## **RESEARCH ARTICLE**

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# In vivo bioluminescence imaging of the spatial and temporal colonization of lactobacillus plantarum 423 and enterococcus mundtii ST4SA in the intestinal tract of mice

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### Abstract

**Background:** Lactic acid bacteria (LAB) are major inhabitants and part of the normal microflora of the gastrointestinal tract (GIT) of humans and animals. Despite substantial evidence supporting the beneficial properties of LAB, only a few studies have addressed the migration and colonization of probiotic bacteria in the GIT. The reason for this is mostly due to the limitations, or lack of, efficient reporter systems. Here we describe the development and application of a non-invasive in vivo bioluminescence reporter system to study, in real-time, the spatial and temporal persistence of *Lactobacillus plantarum* 423 and *Enterococcus mundtii* ST4SA in the intestinal tract of mice.

**Results:** This study reports on the application of the firefly luciferase gene (*ffluc*) from *Photinus pyralis* to develop luciferase-expressing *L. plantarum* 423 and *E. mundtii* ST4SA, using a *Lactococcus lactis* NICE system on a high copy number plasmid (pNZ8048) and strong constitutive lactate dehydrogenase gene promoters (*Pldh* and ST*ldh*). The reporter system was used for in vivo and ex vivo monitoring of both probiotic LAB strains in the GIT of mice after single and multiple oral administrations. *Enterococcus mundtii* ST4SA reached the large intestine 45 min after gavage, while *L. plantarum* 423 reached the cecum/colon after 90 min. Both strains predominantly colonized the cecum and colon after five consecutive daily administrations. *Enterococcus mundtii* ST4SA persisted in faeces at higher numbers and for more days compared to *L. plantarum* 423.

**Conclusions:** Our findings demonstrate the efficiency of a high-copy number vector, constitutive promoters and bioluminescence imaging to study the colonization and persistence of *L. plantarum* 423 and *E. mundtii* ST4SA in the murine GIT. The system allowed us to differentiate between intestinal transit times of the two strains in the digestive tract. This is the first report of bioluminescence imaging of a luciferase-expressing *E. mundtii* strain to study colonization dynamics in the murine model. The bioluminescence system developed in this study may be used to study the in vivo colonization dynamics of other probiotic LAB.

**Keywords:** Lactic acid bacteria, *Lactobacillus plantarum* 423, *Enterococcus mundtii* ST4SA, In vivo bioluminescence imaging, Luciferase, Gastrointestinal tract, Colonization, Intestinal persistence

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### **Background**

Lactic acid bacteria are common inhabitants of a healthy human and animal gastrointestinal tract and they play a major role in keeping the gut microbiota in a balanced state [1–4]. Beneficial properties include the inhibition of enteric pathogens [5], alleviation of constipation [6] and diarrhoea [7], stimulation of the immune system [8], repression of cancer cell and tumor growth [9, 10], and synthesis of essential metabolites such as vitamins [11]. For probiotics to confer their beneficial effects on the host, they must be able to survive stomach acids and bile salts and persist at high levels in the intestinal tract [12]. Some strains have adapted to these harsh conditions by over-expressing specific genes when exposed to acids and bile salts [13–15].

The survival and colonization of LAB in the GIT is usually studied in vitro and ex vivo by using models simulating the GIT [16-19]. Although these studies are valuable in understanding the survival of LAB in the GIT, the findings seldom reflect real-life conditions. More in vivo studies are needed to understand the interactions between probiotic bacteria, pathogens, commensal bacteria and gut epithelial cells. The best approach to study real-time interactions between probiotic bacteria and their mammalian host in the GIT is by labelling the cells with fluorescent or bioluminescent markers [20-26]. A selection of genes, encoding proteins that emit light at specific wavelengths, are available for cloning into plasmids or insertion into the genomes of recipient cells [21, 22]. The most commonly used luciferase labelling systems used in in vivo and ex vivo tracking of bacteria are bacterial luxABCDE from Photorabdus luminescence [26], click beetle luciferase (CBluc) from *Pyrophorus plagiophthalamus* [23] and firefly luciferase (Ffluc) from *Photinus pyralis* [25]. The CBluc and Ffluc luciferases require the exogenous addition of D-Luciferin, whereas the *lux* substrate is synthesized by proteins encoded in the *lux* operon [20]. The half-life of luciferase is only several seconds and does not represent bioluminescence accumulated over a period [28]. Another advantage is that only low levels of background luminescence are emitted by mammalian tissue.

To date, very few studies have used either fluorescence or bioluminescence whole-body imaging to monitor the persistence of LAB in the GIT, and with variable degrees of success. This is mostly due to the weak penetration of photons through muscles and tissue. Furthermore, labelled cells orally administered are often dispersed throughout the GIT or become metabolically inactive and emit bioluminescent signals too weak to detect. Cronin et al. [29] used a bacterial *lux* system to study the persistence of *Bifidobacterium breve* in mice. The bioluminescent signal was, however, not emitted from the GIT of live mice and all imaging had to be done ex vivo after dissecting the GIT. In our own studies [24], fluorescence encoded by the

mCherry gene, transformed into Lactobacillus plantarum 423 and Enterococcus mundtii ST4SA, was also only detected after surgical removal of the GIT. This is not unusual, as also reported by Oozeer et al. [26] and Corthier et al. [30] with studies done on mice. Lee and Moon [31] were one of the first to detect Lc. lactis in the GIT of live mice by using a pMG36e Ffluc plasmid vector, although the strain could only be detected for up to 2 h. Berlec and coworkers [25] successfully used the infrared fluorescent protein IRFP713 to monitor Lactococcus lactis, L. plantarum and Escherichia coli in the GIT of mice. The virulence gene expression and gut persistence abilities of two pathogenic Enterococcus faecalis strains were studied using the bacterial luxABCDE cassete [32].In another study, Daniel et al. [33] successfully monitored the colonization and persistence of Lc. lactis and L. plantarum in live mice using the CBluc luciferase system.

Lactobacillus plantarum 423, isolated from sorghum beer and *E. mundtii* ST4SA, isolated from soybeans both have probiotic properties [34–36]. The strains survive conditions in the human GIT, as shown with studies using a model simulating the intestinal conditions of infants [16]. Both strains adhere to human intestinal epithelial cells [37] and produce antimicrobial peptides [38, 39], active against *Listeria monocytogenes*, *Enterococcus faecalis*, *Clostridium sporogenes* and *Salmonella typhimurium* [34, 35, 40–42]. In a previous report, using the *mCherry* fluorescence gene, we have shown that *L. plantarum* 423 and *E. mundtii* ST4SA were localized in the cecum and colon of mice after a single oral dosage [24].

This study reports on the application of a red-shifted thermostable firefly luciferase-system (Ffluc) to study the spatial and temporal persistence of L. plantarum 423 and E. mundtii ST4SA in the GIT of mice after single and multiple dosages. The use of a red-emitting luciferase with a longer wavelength (620 nm) enabled optimal light penetration through intestinal and skin tissue. The in vitro and in vivo expression of the Ffluc system was optimized using a combination of a high-copy number plasmid vector and strong constitutive promoters. Differences between the two strains in viability and persistence in the GIT of mice were demonstrated by monitoring in vivo and ex vivo bioluminescence, using the Caliper in vivo imaging system (IVIS; Caliper Life Sciences, Hopkinton, MA). The bioluminescence system also allowed tracking of each of the strains in different sections of the GIT.

