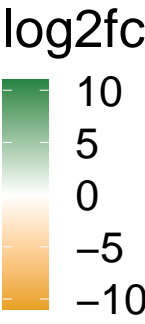
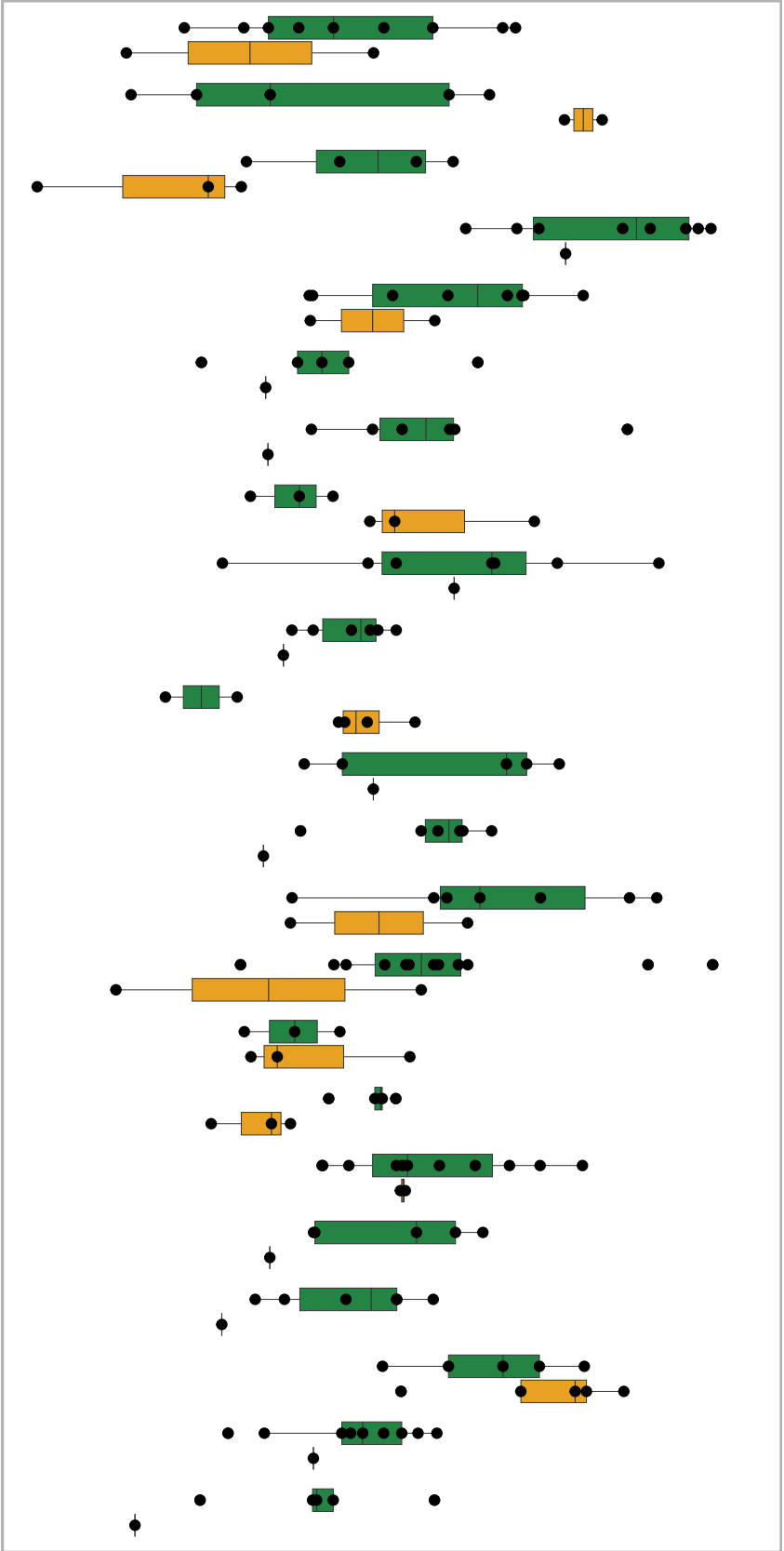
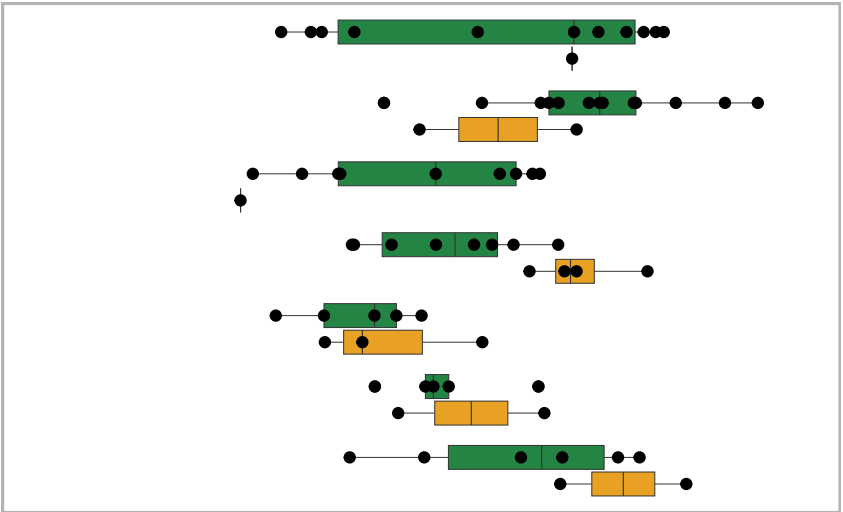


