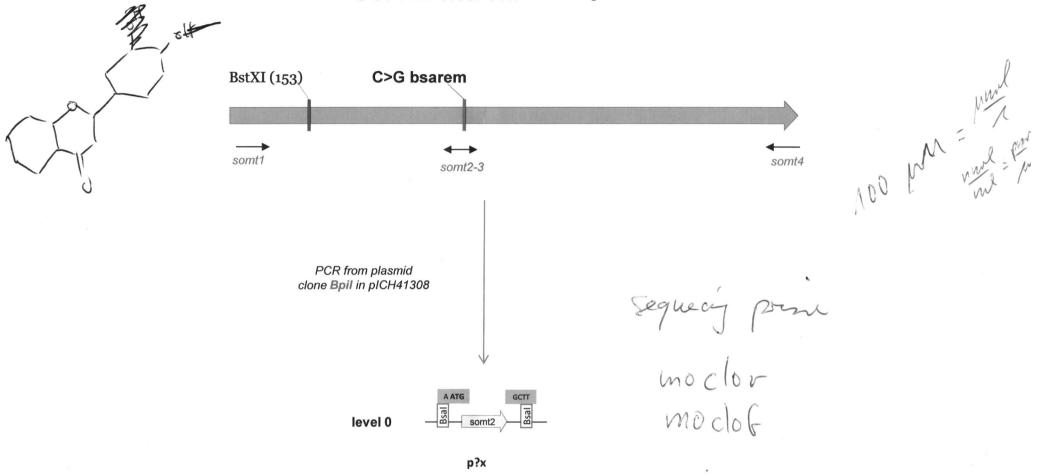
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+2 541 +2 601	ATTTO L N GAACT CTTG V G TGGTO ACCAO CATGO	GACCC CTGGG L TTGGC AACCG G CCACC V L GTGTT	T GCZAA CGTA F G TTTA A ACTT TGZ	ATAT FAGA ATCT G FGGA ACCT R ACGT	TAACA TTGT D GATT CTAA I ATCA TAGT P	ACCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CAAT CAAT A AGCA V FGTT	TTCAA AAGTT W TGGGT ACCCA K AAGAT TTCTA	TG AC	AGG TCC TTG AAC TCT AGA N	CAATG GTTAC E G AGGGA TCCCT C E GTGAG CACTC L S	GCT CGA CTG GAC A GCT CGA GGA	TGT ACA E GAA CTT F TTT AAA AGC	GATT CTAA S TCCA AGGT P CCTA GGAT N AACA	CT GA I TT AA K AG TC	CAG GTC V CAC CTG CTG	ATG TAC GAT CTA K AAA TTT T	TT AA V GT CA C TG AC TT TT
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+2 541 +2 601 +2 661	ATTTC L N GAAC CTTG V G TGGTC ACCA CATGC GTAC F V	GACCC CTGGG L / TTGGC AACCG G C GGTGG CCACC V L GTGTT CACAA G C	T GCAAA CGTAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	ATAT R R FAGA ATCT G FGGA ACCT R ACGT FGCA	CAACA TTGT D CGATT CTAA I ATCA TAGT P CCAA GGTT F	ACC CC CC CC CC CC T N AT AT K	CAAT A AGCA CGTT CGTT CCAAC C C	TTCAA AAGTT W TGGGT ACCCA K AAGAT TTCTA V GTGGA	TG AC CT GA AT TA AAA TT	AGG TCC F TTG AAC I TCT AGA N ATT TAA	CAATG GTTAC E G AGGGA TCCCT C E GTGAG CACTC L S TGTCA ACAGT	GCT CGA L CTG GAC A GCT CGA G G G G A	TGT ACA E GAA CTT TTT AAAA S AGC TCG	GATT CTAA S TCCA AGGT P CCTA GGAT N AACA TTGT	CTI GAA K AG TC N AT TA	CAG GTG CAC L GTG GAC L TTG AAC	ATG TAC GAT CTA K AAA TTT T ACA TGT	TT AA V CA CA TG AC TT AA V TT AA
+2 541 +2 601 +2 661	ATTTO L N GAACT CTTG TGGTO ACCA CATGO GTACO F V TGTTO	GACCC CTGGG L TTGGC AACCG GCACC V L GTGTT CACAA GCGGTGG	T GCAAA CGTA F G TTTA ACTA TGAAC G GAAC	ATAT R R FAGA ATCT G G G G G ACCT R ACGT G CATG	AACA TTGT D GATT CTAA ATCA TAGT P CCAA GGTT F	AC TC C C C C C C C C C C C C C C C C C	N CAAT A AGCA CGTT V CGTT ACAA	TTCAA AAGTT W TGGGT ACCCA K AAGAT TTCTA V GTGGA CACCT	TG AC CT GAA AT TA AA AT TT	AGG TCC F TTG AAC I TCT AGA N ATT TAA	CAATG GTTAC E G AGGGA TCCCT C E GTGAG CACTC L S TGTCA ACAGT A D	GCT CGA L CTG GAC A GCT CGA GCA GCA GGA CCT A	TGT ACA E GAA CTT TTT AAA S AGC TCG V	GATT CTAA S TCCA AGGT P CCTA GGAT N AACA TTGT L CTGC	CTI GAA K AG TC N AT TA	CAG GTC V CGTG CAC L CTG GAC L TTG AAC K	ATG TAC GAT CTA K AAA TTT ACA TGT L	TT AA V CA C TG TG AC TT TT AA V GT
