

## **Bacterial-derived gamma-aminobutyric acid significantly attenuates proliferation of human colon adenocarcinoma cells in vitro**

Florian Kratz<sup>1,2</sup>, Sinead Heuston<sup>1,2</sup>, Lis London<sup>2,3</sup>, Paul Ross<sup>2,3</sup>, Catherine Stanton<sup>2,3</sup>, Aileen Houston<sup>1,2</sup>, Niall Hyland<sup>1,2</sup>. <sup>1</sup>University College Cork, Cork, Ireland, <sup>2</sup>Alimentary Pharmabiotic Centre, Cork, Ireland, <sup>3</sup>Teagasc Food Research Centre, Moorepark, Ireland

Gamma-aminobutyric acid (GABA) is the major inhibitory neurotransmitter in the central nervous system. However, GABA and its receptors have recently emerged as having potential tumour-suppressive effects, with the strongest evidence found in colon cancer. Previously we demonstrated that GABA significantly reduced the proliferation and migration of human and mouse colon cancer cells *in vitro* (Fernandes et al., 2012). Lactic acid bacteria have been described as *cell factories* for GABA (Li & Cao, 2012), and thus have potential as therapeutic tools for the delivery of anti-cancer agents to the colon. However, the ability of GABA-producing bacteria to suppress colon cancer is unknown. Thus, the aim of the current study was to investigate the effect of bacterial-derived GABA on the proliferation of human (SW480) and murine (CT26) colon tumour cell lines *in vitro*. A screen of 91 human intestinally-derived bacteria identified five efficient GABA-producing strains (Barret et al., 2012). Of these, we selected a *Lactobacillus* with the greatest GABA production in the presence of monosodium glutamate (MSG) to prepare GABA-enriched supernatants. Control supernatants were prepared by omitting MSG from the culture media. Lyophilised supernatants from *Lactobacilli* cultured with MSG contained 0.858 mg/mg GABA, while those prepared from *Lactobacilli* cultured in the absence of MSG contained 0.015mg/mg GABA. SW480 and CT26 cells were cultured under standard conditions and treated with GABA (10 $\mu$ M), GABA-enriched (final concentration of GABA, 1 $\mu$ M) and control supernatants for 16 hours, after which cell proliferation was measured by resazurin reduction. To determine whether the anti-proliferative effect of GABA (10 $\mu$ M) on SW480 cells was mediated by GABA<sub>A</sub> or GABA<sub>B</sub> receptors, cells were pre-treated (1 hour) with bicuculline (100 $\mu$ M) or phaclofen (100 $\mu$ M) respectively, and incubated overnight with GABA in the presence of the antagonists. Data are representative of three independent experiments performed in triplicate (supernatant studies) or two independent experiments performed in duplicate/triplicate (antagonist study). Statistical differences were determined using a one-way ANOVA. All tests were performed using GraphPad Prism 5. Both GABA (10 $\mu$ M;  $p$ <0.05) and GABA-enriched bacterial supernatants ( $p$ <0.05) significantly inhibited proliferation of the human adenocarcinoma cell line, SW480. In contrast, supernatants obtained from *Lactobacilli* cultured in the absence of MSG (control) had no significant effect on proliferation. Neither GABA (10 $\mu$ M), nor either of the bacterial-derived supernatants, significantly influenced proliferation of murine, CT26 colon tumour cells. This is consistent with our previous findings with GABA (500nM-100 $\mu$ M) in which GABA had the greatest inhibition of proliferation in the range 2.5 $\mu$ M-5 $\mu$ M (Fernandes et al., 2012). Neither the GABA<sub>A</sub> nor GABA<sub>B</sub> receptor antagonists significantly inhibited the anti-proliferative effect of GABA on SW480 cells. These data demonstrate that bacterial-derived GABA exerts

an anti-proliferative effect on human adenocarcinoma cells *in vitro*, and suggests that further characterisation of GABA production by such bacteria, and their application in preclinical models of colon cancer is warranted.

*Fernandes P et al, Proceedings of the British Pharmacological Society at <http://www.pA2online.org/abstracts/Vol10Issue4abst044P.pdf>, 2012*

*Li H & Cao Y, Amino Acids 39:1107, 2010.*

*Barrett E et al, J Appl Microbiol 113:411, 2012.*