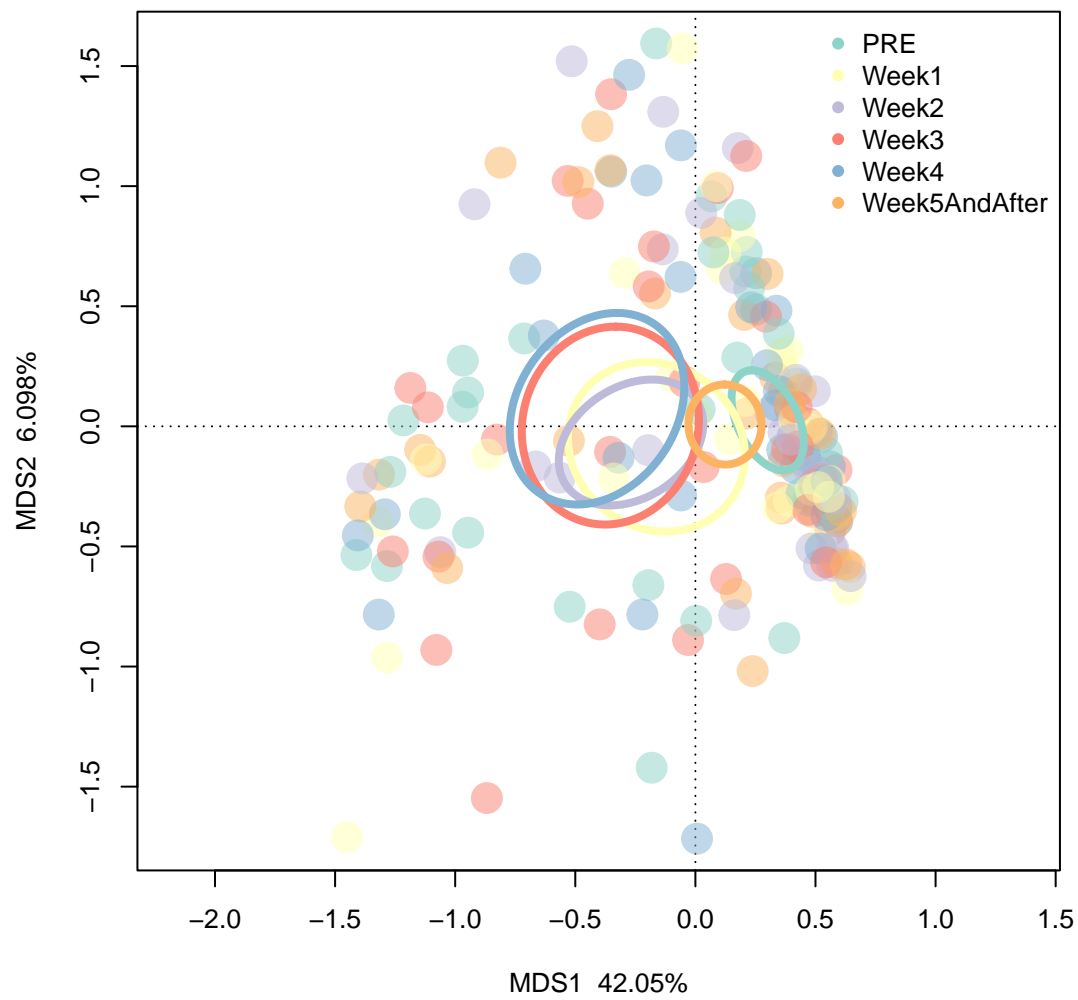
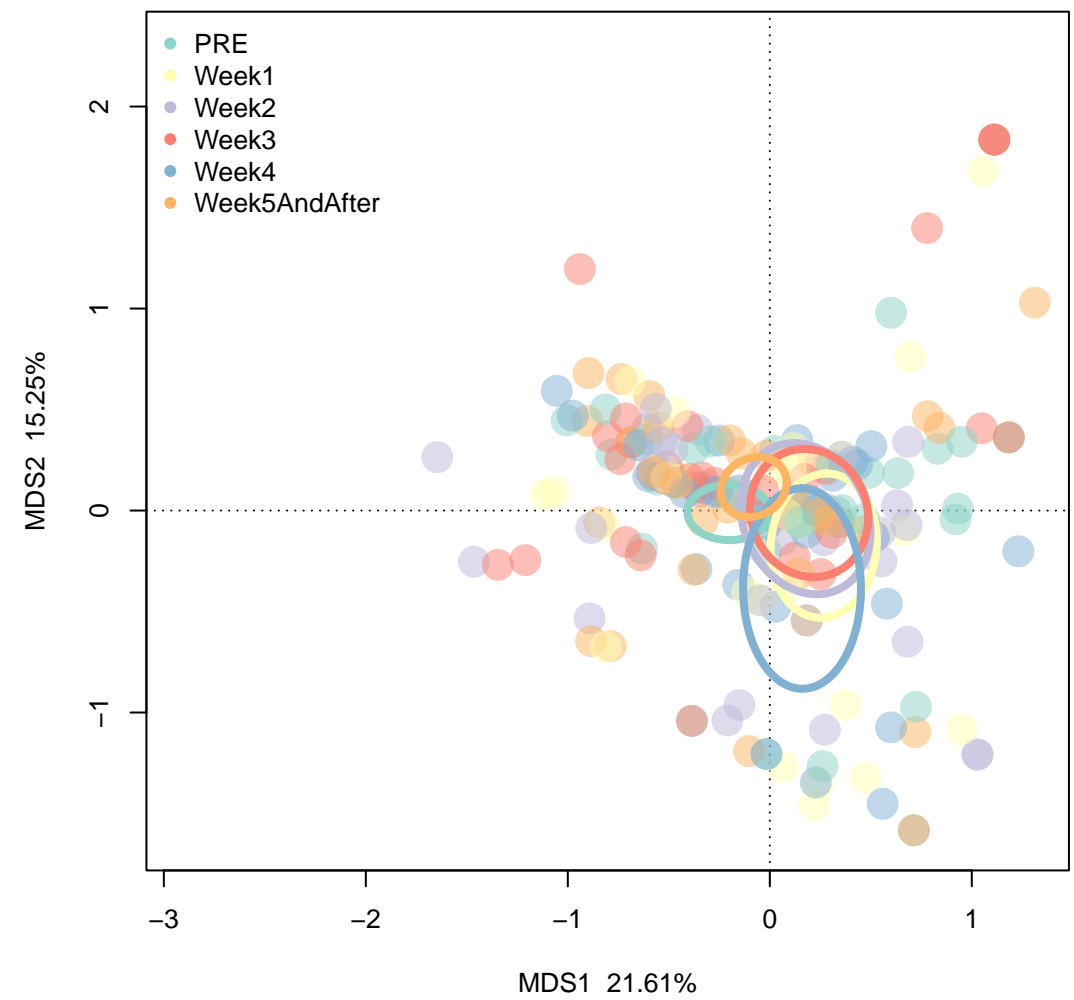


