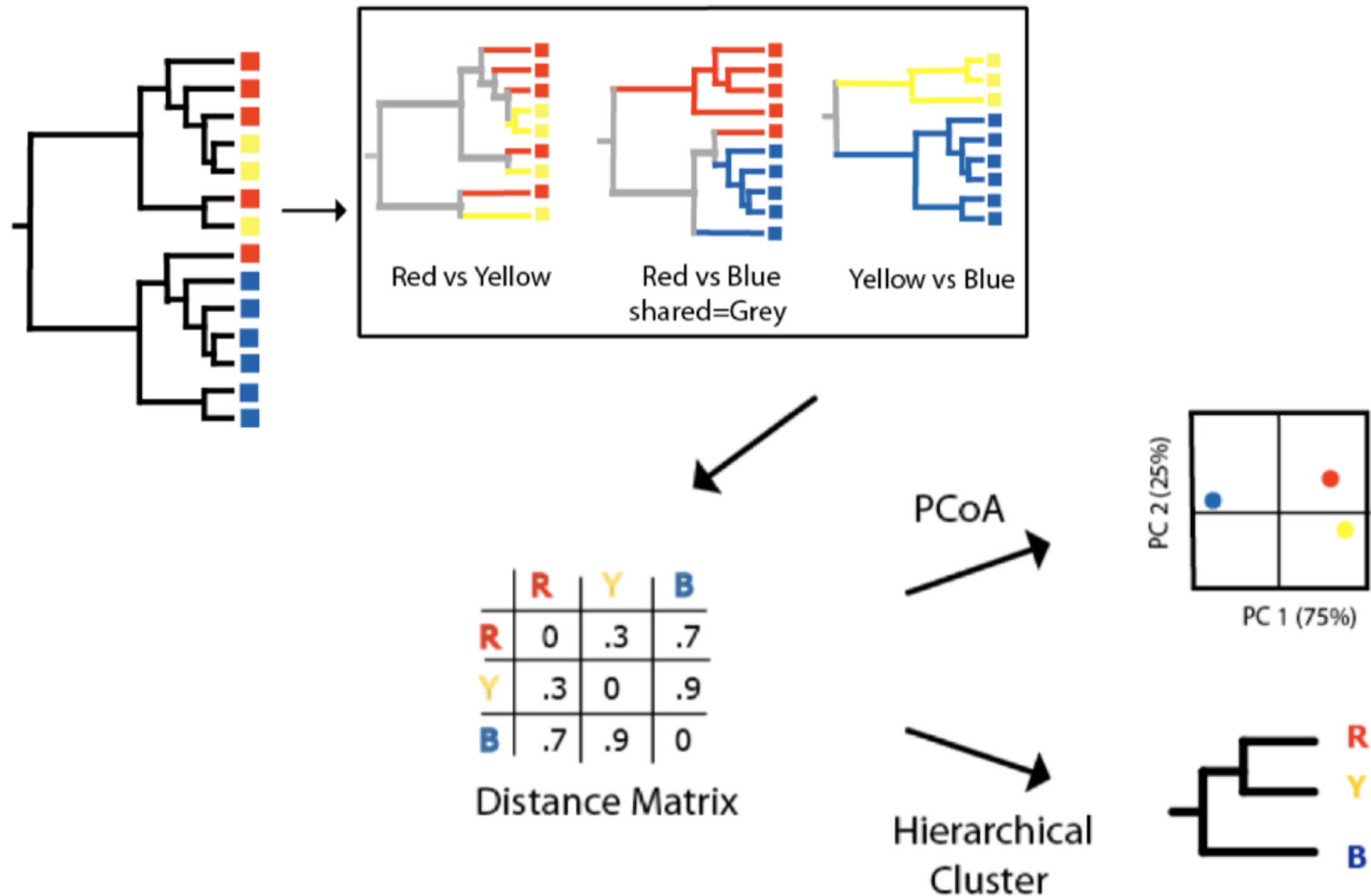


Beta Diversity: UniFrac continued



Do samples contain significantly different microbial communities?

Which taxa have increased abundance compared to another sample?

Are there broad trends that relate many samples? Can these trends be explained by an environmental factor?