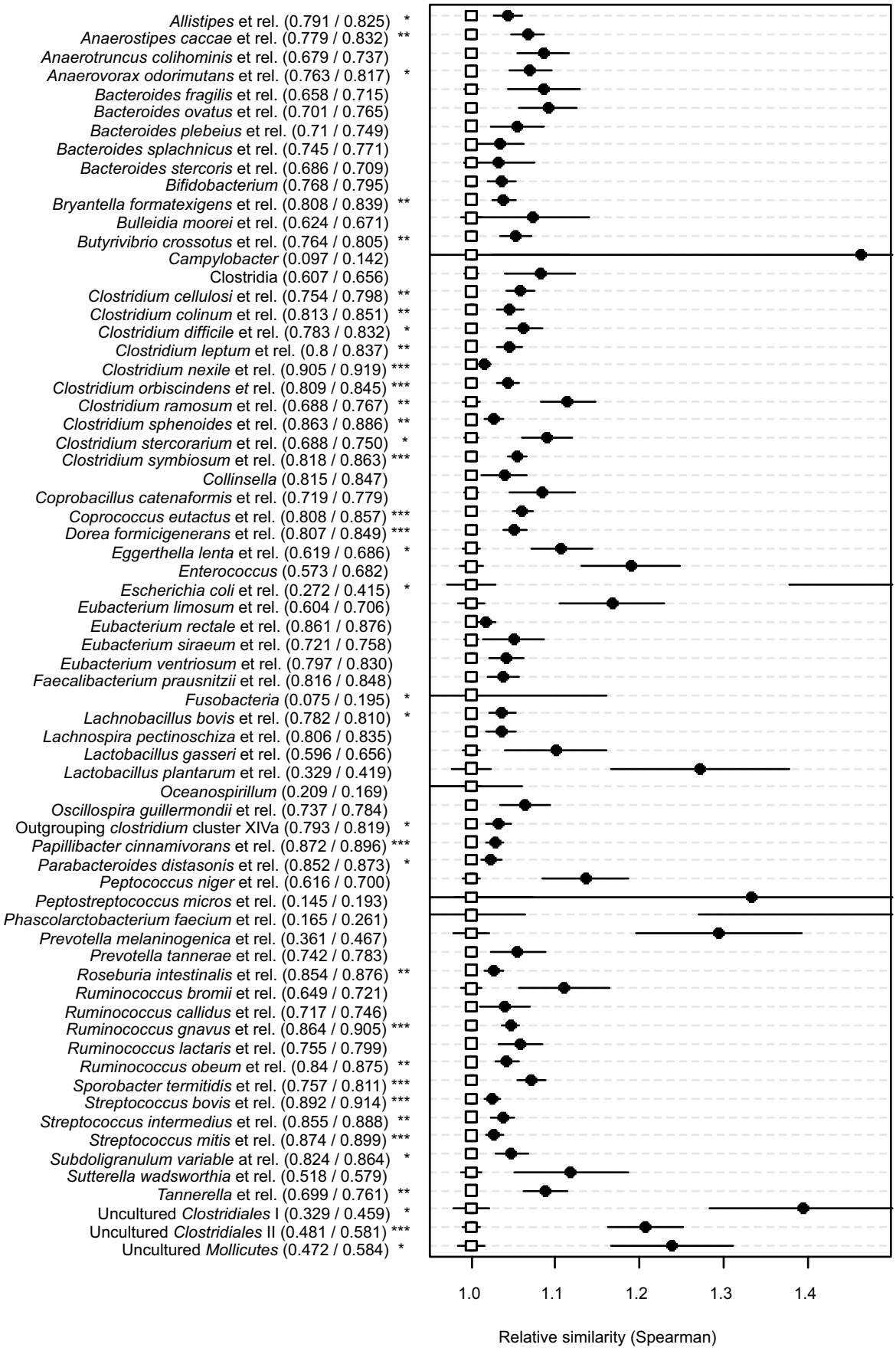
Allistipes et rel. (0.791 / 0.825) *
Anaerostipes caccae et rel. (0.779 / 0.832) **
Anaerotruncus colihominis et rel. (0.679 / 0.737)
Anaerovorax odorimutans et rel. (0.763 / 0.817) *
Bacteroides fragilis et rel. (0.658 / 0.715)
Bacteroides ovatus et rel. (0.701 / 0.765)
Bacteroides plebeius et rel. (0.71 / 0.749)
Bacteroides splachnicus et rel. (0.745 / 0.771)
Bacteroides stercoris et rel. (0.686 / 0.709)
Bifidobacterium (0.768 / 0.795)
Bryantella formatexigens et rel. (0.808 / 0.839) **
Bulleidia moorei et rel. (0.624 / 0.671)
Butyrivibrio crossotus et rel. (0.764 / 0.805) **
Campylobacter (0.097 / 0.142)
Clostridia (0.607 / 0.656)
Clostridium cellulosi et rel. (0.754 / 0.798) **
Clostridium colinum et rel. (0.813 / 0.851) **
Clostridium difficile et rel. (0.783 / 0.832) *
Clostridium leptum et rel. (0.8 / 0.837) **
Clostridium nexile et rel. (0.905 / 0.919) ***
Clostridium orbiscindens et rel. (0.809 / 0.845) ***
Clostridium ramosum et rel. (0.688 / 0.767) **
Clostridium sphenoides et rel. (0.863 / 0.886) **
Clostridium stercorarium et rel. (0.688 / 0.750) *
Clostridium symbiosum et rel. (0.818 / 0.863) ***
Collinsella (0.815 / 0.847)
Coprobacillus catenaformis et rel. (0.719 / 0.779)
Coprococcus eutactus et rel. (0.808 / 0.857) ***
Dorea formicigenerans et rel. (0.807 / 0.849) ***
Eggerthella lenta et rel. (0.619 / 0.686) *
Enterococcus (0.573 / 0.682)
Escherichia coli et rel. (0.272 / 0.415) *
Eubacterium limosum et rel. (0.604 / 0.706)
Eubacterium rectale et rel. (0.861 / 0.876)
Eubacterium siraeum et rel. (0.721 / 0.758)