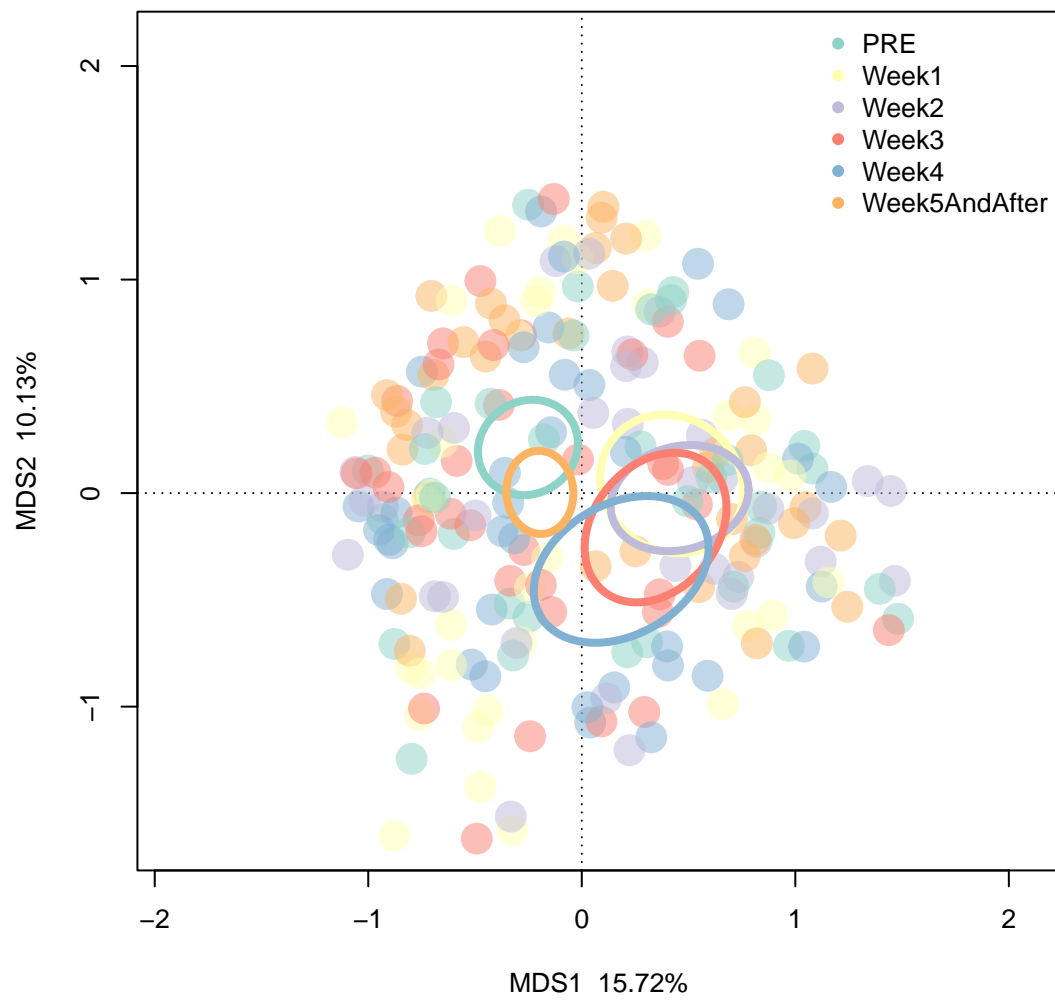
**Bracken-Species n = 200**



**AMR n = 211**



**RGI n = 211**



**vsearch n = 200**

