

Please e-mail completed form to NTR data management: [ntr.datamanagement.fgb@vu.nl](mailto:ntr.datamanagement.fgb@vu.nl)

1. **Applicant information:**

|  |  |  |  |
| --- | --- | --- | --- |
| Data Request Number: | <To be completed by NTR data management> | | |
| Original data request number: | Click or tap here to enter text. | | |
| Original data request title: | Click or tap here to enter text. | | |
| Applicant name: | Click or tap here to enter text. | | |
| Country: | Click or tap here to enter text. | | |
| E-mail: | Click or tap here to enter text. | | |
| Mobile phone or secondary e-mail\* | Click or tap here to enter text. | | |
| Site: | NTR | VU | External: Click or tap here to enter text. |
| *\* to send you the access code for retrieving the data* | | | |

1. **Project (working) title :** Click or tap here to enter text.
2. **Please attach the following:**
   1. Original data sharing request form
   2. Original analysis plan
3. **Changes to the original analysis plan motivating the amendment:**

Click or tap here to enter text.

1. **I would like to request the following new phenotype data:**

I am attaching a .JSON file (created with the NTR Data Showcase) with the requested variables

I need the following additional variables not (yet) incorporated in the NTR repository Data Show Case: *(be as specific as possible, e.g. if survey data are requested please indicate from which survey wave(s).*

Click or tap here to enter text.

1. **I would like a new subject selection** *(Only needed if you require data from participants not in your original dataset)*

Click or tap here to enter text.

1. **I would like to request the following new genotype data** *(Only needed if you did not request these before)*

*Please note that individual level genotype data can only be requested for analyses performed on secure computer systems (including SURFSara) under direct supervision of NTR collaborators.*

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| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| **Standard variables (always needed when working with genotype data):** | | | | | | | | |
| Individual ID | | Father ID | | Mother ID | | Sex | Genotype Platform Dummies | |
| First 10 Genetic Principal Components | | | | Ancestry Outlier dummy | | |  | |
| **Other information:** | | | |  | | |  | |
| Imputation base: | | | 1000 Genomes | | Topmed | | | |
| Genome wide SNP data | | |  | |  | | | |
|  | Format: | | Plink best-guess | | VCF | | | |
| All variants in gene: (default is imputed) | | | | | | | | |
|  | Chromosome: | | start (bp): Click or tap here to enter text. | | end (bp): Click or tap here to enter text. | | | Genome build: 37  38 |
| (Set of) specific SNP(s): (default is imputed) | | | | | | | | |
|  | Chromosome: | | start (bp): Click or tap here to enter text. | | end (bp): Click or tap here to enter text. | | | Genome build: 37 38 |
| *Or provide a .csv file as attachment with Chr, start BP + end BP (or RS id), genotyped only Y/N* | | | | | | | | |

1. **Changes in proposed co-authors, if any (in proposed order):**

The NTR Data Access Request Committee will suggest the appropriate collaborative participation by the relevant NTR researchers. Pre-discussion of appropriate co-authorship with members of the NTR-DAC is strongly encouraged.

|  |  |
| --- | --- |
| (shared) First: | Click or tap here to enter text. |
| (shared) Last: | Click or tap here to enter text. |
| Co-authors (2nd, 3rd , … ): | Click or tap here to enter text. |

1. **Timeline for completion and submission of manuscript:**

Click or tap here to enter text.

|  |  |  |  |
| --- | --- | --- | --- |
| 1. **Permissions** | **Local IRB:** | Yes, IRB reference: | Click or tap here to enter text. |
|  | **VUmc METc:** | Yes [Consult NTR for the VU(mc) IRB reference] | |

**I hereby guarantee to use the data only for addressing the research question described under 3, and not for any other purposes, unless I submit a new analysis plan.**

Signed Date

|  |  |  |
| --- | --- | --- |
|  |  | Click or tap here to enter text. |