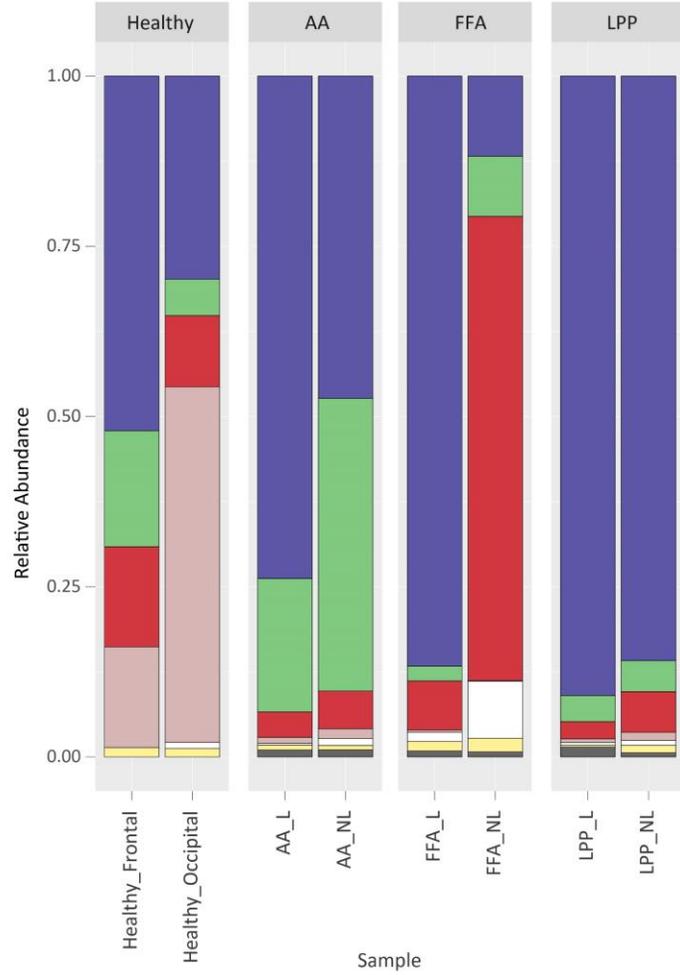
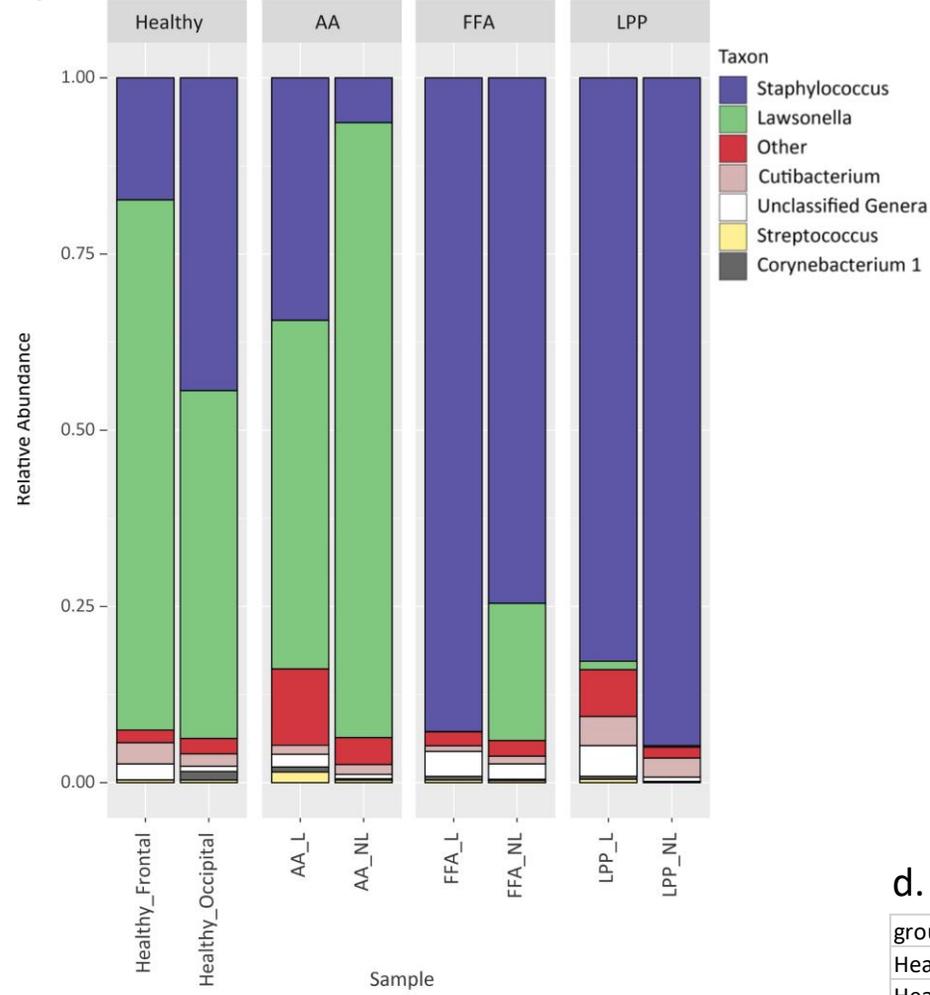


