKHLWSLAIEEQFYLLFPLVITFLL.HRF.K
WP_021723064/1-605	106	SAIFSVQNWWQISQGSS.YFADIAG.A.SPFKHI <b>Y</b> Y <mark>L</mark> SIEGQFFILWP <mark>L</mark> LLIVLL.KFV.K
WP_001220853/1-605	104	GVLGFMTNFYELL.TGG.SYESQFI.P.HLFVHNWSLAVEVHYYILWGLAVWFLS.KQA.K
WP_003687310/1-624	109	TALGFASNLYFAR.GKD.YFDPAQE.E.KPLLHIWS <mark>L</mark> SVEEQFYFVFP <mark>I</mark> LLLLVA.RKS.L
YP_031545/1-343	109	
WP_010975904/1-335	103	ASLFFVPARAGSLFMPSPSSGE.IWPVLVQGWTLNFEMLFYAVFAGSLFMP
WP_011037591/1-364	121	AARWGSHPWWEPIVS.MFTGVGPDL.YVQPPLWFLPVMLVTVIGYVLLR.RWM
AAA86375/1-379	113	SQSWAG
AAA75102/1-357	115	ANIFMIQHTPFNR.IFE.FLPTIPP.L.AQISPLWSILAVEWWLYTLFGLAFFFHK.SSFAN
P08888/1-367	116	QYLFYF.TVYILTAARIWAVYFTYEPYTMSGFYGA.S.DPSGVLWTLTVELTFYLTLPMLLEIWR.RWK.R
Q52778/1-373	124	s.t.evigty <b>w</b> f <b>l</b> waafisfili
P23214/1-333	103	yfshdivrktiss.ifmsqapdaditshlih.a.gingsl <b>w</b>  T <mark>l</mark> pleflcyiitg <mark>W</mark> avahlk.n
WP_009895914/1-394	134	SAVAQVANFHFWS.CYA.FPAKT.C.GIDQVYWSLSLEEQCYILLPVLLYSAS.RRT
WP_004545264/1-394	134	SAVAQVANFHFWA.CYA.FPATT.C.GIDQVYWSLSLEEQCYIVLPVLLYFAN.RRT
NP_706267/1-390	147	TEELFVSIMKWLPFT.ALGMPNIN.DVKDSFT.INAAVTWTLVYEWFFYFSLPVISALIK.RKV.S
EFW62204/1-382	145	PASLTVSLMRWAPFT.ALGMPNIN.GVKDSFT.INAAVTWTLVYEWFFYFSLPVIAALFK.RRV.S
WP_000282635/1-349	111	DIIYGVRN
Q54131/1-367	106	STGLYETVKLAI.KSN.FHN.N.KMFLTGWFLFAYIFVSILSVLIIKSIK.RVV.V
WP_004194788/1-412	157	GPLGYITNNWRLTIGQYGIND.LLRDTTPYGHSISE.S.VFNGSIWTLIYEAKCYVMVGLFAMFGL.LTA.H
WP_001230914/1-332	101	iQ.R.GYFFQF <b>W</b> FFGALILIYLCLP <b>I</b> LRQFLNSKRS
WP_000170108/1-342	108	
P37669/1-331	111	QK.PVFYHL <b>W</b> FFFAIAVIYLVSP <b>L</b> IQVKNV
Q70J69/1-422	131	SHVFLVQSWVPSQWYYL.TMTGIDWSLSCEAFFYLCFPVLLPVLS.RAR.N

WP 000400612/1-640	163	HNRNIVLSCAT.IFIISYA.ISIF.TMASD.GGAGANYYSPA
WP 000639473/1-609	161	.pvgls <b>l</b> sv.ilamsla.itlm.rvtgt.kedifylip
SIU02679/1-640	163	skrnillsca.ifivsyt.isvf.tmaye.gganyyspa
AAX87447/1-622	163	.EIKVLFI <b> I</b>  TL.ILFFILLA.TSFI.PANFYKEVLH.QPN
CAC99369/1-622	169	NPKLLLKI <b>V</b> IG.LGLLSAVWMT.ILYV.PGTD.PSRVYYGTD
WP_002245844/1-622	163	SLRVLRNISI.ILFLILTA.TSFL.PSGFYTDILN.QPN
WP_011101182/1-660	185	KRSRVFWF <b>M</b> MI.AAGISAITMA.MLYDPAN.TNRNRVYYGTD
YP_004888877/1-615	159	RLASGRRALTRFRWGLGL <b>L</b> TTAFAIESFVMMY.FGAV.GLKD.FSPSPVYFSSL
WP_000379821/1-603	174	PRNIIQT <b>L</b> FI.VSLISLGLMI.VIHF.ITGD.NSRVYFGTD
WP_021723064/1-605	162	KRGRIFVI <b>A</b> NL.LALISAIWMA.ILFV.PGAD.PTRTRVYYGTD
WP_001220853/1-605	159	SNGQLKGMVFL <mark>L</mark> SAVAFLISFFSMF.IGSF.LVTTS.YSSVYFSSL
WP_003687310/1-624	164	.RVQFGF <b>L</b> AA.LCALSLA.ASFM.PSALDD
YP_031545/1-343	151	REIEYF <mark>L</mark> IL.WLYASVV.VNLVKYYY.PINFNINIELFYVTN
WP_010975904/1-335	146	RNWRLPVV S GLFLALVIAGRV.VAFFD.D.DAVAVMLTYTR
