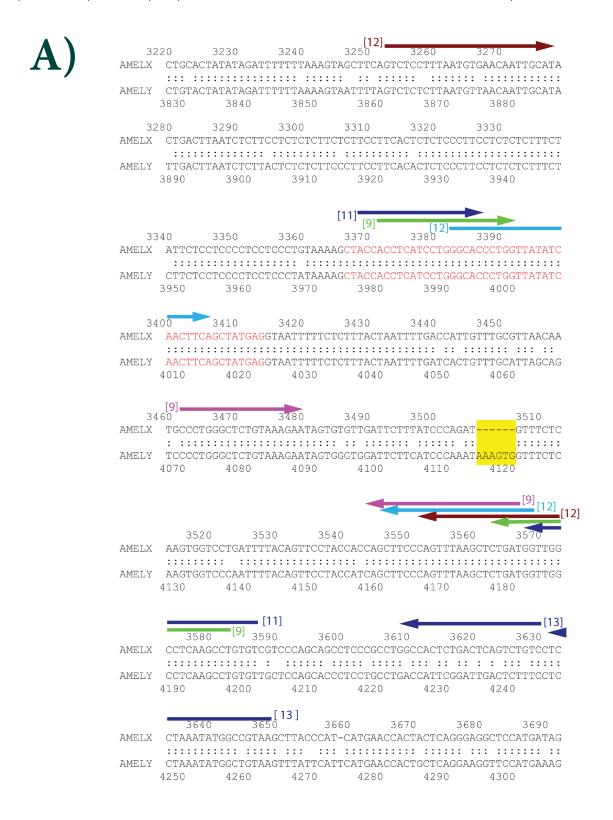
Figure 3: The alignment of AMELX and AMELY genomic sequences mapped with the primers (arrowed lines) used for sex identification. a) intron 3, 6-bp length polymorphism, b) intron 3, 180-bp length polymorphism, c) intron 3, 187-bp length polymorphism, d) exon 5, 3-bp length polymorphism, and e) intron 6, 1-bp length polymorphism. Nucleotide positions are based on AMELX sequence (Ensembl identification number ENSG00000099721) and AMELY (Ensembl identification number ENSG00000125363). "." represents homologous regions; "-" represents length polymorphisms; yellow boxes indicate the polymorphism in question. Bases in red represent exon sequences. Each pair of primers is labeled with its reference in brackets. Additional information about the primers can be found in Table 1.



		[6]			
		360 3			390
AMELX ATTCTCCT					CCTGGTTATATC
AMELY CTTCTCCT 3950	CCCCTCCTCC 3960		TACCACCTCA 3980	TCCTGGGCAC 3990	CCTGGTTATATC 4000
3400 3	410 3	420 3	3430 3	3440 3	3450
AMELX AACTTCAG					TTGCGTTAACAA
AMELY AACTTCAG 4010	CTATGAGGTA 4020		TTTACTAATTI 4040		TTGCATTAGCAG 4060
AMELX TGCCCTGG	GCTCTGTAAA	GAATAGTGTC	GTTGATTCTT		3510 GTTTCTC :::::::
	GCTCTGTAAA		GTGGATTCTT	catcccaaat <mark>f</mark>	AAGTGGTTTCTC 4120
3520	3530				3570
					TCTGATGGTTGG
AMELY AAGTGGTC 4130			ATCAGCTTCO 4160		TCTGATGGTTGG 4180
3580 AMELX CCTCAAGC	3590 CTGTGTCGTC				3630 CAGTCTGTCCTC
	:::::: :				: ::: :::::
		4210			GACTCTTTCCTC 4240
3640 AMELX CTAAATAT	3650 GGCCGTAAGC) 3670 CATGAACCACI) 3690 GCTCCATGATAG
					GTTCCATGAAAG
		4270	4280		4300
3700 AMELX GGCAAAAA					3740 AAAATGTAAGGA
					'AAAATGTAAAGA
				00001100001	
		4330	4340	4350	4360
		3770	3780	3790	4360 3800
AMELX TTAGGTAA	GAT <mark>GTTATTT</mark> ::	3770	3780 ICCAGCTCAAA	3790 AAAACTCCTGA	4360
AMELX TTAGGTAA	GAT <mark>GTTATTT</mark> ::	3770	3780 ICCAGCTCAAA	3790 AAAACTCCTGA	4360 3800
AMELX TTAGGTAA :::::::: AMELY TTAGGTAA 4370 3810	GAT <mark>GTTATTT</mark> :: AAT 3820	3770 AAAACTCTTT 3830	3780 TCCAGCTCAAA 3840	3790 AAAACTCCTG7 	4360 3800 TTCTAAGATAGT