Eubacterium ventriosum et rel. (0.797 / 0.830)
Faecalibacterium prausnitzii et rel. (0.816 / 0.848)
Fusobacteria (0.075 / 0.195) *
Lachnobacillus bovis et rel. (0.782 / 0.810) *
Lachnospira pectinoschiza et rel. (0.806 / 0.835)
Lactobacillus gasseri et rel. (0.596 / 0.656)
Lactobacillus plantarum et rel. (0.329 / 0.419)
Oceanospirillum (0.209 / 0.169)
Oscillospira guillermondii et rel. (0.737 / 0.784)
Outgrouping *clostridium* cluster XIVa (0.793 / 0.819) *
Papillibacter cinnamivorans et rel. (0.872 / 0.896) ***
Parabacteroides distasonis et rel. (0.852 / 0.873) *
Peptococcus niger et rel. (0.616 / 0.700)
Peptostreptococcus micros et rel. (0.145 / 0.193)
Phascolarctobacterium faecium et rel. (0.165 / 0.261)
Prevotella melaninogenica et rel. (0.361 / 0.467)
Prevotella tannerae et rel. (0.742 / 0.783)
Roseburia intestinalis et rel. (0.854 / 0.876) **
Ruminococcus bromii et rel. (0.649 / 0.721)
Ruminococcus callidus et rel. (0.717 / 0.746)
Ruminococcus gnavus et rel. (0.864 / 0.905) ***
Ruminococcus lactaris et rel. (0.755 / 0.799)
Ruminococcus obeum et rel. (0.84 / 0.875) **
Sporobacter termitidis et rel. (0.757 / 0.811) ***
Streptococcus bovis et rel. (0.892 / 0.914) ***
Streptococcus intermedius et rel. (0.855 / 0.888) **
Streptococcus mitis et rel. (0.874 / 0.899) ***
Subdoligranulum variable at rel. (0.824 / 0.864) *
Sutterella wadsworthia et rel. (0.518 / 0.579)
Tannerella et rel. (0.699 / 0.761) **
Uncultured *Clostridiales* I (0.329 / 0.459) *
Uncultured *Clostridiales* II (0.481 / 0.581) ***
Uncultured *Mollicutes* (0.472 / 0.584) *
1.0
1.1
1.2
1.3
1.4
Relative similarity (Spearman)