

Going Beyond the U.S. Haplotype: A Look at Additional Y-STR and Y-SNP Loci in U.S. Populations

John Butler

AAFS Workshop:
Y-STR Analysis on Forensic Casework

February 17, 2004

Commercial Y-STR Kits Available

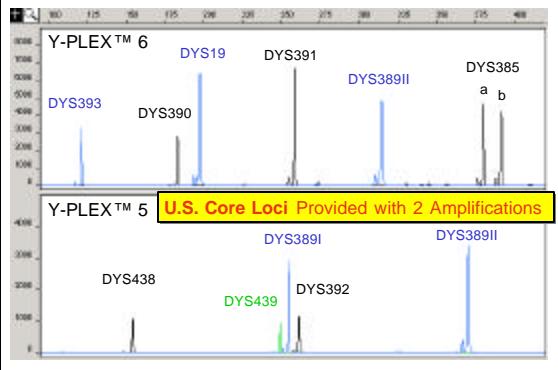
- ReliaGene Technologies (New Orleans, LA)
 - Y-PLEX™ 10: DYS19, DYS389II, DYS390, DYS391, DYS393, DYS385 a/b
 - Y-PLEX™ 5: DYS389II/I, DYS392, DYS438, DYS439
 - Y-PLEX™ 12: DYS19, DYS385 a/b, DYS389II/I, DYS390, DYS391, DYS392, DYS393, DYS438, DYS439, amelogenin
 - Promega Corporation (Madison, WI)
 - PowerPlex® Y: DYS19, DYS385 a/b, DYS389II/I, DYS390, DYS391, DYS392, DYS393, DYS438, DYS439, DYS437
 - Serac (Germany)
 - genRES® DYSplex-1: DYS389II/I, DYS390, DYS391, DYS385 a/b, amelogenin
 - genRES® DYSplice-2: DYS19, DYS389II/I, DYS392, DYS393
 - GKT Inc. (South Korea); *silver-stain kits*
 - GeneKin® Y-STR Systems I: DYS388, DYS19, DYS392
 - GeneKin® Y-STR Systems II: DYS393, DYS390, DYS391, DXY391X
 - GeneKin® Y-STR Systems III: DXYS156X, DXYS156Y, DYS389II/I
 - GeneKin® Y-STR Systems IV: DXYS156X, DXYS156Y, DYS385 a/b

History of Y STR Marker Discovery

Year	Marker	Discoverer	Marker Type
1992	DYS19 ^a	Roewer et al.	Extended Haplotype
1994	YCA1 ab, YCAII ab, YCAIII a/b		Extended Haplotype
1996	DYS389I/II DYS390		Extended Haplotype
1996	DYS391		Extended Haplotype
1996	DYS392		Extended Haplotype
1996	DYS393		Extended Haplotype
1996	DYF371, DYS425, DYS426	Jobling et al.	
1997	DYS288, DYS388	Kayser et al.	
1998	DYS385 a/b ^a	Schneider et al.	Minimal Haplotype
1999	A7.1 (DYS460), A7.2 (DYS461), A10, C4, H4	White et al.	
2000	DYS434, DYS435, DYS436, DYS437, DYS438, DYS439	Ayub et al.)	
2000	G09411 (DYS462), G10123 (de Knijff unpublished)		
2001	DYS441, DYS442 (Iida et al.)		
2002	DYS443, DYS444, DYS445 (Iida et al.); DYS446, DYS447, DYS448, DYS449, DYS450, DYS452, DYS453, DYS454, DYS455, DYS456, DYS458, DYS459 a/b, DYS463, DYS464 a/b/c/d (Redd et al.)		U.S. Haplotype
2002	DYS468-DYS596 (129 new Y STRs; Manfred Kayser GDB entries)		
2003	DYS597-DYS645 (50 new Y STRs; Manfred Kayser GDB entries)		

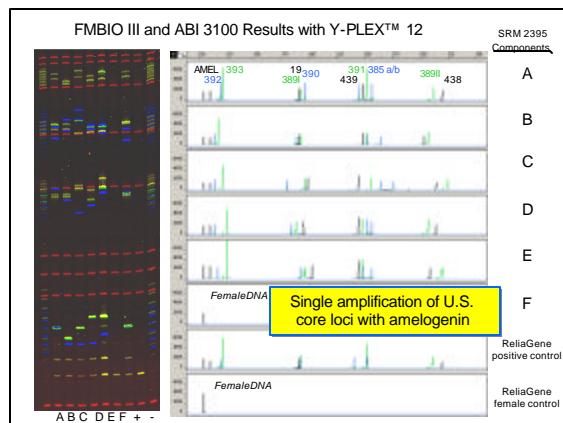
From J.M. Butler (2003) Recent developments in Y-STR and Y-SNP analysis. *Forensic Sci Rev.* 15:91-111