AMELX TTAGGTAA :::::::: AMELY TTAGGTAA 4370 3810 AMELX CACACTCT	GAT <mark>GTTATTT</mark> :: AAT 3820 ATGTGTGTGTCT	3770 AAAACTCTTT 3830 CTTGCTTGCC	3780 TCCAGCTCAAA 3840 TCTGCTGAAA	3790 AAAACTCCTG7 3850 ATATTAGTGAC	4360 3800 TTCTAAGATAGT 3860 TAAGTGGTATAG
AMELX TTAGGTAA :::::::: AMELY TTAGGTAA 4370 3810 AMELX CACACTCT AMELY	GAT <mark>GTTATTT</mark> :: AAT 3820 ATGTGTGTCT	3770 AAAACTCTTT 3830 CTTGCTTGCC	3780 TCCAGCTCAAA 3840 TCTGCTGAAA	3790 AAAACTCCTG7 3850 ATATTAGTGAC	4360 3800 TTCTAAGATAGT
AMELX TTAGGTAA AMELY TTAGGTAA 4370 3810 AMELX CACACTCT AMELY 3870	GATGTTATTT :: AAT 3820 ATGTGTGTGTCT 	3770 AAAACTCTTT 3830 CTTGCTTGCC 3890	3780 ICCAGCTCAAP 3840 ITCTGCTGAAP 3900	3790 AAAACTCCTGA 3850 YTATTAGTGAC 3910	4360 3800 TTCTAAGATAGT 3860 TTAAGTGGTATAG
AMELX TTAGGTAA ::::::::: AMELY TTAGGTAA 4370 3810 AMELX CACACTCT AMELX GAGAGACT AMELX GAGAGACT AMELX GAGAGACT 3930	GATGTTATTT :: AAT 3820 ATGTGTGTGTGT 3880 CCGCAGAACA 3940	3770 AAAACTCTTT 3830 CTTGCTTGCC 3890 GCGGAATGCZ 3950	3780 CCAGCTCAA 3840 CTCTGCTGAA 3900 IIGAGTTTTGC 3960	3790 AAACTCCTGA 3850 ATATTAGTGAC 3910 SACGTCGGGTT 3970	4360 3800 TTCTAAGATAGT 3860 TTAAGTGGTATAG 3920 TGAGGTTCTCCT 3980
AMELX TTAGGTAA 	GATGTTATTT :: AAT 3820 ATGTGTGTGTGT 3880 CCGCAGAACA 3940	3770 AAAACTCTT1 3830 CTTGCTTGCC 3890 GCGGAATGC7 3950 GTGATTTTG	3780 TCCAGCTCAAP 3840 TCTGCTGAAP 3900 ATGAGTTTTGC 3960 GGCAAATCATT	3790 AAACTCCTGA 3850 TATTAGTGAC 3910 3ACGTCGGGTT 3970 TTCCTCTTTC1	4360 3800 XTTCTAAGATAGT 3860 TAAGTGGTATAG 3920 TGAGGTTCTCCT
AMELX TTAGGTAA AMELY TTAGGTAA 4370 3810 AMELX CACACTCT AMELY GAGAGACT AMELX GAGAGACT AMELX CAACCTCT	GATGTTATTT :: AAT 3820 ATGTGTGTGTGT 3880 CCGCAGAACA 3940 TACTAACTTT ::::::::::	3770 AAAACTCTTT 3830 CTTGCTTGCC 3890 GCGGAATGCZ 3950 GTGATTTTGC 51	3780 CCAGCTCAA 3840 CTCTGCTGAA 3900 TGAGTTTTGC 3960 GCAAATCATT SCAATAATT	3790 AAACTCCTGP 3850 ATATTAGTGAC 3910 ACGTCGGGTI 3970 TTCCTCTTTTCT TTCCTCTTTT	4360 3800 XTTCTAAGATAGT 3860 TTAAGTGGTATAG 3920 TGAGGTTCTCCT 3980 GGAACCCTGGTT
