Microhomology scheme: Sense/pCMVΔ/BranchΔ, sgRNA E, forward strand

d	Micr	ohomology scheme: Sense/pCMVΔ/BranchΔ, sgRNA E, forward strand
	5 10 15 20 25 30 35 40 45 50 55 60 65 70	75 80 85 90 95 100 105 110 115 120 125 130 135 140 145 150 155 160 165 170 175 180 185 190 195 200 205 210 215 220 225 230 235 240 245 250
	Primer	Exol Primer
EE1 *		CTACONOGUC CACANCECTON ACTION
EE2 EE3		CALCAGOGGICA CALACCOTA AGCTA AGGTA GOCCAGOCCOTECCCT GOCCAGO GACA ACCORDICACO GOCCAGO GACA GOCCAGO GOCCAGO GACA GOCCAGO GOCCAGO GACA GOCCAGO GOCCA
EE4		CTACGAGGGCACAACACCGTGAAGGTGAACGTAAGGTGACCAAGGGCGGCCCCTGCCCTTGCCTTGGCATGGACATCTGTCCCCCAGTTCCACTACGGTTCAAGTAAGAACCTCTAAGGAACCACCCCCAATCCCCCAAGGGCTTCAAGTAGCACCCCGAATACTGAAGAACCTCTTCCCCCGAAGGCTTCAAGTAGCACCCCGAATACTGAAGAACCTCTTCCCCCGAAGGCTTCAAGTAGCACCCCGAATACTGAAGAACCTCTTCCCCCGAAGGCTTCAAGTAGCACCCCGAATACTGAAGAACCTCTTCCCCCGAAGGCTTCAAGTAGCACCCCGAATACTGAAGAACCTTCCTT
EE5		CTACONOGOCIACANCACCOTONNOCTONNOCTONNOCTOCCCANOGOCCOCCCCCCCCONOCCCCCCCCCCCCCCCCC
EE6 EE7		CALAGRAGICA CALACCETRA ACCTRA ACCTRA ACCTRA ACCTRA CONTROL CON
EE8		TACONOGUECAN AND ACCORDANGE TO ANGERGA ACCORDAGO CONCECTOR CONTROL CONTROL CONTROL CONTROL CONTROL CONTROL CONTROL CONTROL CONTROL AND ACCORD CONTROL AND ACCORD CONTROL AND ACCORD CONTROL CO
EE9		TAGARGACCACAACACCOTGAAGCTGAAGCTGAAGCTGCCCTTCCCCTTCCCCTTCCCCTTCCCCCCAGTCCCCCCAGTTCCAGTACGACCCCCGACGCCCACATCCCCGACGACACCACCAAGAAGCATCCCCGACGACGCTTCAAGAAGCACCCCGACGACGCTTCAAGAAGCACCCCGACGACGCTCAAGAAGCACCCCGACGACGCTTCAAGAAGCACCCCGACGACGCTCAAGAAGCACCCCGACGACGCCTACAAGAAGCACCCGACGACGCTCAAGAAGCACCCGACGACGACCCCGACGACGCTCAAGAAGCACCCGACGACGACCCCGACGACGACCACCCCGACGA
EE10 EE11 *		CHACAGGGGCACAACACCATAAACTAAAGGTACAAAGGGCGGCCCTGCCCTGGGCACATCCCCCGGACAATCCCCCCCGATTCCCAAGGGTAACGGTCAAGGACCCGGGGGGAATCCCCGATTACAAAGAAGTGTCCTTTCCCGAAGGGTTAAAGAACGGTGATACCGAAAGAACTGTCCTTTCCCGAAGGGTTAAAGAACGTGATAAAGAACGGTGATACCGAAGAACGATACAAAGAACTGTCTTTCCCGAAGGGTTAAAGAACCGAAAGAACTGAAAGAACGATACAAAGAACTGTCTTCCCCGAAGGGTTAAAAGAACCCAAAGGTAAACAAAC