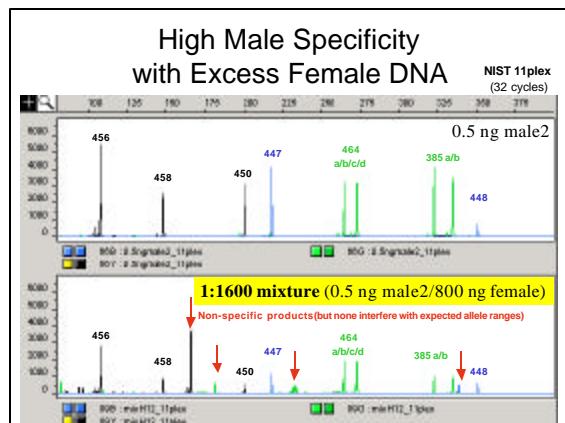
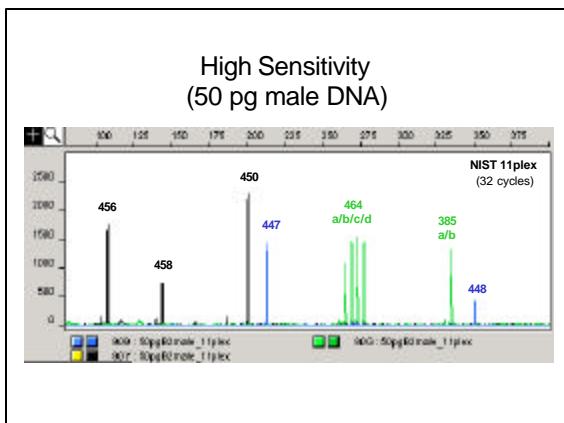
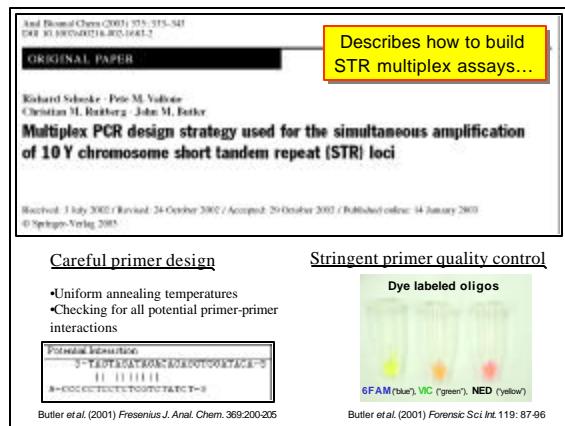
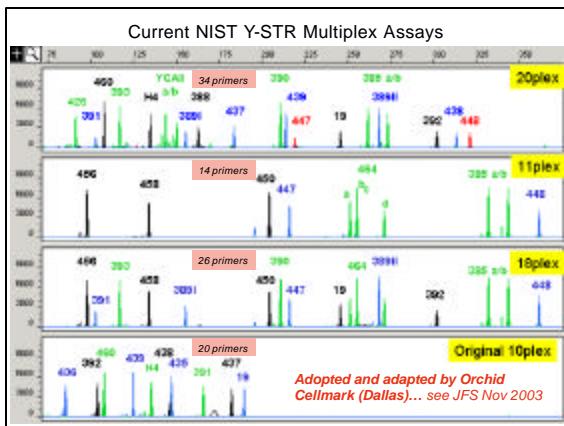
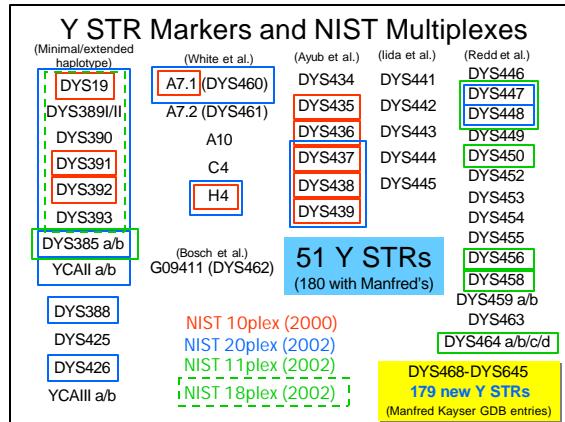
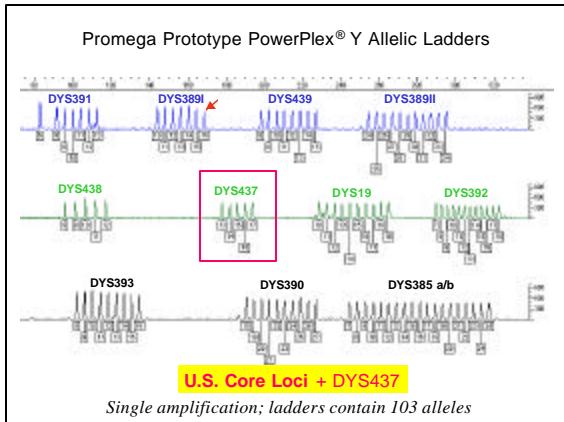
ReliaGene Y STR Kits on ABI 310

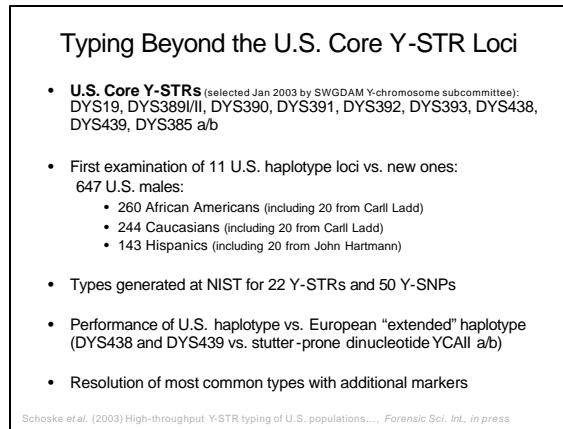
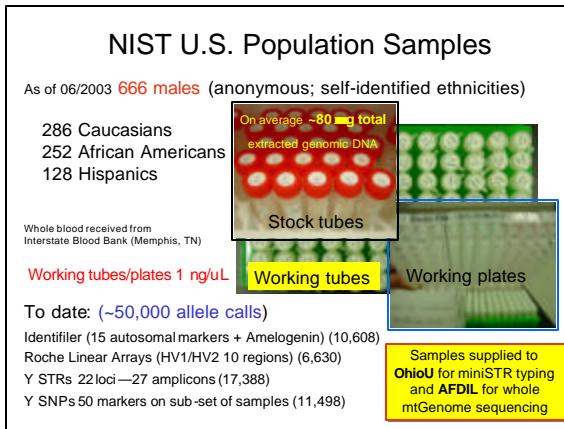
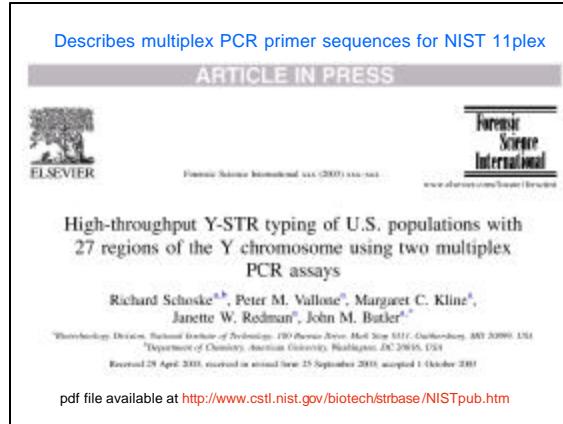
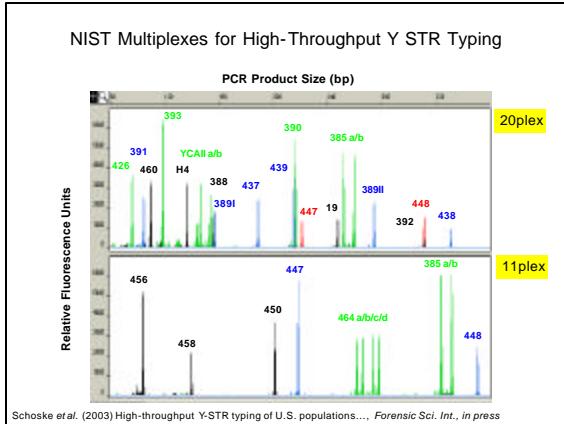