AMELX TTAGGTAA AMELY TTAGGTAA 4370 3810 AMELY CACACTCT AMELY SAGAGACT AMELY SAGAGACT AMELY SAGAGACT AMELY CAACCTCT AMELY 43 3990	GATGTTATTT :: 3820 ATGTGTGTGTCT 3880 CCGCAGAACA 3940 TACTAACTTT 7ACTAACTTT 80 4000	3770 AAAACTCTTT 3830 CTTGCTTGCC 3890 GCGGAATGCZ 3950 GTGATTTTG G	3780 CCAGCTCAAF 3840 CTCTGCTGAAF 3900 ATGAGTTTTGC 3960 GCAAATCATT) 4400 4020	3790 AAACTCCTGP 3850 3910 3970 TCCCTCGGGTT 5000 3977 TCCCTCTTTTC1 4410 4030	4360 3800 XTTCTAAGATAGT 3860 TAAGTGGTATAG 3920 TGAGGTTCTCCT 3980 GGAACCCTGGTT YGGAACCCTGGTT 4420 4040
AMELX TTAGGTAA AMELY TTAGGTAA 4370 3810 AMELX CACACTCT AMELY 33870 AMELX GAGAGACT AMELY 3930 AMELX CAACCTCT AMELY 43 AMELY 43 AMELY 43	GAT GTTATTT :: 3820 ATGTGTGTGTCT 3880 CCGCAGAACA CCGCAGAACA 3940 TACTAACTTT :::::::::: 4000 TGGAGAAAGGG ::::::::::::::::	3770 AAAACTCTTT 3830 CTTGCTTGCC 3890 GCGGAATGC/ 3950 GTGATTTGC GTGATTTGC 4390 4010 AAAATAATTAT	3780 CCAGCTCAAF 3840 TCTGGCTGAAF 3900 ATGAGTTTTGC 3960 3960 3960 3960 3960 3070 3960 4020 TAATAACCATF	3790 AAACTCCTGP 3850 ATATTAGTGAC 3910 3970 TCCCTCTTTCC 100 4030 ATTTCAAAATP	4360 3800 XTTCTAAGATAGT 3860 TAAGTGGTATAG 3920 TGAGGTTCTCCT 3980 GGAACCCTGGTT
AMELX TTAGGTAA AMELY TTAGGTAA 4370 3810 AMELX CACACTCT AMELY 33870 AMELX GAGAGACT AMELY 3930 AMELX CAACCTCT AMELY 43 AMELY 43 AMELY 43	GATGTTATTT :: 3820 ATGTGTGTGTCT 3880 CCGCAGAACA 3940 TACTAACTTT TACTAACTTT 80 4000 TGGACAAGGG	3770 AAAACTCTTT 3830 CTTGCTTGCC 3890 GCGGAATGCA 3950 GTGATTTGC 4390 4010 AAATTAATTAT AAATTACTGT	3780 CCAGCTCAAF 3840 TCTGCTGAAF 3900 TGGAGTTTTGC 3960 3960 3960 3960 3960 3000 TGGAATATCATT 0 4400 4020 PAATAACCATT PAATAACCATT 2.221 2.2111 2.2	3790 AAACTCCTGF 3850 3910 3970 TCCTCGGGTT 3977 TTCCTCTTTTC) 4030 TTTCCAAAATF TTTCCAAAATF	4360 3800 XTTCTAAGATAGT 3860 TAAGTGGTATAG 3920 TGAGGTTCTCCT 3980 GGAACCCTGGTT 3980 GGAACCCTGGTT 4420 4040 XTTCTTTGGAAGAG 2
