## upplemental Table S2: Results of AMOVA (Analysis of Molecular Variance) of selected Asian populations Source of variation Among populations Sum of squares 859.455 Percentage of variation 4.15 3,423 Within populations 20,003.602 5.84388 Vb 95.85 Total 3,435 20,863.057 6.09702 100 b) Population pairwise Fsts Distance method: Pairwise difference Afghanistan Myanmar Uzbekistan Kazakhstan Kyrgyzstan Laos Korea Russians from Uzbekistan Tajikistan Thailand Turkmenistan Vietnam Hong Kong Afghanistan Myanmar Hong Kong 0.08576 0.07269 0.02439 0.03206 Uzbekistan 0.0356 0.04036 Kazakhstan 0.05053 0.04104 0.02778 0.00452 0.05377 0.03228 0.01627 0.00858 0.00525 Kyrgyzstan 0.06886 0.11761 0.06393 0.05454 0.086 0.08941 Korea 0.09431 0.04769 0.01435 0.03642 0.02334 0.01249 0.08284 0 0.11739 0 0.05437 0.05434 Russians from Uzbekistar 0.05619 0.10209 0.09732 0.03454 0.05532 0.13645 0.03759 0.01619 0.02189 0.08081 Tajikistan 0.04877 0.04231 0.02435 Thailand 0.07087 0.01296 0.00572 0.03317 0.03354 0.02376 0.05058 0.03328 0.08905 0.03459 0.04964 0.01904 0.04429 0.03965 0.00333 0.00775 0.01183 0.09269 0.04232 0.03333 0.01982 0.04189 0 0.03203 0.0734 0.04399 0.00542 0.04937 0.00671 0.04005 0.02799 0.03517 0.03961 Vietnam 0.09813 c) Fst p-values (significant p-value after Bonferroni correction 0.000641) Afghanistan Myanmar Hong Kong Uzbekistan Kazakhstan Kyrgyzstan Laos Korea Russians from Uzbekistan Tajikistan Thailand Turkmenistan Vietnam Afghanistan <0.0001 Myanmar Hong Kong < 0.0001 < 0.0001 Uzbekistan Kazakhstan <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 Kyrgyzstan < 0.0001 < 0.0001 < 0.0001 <0.0001 0.00098 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 Korea < 0.0001 Russians from Uzbekistan < 0.0001 < 0.0001 < 0.0001 < 0.0001 < 0.0001 < 0.0001 < 0.0001 Taiikistan <0.0001 <0.0001 <0.0001 <0.0001 < 0.0001 < 0.0001 <0.0001 <0.0001 <0.0001 Thailand <0.0001 <0.0001 0.00098 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 0.00098 <0.0001 < 0.0001 < 0.0001 Turkmenistan < 0.0001 < 0.0001 < 0.0001 < 0.0001 < 0.0001 <0.0001 Vietnam <0.0001 <0.0001 0.00098 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 < 0.0001 <0.0001 0.01660 < 0.0001 d) Population average pairwise differences (16024-16569;1-576; C-insertions around 16193,309,315 and 573 were ignored) Above diagonal: Average number of pairwise differences between populations (PIXY) Diagonal elements: Average number of pairwise differences within populations (PIX) Below diagonal: Corrected average pariwise difference (PIXY-(PIX-PIY)/Z) Distance method: Pairwise difference Afghanistan Myanmar 10.40143 11.87796 Hong Kong 12.20527 Uzbekistan Kazakhstan Kyrgyzstan Laos Korea Russians 11.58257 13.00753 13.36978 11.99111 10.1215 Russians from Uzbekistan Tajikistan Thailand Turkmenistan Vietnam 12.0471 12.17234 10.99285 12.0778 Afghanistan 11.27253 12.07781 Myanmar Hong Kong 10.4168 11.27113 12.00311 12.14755 11.76914 12.12204 11.90342 13.10447 12.98383 11.8268 11.1953 13.32419 11.8696 11.65008 12.38692 11.8717 11.51238 12.78469 12.22844 11.85414 11.82761 12.19479 13.33988 Uzbekistan 0.41329 0.47505 0.38975 11.31704 11.49051 12.81936 13.31868 11.71199 10.40367 12.08397 12.14136 10.99841 12.10871 Kazakhstan 0.60165 0.48765 0.3408 0.05178 11.56041 12.90164 13.51424 11.67823 10.76381 12.31156 12.2749 11.16953 12.28862 0.75288 0.41497 0.21217 0.0675 1.20253 14.10787 14.59292 12.83063 12.24216 13.06301 13.21593 12.68184 13.46264 12.50457 13.29016 13.03417 13.40377 Kyrgyzstan 1.00748 13.08996 1.63756 0.81676 0.71891 1.12866 13.88403 Korea 1.16309 0.56393 0.16852 0.42616 0.27072 0.14939 1.05712 11.25461 11.42748 12.52473 12.10551 11.41973 12.01161 1.43518 8.72998 0.66545 0.63731 11.23428 11.49299 10.01107 12.46396 12.75876 11.7675 11.47951 12.70613

Sources of	data t	aken from	literature:

0.5558

0.61441

0.88683

0 48945

0.90777

1.19475

0.51938

0.15134

0.57413

0.22273

1.21131

0.47893

0.06987

0.47768

0.08184

0.38016

0.19348

0.39805

0.0372

0.48087

Russians from Uzbekistan

Tajikistan

Thailand

Vietnam

Afghanistan	Irwin et al.: The mtDNA composition of Uzbekistan: a microcosm of Central Asian patterns. Int J Legal Med 2010, 124: 195-204.
Hong Kong	Irwin et al.: Mitochondrial DNA control region variation in a population sample from Hong Kong, China. Forensic Sci Int Genet 2009, 3: e119-e125.
Uzbekistan	Irwin et al.: The mtDNA composition of Uzbekistan: a microcosm of Central Asian patterns. Int J Legal Med 2010, 124: 195-204.
Kazakhstan	Irwin et al.: The mtDNA composition of Uzbekistan: a microcosm of Central Asian patterns. Int J Legal Med 2010, 124: 195-204.
Kyrgyzstan	Irwin et al.: The mtDNA composition of Uzbekistan: a microcosm of Central Asian patterns. Int J Legal Med 2010, 124: 195-204.
Laos	Bodner et al.: Southeast Asian diversity: first insights into the complex mtDNA structure of Laos. BMC Evol Biol 2011, 11:49.
Korea	Lee et al.: Mitochondrial DNA control region sequences in Koreans: identification of useful variable sites and phylogenetic analysis for mtDNA data quality control. Int J Legal Med 2006, 120: 5-14.
Russians from Uzbekistan	Irwin et al.: The mtDNA composition of Uzbekistan: a microcosm of Central Asian patterns. Int J Legal Med 2010, 124: 195-204.
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Turkmenistan	Irwin et al.: The mtDNA composition of Uzbekistan: a microcosm of Central Asian patterns. Int J Legal Med 2010, 124: 195-204.
Vietnam	Irwin et al.: Mitochondrial control region seguences from a Vietnamese nonulation sample. Int Llegal Med 2008, 122: 257-259

0.61861

0.29938

0.08664

0.53909

0.4099

0.82323

0.29776

0.32391

0.14794

0.38051

1.78535

1.12054

0.67386

1.19997

0.58913

0.39341 1.04321

0.48974 0.3434

0.41498 1.1452 0.44199

0.23283

0.50483

12.16959 11.8802

0.49271 10.60538

0.06575 0.58001

12.11987

11.85202

11.93865