Commercial Y STR Kits

Y STRs in YCAII				
	(White et al.)	(Ayub et al.)	(Iida et al.)	(Redd et al.)
DYS19				DYS446
DYS389I/II				DYS447
DYS390	A7.1 (DYS460)	DYS434	DYS441	DYS448
DYS391	A7.2 (DYS461)	DYS435	DYS442	DYS449
DYS392	A10	DYS436	DYS443	DYS449
DYS393	C4	DYS437	DYS444	DYS450
DYS385 a/b	H4	DYS438	DYS445	DYS452
YCAII a/b	G09411 (DYS462)	DYS439		DYS453
DYS388				DYS454
DYS425				DYS455
DYS426				DYS456
YCAII a/b				DYS458
				DYS459 a/b
				DYS463
				DYS464 a/b/c/d
				DYS468-DYS645
				179 new Y STRs
				(Manfred Kayser et al. entries)







113 (79) different types in 679 males with DYS464

Y-STR		Overall Population		African Americans		Caucasians		Hispanic	
		ETL	Average	G2L	Average	G2L	Average	G2L	Average
		(N=647) Rank	(N=647) Rank	(N=340) Rank	(N=340) Rank	(N=340) Rank	(N=340) Rank	(N=143) Rank	(N=143) Rank
DYS464	white	0.956	1	0.954	3	0.954	1	0.953	1
	a/b	0.912	2	0.942	2	0.832	2	0.933	2
YCAII a/b		0.790	3	0.797	3	0.701	5	0.772	4
DYS533		0.765	4	0.758	5	0.743	3	0.793	3
DYS390		0.764	5	0.664	10	0.701	5	0.665	13
DYS391		0.747	6	0.767	4	0.683	7	0.748	5
DYS392		0.736	7	0.732	6	0.675	8	0.734	6
DYS393		0.721	8	0.722	6	0.595	11	0.704	8
DYS438		0.700	9	0.671	9	0.731	4	0.695	9
DYS439		0.691	10	0.560	15	0.984	12	0.690	10
DYS385		0.676	11	0.642	8	0.498	19	0.672	12
DYS399		0.656	12	0.636	11	0.659	9	0.717	7
DYS460		0.637	13	0.499	17	0.583	13	0.624	14
H4		0.611	14	0.612	12	0.562	14	0.609	15
DYS299		0.609	15	0.454	20	0.296	10	0.673	11
DYS460		0.570	16	0.568	14	0.555	15	0.556	18
DYS389I		0.549	17	0.531	16	0.538	17	0.596	16
DYS391		0.534	18	0.447	19	0.552	16	0.577	17
DYS392		0.519	19	0.373	21	0.402	20	0.522	19
DYS393		0.489	20	0.487	18	0.177	22	0.414	21
DYS385		0.485	21	0.586	13	0.363	21	0.448	20
DYS388		0.365	22	0.246	22	0.501	18	0.312	22

Schoske et al. (2003) High-throughput Y-STR typing of U.S. populations..., *Forensic Sci. Int.*, in press