AMELX TTAGGTAA AMELY TTAGGTAA 4370 3810 AMELY CACACTCT AMELY GACACTCT AMELY GACAGACT AMELY GACAGACT AMELY CAACCTCT AMELY 43 3990 AMELX TCCTCATC 43 3990 AMELX TCCTCATC 4430 4050	GATGTTATTT :: AAT 3820 ATGTGTGTGTCT 3880 CCGCAGAACA 3940 TACTAACTTT TACTAACTTT 80 4000 TGGAGAAAGGG 4440 4060	3770 AAAACTCTTT 3830 CTTGCTTGCC 3890 GCGGAATGCA 3950 GTGATTTGC GTGATTTGC 4010 AAATAATTATTGT 4450 4070	3780 CCAGCTCAAF 3840 TCTGCTGAAF 3900 ATGAGTTTTGC 3960 GCAAATAATTA) 440(4020 CAATAACCATF) 440(4020 TAATAACCATF TAATAACCATF CAATATCACF) 446(4080	3790 AAACTCCTGF 3850 ATATTAGTGAC 3910 SACGTCGGGTT 3970 TTCCTCTTTTCT 4030 ATTTCAAAATF 1 441C 4030 ATTTCAAAATF 4470 4090	4360 3800 XTTCTAAGATAGT 3860 TAAGTGGTATAG 3920 TGAGGTCTCCCT 3980 GGAACCCTGGTT
AMELX TTAGGTAA AMELY TTAGGTAA 4370 3810 AMELX CACACTCT AMELY 3870 AMELX GACACTCT AMELY 3930 AMELX CAACCTCT AMELY 43 3990 AMELX TCCTCATC AMELY TTCCATAT AMELY TTCCATT 4430 AMELX TAATATAG 	GAT GTTATTT :: AAT 3820 ATCTGTGTGTCT 3880 CCCCAGAACA 3940 TACTAACTTT :::::::::: TGGACAAGGG ::::::::::: TGGACAAGGG 4000 TGGACAAGGG :::::::::::::::::::::::::::::::::	3770 3830 CTTGCTTGCT 3890 GCGGAATGCT 3950 GGGGATTTGC 4010 4010 4010 4010 44450 4070 TGAAAAGTGC 5	3780 CCAGCTCAAF 3840 TCTGCTGAAF 3900 ATGAGTTTGC 3960 3960 3960 3960 3960 3960 300 4000 4020 4000 4	3790 AAACTCCTGP 3850 ATATTAGTGAC 3910 3070 TTCCTCTGTTTC 4030 4030 ATTTCAAAATP 0 4470 4090 FATAATATGAC 500 100 100 100 100 100 100 100	4360 3800 STTCTAAGATAGT 3860 TAAGTGGTATAG 3920 TGAGGTCTCTCCT 3980 GGAACCCTGGTT SISSING GGAACCCTGGTT 4040 SISSING 4040 SISSING 4040 SISSING 4040 SISSING 4040 SISSING 4040 SISSING 4040 SISSING SIS
AMELX TTAGGTAA AMELY TTAGGTAA 4370 3810 AMELY CACACTCT AMELY GAGAGACT AMELY GAGAGACT AMELY GAGAGACT AMELY GAGAGACT AMELY 43 3990 AMELX TCCTCATC 3990 AMELY TTCTCATT 4430 4050 AMELX TAATATAG	GAT GTTATTT :: AAT 3820 ATGTGTGTGTCT 3880 CCCCAGAAACA 3940 TACTAACTTT ::::::::: TGGACAAGGG 4000 TGGAGAAAGG 4400 TTAACAATTA TTAACAATTA	3770 3830 CTTGCTTGCT 3890 GCGGAATGCT 3950 GGGGATTTGC 4010 4010 4010 4010 44450 4070 TGAAAAGTGC 5	3780 CCAGCTCAAF 3840 