### Results

# In vitro functionality and stability of bioluminescent *L. plantarum* 423 and *E. mundtii* ST4SA

Bacterial cultures resuspended in phosphate buffered saline (PBS) were used to image the intensity of bioluminescent signals produced by *L. plantarum* 423

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Fluc and *E. mundtii* ST4SA Fluc (Fig. 1a). No significant difference was observed in the maximum intensities of bioluminescent signals produced by *L. plantarum* 423 Fluc or *E. mundtii* ST4SA Fluc. Maximum bioluminescence was recorded for *E. mundtii* ST4SA Fluc with a mean value of  $2.49 \times 10^8$  photons per second (p/s), while a slightly lower mean value of  $1.94 \times 10^8$  p/s was recorded for *L. plantarum* 423 Fluc. None of the control strains (*L. plantarum* 423 (pNZ8048) or *E. mundtii* ST4SA (pNZ8048)), emitted bioluminescent signals. No bioluminescence was detected in culture supernatants of *L. plantarum* 423 Fluc or *E. mundtii* ST4SA Fluc, indicating that bioluminescent light production was strictly intracellular.

Bioluminescence emitted by cells that expressed firefly luciferase correlated with serial dilutions of total CFUs of cultures of *L. plantarum* 423 Fluc ( $R^2 = 0.9830$ ) and *E. mundtii* ST4SA Fluc ( $R^2 = 0.9870$ ), indicating that photon emission accurately reflects bacterial cell numbers (Fig. 1b and c). The bioluminescence signal corresponded to the detection of bacterial CFU over a broad range, from approximately  $6 \times 10^6$  CFU to  $8.5 \times 10^8$  CFU for *L. plantarum* 423 Fluc (Fig. 1b) and from approximately  $1 \times 10^7$  CFU to  $2 \times 10^9$  CFU for *E. mundtii* ST4SA Fluc (Fig. 1c).

No significant difference in growth was observed between wild-types (WT) L. plantarum 423 and E. mundtii ST4SA and recombinants L. plantarum 423 Fluc and E. mundtii ST4SA Fluc, respectively, after 9 h of growth (not shown). Bioluminescent light production and the presence of the luciferase expressing plasmids in recombinant L. plantarum 423 Fluc and E. mundtii ST4SA Fluc strains had no detectable effect on bacterial growth. The stability of the luciferase-expressing plasmids pNZPldhFfluc in L. plantarum 423 Fluc and pNZSTldhFfluc in E. mundtii ST4SA Fluc was tested in vitro by subculturing for up to 7 days with replica-plating on non-selective and selective media (Additional file 1: Figure S1a). The stability of the autonomous plasmids in L. plantarum 423 Fluc and E. mundtii ST4SA Fluc transformants was indicated by 100% plasmid retention and retained resistance to Cm following culturing for 7 days in the absence of the antibiotic. The bioluminescent signals of the recombinant strains were also imaged with the IVIS in parallel to replica plating (Additional file 1: Figure S1b and c).

# Colonization dynamics of bioluminescent *L. plantarum* 423 and *E. mundtii* ST4SA in the GIT of mice after a single dosage

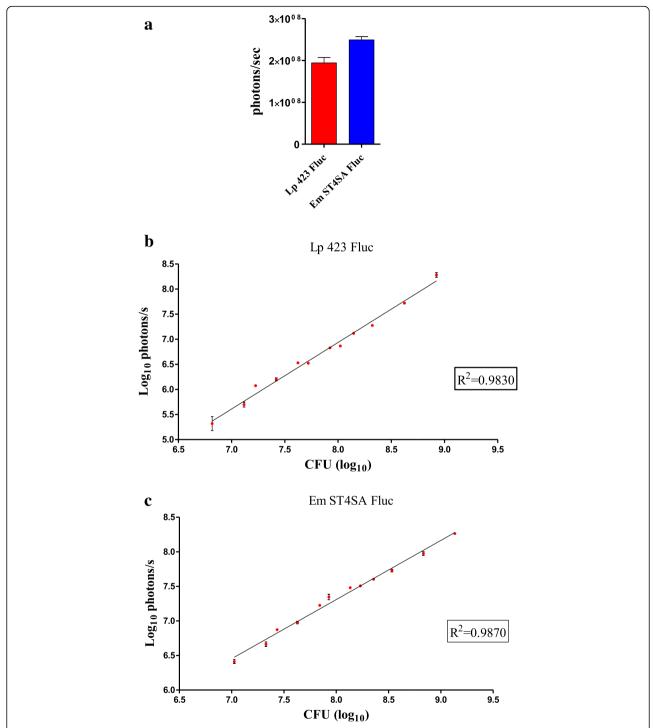
To determine the spatial and temporal colonization of L. plantarum 423 Fluc and E. mundtii ST4SA Fluc after a single oral administration, groups of mice (n = 13, per strain) were monitored over a 24 h period by transcutaneous in vivo bioluminescence imaging (BLI) and ex vivo BLI of GITs and faeces. The viable bacteria numbers in the GIT and faecal samples were also recorded. Three

anesthetised mice (n = 3, per strain) were imaged at 15 and 30 min and, 1, 1.5, 2, 3, 4, 6 and 24 h after the administration of L. plantarum 423 Fluc and E. mundtii ST4SA Fluc, respectively (Fig. 2). The same three mice were used throughout the 24 h trial period. At time zero (before administration) no bioluminescence was recorded (background signal corresponded to approximately  $3 \times 10^4$  p/s). A maximum bioluminescent signal of approximately 2 × 10<sup>8</sup> p/s was detected for *L. plantarum* 423 Fluc and *E.* mundtii ST4SA Fluc at 1 h and 30 min, respectively. The bioluminescent signal of L. plantarum 423 Fluc remained at high levels until 2 h, but declined to lower levels 1 h later (mean value of approximately  $9 \times 10^6$  p/s). The bioluminescent signal of L. plantarum 423 Fluc remained at a plateau (mean value of approximately  $4 \times 10^5$  p/s) until 6 h. The bioluminescent signal of E. mundtii ST4SA Fluc steadily declined between 1 and 6 h and was significantly higher than that observed for L. plantarum 423 Fluc between 3 and 4 h. After 24 h, no bioluminescent signal could be detected for L. plantarum 423 Fluc (background level), whereas the signal of E. mundtii ST4SA Fluc declined to approximately  $2 \times 10^5$  p/s.

Next, the localization of bioluminescent L. plantarum 423 Fluc and E. mundtii ST4SA Fluc in the GITs of mice after oral administration was determined by ex vivo imaging and recording viable cell numbers in the small and large intestinal tracts (Fig. 3). Results showed that 15 min after administration of bacterial strains, both L. plantarum 423 Fluc and E. mundtii ST4SA Fluc survived passage through the stomach by the observation of high cell numbers and bioluminescent cells throughout the small intestine (Fig. 3a and b). After approximately 45 min, bioluminescent cells of *E. mundtii* ST4SA Fluc reached the cecum and colon. From 90 to 240 min after oral administration of bacteria to mice, the majority of bioluminescent L. plantarum 423 Fluc and E. mundtii ST4SA Fluc had travelled through the small intestine and were located exclusively in the cecum and colon. Some of the viable cells of both strains remained in the small intestine after 90 min, but emitted weak or no bioluminescence signals (Fig. 3b). No bioluminescence of either strain was detected in the large intestine 15 min (Fig. 3a) after intragastric administration, suggesting that the viable cells' bioluminescence emission was below the detection limit of the IVIS through the intestinal tissue or the cells were metabolically inactive (Fig. 3c). After 24 h, a significantly higher number of viable E. mundtii ST4SA Fluc was detected in the cecum/colon compared to L. plantarum 423 Fluc.

