



Figure S4: The number of potential inputs of the metabolic networks according to Borenstein method [1].

References

1. Borenstein E, Kupiec M, Feldman MW, Ruppin E: **Large-scale reconstruction and phylogenetic analysis of metabolic environments.** *Proceedings of the National Academy of Sciences of the United States of America / PNAS* 2008, **105**(38):14482–14487.