TCTGCTGAAF 3900 ATGAGTTTGC 3960 3960 3960 3960 3960 3960 300 4000 4020 4000 4	3790 AAACTCCTGP 3850 ATATTAGTGAC 3910 3070 TTCCTCTGTTTC 4030 4030 ATTTCAAAATP 0 4470 4090 FATAATATGAC 500 100 100 100 100 100 100 100	4360 3800 STTCTAAGATAGT 3860 TAAGTGGTATAG 3920 TGAGGTCTCTCCT 3980 GGAACCCTGGTT SISSING GGAACCCTGGTT 4040 SISSING 4040 SISSING 4040 SISSING 4040 SISSING 4040 SISSING 4040 SISSING 4040 SISSING SIS
AMELX TTAGGTAA AMELY TTAGGTAA 4370 3810 AMELY CACACTCT AMELY GACACTCT AMELY GACAGACT AMELY GACAGACT AMELY GACACTCT AMELY 430 AMELX TCCTCATT AMELY TTCTCATT AMELY TAATATAG 4480 4110	GAT GTTATTT :: AAT 3820 ATGTGTGTGTCT 3880 CCCCAGAACA 3940 TACTAACTTT ::::::::: TACTAACTTT 80 4000 TGGACAAGGG 4440 4060 TTAATGAATA :::::::::: TTAACAATTA 4490 4120	3770 AAAACTCTTT 3830 CTTGCTTGCC 3890 GCGGAATGC/ 3950 GGGAATTGC/ 4390 4010 AAATAATTAT ::::::::::::::::::::::::::	3780 CCAGCTCAAF 3840 TCTGCTGAAF 3900 ATGAGTTTTGC 3960 3960 3960 3960 3960 3960 3960 307 3960 307 3960 307 3960 307 307 400 400 400 CTTTGTCAAGT 4080 CTTTGTCAAGT 4510 4140	3790 AAACTCCTGP 3850 ATATTAGTGAC 3910 3970 TTCCTCTCTTTTC 4030 ATTTCCAAAATP 4030 ATTTCCAAAATP 4470 4090 TATAATATGAC 4090 TATAATATGAC 4090 TATAATATGAC 4090 TATAATATGAC 4520 4150	4360 3800 XTTCTAAGATAGT 3860 TAAGTGGTATAG 3920 TGAGGTCTCCCT 3980 CGGAACCCTGGTT CGGAACCCTGGTT 4420 4040 XTTGTTTGGAGAG ::::::::::::::::::::::::::::::::

 AMELY
 ----TTTTTTTTTTTTTGATTACATGCTGTATTACCATATAAAGAATCCCCCAAACCTAAGGTT

 4540
 4550
 4560
 4570
 4580
 4590

AMELY AACTAAG 4600 4230 AMELX TAACAAT :::::: AMELY TAACAAT 460 4290 AMELX GGCAAGA	CTGTGATACTG 4610 4240 CTATGAGTTTA CTGTGAGTTTA 4670 4300 CTTTCTGTTGA CTTTCTGTTGA	TTCAGAAAGC 4620 4250 TTTACTTATA TTACTTACA 4680 4310 ACCGAAAAGA	GAATAAAATTO 4630 4260 AAAAGCTGAAG : :::::: ACAAGCTGCTG 4690 4320	CTTACTTCTC 4640 4270 GACAAATGTT. GACAAATGTT. 4700	TCATAGGTTCAGG 4650 4280 AGTAAGATTTTGA