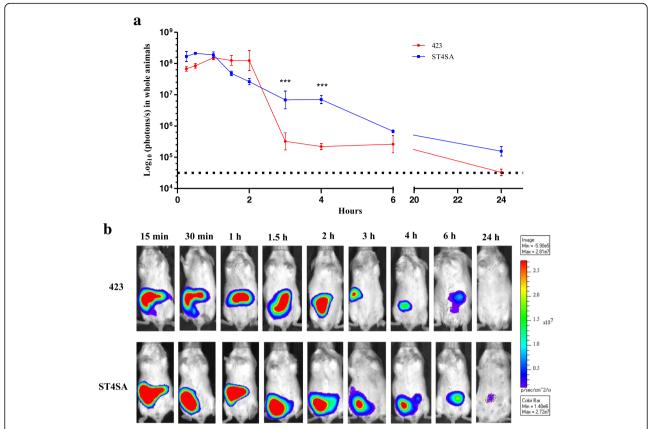
Colonization of *L. plantarum* 423 Fluc and *E. mundtii* ST4SA Fluc in the GIT of mice was also determined by monitoring the number of viable bacterial cells in faeces at different time points after intragastric administration (Fig. 4a). The respective bioluminescent signals of *L.* 

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**Fig. 1** Quantification of bioluminescent *L. plantarum* 423 Fluc and *E. mundtii* ST4SA Fluc. **a** Bioluminescence measured in cultures of *L. plantarum* 423 Fluc and *E. mundtii* ST4SA Fluc distributed in black 96-well microtitre plates. Means from six independent cultures (4 × 10<sup>9</sup> CFU per culture) are shown with standard deviations, and the background from each strain has been subtracted from each respective measurement. Correlation between bioluminescent signals and bacterial cell numbers of **(b)** *L. plantarum* 423 Fluc and **(c)** *E. mundtii* ST4SA Fluc. Cultures of each strain were serially diluted in black microplates and the bioluminescent signals quantified using the IVIS and then correlated with CFUs. Log<sub>10</sub> averages of three cultures are plotted, with error bars indicating standard deviations. The logarithmic trendline and the correlation of determination (R<sup>2</sup>) between bioluminescence measurements and bacterial numbers of each strain are shown

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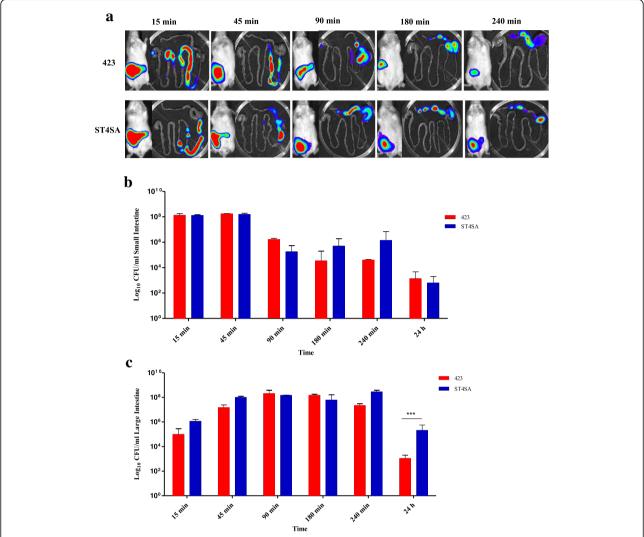
**Fig. 2** Monitoring of *L. plantarum* 423 and *E. mundtii* ST4SA colonization in the digestive tract of mice by bioluminescence imaging in whole animals after one oral administration. *Lactobacillus plantarum* 423 Fluc and *E. mundtii* ST4SA Fluc were fed intragastrically ( $4 \times 10^9$  CFU) to two sets of three mice (n = 3, per strain). The bioluminescent signals in  $\log_{10}$  photons/s measured from whole animals at different time-points over a 24 h period (**a**) are plotted, with standard deviations. Significant statistical differences between the bioluminescence signals of the two groups of mice are indicated with three asterisks (P < 0.001); Mann-Whitney nonparametric test). The background bioluminescence signal (approximately  $4 \times 10^4$  p/s) emitted is represented by a dashed line. **b** Visual images of bioluminescence emission in whole animals by mice fed once with *L. plantarum* 423 Fluc or *E. mundtii* ST4SA Fluc. In each case, one representative image of one mouse is shown. The intensity of the photon emission is represented as a pseudo-color image. One representative scale bar is shown (p/s)

plantarum 423 Fluc and E. mundtii ST4SA Fluc in faeces were also monitored with the IVIS (Fig. 4b). High cell numbers of both strains were excreted in the faeces and were proportionate to the respective bioluminescence signals emitted. The bacterial populations of both strains in faeces increased with time. Both strains reached a maximum number of approximately  $2 \times 10^8$  CFU/100 mg faeces after 4 h and remained at this level for the following 2 h. These peaks correlated excellently with maximum bioluminescent signals of approximately  $4 \times 10^6$ 100 mg of faeces for *L. plantarum* 423 Fluc and  $2 \times 10^7$  p/ s/100 mg of faeces for E. mundtii ST4SA Fluc, from 4 to 6 h. The maximum level of viable E. mundtii ST4SA Fluc cells shed in the faeces was significantly higher in the first 2 h, reaching approximately  $8 \times 10^7$  p/s/100 mg of faeces. Bioluminescence signals emitted by E. mundtii ST4SA Fluc cells in the faeces were higher compared to those of L. plantarum 423 Fluc throughout the 24 h study period. After 24 h, the number of L. plantarum 423 Fluc declined to approximately  $4\times10^4$  CFU/100 mg faeces with no bioluminescent signal (background), while *E. mundtii* ST4SA Fluc cells declined to approximately  $1\times10^5$  CFU/100 mg faeces with a weak bioluminescent signal of  $1.5\times10^4$  p/s/100 mg of faeces.

# Persistence of bioluminescent *L. plantarum* 423 and *E. mundtii* ST4SA in the GIT of mice after five oral dosages

In vivo bioluminescent signals were measured everyday (1 h after administration of bacteria to mice on days 1 to 5) for nine days after two groups of mice (n = 22, per strain) each received daily doses of L. plantarum 423 Fluc and E. mundtii ST4SA Fluc for five consecutive days (Fig. 5). The experimental design is described in Fig. 5a. High intensity bioluminescence was emitted by mice in both groups from days 1 to 5. Highest bioluminescent signals were recorded for E. mundtii ST4SA Fluc (mean =  $4.27 \times 10^8$  p/s) compared to E. plantarum 423 Fluc (mean = E. E vivial the plantarum 423 Fluc (mean = E vivial the plantarum 424 Fluc (mean = E vivial the plantarum 425 Fluc (mean = E vivial the plantarum 425 Fluc (mean = E vivial the plantarum 426 Fluc (mean = E vivial the plantarum 427 Fluc (mean = E vivial the plantarum 428 Fluc (mean = E vivial the plantarum 429 Fluc (mean = E vivial the plantarum 429 Fluc (mean = E vivial the plantarum 429 Fluc (mean = E vivial the plantarum 420 Fluc (mean