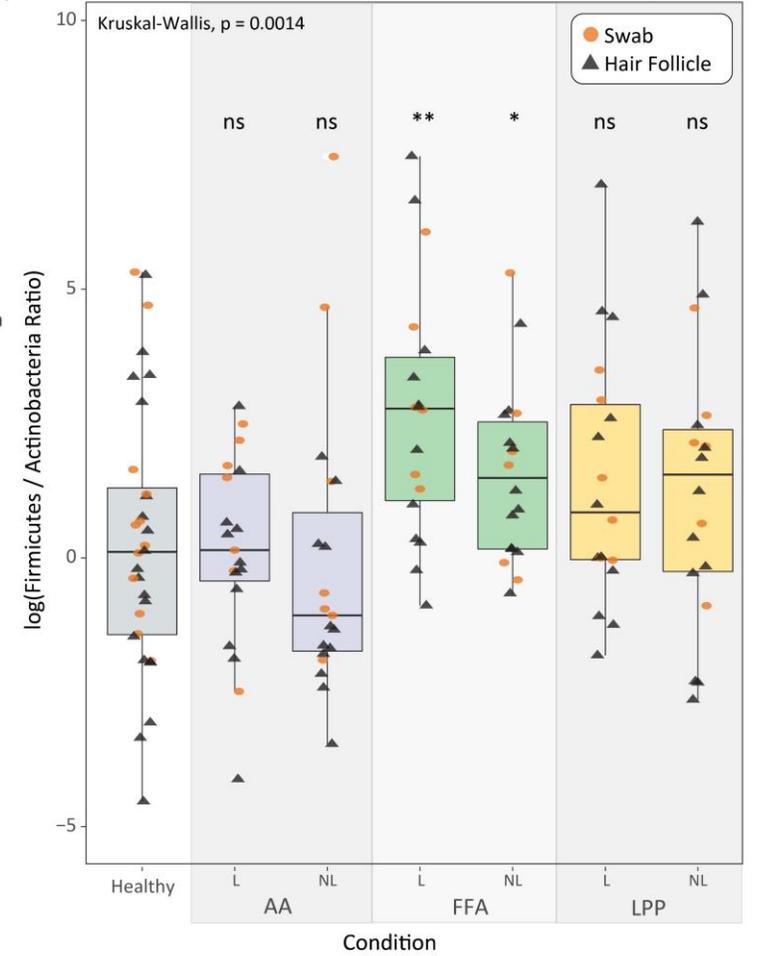
a. Swab community composition at the genus level



b. Hair community composition at the genus level



c. Firmicutes to Actinobacteria Ratio across Groups



d.

| group1 | group2 | p | p.adj | p.format | p.signif | p.adj.signif | method |
|---------|--------|----------|----------|----------|----------|--------------|----------|
| Healthy | AA_L | 0.455391 | 0.464866 | 0.4649 | ns | ns | Wilcoxon |
| Healthy | AA_NL | 0.464866 | 0.464866 | 0.4649 | ns | ns | Wilcoxon |
| Healthy | FFA_L | 0.000553 | 0.003319 | 0.0033 | *** | ** | Wilcoxon |
| Healthy | FFA_NL | 0.00954 | 0.028619 | 0.0286 | ** | * | Wilcoxon |
| Healthy | LPP_L | 0.048541 | 0.097083 | 0.0971 | * | ns | Wilcoxon |
| Healthy | LPP_NL | 0.070443 | 0.105664 | 0.1057 | ns | ns | Wilcoxon |

Supplementary Figure 2: Preliminary data analyses excluding the male population of our control group.

Our metagenomic analyses using only female subjects, healthy (n=6) and patients AAc (n = 6), FFA (n=6), LPP (n=6). (a) Community composition of the scalp surface of healthy subjects and alopecia patients at the Genus level. Relative abundance (%) of the most abundant bacteria collected from scalp swabs, sampled from frontal (F) and occipital (O) scalp in healthy females and lesional (L) and non-lesional (NL) scalp regions in female patients. (b) Hair follicle community composition at the Genus level. Relative abundance (%) of the most abundant bacteria in hair follicles, plucked from frontal (F) and occipital (O) scalp in healthy females and lesional (L) and non-lesional (NL) scalp regions in female patients. (c) The statistical analysis suggests a trade-off between the phyla Firmicutes and Actinobacteria and the diseases under study. log(FAR) pairwise comparisons between healthy and all disease sub-groups (L for lesional and NL for non-lesional scalp sites). FAR comparisons between healthy and lesional sites show a notable increase in all diseases. In cicatricial alopecias, this difference is statistically significant (p-val<0.05). Non-lesional FFA sites also exhibit significantly increased values. Significance is based on adjusted pvalue. Orange circles and black triangles represent swab and scalp HF samples, respectively. ns indicates p-value > 0.05, * <0.05, ** < 0.01 and *** <0.001 from Mann-Whitney U-test comparing each subgroup with the reference healthy group. FAR, actinobacteria and firmicutes ratio; AA, alopecia areata circumscripta; FFA, frontal fibrosing alopecia; ns, not significant. (d) The table includes the significance (*) based on the pvalue (p) before adjustment for multiple testing and adjusted pvalue (p.adj).