EE12 *		CTACONOGGICACANCACCOTONNOCTONACOTONCONAGOCCOCCCCOCCCCTCCCCTTCCCCTGGACACTCCACTC
EE13 EE14		CALCAGGGCCACAACCGTAAACCTGAAGCTAACCTAACCT
EE15 *		TACOMBUCIALA ACCUSTANCE TO MANUFACICA AND SUBJECT CONTROLLED TO SUBJECT TO SU
EE16		CTACOAGGGCCACAACACCGTGAAGGTGAAGGTGACCCAAGGGGGGCCCCCTGGCCTTCGCCTGGGACACTCCCAGTTCCAGTTCCAGTACGACTCCAAGTTCCAGTACGACTCCAAGTACCCCGACATCCCCGACACTACCAAGAAGCTGTCCATCCCCGAGGGCTTCAAGTGGAGCCCGTGAT
EE17 * EE18		CARGAGGGGCCA ARCCGTBAAGTBAAGTBAAGTBAGGACAGGGGCGATTCCCAGAGGACATCTGCCCCAGGGGGTAAT CARGAGGGGCAAAACGGGTAACGGGGGGGAATCCCCGAAGAAAACGGTCTTCCCCGAGGGGTAACGGGGGGGAATCCCCGAAGAAAACGGTCTCCCCCAAGGGACAACGGGGGGAATCCCCGAAGAAAACGGTCCCCCCCAAGGGGGGGG
EE19		TACGAGGGCCACAACACCGGAAGAGCGGAAGACGAAGGCGGC
EE20 *		CTACGAGGCCACAACACCGTGAAGGTGACCAAGGG <mark>CGGCCCCCCCCCGGGGCCTCCCCCTGGGACCCCCCCCAGGGCTTCCAGTACGGTGTACGTGAAGGACCCCGCCGACATCCCCGACTACAAGAAGCTGTCCCCGAGGGCTTCAAGTGGGAGCGCGTGAT</mark>
EE21 EE22		CHACAGOGGICACARACCETANACCTORAGGICACCACCACCACCACCACCACCACCACCACCACCACCAC
EE23	CTTCAAGGTGCGCATGGAGGGC <mark>ACCGTGAACG<mark>GCCA</mark>CGAGTTCGAGATCGAGGGCGAGGGCGAGGGCCGCC</mark>	CTACGAGGGCCACAACACCGTGAAGCTGAAGCTGACCAAGGGCGCCCCTCGCCCTTCGCCTTCGCCTGGACACTCCCCGAGTTCCAGTTCCAGTTCCAGTACAAGAAGCTGTACAAGAAGCTGACCCGACTACAAGAAGATGTCCCTTCCCCGAGGGCTTCAAGTGGGAOCGCGTGAT
EE24 * EE25		CALGRAGGE CALARCESTANACTANAGETA CALAGGE CONCECTE COCCEDE CONCENTRACE CONCENTRA
EE26		TACCAGGGCCAACAACCCGGAAACTGAAACTGAACTGAA
EE27		CTACOMOGGICACANCACCOTGANGCTGANGGIGANGGIGAGGGCCCCTGCCCCTCCCCTTCGCCTGGGACACCTCCCCGATTCCAGTTCCAGTGANGCACCCCGACGCCCCCGACACCACCAAGAAGAAGAAGAAG
EE28 * EE29		CTACOMO GOCO CA ACCOSTO ACCOSTO ACTO CALAGORISTIC CALAGORISTIC CONTROL CONTROL CONTROL CALAGORISTIC CALAGORIS
EE30	CTTCAAGGTGCGCATGGAGGGCAC <mark>CGTGAACGGCCAC</mark> GAGTTCGAGATCGAGGGCGAGGGCGAGGGCCGCC	CTACGAGGGCCACAACACCGTGAAGCTGAAGGTGACCAAGGGCGGCCCCTGCCCTTGCCCTGGCACTTCCCCTGTCCCCCAGTTCCAGTTCCAGTTCCAGTACGGCTGAAGCTGTAAGTGAAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGA
EE31	CTTCAAGGTGCGCATGGAGGGCACC <mark>GTGAACGGCCACG</mark> AGTTCGAGATCGAGGGCGAGGGCGAGGGCCGCC	CTACGAGGGCCACAACACCGTGAAGCTGAAGCTGACCAAGGGCGCCCCTCGCCCTTCGCCTTCGCCTGGACACTCCCCGATTCCAGTTCCAGTACAAGAAGCTGTACTGAAGCACCCCGCCGACTACAAGAAGCTGTCCTTCCCCGAGGGCTTCAAGTGGGAGCGCTGAT
EE32 EE33		TACASSEGUCACAACCETAAGCTAAGCTAAGCTAACCTACAAGAAGCTGTCCCTTCCCGGGGCATCCCCCATTCCCATTGCCATGGCCTGATCCCCATTCCCATTGCATGCGTGATCCCTGATGCATTCCCATTGCATGCGTGATCCCTGATGCATTCCCTGATGCATTCCCTGATGCATTCCCTGATGCATTCCCTGATGCATTCCCTGATGCATTCCCTGATGCATTCCCTGATGCATTCATTCCATTCCATTCCATTCCATTCATTCATTCCATTCCATTCCATTCATTCATTCCATTCCATTC
EE34		CTACGAGGGCACAACACCGTGAAGGTGACCAAGGGCGGCCCCTGCCCTGGGACATCCCGCCAGTTCCACTGGATCCGGTGTAACCCAACAACAACCCGCGCACATCCGCGAAGTGCCGCACATCCGCGAAGGCTCCAAGGGCTCCAAGTGGAACCCGGAAATCCGCGAAGTGCAAGAAGTGTCCTTCCCCGAAGGCTTCAATTGGAACCCGGTGAT
EE35 *	CTTCAAGGTGCGCATGGAGGGCACC <mark>GTGAACGGCC</mark> ACGAG <mark>T</mark> TCGAGATCGAGGGCGAGGGCGAGGGCCGCC	CTACGAG OCC. CAACA CCGTGAAGCTGAAGGTGACCAAGGGCGGCCCCCGCCCTCGCCTCGCCTGGGACATCCCCGCCAGTTCCAGTACGGCTCCAAGGTGTACGCTCCAAGGACACCCCCCCGACTACAAGAAGCTGTCCCCGAGGGCTTCAAGTGGGACGCGCGTGAT
EE36 * EE37		TAGARGGCA CARACGETA AGCTA AGGTA AGGTA ACTA AGGGGGGCCCCTGCCCTG
EE38 *		CTACGAGGGCCACAACACCGTGAAGGTGAAGGTGACCAAGGCGGCGCCCTGCCCTGCCCTGGCCTGGACCTGCCCCCCAATCCCCCAAGGTTCCAATCCCCCAAGGTTCCAAGGAAGCACCCCGCCGACATCCCCGACTACAAGAAGCTGTCCTTCCCCGAGGGCTTCAAGTGGAGCGCGTGAT
EE39		TAGARGEGICA ARCCESTA ACCOMAGNETA ACCOMAGNE
EE40 * EE41 *		CTACAAGGGCCACAAACCGTGAAGCTGAAGGTGACCAAGGGCCCCCTGCCTTGCCTGGACACTCCCCCCAGTCCAGTCCAGTACGGTCCAAGGTACGTAAGGAAGCCCCGCCGACTACCAAGAAACAACTGTCCTTCCCCCAAGGGTCCAATGGAAGCACCCGTCATACGAAGCACCCGGCCAACACCACGTACAAGAACTGTCCTTCCCCCAAGGACTCCAATGGAACCCGTCATACGTAAACAACTGTCCAAGAACTGTCGATCAGTAAGAACTGTCATACGAAGAACTGTCATTCCCCAAGGACCGTCATACGAAGAACTGTCATTCCCCAAGGACCGTCATACGAAGAACTGTCATTCCCCAAGGACCGTCATACGAAGAACTGTCATTCCCCAAGGACCGTCATACGAAGAACTGTCATTCCCCAAGGACCGTCATACGAAGAACTGCATACGAAGAACTGTCATACGAACTACGAAGAACTGTCATACGAAGAACTACAAGAACTACAAGAAACTGTCATACGAAAACTGAAACTACAAGAAACTGTCATACGAAAACTACAAACAA
EE42		CTACOAGGCCACAACACCOTGAAGGTGAAGGTGACCAAGGCGCCCCTGCCCTG