WP_011037591/1-364	171	PPLVIIAA.VAVVLAW.FWMNWF.PLQHMRMRLFWGLD
AAA86375/1-379	163	saav <mark>l</mark> av.csllvvw.awtrwf.pglrlrlrlrlrlpfald
AAA75102/1-357	171	.RLIMSI <mark>L</mark> .I.IPALLVA.GYFT.LKEY
P08888/1-367	182	.AGALVVAVAALG.SWVMAQHF.NITD.KYNPFLSVTAAG
Q52778/1-373	160	.RLSIW
P23214/1-333	162	.GKAFI
WP_009895914/1-394	186	.IMPILCVVVLI.QVFL.PR
WP_004545264/1-394	186	.IAPILCV <b>V</b> VLI.QVFL.PR
NP_706267/1-390 EFW62204/1-382	208 206	IYMVM <b>I</b> SA.ISLFVFILK IHMII <b>I</b> SV.LVLVIFLCK
WP 000282635/1-349	147	NNIVILI <b>I</b> SU.LUSFMLSOT.VLNHN.PLLDPOWFWNID
054131/1-367	156	SNALLILSV.LVAISVL.LITV.SITYL.SPOYILVKDY.K
WP 004194788/1-412	224	RRVLLAVTV.VAWFVLA.IQTINPAFSAQLLPWAGD.RHLVQYVQY
WP_004194788/1-412 WP_001230914/1-332	144	YLYSLSL.LMTIGLI.FELS.NILLQ.MPIQTYVIQTFRL
WP_001230914/1-332 WP_000170108/1-342	152	SLFLVSV.FAYIFTLF.IOID
P37669/1-331	140	GGKMLLVL <b>M</b> AV.IGIIANPNTV.PQKID.GFEWLPINLYING
070J69/1-422	179	GM LYA VSACS.VFLV.FFLPYVVG HTF.AI
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WP_000400612/1-640	199	.SRFWELMACAIISTL.RFI.GI.NTSLSKLM.SLLGI.IL.IALLSIT.M
WP_000639473/1-609	194	TRAWEMLAGGLVYIA.SVR.YK.MPEWIKHC.EVYGI.VL.IV
SIU02679/1-640	199	SRFWELMAGAIIATL.RFM.GI.KTSVSKSM.SLIGV.II.ITLSIA.L
AAX87447/1-622	204	LRFPELLVGSLLAIY.HNL.SA.SKQASKQASNVIAI.LST.LL.LF
CAC99369/1-622	207	TRAFDLLS <mark>G</mark> CALAFV.WPF.NRLSPVVPRKS.KAVLNIAGTISILCF.IL
WP_002245844/1-622	204	.LRFPELLA <mark>G</mark> SLLAVY.GQT.QNGRRQTANGKRQLLSS.LCF.GA.LLACLF.V
WP_011101182/1-660	222	TRMFAILL <mark>G</mark> SGLAFI.WPS.RELSADIANVN.RVTLDILGGASL.IA.II
YP_004888877/1-615	208	THCFPFFVGGLIGVL.SGI.KA.QGPIYRWT.TAHCNPIIAGAVMLISF.IL.LVALGD.R
WP_000379821/1-603	211	TRLQTLLLGCILAFI.WPP.FALKKDISKKI.VVSLDIIGISGF.AV.LMTTLFF.I
WP_021723064/1-605	200	TRFFSLIMGASLAFV.WPL.NKLSHKVNKRAVKIAWQLTIGLSL.LL.LL.LLAYIF
WP_001220853/1-605	201	THVYPFFLGSMLATI.VGV.RQ.TTSLVKQL.DKIWDLRKTLVVFGGGFGF.LV.LL
WP_003687310/1-624	195	.LRACEMLVGSLTAVR.MRY.RQQRN.PAVGKRY.AAVGALFSA.CI.LSAACLF.A
YP_031545/1-343	188	YVGYFLLGYYLSNFDI.SKKWRNIS.YIGG.F.VGF
WP_010975904/1-335	180	.PVILEFVAGMIIGEF.WLK.GRVPPLAVGS.ALFACSLGG.FA.LI
WP_011037591/1-364	203	.VLPVSLCFYALGALLIHV.SPYLPT.SLPGSALVTVVTVVTVVVAALVA
AAA86375/1-379	195	.VLPVALFFIAVGAWLSRF.AERV.R.ALPAVVWVVA
AAA75102/1-357 P08888/1-367	194	valvwfl <b>g</b> sgcayh.f.c.ninrkynnhg.ilmlslitga.af.lv
052778/1-373	217 189	TYPFYCL <b>G</b> FLFAOP.IGWO.NGVIWRYK.WIFV.VLLSIAAA
P23214/1-333	198	LRGLAFFF <b>G</b> ATMAMY.EKS.WNVSNVKITVVSL.LAM
WP 009895914/1-394	211	.URTDALAL <b>G</b> ALLALL.QHR.EW.YAKVEPRTLARGLLSA.IALAGM.CL.IL
WP_009895914/1-394 WP_004545264/1-394	211	.VRTDALALGALLALL.QHR.EW.YAKVEPRPLGRGLISA.YALAGM.CL.YLALALA.G