Extended Haplotype vs. US Haplotype

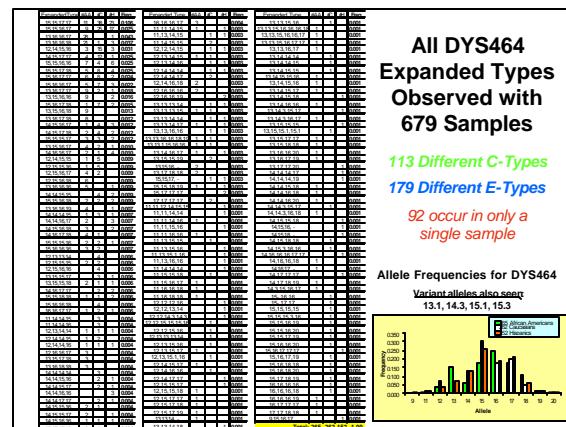
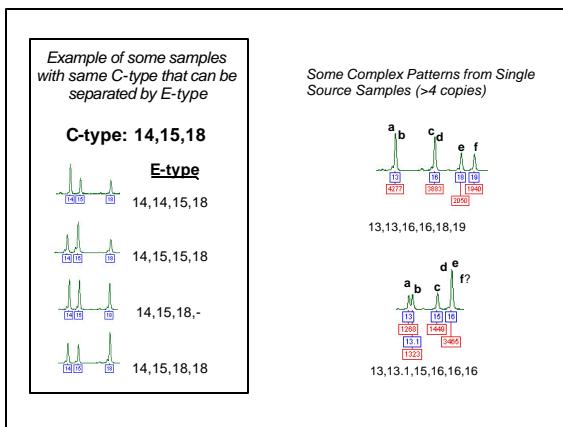
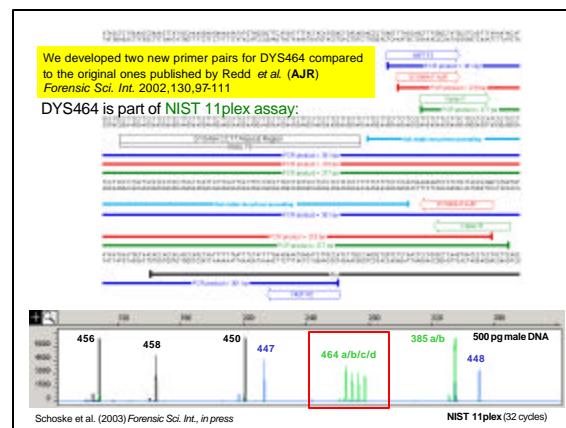
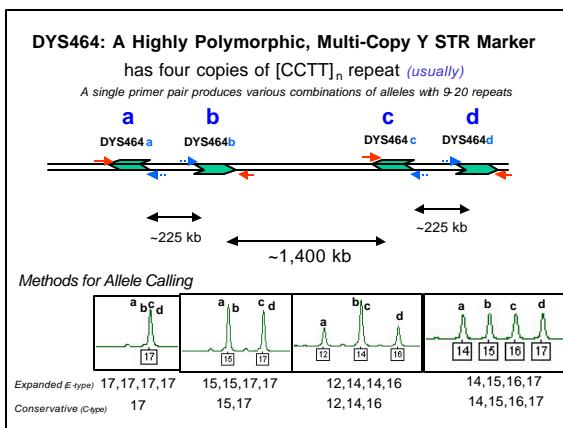
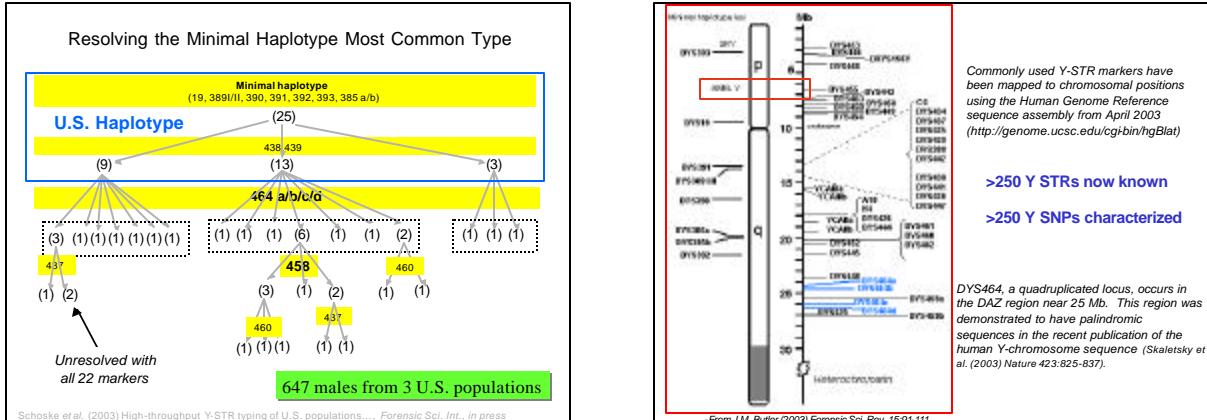
Extended haplotype : 19, 389/II, 390, 391, 392, 393, 385 a/b + YCAII a/b
US haplotype: 19, 389/II, 390, 391, 392, 393, 385 a/b + 438, 439

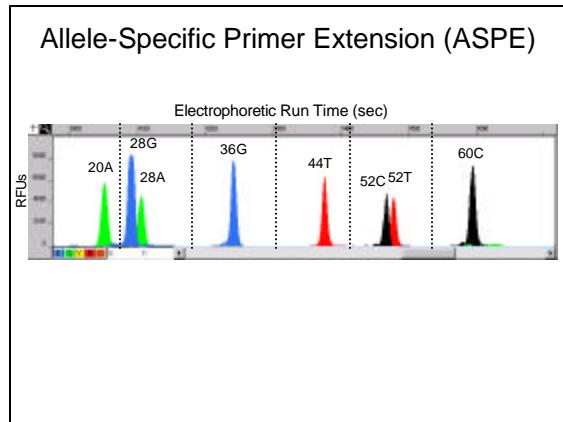
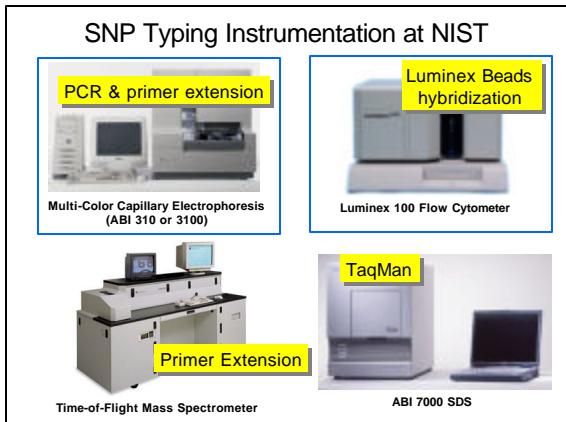
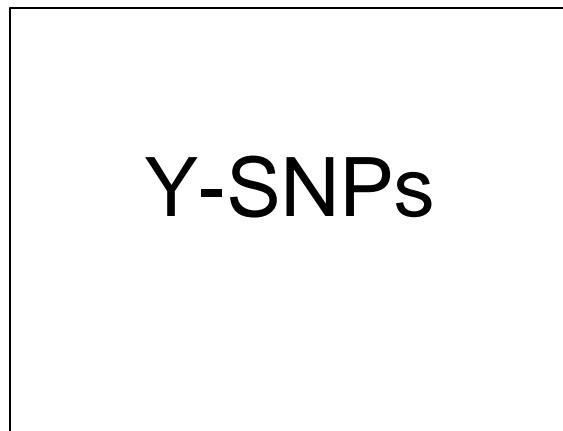
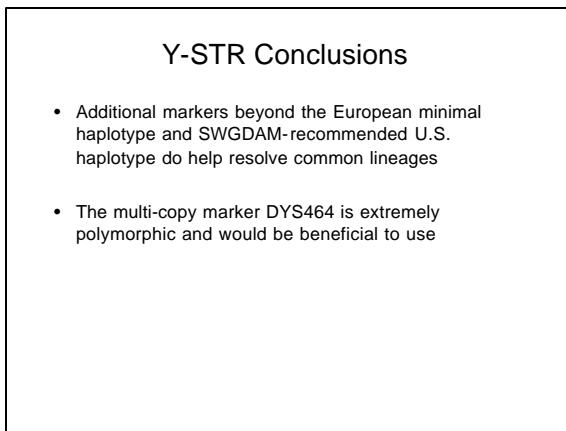
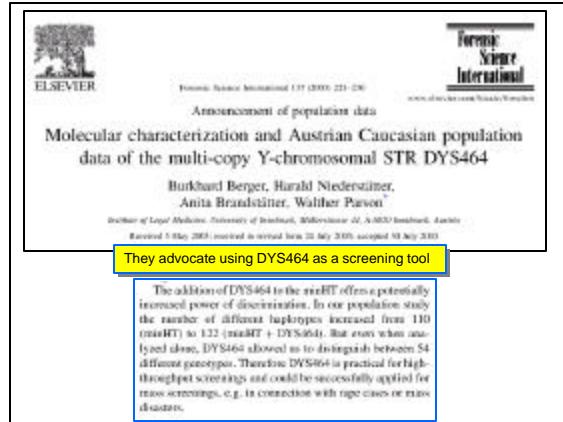
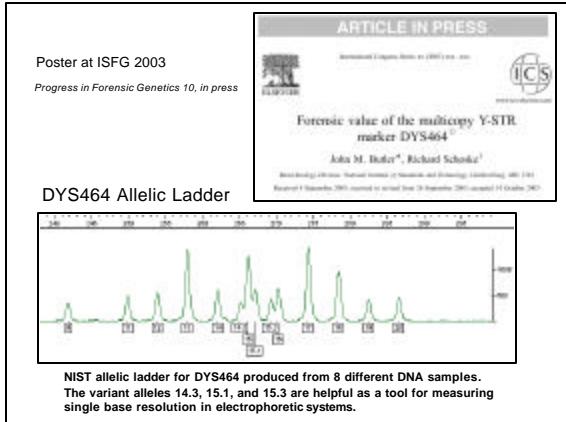
Y-STR Major Combinations		360 African Americans		246 Caucasians		143 Hispanics		
HD	RMP	HD	RMP	HD	RMP	HD	RMP	
extended haplotype	0.9882	0.0013	0.9971	0.0029	0.9975	0.0025		
"U.S. haplotype"	0.2923	0.0007	0.3974	0.0025	0.3986	0.0014		
HD = haplotype diversity								
RMP = random match probability ($1 - \prod_i p_i^{n_i}$)								