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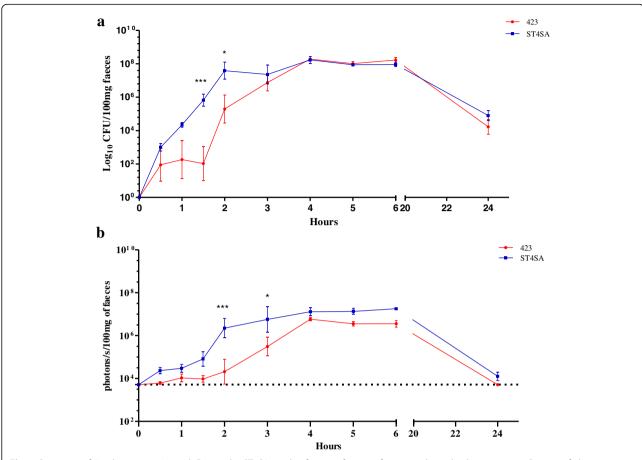
**Fig. 3** Transit of *L. plantarum* 423 and *E. mundtii* ST4SA through the digestive tract of mice after one oral administration. Groups of mice were gavaged once with  $4 \times 10^9$  CFU of *L. plantarum* 423 Fluc or *E. mundtii* ST4SA Fluc and the intestines resected at 15, 45, 90, 180, 240 min and 24 h. At each time point two mice (n = 12, per strain) were sacrificed, and (**a**) one representative image of one mouse and its GIT are shown. Persistence of viable *L. plantarum* 423 Fluc or *E. mundtii* ST4SA Fluc cells in (**b**) the small and (**c**) large intestinal tract of mice sacrificed at time points indicated in A. The limit of detection was approximately  $4 \times 10^4$  p/s. Significant differences between the two groups were assessed using the Mann-Whitney nonparametric test are indicated with asterisks (P < 0.001)

first 5 days. After 6 days the signal for *L. plantarum* 423 Fluc rapidly declined to approximately  $1\times 10^5$  p/s, while the signal for *E. mundtii* ST4SA Fluc declined to approximately  $5\times 10^5$  p/s. From day 6, the *L. plantarum* 423 Fluc bioluminescent signal remained at a plateau until day 9 (approximately  $1\times 10^5$  p/s). The *E. mundtii* ST4SA Fluc bioluminescent signal declined to approximately  $2\times 10^5$  p/s on day 7 and to approximately  $8\times 10^4$  p/s after 9 days.

The intestines of mice were imaged ex vivo to determine which sections of the GIT the bacterial strains colonized after 5 oral administrations (Fig. 6). On day 1, both strains were detected throughout the small intestine 30 min after oral administration of

bacteria. From day 2 to day 5, both strains were detected in the jejunum and ileum sections of the small intestine (jejunum and ileum) and in both the cecum and colon sections of the large intestinal tract. At days 6 and 7 (1 and 2 days after last bacterial dose), *L. plantarum* 423 Fluc colonized the small intestine and cecum readily, but the bioluminescent signal detected from the small intestine progressively declined and by day 9 a very weak or no bioluminescent signal remained (Fig. 6a). On day 9, *L. plantarum* 423 Fluc was predominantly localized in the cecum and colon. From day 6 to day 9, *E. mundtii* ST4SA Fluc colonized the upper section of the small intestine and the cecum/colon most prominently (Fig. 6b).

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**Fig. 4** Presence of *L. plantarum* 423 and *E. mundtii* ST4SA in the faeces of mice after a single oral administration. Groups of three mice each (n = 3, per strain) were administered once with  $4 \times 10^9$  CFU *L. plantarum* 423 Fluc or *E. mundtii* ST4SA Fluc. At each time point,  $\log_{10}$  averages of the **(a)** cell counts per 100 mg faeces and the corresponding **(b)** bioluminescence in  $\log_{10}$  p/s per 100 mg faeces for each group of three mice are plotted with standard deviations. Significant differences between the two groups were assessed using the Mann-Whitney nonparametric test and are indicated with one (P < 0.05) or three (P < 0.001) asterisks. The background bioluminescence signal (approximately  $5 \times 10^3$  p/s) emitted is represented by a dashed line

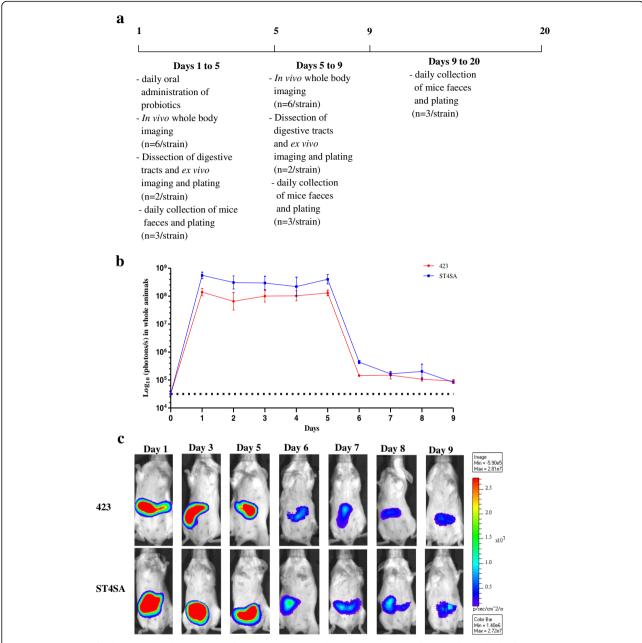
Viable counts of the small and large intestine taken on days 1 to 5 revealed high numbers (approximately 108 CFU) of both L. plantarum 423 Fluc and E. mundtii ST4SA Fluc (Fig. 6c and d). At day 7 (2 days after last bacterial dose), the colonization of the large intestine (mean value of  $3 \times 10^5$  CFU) by L. plantarum 423 Fluc was 10-fold superior to that of the small intestine (mean value of  $3 \times 10^6$  CFU). Two days later, the difference between the amount of viable L. plantarum 423 Fluc cells in the small (mean value of  $9 \times 10^2$  CFU) and large intestine (mean value of  $8 \times 10^4$  CFU) increased to 100-fold. In the case of E. mundtii ST4SA Fluc, the number of viable cells in the small intestine increased from approximately  $1 \times 10^3$  CFU on day 6 to  $2 \times$ 10<sup>4</sup> CFU on day 9. On day 9, mice administered with L. plantarum 423 Fluc harboured approximately 9 × 10<sup>4</sup> CFU in the cecum/colon, while mice administered E. mundtii ST4SA Fluc harboured approximately 7 × 10<sup>4</sup> CFU in the cecum/colon. After 20 days, no L.

*plantarum* 423 Fluc were detected in the intestines of mice. However, the number of *E. mundtii* ST4SA Fluc cells in the small intestine was still approximately  $1.4 \times 10^3$  CFU and  $5 \times 10^3$  CFU in the large intestine.

