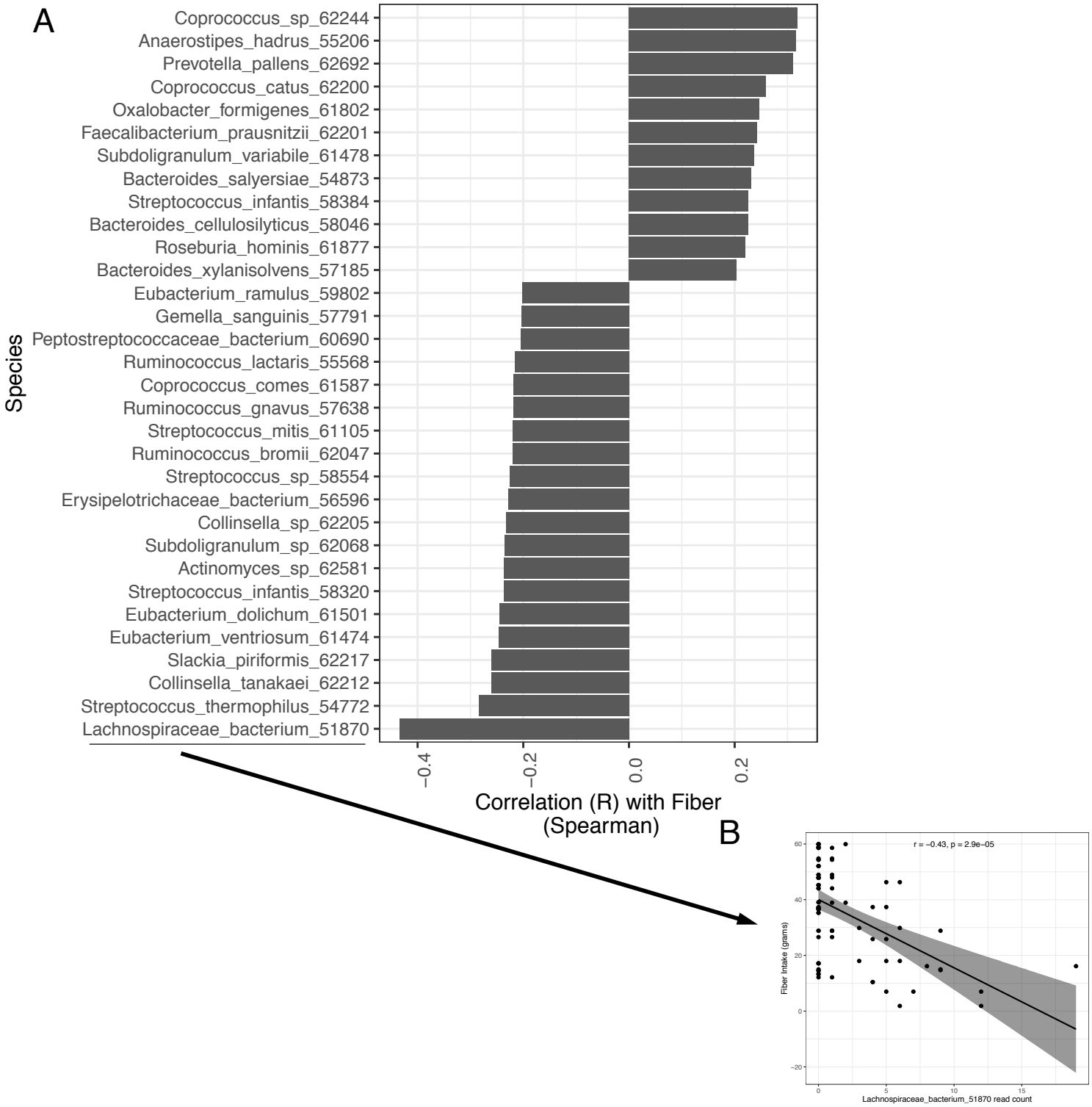
A
Coprococcus_sp_62244
Anaerostipes_hadrus_55206
Prevotella_pallens_62692
Coprococcus_catus_62200
Oxalobacter_formigenes_61802
Faecalibacterium_prausnitzii_62201
Subdoligranulum_variabile_61478
Bacteroides_salyersiae_54873
Streptococcus_infantis_58384
Bacteroides_cellulosilyticus_58046
Roseburia_hominis_61877
Bacteroides_xylanisolvens_57185
Eubacterium_ramulus_59802
Gemella_sanguinis_57791
Peptostreptococcaceae_bacterium_60690
Ruminococcus_lactaris_55568
Coprococcus_comes_61587
Ruminococcus_gnavus_57638
Streptococcus_mitis_61105
Ruminococcus_bromii_62047
Streptococcus_sp_58554
Erysipelotrichaceae_bacterium_56596
Collinsella_sp_62205
Subdoligranulum_sp_62068
Actinomyces_sp_62581
Streptococcus_infantis_58320
Eubacterium_dolichum_61501
Eubacterium_ventriosum_61474
Slackia_piriformis_62217
Collinsella_tanakaei_62212
Streptococcus_thermophilus_54772
Lachnospiraceae_bacterium_51870
Species
−0.4
−0.2
0.0
0.2
Correlation (R) with Fiber
(Spearman)
B
r = −0.43, p = 2.9e−05
Fiber Intake (grams)
Lachnospiraceae_bacterium_51870 read count