EE43 *		CTAL CAAGGGCOC CAAAAC COSTAAAGCTAAAGSTAACCASGGGGGCCCCTGCCCTTGCCCGGGACATCCGTGCCCCCCASTTCCCCAAGTAGCCCCCGACTACAAAAAAACTGTCCCTTCCCCGAGGGCTTAAATGGGAGGCGGTGAATGCCCGAAGAAACTGTCCCCCAATTACAAAAAAACTGTCCCTTCCCCGAGGGCTTAAATGGGAGGCGTGAAT
EE45 *		TACAMBUCCALARACCUTARACCTARACCTARACCTARACCTARACCTARACCTARCCTARACCTA
EE46 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCA <mark>CGAGTTCGAG</mark> ATC <mark>GAGGGCGAGGGCGAGGGCCGC</mark>	CTACGAGGGCCACAACACCGTGAAGGTGACGTAGAGGGGGGCCCCTGCCCTGCCCTGGGAGATCCCCCGAGTTCCAAGTACGGCTCCAAGTGTAAGCACCCCGCCGACATCCCCGACTACCAAGAAGAGCTGTCCTCCCCGAGGGCTTCAAGTGGGAGCGCGTGAT
EE47 * EE48		TACASSEGUCA ARCCETA ASCITA ASCITA ACTACA ASSISTANCE ASS
EE49		CTACGAGGGCACAACACCGTGAAGTGAAGTGACCAAGGGCGGCCCCTGCCCTTGCCTGGCAGGACATCCTGTCCCCCAGTTCAGTACGCTCAAGTGTAAGTAA
EE50		CTACGAGGGCCACAACACCGTGAAGCTGAAGGTGACCAAGGGCGGCCCCTGCCCTTCGCCTGGGACATCCCCCCAGTTCCAGTACGACTCCAAGTACCACCAGCGACATCCCCCAAGAAGCACCCCCCAAGGACACCACCACCACCACC
EE51 * EE52		CHARGESGERON ANGESTANDETS AND AGE ACCANGE GEOCETECCET GOOG ACCANGE AFFORDER CONTROL OF THE CONTR
EE53 *		TACOMOGOCIANA NECTOMA OCTOMA OCTOMA OGGO COCCCTOCCCTTCOCCTTOMA COLOCCCO COLOTACO AND A CALCCCCO COLOTACCA AND A A OCTOCCCO ACT A CAMBA A OCTOCCCO ACT A CAMBA A OCTOCCCO ACT A CAMBA A OCTOCCO ACT A CAMBA A
EE54 *		**************************************
EE56		CHACAGOGUCACARACCOTARACOTORAGOTACCARAGOGOGUCCCOTOCCCTOCCCTGGGCACATCCCCCAGTCCCCAGTCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCAGTCCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCAGTCCCCCCAGTCCCCCCAGTCCCCCAGTCCCCCCAGTCCCCCAGTCCCCCCAGTCCCCCCAGTCCCCCAGTCCCCCCAGTCCCCCCAGTCCCCCAGTCCCCCAGT
EE57 *		CTACOAGGGCCACAACACCGTGAAGGTGAAGGTGACCAAGGGGGCCCCTGCCCTTCCCCTGGGCACATCCCCCGACTTCCAGTTACGGGTCCAAGGTGTACGCCGCCGACATCCCCGACATCCCCAAGAAGCTGTCCTTCCCCCGACATCCCGACATCCCCGACATCCCCAAGAAGCACCCCGACATCCCAAGAAGCACCCCGACATCCCAAGAAGCACCCCGACATCCCAAGAAGCACCCCGACATCCCAAGAAGCAAGC
EE58 * EE59		CHARGESCENCE ALECCETAL ACCETAL AGGREGATIC ALL AGGREGATIC CALL