# Persistence of *L. plantarum* 423 and *E. mundtii* ST4SA in faeces after five oral dosages

The persistence of bioluminescent *L. plantarum* 423 Fluc and *E. mundtii* ST4SA Fluc in faeces and their respective bioluminescent signals were monitored every day for 20 consecutive days (Fig. 7a and b). *Enterococcus mundtii* ST4SA Fluc persisted in the faeces throughout the trial, with significantly higher cell numbers recorded at days 4, 19 and 20 compared to *L. plantarum* 423 Fluc. The maximum number of viable *E. mundtii* ST4SA Fluc cells in faeces was excreted at day 4 (approximately  $2 \times 10^8$  CFU/100 mg faeces). From day 4 to 9 the level of *E. mundtii* ST4SA Fluc cells in faeces declined to approximately  $1 \times 10^4$  CFU/100 mg faeces. After day 9 there is a

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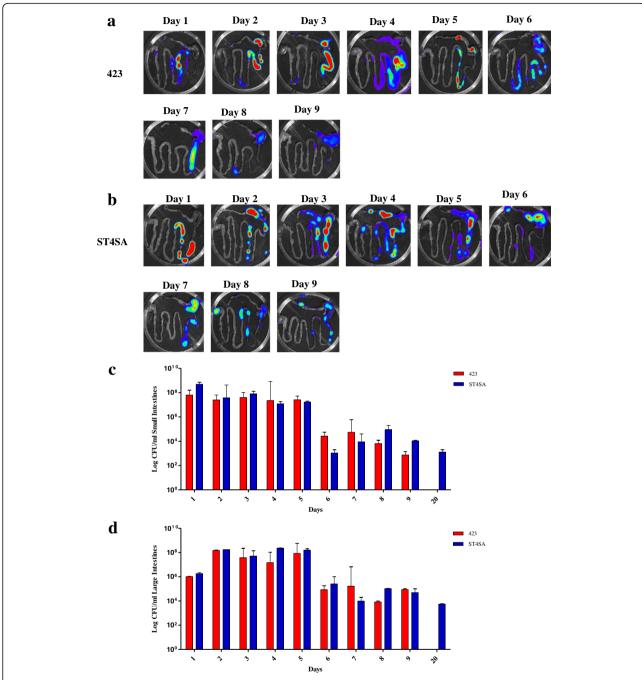
**Fig. 5** Monitoring of colonization and persistence of *L. plantarum* 423 and *E. mundtii* ST4SA in the GIT of mice by bioluminescence imaging in whole animals after five daily administrations. **a** The experimental design. *Lactobacillus plantarum* 423 Fluc and *E. mundtii* ST4SA Fluc were fed once daily by intragastric gavage  $(4 \times 10^9 \text{ CFU})$  to two groups of mice (n = 22, per strain) for five consecutive days (days 1 to 5). The bioluminescent signals in  $\log_{10}$  photons/s measured in whole animals from day 1 to 9 for each set of three mice (**b**) are plotted, with standard deviations. No statistical differences between the bioluminescence signals of the two groups of mice were observed. The background bioluminescence signal (approximately  $4 \times 10^4$  p/s) emitted is represented by a dashed line. **c** Visual images of bioluminescence emission in whole animals by mice fed once daily for five consecutive days with *L. plantarum* 423 Fluc or *E. mundtii* ST4SA Fluc. One representative image of one mouse is shown. The intensity of the photon emission is represented as a pseudo-color image. One representative scale bar is shown (p/s)

slight increase in the number of *E. mundtii* ST4SA Fluc cells in faeces until day 11 after which levels of approximately  $10^4$  to  $10^5$  CFU/100 mg faeces were maintained from days 12 to 20. *Lactobacillus plantarum* 423 Fluc, on the other hand, reached a maximum level of approximately  $2 \times 10^8$  CFU/100 mg faeces at day 1 and then

steadily declined until day 19 (mean value of  $1 \times 10^4$  CFU/100 mg faeces). *Lactobacillus plantarum* 423 Fluc persisted for only 13 days in faeces after the last day of intragastric administration of bacteria (day 5).

The bioluminescent signal of *E. mundtii* ST4SA Fluc cells in faeces was detected throughout the trial period,

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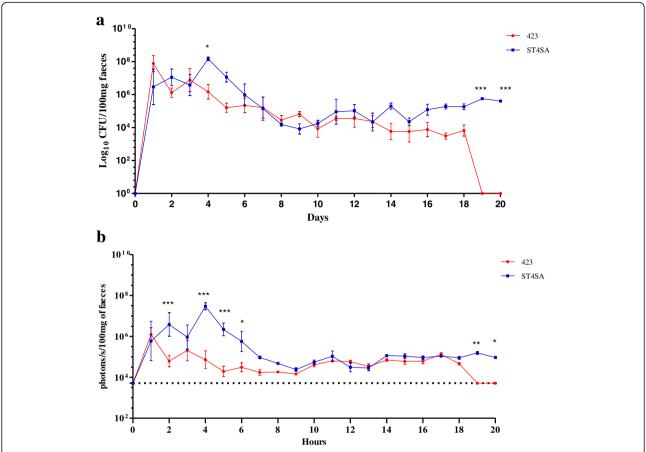


**Fig. 6** Comparison of colonization abilities of *L. plantarum* 423 and *E. mundtii* ST4SA in the GIT of mice after five daily oral administrations. *Lactobacillus plantarum* 423 Fluc and *E. mundtii* ST4SA Fluc were fed once daily by oral gavage  $(4 \times 10^9 \text{ CFU})$  to two groups of mice (n = 18, per strain) for five consecutive days (days 1 to 5). Four mice (two per group) were sacrificed from day 1 to 9, and a representative image of the GIT of one mouse is shown (days 1 to 9) in mice fed with **(a)** *L. plantarum* 423 Fluc or **(b)** *E. mundtii* ST4SA Fluc. Persistence of viable *L. plantarum* 423 Fluc or *E. mundtii* ST4SA Fluc cells in **(c)** the small and **(d)** large intestinal tract of mice sacrificed at time points indicated in A and B and day 20. The limit of detection was approximately  $4 \times 10^4 \text{ p/s}$ 

while the *L. plantarum* 423 Fluc bioluminescent signal declined to the background level at day 19. The peaks of the amount of viable cells per 100 mg faeces of both strains correlated perfectly with the amount of bioluminescent signals emitted at different time points.

The bioluminescent signal of *L. plantarum* 423 Fluc was detected at lower levels between days 2 and 6, but displayed levels similar to those observed for *E. mundtii* ST4SA Fluc from day 7 until 18. The bioluminescent system allowed the detection of both *L. plantarum* 423

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**Fig. 7** Persistence of *L. plantarum* 423 and *E. mundtii* ST4SA in mouse faeces after five daily oral administrations. Groups of three mice (n = 3, per strain) each were administered once daily (days 1 to 5) with  $4 \times 10^9$  CFU *L. plantarum* 423 Fluc or *E. mundtii* ST4SA Fluc for five consecutive days. Mouse faeces were collected daily from days 1 to 20. At each time point,  $\log_{10}$  averages of the **a** cell counts per 100 mg faeces and the corresponding **b** bioluminescence in  $\log_{10}$  p/s per 100 mg faeces for each group of three mice are plotted with standard deviations. Significant differences between the two groups were assessed using the Mann-Whitney nonparametric test and are indicated with one (P < 0.05), two (P < 0.01) or three (P < 0.001) asterisks. The background bioluminescence signal (approximately  $5 \times 10^3$  p/s) emitted is represented by a dashed line

Fluc and *E. mundtii* ST4SA Fluc viable cells in faeces as low as  $10^4$  CFU/100 mg faeces.

### Discussion

A major advantage of using the BLI technique compared to conventional approaches is that it allows for drastic reductions in the number of animals to be sacrificed to establish the precise location of bacteria in mouse or rat models [20]. Moreover, more information is gathered over a shorter period per experiment compared to traditional pre-clinical animal trials [21]. Mouse models provide a complex whole body system for the non-invasive real-time monitoring of bioluminescent probiotic LAB through the GIT of the mammalian host [43]. The murine model is the predominant choice for the in vivo evaluation of probiotic properties and has been used to study the persistence and localization of potential probiotic LAB in several studies [21, 24, 27, 33, 43]. In the current study, an

IVIS system and the BLI technique were used to study the colonization dynamics of the probiotic LAB strains *L. plantarum* 423 and *E. mundtii* ST4SA in the GIT of mice.

