S16264 Proposed Mutation History for groups with 2 or more known haplotypes Rev.2 21-Dec-16			
All known S16264+ have DYS19 >= 15	16 may have been the original ancestral value as it is shared by 2 widely separated lines while only being present	t in less than (	0.5% of L21
Hawks + Manns + Mills + L DYS464c <= 16 DYS444 >= 13 DYS504 = 18	eprovost + Bond + Salmon Turners have 17 modal, 16 in Manns and 41% of L21, 15 in Hawk + Leprovost + Bond + Salmon Turners and 8% of L21. Hawk lii 13 is the likely ancestral value with later mutation to 14 in ancestor of Leprovost + Bond + Salmon Turners Grouazel, Mann, Mills and Leprovost not tested for this marker but Hawk + Bond + Salmon Turners have this valu	kely due to ui ue (as do 5% d	nrelated unusual 464 mutations. of L21)
Hawks have DYS390 = 23 DYS389ii = 28 DYS464 extra DYS576 = 16 DYS413a = 22	frequency 7% of L21 vs. modal 24 or (ii-i = 15) frequency 3% of L21 vs. modal 16. range from 14-15-15-16-17 to 14-15-15-15-16-17. Highly unusual. frequency 2% of L21 vs. modal 18 frequency 20% of L21 vs. modal 23. Also shared with Grouazel but his is most likely an independent mutation.		
DYS534 = 14 Note there may be	frequency 13% of L21 vs. modal 15 nore shared off-modals in markers 68-111 but only one Hawk has tested these, finding 6 off-modals not shared w	ith other grou	ıps.
Manns + Mills + Leprovost DYS391 = 10 DYS385a = 12 DYS458 = 16 DYS460 >= 12 DYS456 = 15 DYS570 <= 16 Y-GATA-A10 = 12 DYS463 = 21 Note there may be Mills and Leprovost	+ Bond + Salmon Turners have frequency 32% of L21 vs. modal 11 frequency 6% of L21 vs. modal 11. Bond has back-mutation to 11. frequency 16% of L21 vs. modal 17 (assuming 455 = 10 mutation has occurred only once with Mills having 458 ba 11 modal, 12 in Manns, Mills and 9% of L21, 13 in Leprovost + Salmon Turners and 1% of L21. Bond, 1 Mann and Mann, Mills, Leprovost and 36% of L21 vs. modal 16. Most parsimonious tree requires a back-mutation to 16 for 17 modal, 16 in Mills + Leprovost + Bond + Salmon Turners and 15% of L21, 15 in Manns and 1% of L21 frequency 15% of L21 vs. modal 13 frequency less than 0.5% of L21 (to date only found in this group in L21) vs. modal 24 more shared off-modals in markers 38-111 but only 1 Mann has tested 11 out of these 74 markers not tested for markers 68-111 so only assumed to match GATA-A10 and DYS463 based on other men's results	ck-mutation † 1 Mills has b Bond and Tu	to 17) ack-mutation to 11. rners.
Manns have DYS439 = 11 DYS576 = 19 CDYa = 37 CDYb = 37 Note there may be	frequency 21% of L21 vs. modal 12 frequency 20% of L21 vs. modal 18 frequency 27% of L21 vs. modal 36 frequency 18% of L21 vs. modal 38 more shared off-modals in markers 38-111 but only 1 Mann has tested 11 out of these 74 markers		
Mills, Leprovost + Bond + DYS455 = 10 DYS520 = 21 Note there may be	Salmon Turners have frequency 1% of L21 vs. modal 11 frequency 9% of L21 vs. modal 20 (no Mann tested for this marker) nore shared off-modals in markers 68-111 but Mills and Leprovost have not tested these		
Mills have DY\$385b = 15 DY\$389i = 14 DY\$449 = 30 DY\$576 = 17 CDYa = 35 CDYb = 39 Note there may be	frequency 17% of L21 vs. modal 14 frequency 14% of L21 vs. modal 13. Hence Mills DYS385ii = 30 with i-ii constant at 16. frequency 41% of L21 vs. modal 29 (29 is P312 modal, 30 is L21 modal) ** frequency 21% of L21 vs. modal 18 frequency 12% of L21 vs. modal 36 frequency 26% of L21 vs. modal 38 more shared off-modals in markers 68-111 but Mills have not tested these		
Leprovost + Bond + Salmon Turners have DYS464d = 15 frequency 1% of L21 vs. modal 17 Note there may be more shared off-modals in markers 68-111 but Leprovost has not tested these			
Bond + Salmon Turners ha DYS393 = 12 DYS439 = 13 DYS464a = 14 CDYb = 36 DYS446 = 14 DYS710 = 34 DYS540 = 13 DYS540 = 13 DYS5714 = 25 DYS505 = 13 DYS505 = 13 DYS522 = 11 DYS650 = 20 DYS561 = 16 ** Mann and Lepro	ve frequency 2% of L21 vs. modal 13 frequency 13% of L21 vs. modal 12 frequency 11% of L21 vs. modal 15 frequency 16% of L21 vs. modal 15 frequency 6% of L21 vs. modal 13 frequency 24% of L21 vs. modal 13 frequency 24% of L21 vs. modal 37 (37 is P312 modal, 34 is L21 modal - a highly variable SNP) ** frequency 4% of L21 vs. modal 12 ** frequency 41% of L21 vs. modal 12 (6 (26 is P312 modal - another highly variable SNP) ** frequency 79% of L21 vs. modal 12 ** frequency 79% of L21 vs. modal 10 (10 is P312 modal - another variable SNP) ** frequency 12% of L21 vs. modal 19 ** frequency 15% of L21 vs. modal 15 ** vost not tested for these markers		
Salmon Turners have DYS390 = 23 CDYa = 34	frequency 7% of L21 vs. modal 24 frequency 3% of L21 vs. modal 36		