EE60		TA COASGGCANA NACOTTAN AGGTACCAN AGGGCGCCCTGCCCTTGCCTGGGCACATCCCCCAGTTCCAGTTTCAGTTCAGTTTCAGTTTCAGTTCAGTTCAGTTTCAGTTTCAGTTTCAGTTTCAGTTTCAGTTTCAGTTTCAGTTTCAGTTTCAGTTTTC
EE61 *		CTACGAGGGCCACAACACCGTGAAGGCTGACGAAGAGGCGGCCCCTGCCCTGCCCTGGGACATCCCAGTGCCCCAGTTCCAGTACGGGTGTACGTGAAGCACCCCCCCGACATCCCCGACAAGAAGAAGCTGTCCTTCCCCGAGGGCTTCAAGTGGGAGCCCGGATACA
EE62 EE63		CALAGRIGUE CA CALAC COSTA AGOSTA CONTRA AGOSTA CONTRA CONT
EE64 *		CTACOAGGCC REARCACCGTGAAGCTGAAGCTGACCAAGGCGGCCCCTCCCCT
EE65 *		CTACGAGGCCACAACACCGTGAAGGTGAACCAAGGACGGCGCCCCTCCCCTTCCCCTGGCACGTCCCAGTACCAGTACCAGTACCGGTGAAGCACCCCCCGACATCCCGACTACAAGAAGCTGTCCCCGAGGCCTTCAAGTAGGAGCCCGATACA
EE66 *		CALAGRAGICA CARACCETRA ACCTRA ACTRA ACTRA CARACTERICA CONTROL
EE68 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCG <mark>AGGGCC</mark> AGGGCCGCC	CTACGAGGGCCACAACACCGTGAAGCTGAAGCTGACCA <mark>AGGGCG</mark> CCCCCTCCCCTTCGCCTTCGCCTTCGCACTACCAGTTCCAGTTCCAGTTCCAGTACAAGGAGCCCCCAACTACAAGAAGCTGTCCTACAAGAAGCTGTCCTTCCCCGAAGGAGCGCTTCAAGTAGGAGCAGCCGTGAT
EE69 EE70		TACAMAGGCCACAACACCCTGAAGCTGAAGCTGAAGCTGAAGAGGGCCCCCTGCCCTGCCCTGCCCTGGGCAACACGTGTCCCCCCAGTTCCAGGTGCAAGGGCTTCAAGTGTAAGAGCCCGGAGTGAAGAAGCTGTCCTTCCCCGAGGGCTTCAAGTGGAAGCCCGTGAT
EE70 EE71 *		CALAGOGGICA CALAGOGGIA AGOTA AGOTA CALAGGGIGGICCOCTUDOCOTT COCCAGAGGIT CALAGOGGICA CALAGOG
EE72	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAG	CTACGAGGGCCACAACACCGTGAAGCTGAAGCTGACCAAGGGCGGCCCCTCGCCTTCGCCTGGGACACCCCCCCC
EE73 * EE74 *		CHACAGOGGICACAAACGETAAGCTAAGGTAAGCTAAGGGGGGCCCCTGCCCTACCCGACTACAGTGGGACATCCCGACTACAAGAAGCTGTCCCCGAGGGCTCAAGTGGGACGGGTGATCCGACTACCGACTACAAGAAGCTGTCCCTTCCCCGAGGGTTAAGTGGAGGGGGTGATCCCGAGGGCTCAAGTGGACGGGGTTAAGTGGAGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGTTAAGTGGAGGGGTTAAGTGGAGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGTTAAGTGGAGGGGGG
EE75 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCGAGGGCGAGGGCCAC	C TACGAGGGCACAACACCGTGAAGCTGAAGGTGACCAAGGGCGCGCCCCTGCCCTGCCCTGGCCTTGCCCTGGCACTCCCCGATTCCCAGTACTACTAAGAAGACCCCGCCGACATCCCCGACTACAAGAAGACCTCTCCCCGAGGGCTTCAAGTGGGAGCCCTGAT