The stable expression of reporter genes during in vivo imaging is an absolute prerequisite for the detection of luminescent light through animal tissues. Stable expression of reporter genes depends on the nature of the expression vector used, including the promoters used to drive expression and plasmid copy numbers. With this in mind, expression of the firefly luciferase gene from P. pyralis (ffluc) was optimized for use in L. plantarum 423 and E. mundtii ST4SA. Results in this study demonstrate that highest bioluminescence signals were achieved using the pNZ8048 high-copy number plasmid [44] for luciferase gene expression (Fig. 1). Attempts to use a low-copy number plasmid and chromosomally integrated ffluc genes in L. plantarum 423 and E. mundtii ST4SA resulted only in the emission of weak and inconsistent bioluminescent signals (not Van Zyl et al. BMC Microbiology (2018) 18:171 Page 11 of 16

shown). The former results are most likely linked to the presence of multiple copies of the pNZ8048 plasmid that in turn leads to higher expression levels of the ffluc gene within the respective LAB hosts. The expression of reporter genes in bacteria using plasmids is widely used. Bacterial plasmids play an important role in the ability of bacteria to adapt in diverse environments [45]. However, when introducing a new recombinant plasmid harbouring a reporter gene, it is critical to evaluate its stability and persistence within the bacterial host with and without antibiotic pressure. This study demonstrates that in vitro ffluc expression plasmids were remarkably stable in L. plantarum 423 Fluc and E. mundtii ST4SA Fluc and that bioluminescence did not affect the growth of the respective host strains compared to the WT derivatives (see Additional file 1). We have previously demonstrated plasmid stability in E. mundtii ST4SA [24]. In terms of promoter selection, the lactate dehydrogenase gene (ldh) promoters of L. plantarum 423 and E. mundtii ST4SA, respectively, drove highest ffluc gene expression. The L. plantarum ldh gene promoter has been used for the constitutive expression of genes in LAB in several studies [24, 33, 46, 47]. These results demonstrated that both *L. plan*tarum 423 Fluc and E. mundtii ST4SA Fluc could produce bioluminescence with no loss of the firefly luciferase-expressing plasmids.

The current study was designed to study the colonization (spatial and temporal) dynamics of L. plantarum 423 and E. mundtii ST4SA orally inoculated once or for five consecutive days over a period of 24 h and 20 days, respectively. Strong bioluminescence signals emitted by L. plantarum 423 Fluc and E. mundtii ST4SA Fluc could be detected as soon as 15 min after oral administration with a single bacterial dosage of the respective strains (Fig. 2). The bioluminescence signals of both strains as recorded with whole body imaging showed an overall decline after a single oral administration to mice. This is an indication that the probiotic strains only transiently colonized the GIT of mice after a single oral administration. In a previous report, using the mCherry fluorescence gene and a conventional culture-based method, we showed similar transit dynamics of viable L. plantarum 423 and E. mundtii ST4SA after a single dose of each bacterial strain [24]. However, we could not obtain fluorescence signals from in vivo whole body imaging for real-time analysis and could only locate the bacterial strains in the GIT after killing the animals. Interestingly, in the current study bioluminescence signals were detected up to 6 h in whole animals fed with either L. plantarum 423 or E. mundtii ST4SA and up to 24 h in mice fed with E. mundtii ST4SA. Eom et al. [27] could not detect a bioluminescence signal after 3 h from mice administered bioluminescent L. casei CJNU 0588. Based on whole body bioluminescence imaging in real-time, these results clearly reflect the superior active replication and colonization abilities of *L. plantarum* 423 and *E. mundtii* ST4SA in the mouse GIT compared to *L. casei* CINU 0588.

The bacterial transit time in the GIT for viable L. plantarum 423 and E. mundtii ST4SA varied. Lactobacillus plantarum 423 showed a slower transit and was detected in the lower part of the jejunum and in the ileum 45 min post gavage, while E. mundtii ST4SA was detected in the ileum, cecum and colon (Fig. 3a). Similarly, Karimi et al. [48] demonstrated that the gastrointestinal transit time differs between strains of L. reuteri, using fluorescence and bioluminescence imaging techniques. However, the authors did not selectively monitor the bacterial cell numbers of the LAB strains present in gastrointestinal tissue or stool samples. Although it was shown in the current study that there is excellent correlation between bioluminescence and bacterial load, it is important to verify that a certain level of bioluminescence correlates to a certain number of cells. Enumeration of viable counts in dissected intestines revealed that both bioluminescent LAB strains were located predominantly in the large intestine (cecum/colon) 3 h post gavage (Fig. 3b). No bioluminescent cells of L. plantarum 423 or E. mundtii ST4SA were detected in the stomach or duodenum compartments 45 min after intragastric administration of the respective bacteria (Fig.3a). These results are indicative of the harsh conditions the LAB strains are exposed to in those compartments of the GIT and may be the reason for the rapid transit in those sections of the gut. The CFU counts revealed the presence of viable cells of L. plantarum 423 and E. mundtii ST4SA in the small intestine at 4 and 24 h post gavage, but no bioluminescent signals could be detected. This might be caused by reduced cell activity or the inhibition of protein synthesis during passage in the stomach and duodenum compartments of the gut, which leads to inefficient bioluminescence emission. Bioluminescent light can only be produced in metabolically active cells [20]. Both strains could be detected in the cecum and colon after 4 h, indicating the metabolically active state of the bacteria in the large intestine (Fig. 3a).

After 24 h, no bioluminescent signal could be detected from mice administered with *L. plantarum* 423, while a detectable bioluminescent signal was still observed from mice fed with *E. mundtii* ST4SA (Fig. 2). This indicates that *L. plantarum* 423 was eliminated from the GIT of mice more rapidly than *E. mundtii* ST4SA, and is associated with a significantly higher amount of viable *E. mundtii* ST4SA in the large intestine compared to *L. plantarum* 423 (Fig. 3c). These findings are in agreement with previous studies [33, 48] reporting on the gastrointestinal transit of *Lactobacillus reuteri*, *L. plantarum* and *Lactococcus lactis*.

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The authors showed that the intestinal transit times differed between two strains of *L. reuteri*, and that a *Lc. lactis* strain had shorter survival times in the GIT compared to a *L. plantarum* strain. *Lactobacillus plantarum* 423 and *E. mundtii* ST4SA were excreted in high numbers in the faeces of mice over the 24 h period, but it is clear that a small amount of each LAB strain persisted in the GIT after a single oral dose as demonstrated with intestinal tissue CFU counts (Figs. 3 and 4). Bioluminescence signals from bacteria in faecal material accurately reflected CFU data and this serves as an indication that the plasmids are retained in vivo to a large extent.