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+2 541 +2 601 +2 661 +2 721 +2 781	ATTTO L N GAACT CTTG TGGTO ACCAO CATGO GTACO F V TGTTO ACAAO V L TTTAO AAATO	GACCC CTGGG L TTGGC AACCG GCACC V L GTGTT CACAA GCACC CATAA GTATT G L GGTGA	T GCA A CGT C AAA T TGA C CTT G G GA C CTT G	ATAT FATA R R FAGA ATCT G FGGA ACCT R ACGT FGCA M CATG GTAC N GAAT CTTA	AACA TTGT D GATT CTAA I ATCA TAGT P CCAA GGTT F TAAAT D CGACA CTGT T	A(C) T(C) G(C) T T T T T T T T T T T T T T T T T T T	CAAT CAACAA AGCAA AGCAA ACAAC C ATGCA ACAAC C C ATGCA ACAAC ACAAC C C C C C C C C C C C C	TTCAA AAGTT W TGGGT ACCCA K AAGAT TTCTA V GTGGA CACCT TAGGG C TGCAT ACGTA V GTAGT	TG AC CT GAA TA AT TT CA GT GA CT TT TT TT TT TT TT TT TT T	AGG TCC F TTG AAC I TCT AGA K ATT TAA K AGG TCC K AGA TCC V TCA	CAATG GTTAC E G AGGGA TCCCT C E GTGAG CACTC L S TGTCA ACAGT A D CTGAT GACTA TATTA ATAAT I D TAGAT	GCT CGA GCT CGA GCT A GCA CCT A GCA CCT T A CTT	TGT ACA E GAA CTT F TTTT AAA S AGC V GTT CAA N AAT TTA	GATT CTAA S TCCA AGGT P CCTA GGAT N AACA TTGT L CTGC GACG C TGTA ACAT I ATAA	CT GAA K AAA TT AAA AAA TT N AAA	CAGGGTC V CGTG CCTG CGAC L TTG AACC K CAACC CTTC CGAA CCTT E	ATG D GAT CTA K AAAA TTT T ACA TGT L TTG AAC A GCT N AAAC	CA CA CA CA CA CA CA CA CA CA CA CA CA C
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+2 541 +2 601 +2 661 +2 721 +2 781 +2 841	ATTTO L N GAACT CTTG V G TGGTO ACCAO CATGO GTACO F V TGTTO ACAAO V L TTTAO AAATO AAATO AAGTO	GACCC CTGGG L TTGGC AACCG GCACC V GGTGT CACAA GCACC H CATAA GTATT G GGTGA CCACCT E F	T GCA A CGT C AAA T TGA A ACT T TGA C CTT W T TGGA A ACC I C CTC I W	ATAT FATA R FAGA ATCT G FGGA ACCT R ACGT FGCA M CATG GTAC V CAAA GTTT	AACA TTGT D GATT CTAA TAGT P CCAA GGTT F GACA CTGT T ACAG TGTC T	A(C) T(C) T(C) T(C) T(C) T(C) T(C) T(C) T	CAAT CAAT CAAT CAAT CAAT CAAT CGTT CAACA CCATGCA CCATG	TTCAA AAGTT W TGGGT ACCCA K AAGAT TTCTA V GTGGA CACCT I TGCAT ACGTA