EE76 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCGAGGGCGAGGGCCGCC	CTACGAGGGCCACAACACCGTGAAGGTGACCAAGGGCGCCCCTGCCCTTGCCCTGGGACATCCCGTTTCCAGTTCCAGTACGGCTCCAAGGTGTACGCGCGACATCCCCGACACACAAGAAGCTGTCCTTCCCCGAGGCTTCAAGTGGAAGCCGTGAT
EE77 *	CTTCANGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGGACCGAGGGCCACGCC	TARAGGICÀ CALA COGTÓ A LOCA TORA A GOGGICA COGGICA COCCUTA COGGICA TO COGGICA
EE79 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCGAGGGCGAGGGCCACG	TACGAGGGCAACAACCCGTAAAGTAACGTAAAGTACCAACAACAACGGGCCCCTGGCCTTCGCCTGGGACATCCCGGATCCAGTTCGAGTACGGCTCAAGTACGCCCCCCACATCCCAGAACAACACCCGCCCAACATCCCAGAACACACAC
EE80 *		CTACONOGUECACANCACCOGONNOCTONAGOTONCONAGOCOGOCCCCTCCCCTTCQCCTCONCOCTOCCCCCONCOCTOCCCCONCOCTOCCCCONCOCTOCCCCONCOCTOCCCCCONCOCTOCCCCCONCOCTOCCCCCONCOCTOCCCCCONCOCTOCCCCCONCOCTOCCCCCCCC
EE81 * EE82 *		TACAMAGGERIANA CONTINUED AND TANACTAN AGGING CONTINUED ACCAMAGGING CONTINUED ACCAMAGGING CONTINUED ACCAMAGGING CONTINUED ACCAMAGGING CONTINUED ACCAMAGA ACCORDANCE ACCAMAGA ACCORDANCE ACCAMAGA ACCORDANCE ACCAMAGA ACCORDANCE ACCAMAGA ACCORDANCE ACCAMAGA ACC
EE82 * EE83		TAGABBECACHARACGGANAGGTAAAGTAAAGTACAAAGGGGCCCCTTBCCCTGGCTGACTACCCGGATCAATTCCAGTTCCAGTACGGATCAGTTCCAGTTCCAGTGAAAGGACCCGATCAATTCCAGTTCCAGTGAAAGGACCCGATCAATTCCAGTTCCAGTGAAAGGACCGATCAATTCCAGTGAAAGCACCGATTCCAGTTCCAGTGAAAGCACCGAATTCCAGTTCCAGTGAAAGCACCGAATTCCAGTTCCAGTGAAAGCACCGAATTCCAGTAAAGCACCGAATTCCAGTGAAAGCACCGAATTCCAGTTCCAGTGAATTCCAGTGAATTCCAGTTCAGTTCAGTTCCAGTTCAGTTCCAGTTCAGTTCAGTTCCAGTTCAGTTCCAGTTCAGTTCAGTTCAGTTCAGTTCCAGTTCAGTTCCAGTTCA
EE84		CTACGAGGCCACAACACCOTGAAGGTGAAGGTGAAGGTGACCAAGGGCCCCCCCCCC
EE85 *		CHACAN SOCK ALACCOS ALACCOS ALACOS ALACOS ALACOS ALACOS ALACAN ALACOS ACCORDANCE ALACAN ACCORDANCE ALACAN ALACAN ACCORDANCE ALACAN ALACAN ACCORDANCE ALACAN ALACAN ACCORDANCE ACCORDANCE ALACAN ACCORDANCE ALACAN ACCORDANCE ACCOR
EE86 * EE87 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCG <mark>AGGGCGAGGGCCC</mark>	CTACGAGGGCCACAACACCGTGAAGGTGACCAAGGGCGACCCACCC