NP 706267/1-394	228	.IHIASFLFGL.LAFL.LNK.SKIVNGIAKAKVTPIIITAIM.V
EFW62204/1-382	227	.IHVISFLFGL.LASY.SNNYVHRVAKAKITPIMIIALL.SY.E
WP 000282635/1-349	182	.SAMAYWWLLPLCRCMF.LELT.RDRFFGKSKIGF.IV.FSITAIMSAYQLLNQKPLLFKIISI
054131/1-367	191	LNFICOVLTCOVLTCCNS.FYI.FGY.VI.RNOIYNLL.NF
WP 004194788/1-412	264	GTĪF LI <b>G</b> SSAAAY.SKS.L PISDK LGA FAV.VV.YL
WP 001230914/1-332	180	WTWFFYYLLCGYIAOF.TIE.EI.ESRFKNWM.KIVSI.LL.LL
WP 000170108/1-342	178	LVWGICFFL@SVLSEI.HFD.KINLK.KFLFF.FV.LFDFIYMFA
P37669/1-331	179	.DTFYYILY <b>G</b> MLGRAI.GMM.DT.OHKAL.SWVSA.AL.FA
Q70J69/1-422	234	.MRMAEFVIGISLAVL.VRR.GA.WRGPGVAAC.LV.LCAIGWGVN.L

WP_000400612/1-640	241	IDEKMIASN.GNDLVVSKLLSVRPV
WP_000639473/1-609	235	ILHSNGILAN.KQ.NSLFTSNRIA
SIU02679/1-640	241	INEKMIASN.GNDWIASKILSFKPI
AAX87447/1-622	250	MNNDIA
CAC99369/1-622	257	VSEYQPFATIS.HPASYLSKIFSFKPL
WP_002245844/1-622	251	IDKHNPIRSM.QYGTLPTRILSASPI
WP_011101182/1-660	272	MSGQSDFATVA.HPASHLNRVLTNPLF
YP_004888877/1-615	262	MKFDALQ
WP_000379821/1-603	261	VGDQDQWAIAV.HPSSLFAKFLSMKPL
WP_021723064/1-605	250	MPAQGTFALVA.HPSLPTNKLFSNPVF
WP_001220853/1-605	255	VKFTYLFLAAR.VLHEKTHHIQESKII
WP_003687310/1-624 YP 031545/1-343	243 226	YSEQTAIYFN.HYEHPLKKFFQWKIT
WP 010975904/1-335	225	.TYFYTVKA.NGQLEQFWYGYFAPGVVLMAIGLFIFFKYAFQKSERELPLLF LPFDELTV.IGVLS.LEANGCVRALSLP
WP_011037591/1-364	244	WLAGVNG
AAA86375/1-379	235	WGGV.AAMNGOVDVNNLOFGKSSLLFLIASL.LGTA.MTLLCI.AYFMOGWRWL
AAA75102/1-357	238	LK. HSL M
P08888/1-367	260	. TS. AAFI
052778/1-373	226	.FICFLGWG.KE.TYAYNNLVLIHDEQSAKQVFLMFSGSLAASAV.AMQSMFQCWRLVYSTRVARFVA
P23214/1-333	235	A
WP 009895914/1-394	262	RLASIPIASY.GKSYTMPDGAVKRVL
WP_004545264/1-394	262	YLARIPIASY.GRGYTMRDGAVKRVL
NP_706267/1-390	269	MTY.FKTIIAS.GCDLYGILRLNIT
EFW62204/1-382	266	MIY.FPTLIAS.GCDFYGILRLNLT
WP_000282635/1-349	241	FNVGLIIFNI
Q54131/1-367	231	YVSKSYGFSTQTIMSWSYYPDGLIMSVI.NALI.GIYAVFFISLLI.TRGMKEIKLL
WP_004194788/1-412	297	SYAIL.WLACRLPRWARRI
WP_001230914/1-332	222	LFVKVILFFIAKTIYHNLFAEYFYDTLFVKVKV.STLG.IFLTILMLTLNESR.RESI
WP_000170108/1-342	218	WFLFYEVGS.KK.DYVSYINPGLWGIAF.IVCV.LVAFAIFPK.ME.KNFPKTFLYF
P37669/1-331	218	ISRGTLYELQW.RLVKNTLDTRTIRGL
Q70J69/1-422	275	HL.DSYLALST.ADVNSAWSPLRWKPL

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WP_000400612/1-640	282	VFFGLISYPLYLWHWPIYSF.YRSIFAG	S.P	DY.HEL	TLLLL.LSFFLAIL
WP_000639473/1-609	274	QWVGKI $ \mathbf{S} $ YS $ \mathbf{V} $ YLWHWPVIVA.MKHYDIEF			
SIU02679/1-640	282				LILMA.LALVLAIL
AAX87447/1-622	292	VFIGKI <b>S</b> YS <b>L</b> YLYHWIFIAF.AYYITGE	KQI	NN.QSI	AIVII.LTIIFSVL
CAC99369/1-622	300	RWIGTRSYGIYLWHYPIITL.TTPVLEI	TQP	NI.W	ILQVA.ATFIIAEL
WP_002245844/1-622	293	VFVGKI <b>S</b> YS <b>L</b> YLYHWIFIAF.AHYITGD	KQL	GL.PAV	sAVAA.LTAGFSLL