V GTAGT CATCA K	TG AC CT TG AC CT TG AC CT	AGG TCC F TTG AAC I TCT AGA N ATT TAA K AGG TCC K AGG TCT V TCA AGT L	CAATG GTTAC E G AGGGA TCCCT C E GTGAG CACTC L S TGTCA ACAGT A D CTGAT GACTA I L TATTA ATAAT I D TAGAT ATCTA M D	GCT CGA GCT CGA GCT CGA CCT A GCA CCT T TGA	TGT ACA E GAAA CTT AAA S AGC V GTT CAA N AATT TTA V GTG CAC	GATT CTAA S TCCA AGGT P CCTA GGAT N AACA TTGT L CTGC GACG C TGTA ACAT I ATAA TATT	CTGAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	CAG GTC V GTG CAC L GCTG GAC L TTG AAC K AAC CTT E GAA CCTT C C	ATG D GAT CTA K AAAA TTT I ACA TTG A GCT N AAC TTG	TT AA V GT CA C TG AC F TT AA V GT TT AA TT TT TA TT TT TT TT TT TT TT TT
+2 541 +2 601 +2 661 +2 721 +2 781 +2 841	ATTTO L N GAACT CTTG TGGTO ACCA CATGO GTACO F V TGTTO ACAA V L TTTAO AAATO AAATO AAGTO AGATO AGATO	GACCC CTGGG L TTGGC AACCG GCACC V L GTGTT CACAA GCACC H N CATAA GTATT G L GGTGG CCACC H N CATAA GTATT G L GGTGG	T GCA A CGT C AAA T TGA A ACT T TGA C CTT O T TGA T TGA C CTT T TGC T TCC T TC	ATAT FATA R FAGA ATCT G FGGA ACCT R ACGT FGCA M CATG GTAC K CAAA GTTT V AGTT	AACA TTGT CTAA ATCA TAGT CCAA CGGTT F CGACA CTGT ACAG TGTC TACTG	A(C) T(C) T(C) T(C) T(C) T(C) T(C) T(C) T	CAAT CAAT CAAT CAAT CAAT CAAT CCAT CCAT	TTCAA AAGTT W TGGGT ACCCA K AAGAT TTCTA V GTGGA CACCT TAGGG C TGCAT ACGTA V GTAGTA K AAGCT	TG AC CT TA AT TT TG GA CT TG AC CT TG AC CT	AGG TCC F TTG AAC I TCT AGA N ATT TAA K AGG TCC V TCA AGA TCT L	CAATG GTTAC E G AGGGA TCCCT C E GTGAG CACTC L S TGTCA ACAGT A D CTGAT GACTA I L TATTA ATAAT I D TAGAT ATCTA	GCT CGA GAC GGA GCT A GCA CCT A GCA CGT T GAA CTT TGA	TGT ACA E GAAA CTT TTT AAA S AGC V GTT CAAA V GTG CAC H CCAC	GATT CTAA S TCCA AGGT P CCTA GGAT N AACA TTGT L CTGC GACG C TGTA ACAT I ATAA TATT M ATGG	CTGAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	CAG GTC V GTG CAC L CTG CAC L CTTG AAG CTTC E CGAA CTTT C C TTGT	ATG TAC GAT T AAAA TTT TACA A GCT CGA N AAC TTG AAC TTG AAC TTG AAC AAC	TT AA V GT CA C TG AC F TT TA CA TT

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> A ~ 6.56 mg B ~ 6.66 mg C ~ 7.08 mg

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