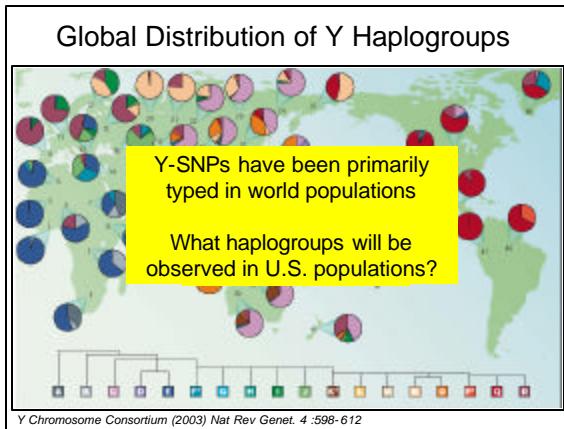
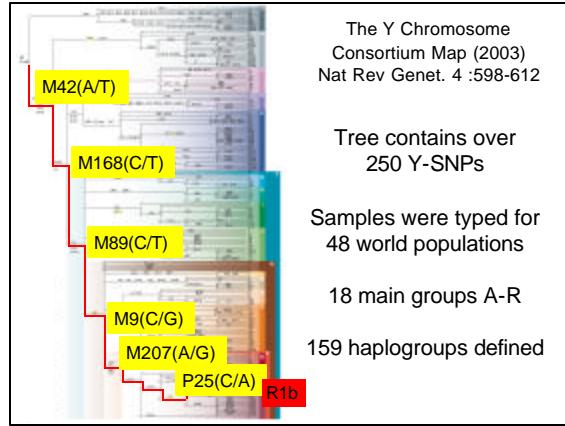
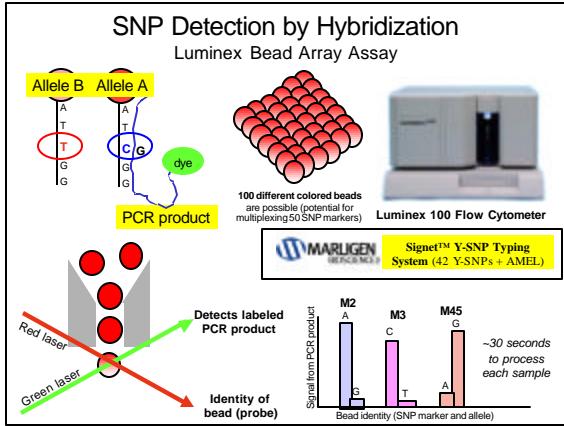
U.S. haplotype is as good as extended haplotype in all major U.S. populations

Y-STR Major Combinations		360 African Americans		246 Caucasians		143 Hispanics		
DC	UH	DC	UH	DC	UH	DC	UH	
"original" haplotype	99.9%	0.1%	99.9%	0.1%	99.9%	0.1%	99.9%	0.1%
"extended" haplotype	91.9%	2.2%	83.0%	18.4%	83.5%	16.0%	83.5%	16.0%
"U.S. haplotype"	91.9%	2.2%	83.7%	17.6%	91.3%	12.1%		
DC = discrimination capacity (number of haplotypes/number of samples)								
UH = unique haplotype (occurs only once in a given population)								

Schoske et al. (2003) High-throughput Y-STR typing of U.S. populations..., *Forensic Sci. Int.*, in press