WP_011101182/1-660	315	KYVGQR <b>S</b> YG <b>I</b> YLYQFPVMIF.YETKVKNI			
YP_004888877/1-615	305	TFLADV <b>S</b> YS <b>V</b> YLYHWPLYVI.FSHMMV			
WP_000379821/1-603	304	LIIGKR <mark>s</mark> ys <b>l</b> ylwhyPIIVF.VNSYYVQ	GQI	PV.YVY	IIEIL.LTALMAEI
WP_021723064/1-605	293	EYIGSR <mark>S</mark> YG <b>I</b> YLWQLPVFAF.AEAKVL	APT	AW.YNL	IWQLA.LILILTEL
WP_001220853/1-605	298	SFLADT $\mathbf{S}$ YA $\mathbf{V}$ YLFHWPFYII.FSQLTS	N.L	L	LLTLI.CSYGFASL
WP_003687310/1-624	285				AAAIV.LTLAFSLI
YP_031545/1-343	276	RFINQASLGIYILHFFLLNN.LLYMVFPK.V.			
WP_010975904/1-335	259	GLLGDA <b>S</b> YS <b>I</b> YLWHTFAISV.VAKAGLAI			
WP_011037591/1-364	291	QWIGRNTLLTLCTHMLVFFV.LSGVAALA			
AAA86375/1-379	284	RWIGANTLLILGTHTLVFLV.VTSVVVRT			
AAA75102/1-357	279	AFLAFI <b>S</b> YT <b>L</b> YLSHEPIRRV.VSTFIEPT			
P08888/1-367	298	LRRQDL SYG I YLYHMLVMHT.LIAIGW.V			
Q52778/1-373	290	VQLGQS <b>T</b> LL <b>L</b> YLVQGAVFRL.MDLIQFGE.V.			
P23214/1-333	268	KGRFDYSYGVYIYAFPVQQV.VINTL			
WP_009895914/1-394	302	VYFGSR <mark>S</mark> YA <b>I</b> YLIHVPVYRL.TREIWERVATI			
WP_004545264/1-394	302	VYFGSR <b>s</b> ya <b>i</b> ylihvpvyrl.treiwerivij			
NP_706267/1-390	307	RKLGET $\mathbf{T}$ YS $\mathbf{V}$ YLLHGIFLYCLMTWIIPN			
EFW62204/1-382	304	RKLGET $\mathbf{T}$ YS $\mathbf{I}$ YLLHGVFLYCVMTWVIPS			
WP_000282635/1-349	281	VRAGRN <b>T</b> LN <b>I</b> CGMEYITKLFIPMA.LAIIGFSV			
Q54131/1-367	285	KMIGQN $\mathbf{S}$ RA $\mathbf{I}$ MAYHLLVYVI.LDIIAS	IL	GD.YSLSGTDVYDN	HFITK.WSVPVY.IALGLLLP
WP_004194788/1-412	330	GSRNDY <b>S</b> YG <b>I</b> YVFGFLVQQV.LAYVGAY			
WP_001230914/1-332	270	VSLSNQ $\mathbf{T}$ MG $\mathbf{V}$ FIIHTYIMKV.WEKVLG.F			
WP_000170108/1-342	269	TKYGKD <b>S</b> LG <b>I</b> YILHAPICSM.IRILMLKV			
P37669/1-331	268	GLISRH $\mathbf{S}$ LG $\mathbf{I}$ YGFHALIIHA.LRTRGIE			
Q70J69/1-422	315	VWFGEI <b>S</b> LSFYVSHLLVQEE.LFSRLWNL	GMRAGLLPAPL	PVLSW.WVAVLSF	LAQFA.LAVLAAWL

		. *
WP_000400612/1-640	329	TYYLTEKPLRNSRS
WP_000639473/1-609	320	SYRT I ENTLRKRVKFRF.SDLQF.NIVLFSST.LALCLFVMFTKGVSFRF.SD
SIU02679/1-640	329	TYFL <mark>L</mark> EKPLRHSRKKS.ITTIILAVV.VFG.SGIFGIVTYSM.NGIKERSVNKS.AGEYA
AAX87447/1-622	340	SYYL <mark>I</mark> EQPIRKSKLNF.KQSFLYIYF.IPS.LLLLGFNLYKRQT
CAC99369/1-622	348	SFRF I ETPIRKNGFINYFKGFKDKNYFIWKNKPVGKW.LSIAGVVA.VLAIFTLGMSNVLSVNT.NAEKQQ.TSVKTTTSTPDEK
WP_002245844/1-622	341	SYYL <mark>I</mark> EQPLRKRKM
WP_011101182/1-660	364	SYRF I ENPMRHYDYSRLLVDFKDFLRKPKF.NRVTTAIVAL.TTVLFVITAVGFVQQPS.KAEANKKTELQKT.IAANSKAAD
YP_004888877/1-615	349	SYYV <mark>L</mark> EPVIAGKRAHVFGHTVTWRQL.QLP.VITAGVLLAVNTGVVVKNAPQLSTLEQN.L
WP_000379821/1-603	352	SYRF I ETPIRKKGFKAFAFLPKKKGQF.ARTVLVILLLVPSIVVLSGQFDALGKQH.EAEKKEKKT.EFKTTKKKVV
