**Colonic mucosa-associated Diffusely-Adherent** *afaC*+ *Escherichia coli* expressing *lpfA* and *pks* **are increased in inflammatory bowel disease and colon cancer.** Maelle Prorok-Hamon, Melissa K. Friswell, Abdullah Alswied, Fei Song, Carol L. Roberts, Paul Flanagan, Paul Knight, Caroline Codling, Julian R. Marchesi, Craig Winstanley, Neil Hall, Jonathan M. Rhodes and Barry J. Campbell.

## Supplementary file S4

Table S4: *E. coli* uptake and replication of fosmid library clones within J774-A1 murine macrophages

| Strain               | Uptake of bacteria <sup>a</sup><br>CFU/wells x10 <sup>4</sup> | Fold replication<br>6h/3h <sup>b</sup> | Fold replication<br>24h/3h <sup>b</sup> |
|----------------------|---|--|---|
| HM358                | 5.36± 1.1   | $17.66 \pm 3.80^{\circ}$               | <b>11.26± 2.67<sup>c</sup></b>          |
| HA+(afaC+)           |   |  |   |
| 1a12                 | $5.14 \pm 2.3$  | $2.37{\pm}0.51$                        | $0.77{\pm}0.19$                         |
| 2f4                  | $5.52 \pm 3.8$  | $1.94 \pm 0.40$                        | $0.19 \pm 0.03$                         |
| 2f9                  | $6.43 \pm 2.7$  | $1.84 \pm 0.18$                        | $0.23 \pm 0.05$                         |
| 3b9                  | $3.99 \pm 2.3$  | $1.53 \pm 0.28$                        | $0.20 \pm 0.01$                         |
| 4b6                  | $6.26 \pm 2.8$  | $1.89 \pm 0.09$                        | $0.49 \pm 0.06$                         |
| 4g9                  | $10.86 \pm 1.5$   | $2.29{\pm}0.51$                        | $1.54 \pm 0.25$                         |
| 8h8                  | $5.08 \pm 2.7$  | $1.19 \pm 0.08$                        | $0.51 \pm 0.28$                         |
| 9b12                 | $7.75 \pm 3.4$  | $1.62 \pm 0.05$                        | $0.14 \pm 0.01$                         |
| HA- ( <i>afaC</i> -) |   |  |   |
| 1c2                  | $7.75 \pm 3.4$  | $1.81 \pm 0.15$                        | $0.92 \pm 0.10$                         |
| 3h5                  | $2.38 \pm 1.0$  | $2.3 \pm 0.48$                         | $1.18 \pm 0.01$                         |
| 4c11                 | $6.45 \pm 2.7$  | $2.15 \pm 0.35$                        | $2.63 \pm 0.60$                         |
| 5b9                  | $5.62 \pm 4.1$  | $1.69 \pm 0.31$                        | $0.18 \pm 0.008$                        |
| 7c11                 | $7.76 \pm 6.6$  | $1.66 \pm 0.39$                        | $0.24 \pm 0.05$                         |
| 8d8                  | $6.29 \pm 3.09$   | $2.19 \pm 0.16$                        | $0.44 \pm 0.11$                         |
| 9b7                  | $4.73 \pm 2.0$  | $1.61 \pm 0.26$                        | $0.23 \pm 0.06$                         |

Haemagglutinin positive (HA+) and negative (HA-) library clones

Data expressed as means  $\pm$  SEM, determined from N=2-7 independent experiments, with each experiment performed with n=2-3 replicates.

<sup>a</sup> CFU recovered from lysed macrophages after 3h (2h infection followed by 1h gentamicin treatment)

<sup>b</sup> recovered intracellular bacteria from lysed macrophages after 6h or 24h, relative to intracellular numbers at 3h **Statistical analysis was performed using Mann-Whitney U with Bonferroni correction.** 

<sup>c</sup> significantly different from EPI300<sup>™</sup>-T1<sup>R</sup>/pCC1FOS<sup>™</sup>; p<0.0001.

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## Methods S4: Assessment of TNFa release from macrophages.

Macrophages were infected with *E. coli*, library clones or *E. coli* constructs in antibiotic-free media for 2h, at MOI 10. Media was removed, macrophages washed and replaced with media containing 20µg/mL gentamicin. Followed 4h incubation, supernatant was collected and assayed using a commercial mouse TNFα Quantikine<sup>®</sup> ELISA (SMTA00: R&D systems, Abingdon; UK).

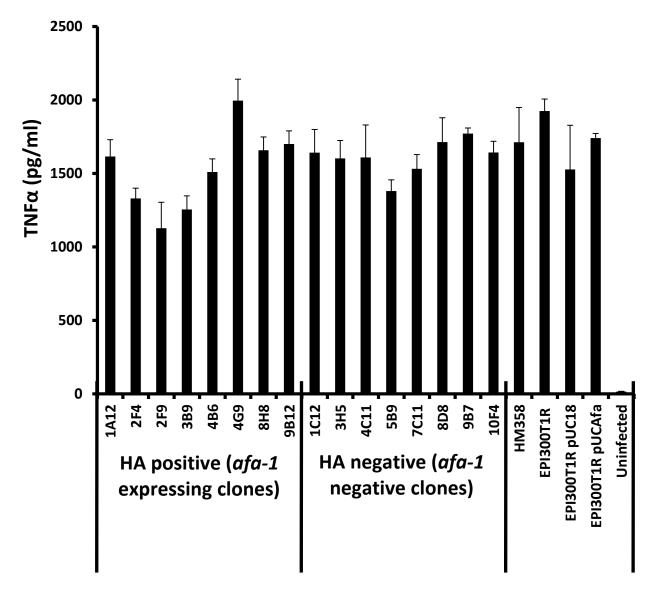


Figure S4: TNF $\alpha$  release from macrophages by *afa-1* positive clones, wild type *E. coli* HM358 and *E. coli* expressing the full *afa-1* operon. Murine J774.A1 macrophages were infected with *E. coli* for 2h, at MOI 10. The 8 HA positive fosmid library clones possessing the *afa-1* gene cluster, exhibiting no significant ability above HA negative clones to release TNF $\alpha$  to culture supernatant (6h post-infection) as assessed by ELISA. Similar TNF $\alpha$  levels were released by the wild-type strain HM358 and *E. coli* EPI300<sup>TM</sup>-T1<sup>R</sup>/pCC1FOS<sup>TM</sup> expressing Afa-1 from pUCAfa. Results are presented as mean  $\pm$  SEM (n=4 for clones; n=3 for other *E. coli*).