Y-SNPs in U.S. populations

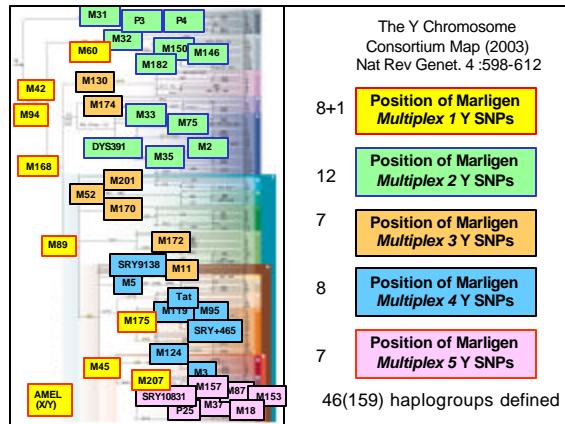
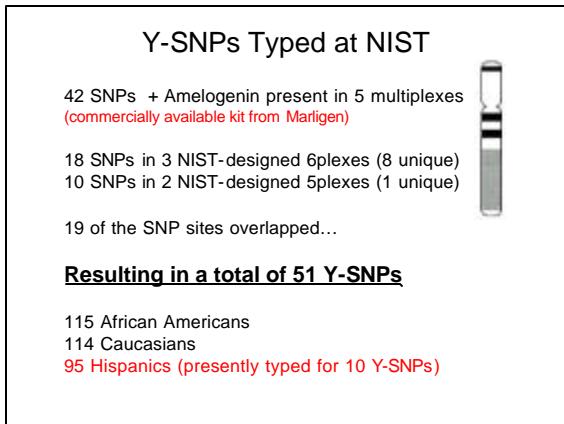
What haplogroups will be observed?

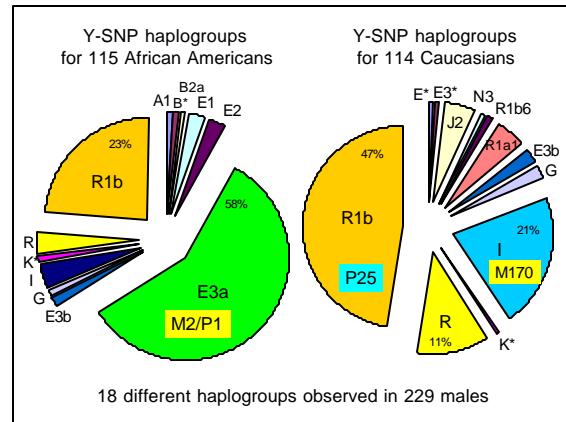
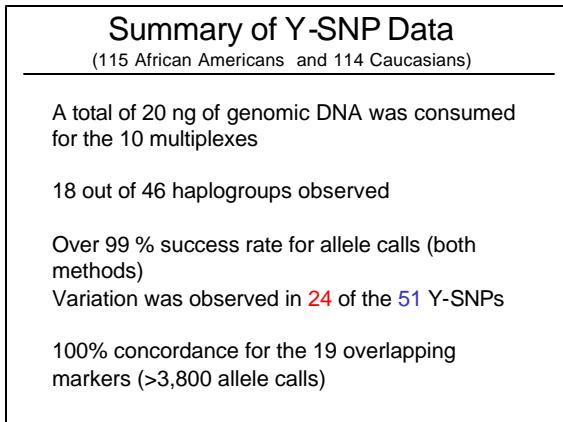
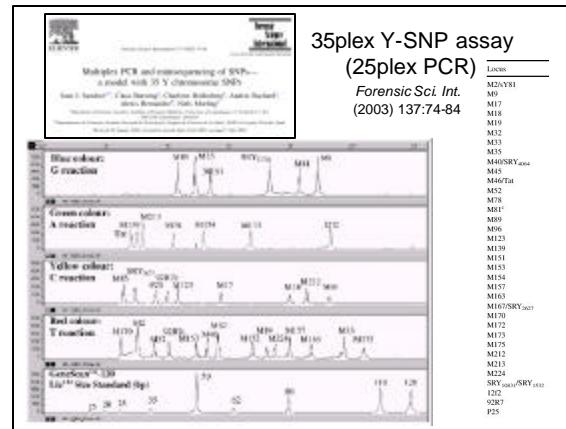
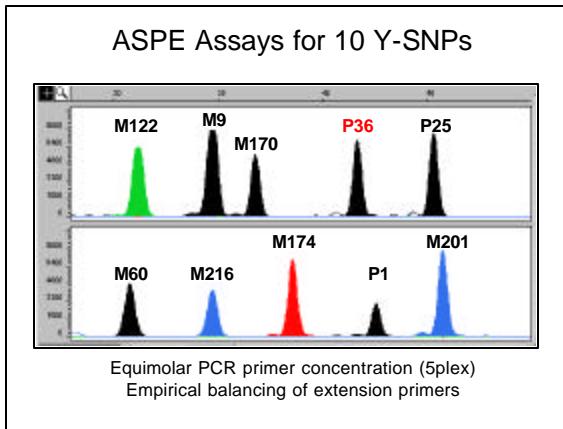
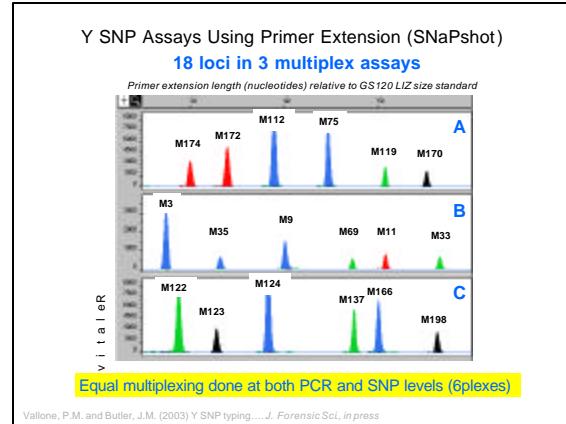
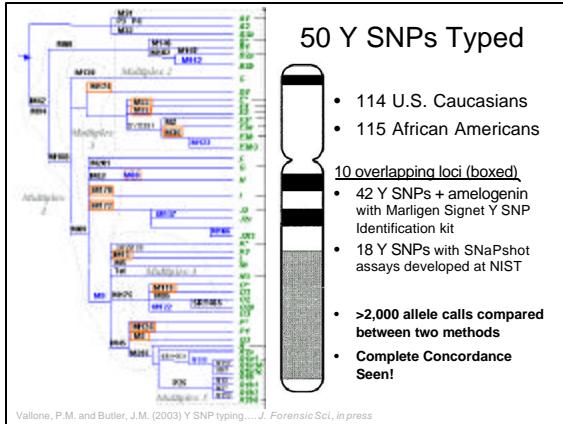
How specific will certain Y-SNPs be for a U.S. population group?