WP_021723064/1-605	340	SYRL <mark>I</mark> ELPTQRFDYSNILGILQNF.VREKGWKL.KKNILPMLIS.GLALISLGFIIFSP.PSPHDQ.RVIEEK.IMAQQV
WP_001220853/1-605	342	SFYV <mark>L</mark> EPWIAGKNTPIV.QTLRPLPY.IHA.ILAAGTGILTIIVCTV.TLLAPQVGAFETD.L
WP_003687310/1-624	332	SYHC  <mark>I</mark>  EKPFKKWKGSF.AQSVLWIYA.LPM.LVLGAGSFFAMR
YP_031545/1-343	328	ITLVL Q RIPVVKK
WP_010975904/1-335	308	AYMM <mark>L</mark> ERPLLRRGRAR
WP_011037591/1-364	345	LRWFL <b>M</b> .RFA
AAA86375/1-379	339	MRAVL <mark>V</mark> RRALDVGIE
AAA75102/1-357	325	IAYL <mark>I</mark> ENKHLVVR
P08888/1-367	344	SWAL <mark>I</mark> EQPAMKLRT
Q52778/1-373	342	IRSIARNLGYVSR
P23214/1-333	314	SWNL <mark>V</mark> EKRFLTRS
WP_009895914/1-394	359	NYRF <b>V</b> ETPLRLHGTRV.ARV.A
WP_004545264/1-394	359	NYRF <b>V</b> EAPLRVYGARL.A
NP_706267/1-390	359	TFKL <mark>I</mark> ETPFIKLTKQTT.TLL
EFW62204/1-382	356	TFKLIELPFINITKQTA.MK
WP_000282635/1-349	334	IGHWLSRTVGGPFLIK
Q54131/1-367	345	LIFSI
WP_004194788/1-412 WP 001230914/1-332	378 316	SWHL <mark>I</mark> EKRALALKD IVGML <b>M</b> KIPYFNR
WP_001230914/1-332 WP_000170108/1-342	319	ATYTIKKIPIPNK
P37669/1-331	319	LLSM <b>L</b> VORI
070J69/1-331	379	LHRL <b>V</b> EQPLVRRLRPKD.VP
Q10069/1-422	319	purpla på praktru

WP 000400612/1-640	384	
WP_000639473/1-609	364	YS.AFSKCQDKMTEKSFVVWG
SIU02679/1-640	384	
AAX87447/1-622	381	EQ.SIPVSNENHYPAKVIILG
CAC99369/1-622	429	KDDKKEDKATKDKEADSNKAS.EQKETQKPDNKNKSAATPKTIITQTVAIG
WP_002245844/1-622	382	LP.GAPLAAENHFPETVLTLG
WP_011101182/1-660	443	KKNAAALKRQKAAQAAAASSKKVATEKMQTKQAEAKLNSKQKQVEKEY
YP_004888877/1-615	407	LAAVTPKKKQ.ATPKGDQEKAPGVSIIG
WP_000379821/1-603	426	
WP_021723064/1-605	414	ALQKKQLAVVAEKASQMNVLALG
WP_001220853/1-605	400	ERAD.ANSLGIADGTMLIG
WP_003687310/1-624	372	
YP_031545/1-343		
WP_010975904/1-335		
WP_011037591/1-364		
AAA86375/1-379		
AAA75102/1-357		
P08888/1-367		
Q52778/1-373 P23214/1-333		
WP 009895914/1-394		
WP_009895914/1-394 WP 004545264/1-394		
NP 706267/1-390		
EFW62204/1-382		
WP 000282635/1-349		
054131/1-367		
WP 004194788/1-412		
WP 001230914/1-332		
WP 000170108/1-342		
P37669/1-331		
Q70J69/1-422		

WP_000400612/1-640			
WP_000639473/1-609 41D DSHAARHLMPGLKSVFGNSL.NIT.QRTA.SLCPPIIGL.QKDDRP.YCKDINDMVAKEISDNKPTTVLMSALWPVYPMRDYLPE AX87447/1-622 409 DSHASHLEAPLNYV.GNKE.GWKAD.IFKDKFECSFIVNE.QYOLDP.NCQSVWQKDSQY.KAIFISAFYDL.RMGGQPVPRFR.PETIEP.DQ WP_002245844/1-622 410 DSHASHLEAPLNYV.GNKE.GWKAD.IFKDKFECSFIVNE.QYOLDP.NCQSVWQKDSQY.KAIFISAFYDL.RMGGQPVPRFR.PETIEP.DQ WP_01101182/1-660 510 DSVLLDUSDLUPYLKEAPUPNITIDGLUGRQLURD.AITTATG.YKKFNSENSSVILE.L.GINGPF.T.EDQ WP_01101182/1-660 510 DSVLLDVSSDLQDVIPGTVVQGRVQRQVTE.VPGIINS.LKSGQLAHNVLLN.I.GTNGTI.T.DDQ WP_00379821/1-603 510 