Figure: Bar chart of Average Importance Across 25 Models versus Aggregate Stability Important Features.

The y-axis is labeled "Average Importance Across 25 Models" with values 0.000, 0.005, and 0.010. The x-axis is labeled "Aggregate Stability Important Features" with the following categories (left to right):

ClimateZ, Roseisolibacter_agri, Angustibacter_luteus, Bradyrhizobium_jicamae, Modestobacter_multiseptatus, Geodermatophilus_normandii, Chelatococcus_reniformis, Gemmatirosa_kalamazoonesis, Geodermatophilus_siccatus, Rhodoplanes_sp__Z2_YC6860, Geodermatophilus_pulveris, Microvirga_makkahensis, Gaiella_occulta, Nocardioides_jensenii, Gemmatimonas_phototrophica, Urbifossiella_limnaea, Blastococcus_deserti, Eggerthella_sp__YY7918, Blastococcus_colisei, Roseomonas_aquatica, Nocardioides_sp__603, Rhodoligotrophos_appendicifer, Rhodoplanes_piscinae, Bradymonadales_bacterium_V1718, Schlegelella_aquatica, Bradyrhizobium_sp_, Sulfurisoma_sediminicola, Angustibacter_peucedani, Slackia_equolifaciens, Pseudarthrobacter_phenanthrenivorans, Herpetosiphon_aurantiacus, Thermomarinilinea_lacunifontana, Adhaeribacter_aerolatus, Sphingomonas_rhizophila, Neobacillus_drentensis, Angustibacter_aerolatus, Streptomyces_chartreusis, Arthrobacter_sp__PAMC25564, Terrabacter_terrigena, Kitasatospora_kazusensis, Mycolicibacterium_phocaicum, Tepidisphaera_mucosa, Methylobacterium_tardum, Streptomyces_canus, Betaproteobacteria_bacterium_GR16_43, Longimicrobium_terrae, Thermaerobacter_composti, Ilumatobacter_fluminis, Bellilinea_caldifistulae, Cryptosporangium_aurantiacum