4230 AMELX TAACAAT(:::::: AMELY TAACAAT(4660 4290 AMELX GGCAAGG :::::: AMELY GGCAAGG	4240 CTATGAGTTTA CTGTGAGTTTA 4670 4300 TTTTCTGTTGA TTTTCTGTTAA	4250 TTTACTTAT ::::::::::::::::::::::::::::::::::	4260 AAAAGCTGAAG : ::::: ACAAGCTGCTG 4690 4320	4270 GACAAATGTT GACAAATGTT GACAAATGTT 4700	4280 AGTAAGATTTTGA : ::::: : ::: AATAAGAATCTGA
AMELY TAACAAT 4660 4290 AMELX GGCAAGA :::::: AMELY GGCAAGG	CTGTGAGTTTA 4670 4300 ITTTCTGTTGAJ IIIIICTGTTAAJ	ITTACTTACA 4680 4310 ACCGAAAAGA	ACAAGCTGCT(4690 4320	GACAAATGTT 4700	AATAAGAATCTGA
AMELX GGCAAGA :::::: AMELY GGCAAGG	FTTTCTGTTGA FTTTCTGTTAA	ACCGAAAAGA			
AMELY GGCAAGG	ITTTCTGTTAA				
		ACCTAAAAGA 4740			
4350 AMELX TAGGATGA ::::: AMELY TAGGTTGA 4780			: :::: :::		::::: ::: : <mark></mark> :
		:::::	: :::::::	IGTGTTTGTT : :::::::	TTGCTCTTACAAT
4470 AMELX GAAAAGAA					4520 TCAGTCAAGTTAA
			GATTTACTCC	CTTTGTAACT	TCAGTCAAGTTAA 4950
			ATCTAAAAAG	IGAGAGTAAT.	4580 AATACTTGCCTCC
					AATACTTGCCTCC 5010
					4640 GTGCCTGTCACAC
					GTGCCTGACACAC 5070
			FCTTCTTTCT	rttgtag <mark>aac</mark> '	4700 TCACATTCTCAGG
					TCACATTCTCAGG 5130
					4760 CATCAGTGAGTTT
					CATCAGTGAATTT 5190
:: :: AMELY CTGCATG	GGATGAAAGTA	AATTAAATC	AATGGATTC	INTATATATCT	4820 TTCTCTTAAGGTG TTCTCTTAAGGTG
5200	5210	5220	5230	5240	5250
:: ::::		4850 IACCAGAGCA			4880 TAGACATTTTGTT :::::::::: TAGACATTTTGTT 5310

C)

D)

AMELX	4770 CTATATTGO	4780 Gatgaaagta <i>f</i>	4790 АТТАААТСА	4800 AATGGGTTCTAJ	4810 ATATCTTTTT	4820 CTCTTAAG <mark>GTG</mark>
AMELY	CTGCATGGG 5200	GATGAAAGTAA 5210	ATTAAATCA 5220	AATGGATTCTA 5230	ATATATCTTT 5240	CTCTTAAG <mark>GTG</mark> 5250
	[14] [15]					
	4830	4840	4850	4860	4870	4880
AMELX	CTTACCCCI	TTGAAGTGGI	ACCAGAGCA	T <mark></mark> AAGGCCA	CCGGTATGTA	GACATTTTGTT
AMELY	CTCACCCCT 5260	TTGAAGTGGT 5270	ACCAGAGCA 5280	T <mark>GAT</mark> AAGACCA 5290	CCAGTATGTA 5300	GACATTTTGTT 5310
Г	15]					
L	4890	4900	4910	[14] 4920	4930	4940
AMELX	CCTTATTCC	ССТБААААТАТ	TAGGCATGC	ATTAAAATTCC	CATATTAAGT	GAAATATCATG
AMELY	: :::::: CTTTATTCC 5320	CCTGAAAATAT 5330	TAGGCATGC. 5340	ATTTAAATTCC 5350	::: :::: CATTTTAAGA 5360	::::: :::: -AAATACCATG 5370

E)