DSVLLDVSSDLQDVIPGTVVQGRVQRQVTE.VPGIINS.LKSGQLAHNVLLN.I.GTNGTI.T.DDQ WP_0112306471-605 450 DSVNUDIGNVFTKKIPNAQIDGRVGRQVVTE.VPGIINS.LKSGQLAHNVLLN.I.GTNGTI.T.DDQ WP_0112306471-605 450 DSVNUDIGNVFTKKIPNAQIDGRVGRQLVD.ATFIVKS.QYND.YAKKGGKVVVE.L.GTNGTI.T.DDQ WP_0112306471-605 450 DSVNUDIGNVFTKKIPNAQIDGRVGRQLVD.ATFIVKS.QYND.YAKKGGKVVVE.L.GTNGTI.KGE WP_00156731017-605 437 DSVADRANTALQTALPGAQINAQVSVTIKT.ANEIMIN.NSQNKFIPKTVVIA.T.GVNNPE.NYRDD WP_00379821/1-303 341 WP_010975904/1-335 344 WP_011037591/1-364 AAR8637511-379 AAA75102/1-337 355 P23214/1-333 357 P08888/1-367 QS2778/1-373 357 P08888/1-367 QS2778/1-373 355 IRGIIK WP_009895914/1-394 WP_004545264/1-394 WP_004545264/1-39	WP 000400612/1-640	429	DSYAAALFNGLSHYIDNKGSDYIIS.OMTD.GNAPPLFVDGKDDLORSVITLNNNRINEIKRVOPEVVLLTWSVRGTNGVHD.KKLAIDALSL
AAX87447/1-622	WP_000639473/1-609	411	
CAC99369/1-622	SIU02679/1-640	429	DSYAAALYNGLSSYIKNNNEKYVIS.QMTD.GNAPPLFVSGKDDLQRDVGSINADRIKEIGIVKPEIVLLTWSVRGSNGVHD.KKLAIEALSL
WP_002458844/1-622         410         DSHAGHLRGFLDYV.GSRE.GWKAK.ĪLSLDSECLWWVDEKLADNP.LCRKYRDEVEKA.EAVFIAQFYDL.RMGGQPVPRFE.AQSFLIPG           WP_01101182/1-660         510         DSVLLDVSSDLQDVIPGTVVQGRVGRQVTEVPGIINSLKSQGQLAHNVLIN.IGTNGTI.T.DDQ           WP_000379821/1-603         453         DSVVLDGTRSYLGDHVANSNIDABEDRTMNL.AYKVMMN.QQRSHTLKQYVVCIGTNADL.DYEEQ           WP_021223064/1-605         452         DSVWNDIGNVFTKKIPNAQIDGRVGRQVIV.ATFIVKS.QYKDYAKKGGKVVVE.LGTNGAF.T.KDQ           WP_003687310/1-605         458         DSVWNVAASTNIQEVFPHMYIDAAVGRQAES.LDTDLTN.AKSKMPNPDAYLIG.LGTNGAT.T.KDQ           WP_003687310/1-624         41         DSVALRANTALQTALPGAQINAQVSVTTKTANEIMLN.NSQNKFLPKTVVIA.TGVNNPE.NYKDD           YP_031545/1-343         341         L.SHADHYKTFDAV.GKKE.KWSAT.MVSA.DACAYVEGYASRVFQNWA.ACRAVYRYAEEH.LPRYPKVVLAM.RWGS.QMPE.NS.RSLAYDAG           YP_010375904/1-335         324         RVTAGLAGR	AAX87447/1-622	409	DSHSSHLEAFLNYV.GNKE.GWKAD.IFKDKFECSFIVNEQYQLDP.NCQSVWQKDSQY.KAIFISAFYDL.RMGGQPVPRFR.PETFIEPD
WP_011001182/1-660 510 DSVLLDVSSDLQDVIPGTVVQGRVGRQVTEVPGIINSLKSQGQLAHNVLIN.IGTNGTI.T.DDQ YP_004888877/1-615 453 DSVTLDTRSYLGDHVANSNIDAEGDRTMNLAYKVMMNQQRSHILRQYVVIC.IGTNALD.DYEEQ WP_001739821/1-603 452 DSVMVDIGNVFTKKIPNAQIDGKVGRQLVDATPIVKSQYKDYAKKQKVVVE.LGTNGAF.T.KDQ WP_01120853/1-605 458 DSVMVAASTNLQEVFPHMYIDAAVGRQAESLDTDLTNAKSKMPNPDAYLIG.LGTNGTI.K.EGE WP_00120853/1-605 437 DSVALRANTALQTALPGAQINAQVSVTKTANEIMLNNSQNKFLPKTVVIA.TGVNPFE.NYKDD WP_03687310/1-624 414 DSHADHYKTFFDAV.GKKE.KWSAT.MVSA.DACAYVEGYASRVFQNWA.ACRAVYRYAEEH.LPRYPKVVLAM.RWGSQMPE.NS.RSLAYDAG YP_031545/1-343 341	CAC99369/1-622	479	DSVMLDIEPYLKEAVPNITIDGLVGRQLRDAITTATGYKKFNSENSSVILELGTNGPFT.EDQ