MET Progression categc

Species

<i>Streptococcus mitis</i>	2.36		2.76
<i>Prevotella histicola</i>	2.78		3.42
<i>Oribacterium asaccharolyticum</i>	2.03		8.59
<i>Granulicatella adiacens</i>	-5.53	-3.09	-3.24
<i>Cardiobacterium valvarum</i>	-3.49		-2.7
<i>Bacteroidetes oral taxon 274</i>	-2.59		-2.32
<i>Actinomyces SGB17168</i>	-2.52		-2.45
<i>Prevotella melaninogenica</i>	3.08		3.1
<i>Veillonella parvula</i>	2.77		2.44
<i>Ruminococcaceae unclassified SGB15234</i>	-2.04		-5.58
<i>Ruminococcaceae bacterium AM07 15</i>		4.16	3.99
<i>Phocaeicola dorei</i>	4.77		4.49
<i>Parasutterella excrementihominis</i>	2.24		3.72
<i>Massilistercora timonensis</i>	2.37		3.15
<i>Lactobacillus gasseri</i>	3.26		8.91
<i>Lachnospira sp NSJ 43</i>	-3.71		-6.49
<i>Klebsiella pneumoniae</i>	3.25		4.16
<i>Gordonibacter urolithinfaciens</i>	2.7		2.96
<i>Firmicutes bacterium</i>	-5.22		-7.82
<i>Faecalicatena fissicatena</i>	2.29		4.18
<i>Eubacteriaceae bacterium</i>	3.29		5.56
<i>Enterocloster clostridioformis</i>	2.09		2.95
<i>Clostridiales bacterium 1 7 47FAA</i>	5.22		8.42
<i>Candidatus Paralachnospira caecorum</i>	-2.04		-3.43
<i>Candidatus Avimicrobium caecorum</i>		2.65	2.31
<i>Blautia caecimuris</i>	3.81		2.6
<i>Anaerotignum faecicola</i>	2.5		4.92
<i>Alloscardovia omnicoles</i>	2.95		5.87
<i>Alistipes shahii</i>	-7.9		-3.05
<i>Agathobaculum butyriciproducens</i>	4.17		3.42
<i>Actinomyces SGB17154</i>	2.59		6.65

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Method



Positive log2fc values indicate higher abundance in Y progression group