

Application for the Walk Through The Y Project

Project Goals

- Sequencing of 3 plates of 96Y chromosome PCR segments each.
- We're sequencing in both the forward and the reverse directions.
- The average number of total unique bases sequenced on the ChrY will exceed 100 kB.
- No re-runs (unless sample failed completely or < 50 kB).
- Segments can be considered as randomly chosen regions on the Y chromosome which are unlikely to recombine.
- All known derived SNPs and all differences from the HUGO reference sequence will be scored.
- Participants can opt for public presentation of their results. This decision cannot be reversed.
- If a new SNP is found the participant can decide whether FTDNA makes it public. In the event that multiple participants have this SNP, only one need submit their decision for FTDNA to make the SNP public. This decision cannot be reversed.
- Only public markers will be available to order as single SNPs from the advanced orders page.
- FTDNA will try to submit new public SNPs to NCBI (for a rs number). We cannot guarantee that they will be accepted.
- If a customer publishes a new marker all involved GRC laboratory personal must be added as co-authors.
- Each participant must add all previous Y-STR and Y-SNP results to his application.
- Please indicate the testing company for non-FTDNA results.
- Collective submission of similar participants by group is strongly encouraged.
- FTDNA reserves the right to limit the number of participants according to our production capacity.
- The GRC laboratory director decides if and when an application will be accepted based on the scientific importance.
- Public groups of well-selected individuals will be preferred. This decision cannot be disputed.
- **Cost: US\$ 750 per participant.**
- Turnaround time: 6 weeks after fresh swabs have been received at FTDNA.

Information supplied by the participant:

Name: _____ Surname: _____ FTDNA kit # _____

Name of the WalkThrough the Y group: _____

My raw data should be handled: private public

I agree that the following persons will have full access to my raw data and results:

Origin of the Y chromosome line ~400 years ago:

Can this origin be genealogically proven? Yes No

Continent: _____ Country on the current world map: _____

City on the current world map: _____

Country and/or city names ~400 years ago: _____

Ethnicity of the Y chromosome line:

Please precisely indicate the ethnicity: _____ (e.g. Askenazi, Inuit, Bantu, Pashtun etc.)

Can this ethnicity be genealogically proven? Yes No

FTDNA Y-STR results: (No details necessary. We'll look up your kit #)

No Y-STR 12 Y-STR 25 Y-STR 37 Y-STR 67 Y-STR

FTDNA/DNA-FP Advanced Y-STR results (No details necessary. We'll look up your kit #)

Please give the approximate number beyond Y67: _____

Y-SNPs tested at FTDNA: (No details necessary. We'll look up your kit #)

Shorthand Haplogroup at deepest tested level: _____ if untested write "untested"

Approximate # of SNPs _____ or email kit # and text file to: thomas@familyreedna.com

Are there any known Y chromosome deletions or duplications? Yes No

If yes, please specify: _____

Y-STR & Y-SNP results from other companies:

List [company, marker , result]: _____

or send an e-mail with your kit # and an attached text file to: thomas@familyreedna.com