YP_004888877/1-615       453       DSVTLGTRSYLGTHVANSNIDAEGDRTMNLAYKVMMNQQRSHTLRQYVVICIGTNALDDYEEQ         WP_001723064/1-605       452       DSVWVD1GNVFTKKIPNAQIDGKVGRQLVDATPIVKSQYKDYAKKGQKVVVE.LGTNGAFTKDQ         WP_001202853/1-605       458       DSVWVAASTNLQEVFFHMYIDAAVGRQAESLDTDLTNAKSKMPNDAYLIG.LGTNGTI.K.EGE         WP_001364571-624       414       DSVALRANTALQTALPGAQINAQVSVTTKTANEIMLNNSONKFLPKTVVIATGVNNPENYKDD         WP_031545/1-343       341       L	<b>-</b>	410	
WP_000379821/1-603			
WP_021723064/1-605			
WP_001220853/1-605 WP_003687310/1-624 WP_003687310/1-624 WP_00154571-343 WP_010975904/1-335 WP_011037591/1-364 AAA86375/1-379 AAA75102/1-357 P08888/1-367 Q52778/1-373 P23214/1-333 WP_009895914/1-394 WP_009895914/1-394 WP_009895914/1-394 WP_004545264/1-394 WP_006267/1-390 SFW 002826355/1-349 Q54131/1-367 WP_00194788/1-412 WP_0049885/1-412 WP_004988675/1-390 WP_002826355/1-349 Q54131/1-367 WP_0049888/1-412 WP_0049888/1-412 WP_0049886/1-412 WP_0049865914/1-392 WP_0049865914/1-393 WP_0049865914/1-394 WP_00494788/1-412 WP_0049865914/1-394 WP_0049865914/1-394 WP_0049865914/1-394 WP_0049865914/1-394 WP_0049865914/1-394 WP_0049865914/1-394 WP_0049865914/1-394 WP_0049484784/1-394 WP_004			
WF_003687310/1-624 414 DSHADHYKTFFDAV.GKKE.KWSAT.MVSA.DACAYVEGYASRVFQNWA.ACRAVYRYAEEH.LPRYPKVVLAM.RWGSQMPE.NS.RSLAYDAG  YF_031545/1-343 341			
YP_031545/1-343			
WP_010975904/1-335 324 RVTAGLAGR. WP_011037591/1-364 AAA86375/1-379 AAA75102/1-357 P08888/1-367 Q52778/1-373 355			~
WP_011037591/1-364 AAA86375/1-379 AAA75102/1-357 P08888/1-367 Q52778/1-373	<u> </u>		
AAA86375/1-379 AAA75102/1-357 P08888/1-367 Q52778/1-373		324	
AAA75102/1-357 P08888/1-367 Q52778/1-373 355 I P23214/1-333 327 SPK WP_009895914/1-394 WP_004545264/1-394 NP_706267/1-390 378 VKELIP EFW62204/1-382 375 IRGIIK WP_000282635/1-349 Q54131/1-367 358 I.K WP_004194788/1-412 392 WGPG			
P08888/1-367 Q52778/1-373			
Q52778/1-373			
P23214/1-333 327 SPK  WP_009895914/1-394  WP_004545264/1-394  NP_706267/1-390 378 VKELIP  EFW62204/1-382 375 IRGIIK  WP_000282635/1-349  Q54131/1-367 358 I.K  WP_004194788/1-412 392 WGPG		255	
WP_009895914/1-394 WP_004545264/1-394 NP_706267/1-390			
WP_004545264/1-394 NP_706267/1-390		321	
NP_706267/1-390			
EFW62204/1-382 375 IRGIIK	·	378	
WP_000282635/1-349 Q54131/1-367	<u> </u>		
Q54131/1-367 358			
WP_004194788/1-412 392WGPGWGPG	_ · · · · _ · · · · · · · · · · · · · ·	358	I.K
WP 001230914/1-332 329		392	WGPG
ma_vvamovvas/a vva   vmv	WP_001230914/1-332	329	
WP_000170108/1-342 332	WP_000170108/1-342	332	
P37669/1-331			
Q70J69/1-422	Q70J69/1-422		

WP_000400612/1-640	519	TIKKIKEASPDSRIIFI.GPVPEWNANLVKIISNYLSEFKKT.PPLYMTYGLNSE.ISE.WDSYFSNNVPKMGI.EYISAYKAL
