

Figure S1 Plot of scores for multidimensional scaling (MDS) analysis based on genetic distance. The groups of dots identify genetic outliers. The slightly larger cluster (circled red) represents Section A&D group.

Each cluster contained following individuals:

> cl1

[1] "F64" "F66" "F79" "F84" "F40" "F41" "F25" "F62" "F75" "F77" "F37" "F13" "F69" "F86" "F42" "F51" "F52" "F83" "F67" "F71" "F82" "F53" "F68"

> cl2

[1] "H34" "H36" "H14" "H18" "H32" "H33" "H6" "H12" "H47" "H22" "H5" "H7" "H26" "H29" "H2" "H30" "H43"

> cl3

"A33" "A41" "A76" "A78" "D19" "D89" "D96" "A63" "A66" "D75" "D77" "A50" "A52" "A79" "A86" "D48" "D62" "D100" "A69" "A72" "D78" "D88" "D13" "D15" "D56" "D63" "A46" "A85" "D52" "D53" "A29" "A36" "A73" "A74" "D79" "D104" "A44" "A84" "A59" "D51" "D87" "A67" "A70"

> cl4

[1] "N12" "N17" "N1" "N2" "N30" "N31" "N18" "N21" "N8" "N32" "N44" "N36" "N37" "N9" "N19" "N41" "N23" "N33" "N6" "N29" "N38" "N39"