To study more thoroughly the persistence of the bioluminescent strains in the GIT, mice were administered orally with either L. plantarum 423 or E. mundtii ST4SA for five consecutive days. Our results demonstrate that L. plantarum 423 and E. mundtii ST4SA had similar GIT transit dynamics during the first 5 days of administration of the respective strains to mice, despite the detection of higher bioluminescent signals for E. mundtii ST4SA compared to L. plantarum 423 (Fig. 5). At day 6 (one day after last bacterial dosage), the in vivo bioluminescent signals of both strains declined to low levels but were maintained at similar levels for the next 3 days. This suggests that while most of the administered bacteria transited the GIT of mice, small populations of both strains persisted until day 9 (4 days after last bacterial dosage). Overall, the gastrointestinal persistence of L. plantarum 423 and E. mundtii ST4SA compared well to several other commercial probiotic strains, including Lactobacillus rhamnosus GG, Bifidobacterium lactis LAFTI B94, L. plantarum 299v and Lactobacillus gasseri SBT2055 [49-52]. These observations were confirmed by ex vivo imaging of dissected intestines (Fig. 6a and b). At day 9, bioluminescent cells of both strains were predominantly localized in the cecum and colon. It has been suggested that the murine cecum may be the site where microorganisms adapt to the gastrointestinal environment and where the activation of genes required for colonization of the colon occur [29]. Interestingly, the cecum and colon have also been shown to be the major sites of colonization of several enteric pathogens including E. coli O157:H7, Citrobacter rodentium, Yersinia enterocolitica, L. monocytogenes and S. typhimurium [42, 53-56]. Both L. plantarum 423 and E. mundtii ST4SA have been demonstrated to exclude pathogens such as L. monocytogenes, S. typhimurium, C. sporogenes and E. faecalis in in vitro or in vivo competitive exclusion experiments [40–42]. The presence of persistent populations of L. plantarum 423 and E. mundtii ST4SA in the murine cecum/colon as demonstrated in this study suggest that their presence may have prevented the pathogenic bacteria from becoming established. It is also interesting to note that the population of E. mundtii ST4SA in the small intestine increased until day 9 compared to that of L. plantarum 423 (Fig. 6c). This could explain why mice pre-colonized with E. mundtii ST4SA showed a more rapid decline in L. monocytogenes EGDe cell numbers compared L. plantarum 423 in a competitive exclusion experiment [42]. Enterococcus mundtii ST4SA cells were able to persist in the faeces of mice throughout the trial period (Fig. 7). In contrast, L. plantarum 423 could not be detected in faeces after 13 days after the last oral administration to mice and in lower numbers compared to E. mundtii ST4SA. Based on bioluminescence, the amount of bacteria administered to mice per day  $(4 \times 10^9 \text{ CFU})$ and the amount of E. mundtii ST4SA cells shed in the faeces per day, there is a clear indication that E. mundtii ST4SA persists better than L. plantarum 423 in the murine GIT of mice. Since the same dosage and administration methods were used for both strains, the difference in the intestinal persistence between the two probiotic strains might be due to differences in the physiological and genotypic properties of the strains.

### **Conclusions**

The construction and optimization of LAB reporter strains is an important step towards a better understanding of the route and destination of orally administered probiotics in the GIT, and the interactions between probiotics and the host. This study demonstrates the application of the firefly luciferase system to compare the colonization dynamics of L. plantarum 423 and E. mundtii ST4SA in mice. The in vivo BLI system revealed the precise location of the bacterial strains within the murine GIT after single or multiple doses. Both strains prominently colonized the cecum and colon. Enterococcus mundtii ST4SA persisted in the GIT and faeces of mice throughout the trial period and also actively colonized the small intestine. This is the first report of bioluminescence in vivo imaging of E. mundtii ST4SA in a mouse model. The bioluminescence system developed here has the potential to allow the study of in vivo colonization dynamics of other important probiotic LAB species.

### Methods

# Bacterial strains, plasmid construction and culture conditions

Escherichia coli MC1061 (Mobitech) was used as a cloning host for construction of pNZ8048-derived bioluminescence expression vectors and was cultured aerobically at 37 °C in Luria-Bertani (LB) broth, or brain heart infusion (BHI) and streaked onto the same media, supplemented with 1.5% (w/v) agar (all from Biolab Diagnostics, Midrand, South Africa). Lactobacillus plantarum 423 and E. mundtii ST4SA were grown without shaking at 30 °C in MRS broth and streaked onto MRS agar (both from Biolab Diagnostics). Where appropriate, Cm was added at

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10 μg/ml to growth media of *E. coli* MC1061 and *L. plantarum* 423 and 5 μg/ml to media of *E. mundtii* ST4SA.

Lactobacillus plantarum 423 and E. mundtii ST4SA were labelled by transformation with plasmids encoding the red-shifted thermostable firefly luciferase gene from P. pyralis (ffluc) [57]. The bioluminescence expression vectors are based on the pNZ8048 Lc. lactis NICE system high copy number plasmid (Mobitech, Goettingen, Germany). The vector contains the cat gene for chloramphenicol (Cm) resistance, the nisA gene promoter region (PnisA), a multiple cloning site (MCS), replication genes (repC and repA) for replication in LAB/E. coli and the termination (T) sequence of the *Lc. lactis pepN* gene [44]. Primers used for PCR amplification are listed in Additional file 2: Table S1 and were from Ingaba Biotechnical Industries (Pretoria, South Africa). DNA restriction enzymes and PCR polymerase were from New England Biolabs (NEB, Ipswich, MA, USA). The construction of pNZPldhFfluc and pNZSTldhFfluc luciferase expression vectors is shown in Additional file 3: Figure S2. Plasmid pNZPldhFfluc carried the ffluc gene under control of the strong constitutive L. plantarum 423 lactate dehydrogenase gene promoter (Pldh). In the pNZSTldhFfluc construct, the ffluc gene was cloned under control of the strong constitutive E. mundtii ST4SA lactate dehydrogenase gene promoter (STldh). The Pldh and STldh promoters were amplified from L. plantarum 423 and E. mundtii ST4SA genomic DNA, using primer pairs Pldh1/ Pldh2 and ldhS1/ldhS2, respectively. The ffluc bioluminescence gene was amplified from plasmid pMV306G13 + FflucRT using primers FlucFor and FlucRev. Briefly, the Pldh (520 bp), STldh (166 bp) and ffluc (1.6 kb) PCR fragments were cloned into pNZ8048 after digestion of Pldh and STldh with BglII/NcoI, digestion of ffluc with NcoI/ XbaI and digestion of pNZ8048 with BglII/XbaI (resulting in the removal of the PnisA promoter), yielding plasmids pNZPldhFfluc and pNZSTldhFfluc, respectively.

The two bioluminescence expression vectors were introduced into *L. plantarum* 423 and *E. mundtii* ST4SA by electro-transformation as described by Van Zyl et al. [24] and were named *L. plantarum* 423 Fluc and *E. mundtii* ST4SA Fluc. *Lactobacillus plantarum* 423 and *E. mundtii* ST4SA containing the empty pNZ8048 vector were used as controls and were labelled *L. plantarum* 423 (pNZ8048) and *E. mundtii* ST4SA (pNZ8048). Plasmid stability in *L. plantarum* 423 Fluc and *E. mundtii* ST4SA Fluc, and growth comparison between WT and recombinant strains was tested by standard methodology as described previously [24].

# Correlation between in vitro bioluminescence measurements and viable cell numbers

Lactobacillus plantarum 423 Fluc and E. mundtii ST4SA Fluc were grown for 12 h at 30 °C in MRS broth,

supplemented with Cm as mentioned elsewhere. From these cultures, 1 ml was inoculated into freshly prepared MRS broth and incubated at 30 °C to an optical density (OD<sub>550nm</sub>) of 2.5 (for *L. plantarum* 423) and 2.3 (for *E.* mundtii ST4SA). Viable cell numbers were determined by plating onto MRS agar containing Cm. The bacterial suspensions were harvested (3 min at 8000 x g), washed twice with sterile PBS, resuspended in gavage buffer (0.2 M NaHCO<sub>3</sub> with 1%, w/v, glucose, pH 8.0) and serially diluted to 1/128 in the same buffer. Two-hundred microliters of each dilution was added in triplicate to black 96-well microtitre plates and bioluminescence measured after the addition of 5 µl of D-Luciferin potassium salt (Anatech Instruments, Bellville, South Africa) at 470 µM. Bioluminescent readings were recorded using the IVIS and the photons emitted from regions of interest (ROI) calculated using the Living Image® software, version 3.0 (Caliper Life Sciences). The ROI of each well were manually selected. Exposure times ranged from 30 s to 2 min, depending on the intensity of the signal. Bacterial cell numbers were plotted against bioluminescence emitted, recorded as p/s. Non-bioluminescent L. plantarum 423 (pNZ8048) and E. mundtii ST4SA (pNZ8048) were used to set the background bioluminescence. A modified version of the method by Rhee et al. [58], was used for the in vitro plasmid stability experiment.

