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**Abstract**

**Background**

Describe the context and purpose of the study.

**Methods**

Write how the study was performed and statistical tests used.

**Results**

Write the main findings.

**Conclusions**

Present a brief summary and potential implications.

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**Methods**

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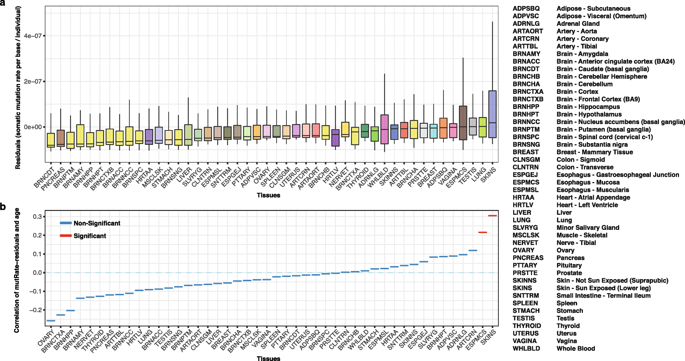
* the aim, design and setting of the study,
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* the type of statistical analysis used, including a power calculation if appropriate.

**Results**

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**Fig. 1**



Rate of somatic mutations varies significantly across the 46 tissues of the GTEx cohort (ignoring kidney, cell-EBV-transformed lymphocytes, and cell-transformed fibroblasts for technical reasons, see the “[Methods](https://genomemedicine.biomedcentral.com/articles/10.1186/s13073-020-00746-1#Sec2)” section). **a** Distribution of the somatic mutation rate per base and individual residuals (mutRate residuals) across analyzed tissues. mutRate residuals represent the somatic mutation rates corrected for non-biological confounders such as PCR duplication rate, RIN, cohort, and read coverage. **b** Spearman correlation between mutRate residuals and age per tissue. Colors show the significance of the correlation test after FDR correction (q value < 0.05 in red).

**Table 1 Table title (max 15 words). Legend (max 300 words) should be included underneath the table**

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Characteristics** | **Group A** | **Group B** | **Group A+B** | **Group C** |
| **Age (years)** | 3.0 (0.9–7.5) | 1.0 (1.0–4.5) | 1.0 (1.0–4.5) | 1.0 (1.0–4.5) |
| > 2.5 | 22/43 (51%) | 21/43 (49%) | 21/43 (49%) | 21/43 (49%) |
| ≤ 2.5 | 21/43 (49%) | 4/7 (57%) | 4/7 (57%) | 4/7 (57%) |
| **Sex** | | | | |
| male | 23/43 (53%) | 5/7 (71%) | 5/7 (71%) | 5/7 (71%) |
| female | 20/43 (47%) | 2/7 (29%) | 2/7 (29%) | 2/7 (29%) |
| **Clinical status** | | | | |
| Asymptomatica | 0/43 (0%) | 2/7 (29%) | 2/50 (4%) | 0/26 (0%) |
| Very mild | 0/43 (0%) | 5/7 (71%) | 5/50 (10%) | 0/26 (0%) |
| Mild | 41/43 (95%) | 0/7 (0%) | 41/50 (82%) | 26/26 (100%) |
| Severe | 0/43 (0%) | 0/7 (0%) | 0/50 (0%) | 0/26 (0%) |
| Critically ill | 2/43 (5%) | 0/7 (0%) | 2/50 (4%) | 0/26 (0%) |
| **Symptoms** | | | | |
| Fever | 29/43 (67%) | 3/7 (43%) | 32/50 (64%) | 21/26 (81%) |
| Cough | 21/43 (49%) | 1/7 (14%) | 22/50 (44%) | 19/26 (73%) |
| Myalgia/fatigue | 2/43 (5%) | 0/7 (0%) | 2/50 (4%) | 0/26 (0%) |
| Sore throat | 1/43 (2%) | 0/7 (0%) | 1/50 (2%) | 0/26 (0%) |
| Diarrhea | 3/43 (7%) | 0/7 (0%) | 3/50 (6%) | 0/26 (0%) |

aNo clinical symptoms and no abnormal CT findings

Results that would otherwise be indicated as "data not shown" should be included as additional files. BioMed Central requires that supporting data are included as additional files or deposited in a recognized repository. The maximum file size for additional files is 20 MB each. Each additional file should be cited in sequence within the main body of text. For more information on additional files, visit: [here](https://genomemedicine.biomedcentral.com/submission-guidelines/preparing-your-manuscript?utm_source=other_website&utm_medium=display&utm_content=mpu&utm_campaign=BSCN_3_CZ01_CN_GM_Template#preparing+additional+files).

**Discussion**

For research articles, this section should discuss the implications of the findings in the context of existing research and highlight limitations of the study.

**Conclusions**

This should state clearly the main conclusions and provide an explanation of the importance and relevance of the study to the field.

**List of Abbreviations**

If abbreviations are used in the text, they should be defined in the text at first use, and a list of abbreviations should be provided as below:

**3Di:** Developmental, Dimensional and Diagnostic Interview

**ABC:** Antecedents, Behaviours and Consequences (chart)

**ACE:** ADHD Child Evaluation

**References**

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