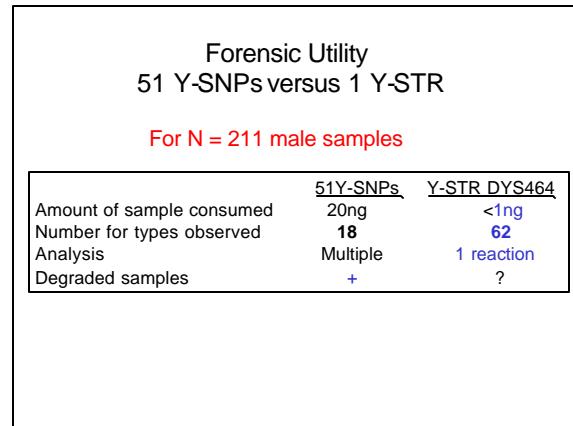
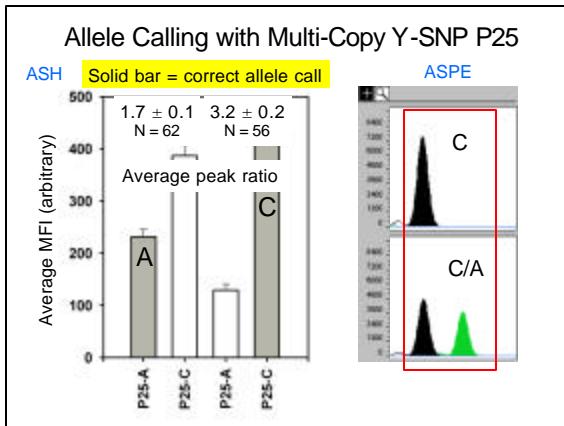
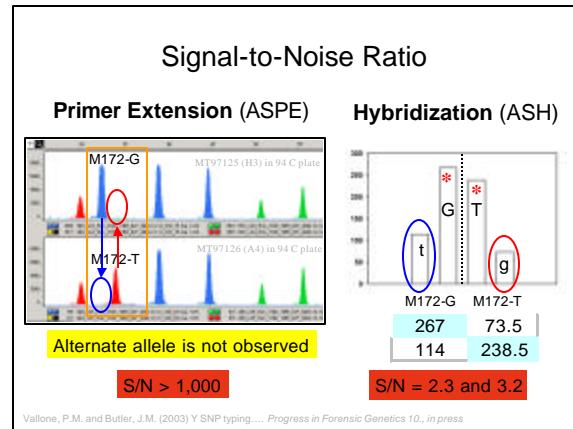
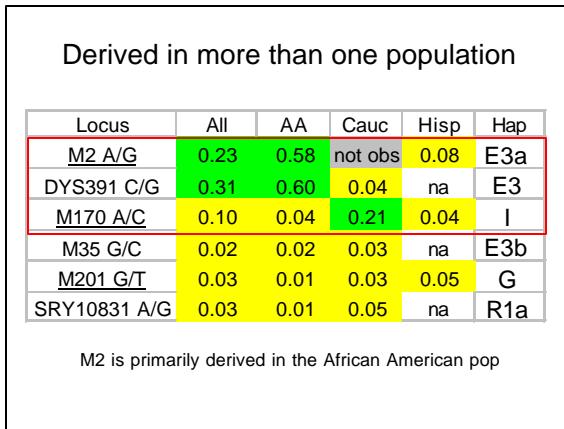
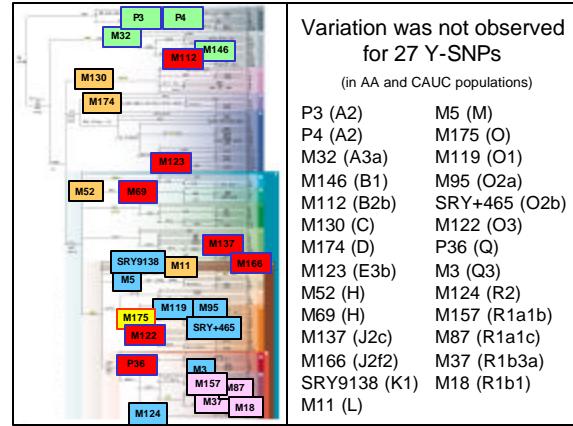
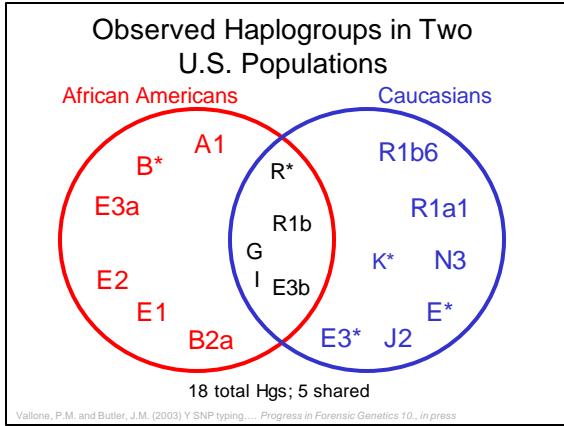
Forensic utility in comparison/addition to Y-STRs

Commercial kit (Marligen) 42 Y-SNPs

Medium sized multiplexes developed in-house (CE or MS)







Y-SNP Conclusions

- Full concordance was observed between hybridization and primer extension technologies on 18 different Y-SNPs (>3,800 allele calls)
- Caucasian admixture was observed with our African American population (Hg R and R1b in ~30%)—agrees with Kayser *et al.* (2003) *Genome Res.* 13:624-634 done with 9 Y-STRs
- Y-SNPs may have limited value for ethnic differentiation in U.S. populations
 - One exception: M2 not in Caucasians
- Y-SNPs are not a useful stand-alone assay for forensic purposes, but may be helpful in combination with Y-STRs