### Animals used

Ethical approval for in vivo experiments was granted by the Ethics Committee of Stellenbosch University (reference number SU-ACU-2017-0206-454). Eightweek-old female BALB/c mice were used in all experiments and were obtained from South African Vaccine Producers (Pty.) Ltd. (Sandringham, Pretoria, South Africa). Animals were housed in separate cages under controlled environmental conditions (12 h dark/light cycles, 20–22 °C). Water and a standard rodent feed was provided ad libitum and changed daily. Animal procedures were performed according to the Stellenbosch University ethical guidelines.

### Preparation of bacterial strains and dosing of mice

Lactobacillus plantarum 423 Fluc and E. mundtii ST4SA Fluc were grown at 30 °C for 12 h, whereafter 1 ml of each culture was inoculated into freshly prepared 10 ml MRS broth. Lactobacillus plantarum 423 Fluc was grown to an  $\mathrm{OD}_{550}$  of 2.5 and E. mundtii ST4SA Fluc to an  $\mathrm{OD}_{550}$  of 2.3. The cells were harvested (3 min at 8000 x g), washed twice with sterile PBS and resuspended in gavage buffer at a final concentration of  $4 \times 10^9$  CFU. Mice in each group were then gavaged with 200  $\mu$ l ( $4 \times 10^9$  CFU) of each strain.

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# In vivo gastrointestinal persistence of LAB in the murine model

Groups of mice each received a daily dose of 200 µl (4× 10<sup>9</sup> CFU) of live L. plantarum 423 Fluc or E. mundtii ST4SA Fluc for one (n = 13, per strain) or five (n = 22, per strain) consecutive days by intragastric gavage. Control mice (n = 4, per strain) received 200  $\mu$ l ( $4 \times 10^9$  CFU) of non-bioluminescent L. plantarum 423 (pNZ8048) or E. mundtii ST4SA (pNZ8048) in all experiments. Faeces (100 mg) were collected at different time points and vortexed in 1 ml sterile PBS for 5 min, followed by serial dilution in sterile PBS, and plating onto MRS agar supplemented with Cm and incubated, as described elsewhere. Viable cell numbers were expressed as CFU per 100 mg faeces. Two mice per strain were sacrificed by cervical dislocation at predetermined time points, the intestines surgically removed and immediately separated in a sterile Petri dish. The lumen of all intestinal sections was injected with air using a 27-gauge needle and syringe, as described by Rhee et al. [58], and bioluminescence recorded using the IVIS. The complete duodenum, jejunum, ileum and large intestine (cecum plus colon) were homogenized, separately, in 3 ml sterile PBS, serially diluted and plated (in duplicate) onto MRS agar supplemented with Cm as mentioned elsewhere. The plates were incubated and cell numbers determined as described elsewhere.

### In vivo bioluminescence measurements

In vivo BLI was recorded using the IVIS, equipped with a cooled-charged-device camera mounted on a light-tight specimen chamber (dark box) and a Windows computer system. Mice were gavaged with 200 µl of a D-Luciferin potassium salt suspension (30 mg/ml) 30 min before gavage with L. plantarum 423 Fluc and E. mundtii ST4SA Fluc. Mice were anesthetized with 2% (vol/vol) isoflurane in an oxygen-rich induction chamber before administering the D-Luciferin and bacteria. Mice were kept subdued during bioluminescent readings with a mixture of isoflurane (1.5%, vol/vol) and oxygen. Mice in ventral position were imaged for quantification of bioluminescent photon emission with exposure times ranging from 1 to 5 min, depending on the signal intensity. Pseudo-color images superimposed over grayscale reference images representing light intensity (red, most intense and purple being the least intense) were generated using the Living Image® software program. ROIs were manually selected and bioluminescence expressed as photons emitted per second.

### Statistical analysis

All data were analysed using GraphPad Prism (version 6.05) and statistical differences between groups were

determined using the Mann-Whitney nonparametric test. Statistical differences are shown for each data set. Error was calculated as standard error of mean (SEM). The number of animals required for each experiment was calculated (power analysis) using the resource equation for the sample size.

### **Additional files**

**Additional file 1: Figure S1.** In vitro stability of bioluminescence. (a) Stability of plasmid pNZPIdhFfluc in *L. plantarum* 423 Fluc and plasmid pNZSTIdhFfluc in *E. mundtii* ST4SA Fluc after subculturing for 7 days with replica plating on non-selective (antibiotic-free) and selective (Cm) media. The percentages of Cm-resistant colonies of three independent cultures of each respective strain are shown. Bioluminescent colonies of (b) *L. plantarum* 423 Fluc and (c) *E. mundtii* ST4SA Fluc after 7 days of subculture in antibiotic-free MRS media. (PDF 515 kb)

Additional file 2: Table S1. Primers used in this study. (PDF 497 kb) Additional file 3: Figure S2. Schematic representing the construction of the pNZPIdhFfluc and pNZSTIdhFfluc luciferase expression plasmids. Relevant features are indicated, including restriction sites and PCR primers used for cloning; the *E. coli/LAB repA* and *repC* replication genes; the chloramphenicol acetyltransferase (*cat*) gene conferring resistance to chloramphenicol; the *Pldh* promoter from the *L. plantarum* 423 lactate dehydrogenase gene and the *STldh* promoter from the *E. mundtii* ST4SA lactate dehydrogenase gene. (PDF 507 kb)

### **Abbreviations**

BLI: Bioluminescence imaging; CBluc: Click beetle luciferase; CFU: Colony forming units; Cm: Chloramphenicol; DNA: Deoxyribonucleic acid; Ffluc: Firefly luciferase; GIT: Gastrointestinal tract; IVIS: In vivo imaging system; LAB: Lactic acid bacteria; LB: Luria Bertani; MCS: Multiple cloning site; OD: Optical density; p/s: Photons per second; PBS: Phosphate buffered saline; Pldh: Lactate dehydrogenase gene promoter of Lactobacillus plantarum 423; repA: Replication gene A; repC: Replication gene C; ROI: Region of interest; STldh: Lactate dehydrogenase gene promoter of Enterococcus mundtii ST4SA; T: Termination sequences

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### Availability of data and materials

All data analysed during this study are included in this published article and its supplementary information files. Additional raw data from sequencing datasets are available from the corresponding author on reasonable request.

### Authors' contributions

WvZ is the principle investigator; designed the project; designed experiments; analysed sequencing data; carried out the animal studies; analysed the data; produced figures and tables and drafted the main manuscript text. SD participated in the design of the project; helped to interpret data and reviewed the manuscript. LD participated in the design of project and was a major contributor in writing of the manuscript. All authors have read and approved the final manuscript.

### Ethics approval and consent to participate

Ethical approval for in vivo experiments on mice was granted by the Ethics Committee of Stellenbosch University (reference number SU-ACU-2017-0206-454).

### Consent for publication

Not applicable.

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### Competing interests

The authors declare that they have no competing interests.

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