SUPPLEMENTARY INFORMATION

A gut-oral microbiome-driven axis promotes mucosal immunity to *Candida albicans* through retinoic acid

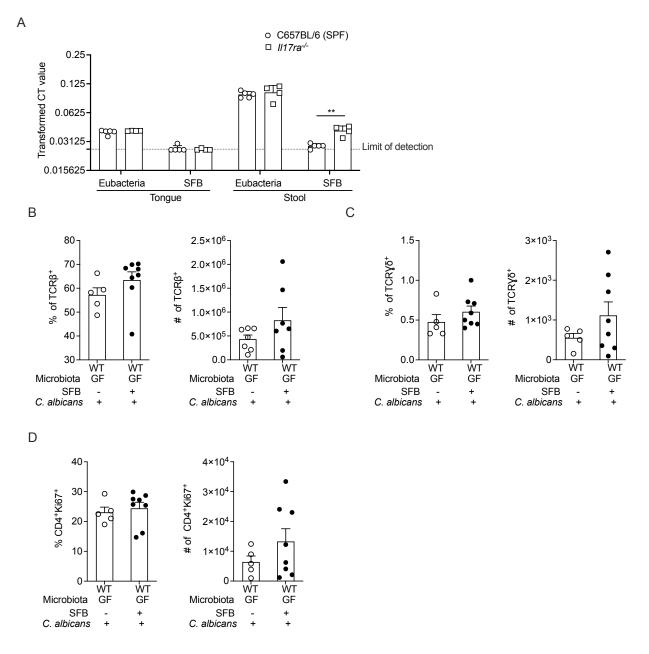
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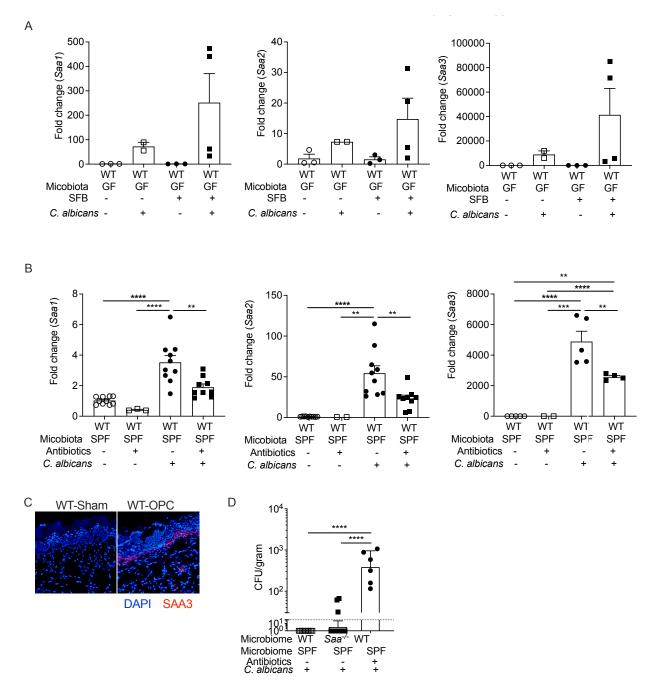
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Supplementary fig. 1: SFB is not detected in murine tongue

(A) Tongue and stool samples from WT C56BL/6 mice (JAX) or $Il17ra^{-/-}$ mice were evaluated for SFB or total Eubacteria by qPCR. Dashed line = limit of detection (LOD). Percentage and numbers of tongue TCR β^+ (C) and $\gamma\delta^+$ T cells (D) from *C. albicans*-infected GF or SFB mice harvested day 2 p.i. (E) Percentage and numbers of proliferating Tongue T cells (CD4⁺Ki67⁺) from *C. albicans*-infected GF and SFB mice harvested day 2 p.i.



Supplementary fig. 2: Serum amyloids are induced during OPC but are not required for fungal clearance.

(A) WT-GF mice were gavaged with PBS or SFB, and after 14 days infected orally with *C. albicans*. Tongue homogenates prepared on day 2 p.i. were evaluated for the indicated mRNAs by qPCR, normalized to *Gapdh*. (B) WT-SPF mice were given antibiotics in drinking water for 7 days and then infected orally with *C albicans*. Tongue homogenates prepared on day 2 p.i. were evaluated for the indicated mRNAs by qPCR, normalized to *Gapdh*. (C) Frozen tongue sections from Sham (PBS)-infected or *C. albicans*-infected mice day 2 p.i. were stained for DAPI and

SAA3. (D) The indicated mice were given antibiotics in drinking water for 7 days and then infected orally with *C albicans*. After 5 d, fungal loads in tongue were evaluated by CFU enumeration of tissue homogenates. Graphs show geometric mean \pm SD analyzed by ANOVA with Mann Whitney U. Data from 2 independent experiments.