NIST SRM 2395 Human Y-Chromosome DNA Profiling Standard

NIST SRM 2395 Human Y Chromosome DNA Profiling Standard

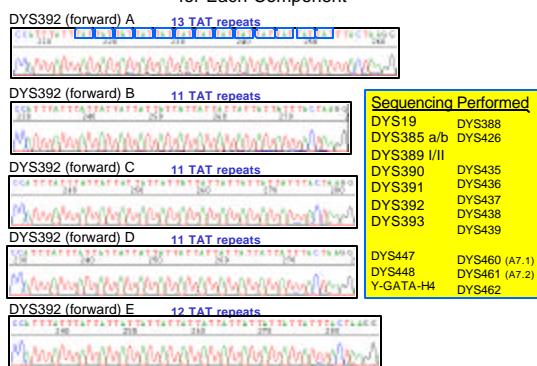
- 5 male samples + 1 female sample (neg. control)
- 100 ng of each component (50 µL at ~2 ng/µL)
- 22 Y STR markers sequenced to provide certified values (number of repeats)
- 9 additional Y STR markers typed
- 42 Y SNPs typed with Marligen kit
- Now available from NIST Standard Reference Material office (www.nist.gov/srm)
- Can be used to verify results with *any* primer sets
- Will help U.S. labs meet DAB/FBI Standards

SRM 2395 Now Available...



<http://www.nist.gov/srm>

Sequence Determination of Y STR Repeat Region for Each Component



Y SNP Results on SRM 2395 from Marligen Signet™ Multiplexes (Luminex bead assay)

SRM 2395	AMEL	M207 (A/G)	M45 (A/G)	M89 (C/T)	DYS391 (C/G)	M2 (A/G)	M170 (A/C)	M172 (G/T)	M201 (G/T)
Component A	XY	G	A	T	C	A	A	T	G
Component B	XY	A	G	T	C	A	A	G	G
Component C	XY	A	G	C	G	G	A	T	G
Component D	XY	A	G	T	C	A	A	T	T
Component E	XY	A	G	T	C	A	C	T	G
Component F	XX								

SRM components are all distinguishable from one another with these Y SNPs

42 Y SNPs measured across all samples

STRBase
Short Tandem Repeat DNA Internet Database

working with industry to develop new applications for forensic science and continuing to support the needs of law enforcement and scientists

General Information	Forensic Interest Data	Supplemental Info
• Intro to STRs (downloadable PowerPoint)	• FBI CODIS Core Loci	• Reference List
• STR Fact Sheets	• DAB Standards	• Technology Review
• Sequence Information	• NIST SRM 2391	• Addresses for Scientists
• Multiplex STR Kits	• Published PCR Primers	• Links to Other Web Sites
• Variant Allele Reports	• Y-Chromosome STRs	Standardized information formats
	• Population Data	
	• Validation Studies	

<http://www.cstl.nist.gov/biotech/strbase>

STRBase
Short Tandem Repeat DNA Internet Database

<http://www.cstl.nist.gov/biotech/strbase>

Y-Chromosome STR Information Available

- Over 200 publications on Y STRs & SNPs cataloged
- Allele information on over 20 Y STR loci
- Downloadable PowerPoint on Y STRs and Y SNPs
- Links to other Y-chromosome sites
- Information on new Y STR multiplexes developed at NIST (published 20plex primers)
- Y STR mapped positions along chromosome

Example Y STR Fact Sheet from STRBase

www.cstl.nist.gov/biotech/strbase

We would like to collect variant alleles for Y STRs as they are discovered...

PCR Product Sizes of Observed Alleles					
	Allele	Set 1	Set 2	Set 3	Repeat Structure
DYS391	288	287	295	297	(TAT) ₂₈
DYS392	348	348	346	346	(TAT) ₃₄
DYS393	346	346	346	346	(TAT) ₃₄
DYS394	348	348	347	347	(TAT) ₃₄
DYS395	252	252	248	248	(TAT) ₂₅
DYS396	255	255	252	252	(TAT) ₂₅
DYS397	258	258	254	254	(TAT) ₂₅
DYS398	361	361	357	357	(TAT) ₃₆
DYS399	364	362	360	360	(TAT) ₃₆
DYS400	387	387	385	385	(TAT) ₃₈

Allele Callers: alleles 18-19 present in T-FLX; 20 from Profiler Plus

Summary of NIST Y Chromosome Work

- Development of new Y STR multiplex assays (**Y STR 20plex, etc.**)
- Evaluation of SNP typing methodologies and development of **Y SNP assays** involving primer extension and the SNaPshot kit
- Creation of a Y Chromosome Standard Reference Material (**SRM 2395**)
- Standardization of information on Y chromosome markers with internet accessibility (**STRBase**)

>30,000 Y chromosome allele calls generated to aid studies on optimal markers for U.S. populations

Publications from Our Group on Y Chromosome Assays

- Schoske, R., Butler, J.M., Vallone, P.M., Kline, M.C., Prinz, M., Redd, A.J., Hammer, M.F. (2001) Development of Y STR megaplex assays. *Proceedings of the Twelve International Symposium on Human Identification 2001*, Promega Corporation. <http://www.promega.com/genetid/proc/usymp12/proc/content/butter.PDF>
- Butler, J.M., Schoske, R., Vallone, P.M., Kline, M.C., Redd, A.J., Hammer, M.F. (2002) A novel multiplex for simultaneous amplification of 20 Y chromosome STR markers. *Forensic Sci Int.* 129: 10-24.
- Schoske, R., Vallone, P.M., Ruitberg, C.M., Butler, J.M. (2003) General strategy for the design of multiplex PCR assays involving Y STR markers. *(2003) Anal Bioanal Chem.*, 375: 333-343.
- Butler, J.M., Schoske, R., Vallone, P.M. (2003) Highly multiplexed assays for measuring polymorphisms on the Y-chromosome. *Progress in Forensic Genetics 9* (Brinkmann, B. and Carracedo, A., eds.), Elsevier Science: Amsterdam, The Netherlands, International Congress Series 1239, pp. 301-305.
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NIST Project Team:
Pete Vallone
Margaret Kline
Jan Redman
Rich Schoske (American U)
Dave Duewer

Collaborators:
Mike Hammer and Alan Redd (U. AZ)
for Y STR sequences

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