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We use this collection and it's working

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Human metagenomics protocols Payami lab

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ABSTRACT

This is a collection of protocols that details the entire process of conducting a human microbiome study from start to end. It includes (1) consent form, to enroll subjects with permission to store and share data (2) data collection protocols designed to be self-administered by subjects (Environmental & Family History Questionnaire, Gut Microbiome Questionnaire, instructions for saliva collection, and instructions for stool collection), (3) codes for bioinformatics and statistical data analysis, (4) workflow at a glance created according to STORM guidelines and (5) final product, the manuscript, which describes the whole process from start to end using metagenomics of Parkinson's disease implicates the gut microbiome in multiple disease mechanisms.

ATTACHMENTS

1. Informed Consent Form used for enrolling subjects into study. This consent allows long-term storage of samples and sharing data.pdf	2a. Data collection protocol_Environment & Family History Questionnaire_Self administered by subjects.pdf	2b. Data collection protocol_Gut Microbiome Questionnaire_Self administered by subjects (1).pdf	2c. Data collection protocol_Saliva Collection Instruction_Self administered by subjects.pdf
2d. Data collection protocol_Stool Collection Instruction_Self administered by subjects.pdf	3a. Data analysis code_bioinformatics and statistics.pdf	3b. Data analysis code_bioinformatics and statistics.Rmd	4. Workflow at a glance_STORM chart.pdf
5. Final product_describes the whole process from start to end_Nat Comm PMID 34408160 .pdf			

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4. Workflow at a glance_STORM chart.pdf	5. Final product_describes the whole process from start to end_Nat Comm PMID 34408160 .pdf					

FILES

Protocol

NAME



1. CONSENT FORM

VERSION 1

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Protocol

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2. Data collection protocols

VERSION 1

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Protocol

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 3. Codes for bioinformatics and statistical data analysis

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Protocol

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 4. Workflow at a glance_ STORM chart

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Protocol

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 5. Metagenomics of Parkinson's disease implicates the gut microbiome in multiple disease mechanisms

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