EE88 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCGAGGGCCGAGGGCC	CTACGAGGCCACAACACCGTGAAGGTGACCAAGGCCGCCCCCCCC
EE89 *		CHARAGEGERA ARRICOSTA ARGESTA ARGESTA ARGESTA ARGESTA ARGESTA CONTROL ARGESTA ARGESTA CONTROL ARGESTA
EE90 * EE91 *		TREBARGUECACALACEGRANGETRANGETRANGETRACCANGGGGGGCCCCTGCCCTTGCCTGGCTGGACATCCCCCAGTTCCATACGGCTCATCCCTGAACACGCTCATCCCCAACTACAACAACTCCCCAACTACAACAACTTCCCCCAGGGCTTCAACTGCAGCGCTGATCCCTGAACAACTGCCCTGACTACAACAACTGCAGCGCTGATCCCAACTGCCAGGGCTTCAACTGCAGCGCTGATCACAACAACTGCCAGGGCTTCAACTGCAGCGCTGATCACAACAACAACAACAACAACAACAACAACAACAACAA
EE92 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCGAGGGCGAGGGCCG <mark>CC</mark>	TREGRIGGE CANADA COTERA GOTERA GOTERA CONSEGUE CONCESSE CONTROLLES OF A CONCESSE OF A CONCESS
EE93 *		CHARLES CONTROL AND TRANSPORT
EE94 * EE95 *		CHACAGGGCACAAACGGTAAAGGTAAAGGTAACGTAAAGGGGGGCCCCTGCCCTGGCCTGGGCATCCCCCCATTCCCATACGGCTCATCCCGATACAAACAA
EE96 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCGAGGGCGAGGGCCG	TACGAGGGCCACAACACCGTGAAGCTGAAGCTGAACGAGGGCGCCCCTCGCCCTCGCCTTCGCCTGGGACACCCCCCCC
EE97 * EE98 *		CHACAGGGGCCACAAACGGGAAGGTAAGGTAAGGTAACGTAAGGGAGGCCCCTGCCCTGACCTACAGTAGGAGGACGTAAAGAAGTTACAGTCCAAGAAGAGTTAACGGGCACAAAAAAGTTACCTTCCCCAAGGACTAAAGAGGTTAACGGACGACAACAAGAAGTTACAAGAAGAGGTTAAAGTGAAGAGCGTTAACGGACGACTAAAAAAGTTACCTTCCCCCAAGGTTAAAGTGAAGACGCTGAA
EE98 * EE99 *		TAGABBECACHARACGTBANGGTBANGGTBACCANBBECBGCCCGCCTCGCCCTBGACCATCCCABGACTTCCACCCABTCCABT
EE100 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCGAGGGCCAGGGCCGCCC	TACOAGGGCCACAACACCGTGAAGGTGACCAAGGGCGCCCCTGCCCTTCGCCTTCGCCTGGGACACCCCCGATCCCCAGTTCCAGTACGGCTGTACGGACGCTGAAGCACCCCGCCGACTACAAGAAGCTGTCCTTCCCCCAAGGACGCTTCAAGTAGCACCGTGAT
EE101 *	CTTCAAGGTGCGCATGGAGGGCACCGTGAACGGCCACGAGTTCGAGATCGAGGGCGAGG	CTACOAGGGCCACAACACCCGTGAAGGTGACCAAGGGCGCCCCTGCCCTTCACCTGGGACATCCCGGTTCCAGTTCCAGTTCCAGTTACGGTGAAGCACCCCGCGACATCCCGACTACAAGAAGCTGTCCTTCAAGTAGCGTACCGGACGCTTCAAGTAGCGACGCTTCAAGTAGCGACGCTTCAAGTAGCGACGCTGAT