

Modulatory effect of faecal microbiota transplantation on procarcinogenic colibactin-producing bacteria in patients with recurrent *Clostridioides difficile* infection

Introduction

Faecal microbiota transplantation (FMT) is an established treatment for recurrent *C. difficile* infections (rCDI). Recently, it was shown that *Escherichia coli* may produce genotoxic colibactin, which is encoded on the *pks* gene island that holds 19 *clb* genes. Also, *pks*⁺ *E. coli* may be transmitted or cleared after FMT. We studied deep sequenced faecal microbiomes of healthy faeces donors and rCDI patients for *pks* genes and the possibility of transmission to patients and clearance by negative donor faeces after FMT.

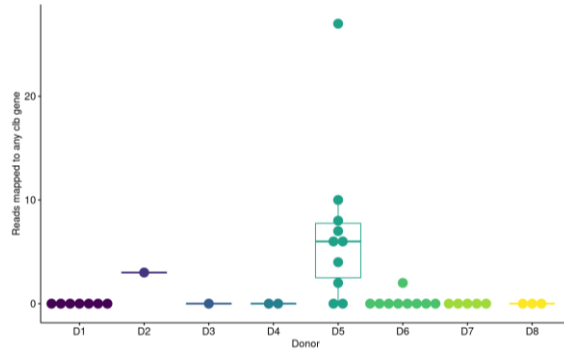


Figure 1, *pks* is present in faecal samples of three out of eight donors. Five donors never have *pks* genes in their faeces, three donors did have at least one *pks*-positive faecal sample. *Pks* values are measured as absolute number of reads matched to the reference.

Materials and methods

CDI patient faeces



× 49

Healthy donor faeces



× 38 (from 9 donors)

Patient faeces after FMT



× 49

49 sample triplets → 136 deep sequenced metagenomes

Method details

Thirty-eight donor faecal suspensions were used to treat 49 patients (some suspensions were used for multiple patients). Metagenomes were deep sequenced by Vedanta Biosciences, Inc. (Cambridge, MA, USA) to 20M+ Illumina 150-PE reads. Metagenomes were analysed with Jovian (<https://doi.org/10.5281/zenodo.3666156>) and *clb* reads were detected by mapping cleaned reads to a reference sequence of the *pks* island: accession ID AM229678 from GenBank.

Results

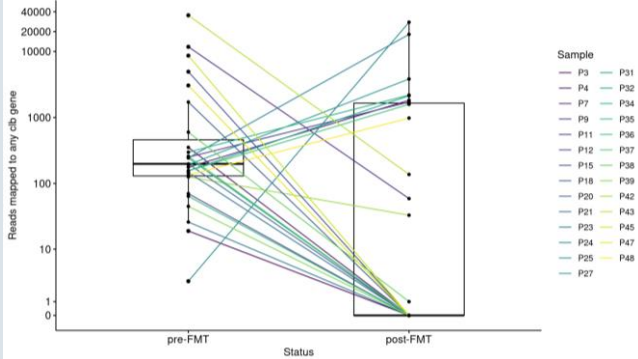


Figure 2a, patients with *pks* before FMT may lose *pks* after FMT. Twenty-seven patients were *pks*-positive before FMT, of which 13 are still *pks*-positive after FMT and 14 patients lose *pks* after FMT. All but one of the patients who lose *pks* were treated with *pks*-negative donor material. Patients whose *pks* levels decrease were mostly treated with *pks*-negative donor material and patients whose *pks* levels rise were mostly treated with *pks*-positive donor samples.

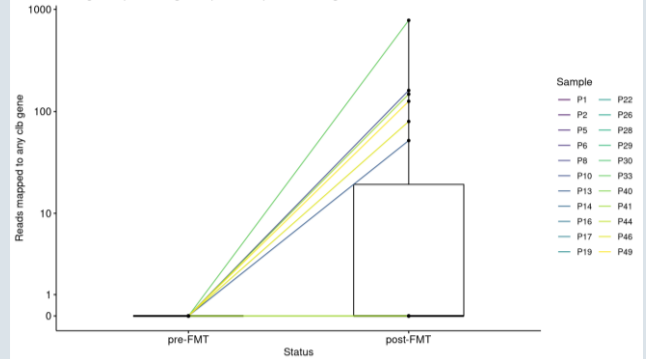


Figure 2b, patients with no *pks* before FMT may obtain *pks* after FMT, although they often remain *pks*-negative. Twenty-two patients had no *pks* prior to FMT. Sixteen of them remain negative after FMT, whereas six patients have detectable *pks* after FMT. All six were treated by negative donor samples. *Pks* values are measured as absolute number of reads matched to the reference – scales are non-linear.

Conclusions

- *Pks* is often absent in healthy donors (74% of samples)
- *Pks* is prevalent in rCDI patients (55% of patients)

- In rCDI patients, *pks* generally decreases after FMT
- Donor *pks* status is not always predictive for *pks* status in recipients after FMT