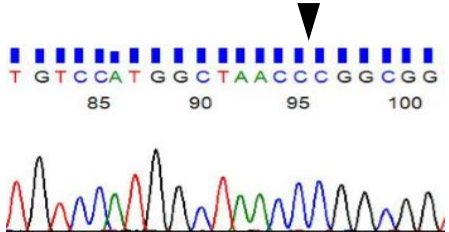
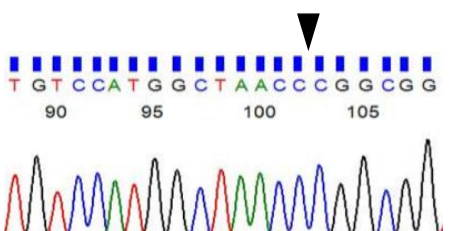
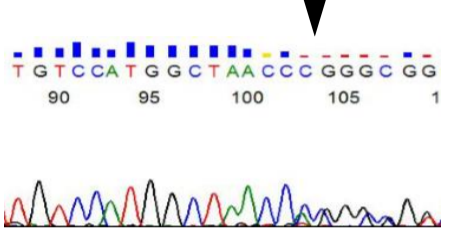
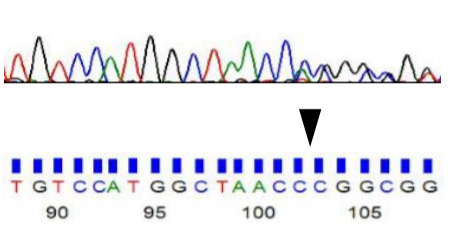
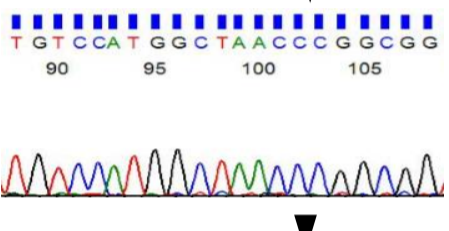
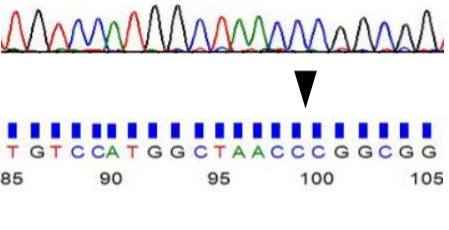
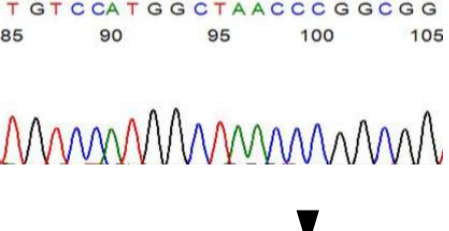
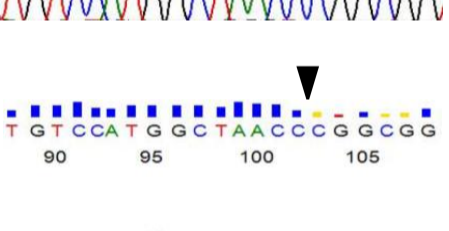
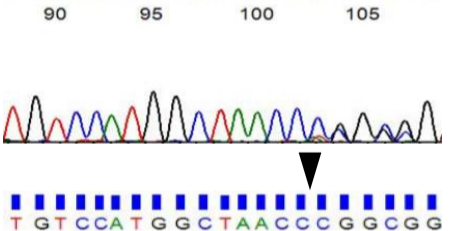
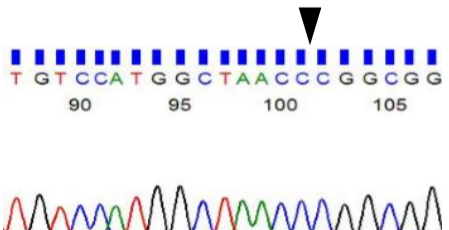
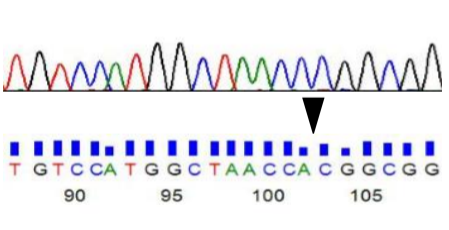


Kitaake		WT: TGTCCATGGCTAACC-CGGCGG WT: TGTCCATGGCTAACC-CGGCGG	Wild-type
OsU6a <i>xa13m</i> /Kit T ₂	1A-5-5 	WT: TGTCCATGGCTAACC-CGGCGG WT: TGTCCATGGCTAACC-CGGCGG	Wild-type
	1A-5-6 	WT: TGTCCATGGCTAACC-CGGCGG M2: TGTCCATGGCTAACCTCGGCGG	Heterozygote
	1A-8-9 	WT: TGTCCATGGCTAACC-CGGCGG WT: TGTCCATGGCTAACC-CGGCGG	Wild-type
	1A-16-2 	WT: TGTCCATGGCTAACC-CGGCGG WT: TGTCCATGGCTAACC-CGGCGG	Wild-type
	3A-6-1 	WT: TGTCCATGGCTAACC-CGGCGG M2: TGTCCATGGCTAACCTCGGCGG M1: TGTCCATGGCTAACCAACGGCGG	Chimera
	3A-6-3 	WT: TGTCCATGGCTAACC-CGGCGG WT: TGTCCATGGCTAACC-CGGCGG	Wild-type
	4A-1-6 	M1: TGTCCATGGCTAACCAACGGCGG M1: TGTCCATGGCTAACCAACGGCGG	Homozygote
	4A-1-7 	M1: TGTCCATGGCTAACCAACGGCGG M1: TGTCCATGGCTAACCAACGGCGG	Homozygote
	4A-3-3 	M1: TGTCCATGGCTAACCAACGGCGG M1: TGTCCATGGCTAACCAACGGCGG	Homozygote
	4A-3-5 	M1: TGTCCATGGCTAACCAACGGCGG M1: TGTCCATGGCTAACCAACGGCGG	Homozygote