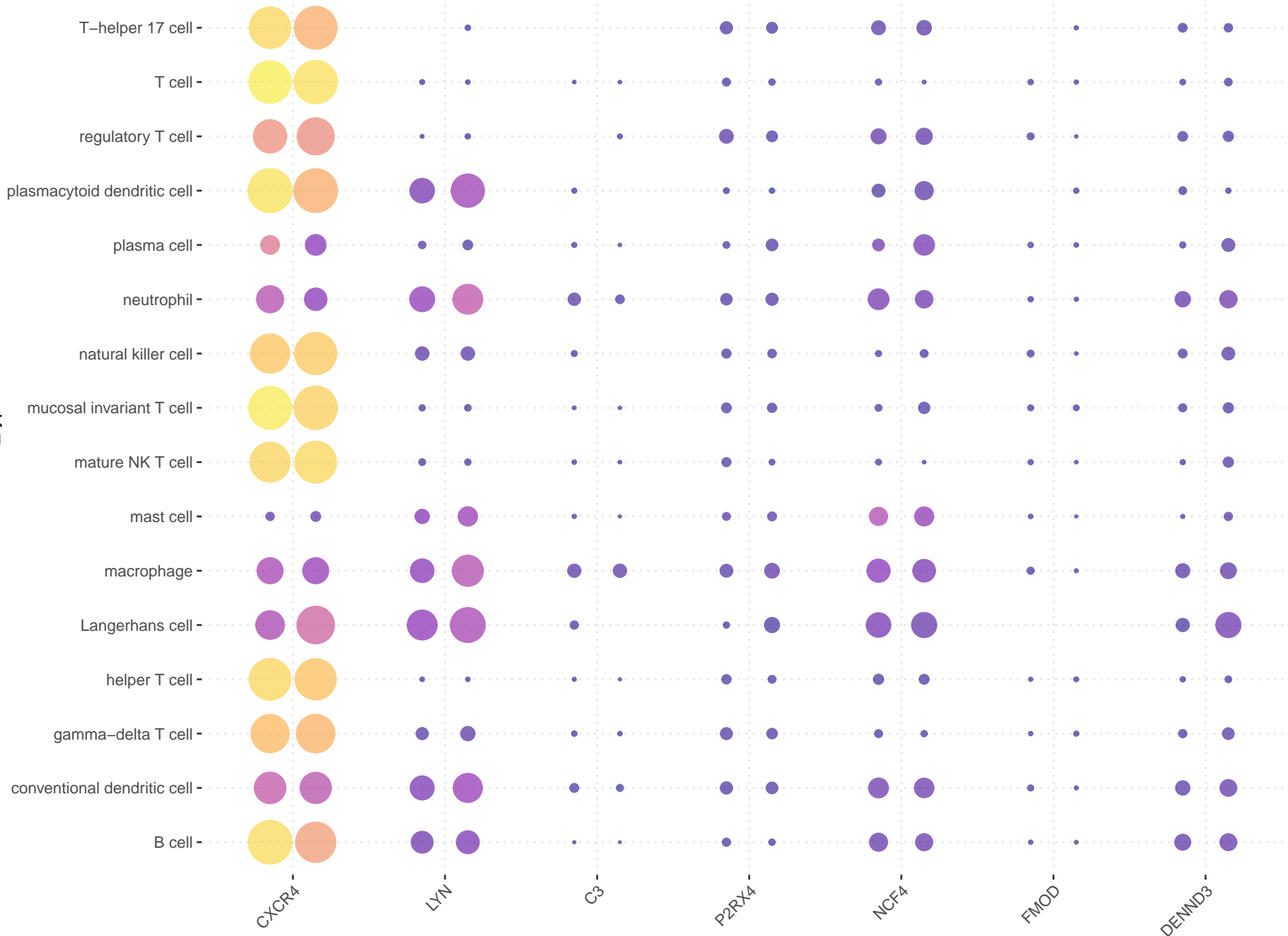


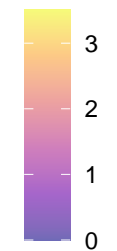
# normal vs periodontitis

cell\_type



factor(gene, levels = input\_gene)

Average Expression



Percent Expressed