WP_000639473/1-609	490	TIKFLKDNK.VKNIIIV.GPFPVWKKTMIDTIEDMGINSGRT.VPWSMT.DETRN.LRD.NDKYLRELAKEHSLTYISPLETMC
SIU02679/1-640	519	TIKEIKKVSPQSRLIVV.GPVPEWNANLVKVISNYTSEFKKT.PPIYMSYGLNDE.IKG.WDKFFEENVPKLGA.EYISAYSALG
AAX87447/1-622	494	FKARFKNTVKQLAMQKP.VYVFANNSSVSRSPLRGYLLENYGLEKYLTPI.HRMGD.IDA.SNKIIHDLVKDIPNVYWVDAQQYLI
CAC99369/1-622	541	LNDLLDQFDKA.TIYLV.NTRVPRGWQSD.VNKSIANAA.SRPNVTVVDWYSRS
WP_002245844/1-622	495	FPARFRETVKRIAAVKP.VYVFANNTSISRSPLREEKLKRFAANQYLRPI.QAMGD.IGK.SNQAVFDLIKDIPNVHWVDAQKYLI
WP_011101182/1-660	572	AEQVVKLIGKDRQIFWV.TAHVPTQSWQNQ.VNAQIAKTAKKHANVHVIDWHGRAQ
YP_004888877/1-615	516	TMKIIHDLEPGHKLILM.TPYNARADADWNSSKLAVLERRLPAKYKFITVADWGKIA
WP_000379821/1-603	515	LNELLDSFGKA.DIYLV.SIRVPRDYEG
WP_021723064/1-605	520	IDAAMKVAGNK.PVYWI.NVHADRVWAK
WP_001220853/1-605	500	WDSIVKNLPKGHHMILV.TPYEGDKTKETYAIVE
WP_003687310/1-624	500	FFQKFDRMLHKLSSEKQAVYLMADNLASSYNVQRAYILSSRIPGCRQTLRPDDESTLK.ANARIRELAAKYPNVYIIDAAAYI
YP_031545/1-343	342	VV
WP_010975904/1-335		
WP_011037591/1-364		
AAA86375/1-379		
AAA75102/1-357	338	
P08888/1-367	362	
Q52778/1-373	356	VV
P23214/1-333		
WP_009895914/1-394	376	
WP_004545264/1-394	376	DR
NP_706267/1-390		
EFW62204/1-382		
WP_000282635/1-349		
Q54131/1-367	360	FK
WP_004194788/1-412	396	
WP_001230914/1-332	330	VKVK
WP_000170108/1-342	334	LPQKYIKL
P37669/1-331		
Q70J69/1-422	398	

WP 000400612/1-640	599	NESGCLTRVGNGPDFI.TAVDWGHLTKPGSDFLFNKIGNKIIK
WP_000400612/1-640 WP_000639473/1-609	568	TESYCKAIIGNRIAYP.IOYDNAHLTPEGSGWFIEEVKKOISK
SIU02679/1-640	599	NESGCLTRVGDGPDFV.TAVDWGHLTKPGSDFLMKKIGHLIIR
AAX87447/1-622	576	KDSVMAEGKY.LYGDODHLTNFGAYYMAKEFSKYORVMTPEOV.KKLYE
CAC99369/1-622	592	GQSQY.FAPDGVHLTKAGAQAYVAMLTSVMNK
WP 002245844/1-622	577	KNTVEIYGRY.LYGDODHLTYFGSYYMGREFHKHERLLKSSRD.GAL.O
WP 011101182/1-660	626	NOSGW.FADDNVHPNTTGNROLTNLIANRIAEVNNN.
YP 004888877/1-615	573	OHP.EVFKGTDGVHFGGIRAGDILYAKVINOALTAAKOTPAKPA.
WP 000379821/1-603	567	GHPEY.FAYDGIHLEYAGSKALTDLIVKTMETHATNKK
WP_021723064/1-605	572	GQSSW.FYSDNIHPKGTGAEKYAALVANSLTDVEK
WP_001220853/1-605	559	EHPEIWAGTDQVHFGSESSTIEAGAKLYADTIATALQTAQDKPVKSK.
WP_003687310/1-624	583	ADFQIGGLP.VYSDKDHINPYGGTELAKRFSEKQRFLDTRHN.H
YP_031545/1-343	343	P
WP_010975904/1-335	333	AAE
WP_011037591/1-364	354	PWTLGARPVSA
AAA86375/1-379	354	HQSMWRVRVSHRQRRSAQ
AAA75102/1-357	347	KNTSNKLAYST
P08888/1-367	363	RLSVA
Q52778/1-373	369	VIN
P23214/1-333	330	LSLD
WP_009895914/1-394	379	
WP_004545264/1-394	379 384	
NP_706267/1-390 EFW62204/1-382	384 381	
WP 000282635/1-349	201	
054131/1-367	367	N
WP 004194788/1-412	403	ARLTMKK EGV
WP 001230914/1-332	332	L
WP 000170108/1-342	342	K
P37669/1-331	325	DRNRLVS
Q70J69/1-422	408	APPA
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