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Edited provisional title

Metagenomic and metabolomic analyses show correlations between
intestinal microbiome diversity and microbiome metabolites in ob/ob
and ApoE-/- mice

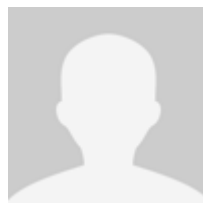
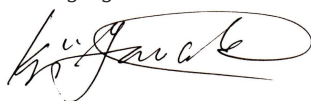
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