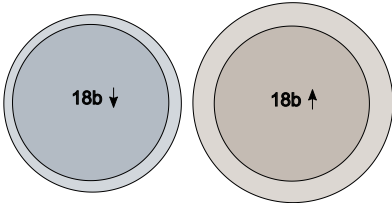
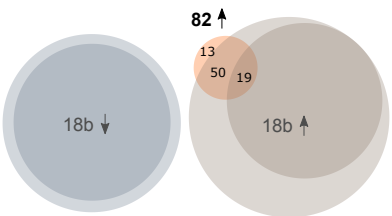


636 genes (2 fold change)
493 genes (2 fold change, FDR 0.05)

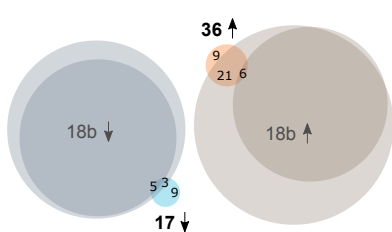
808 genes (2 fold change)
486 genes (2 fold change, FDR 0.05)



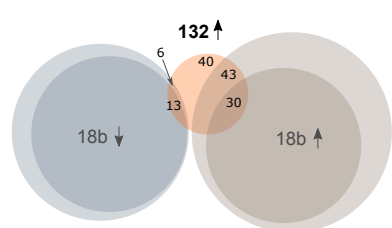
Mtb 18b, streptomycin depletion, 2 weeks



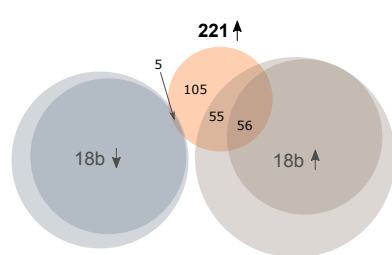
"Dormancy regulon" (Voskuil *et al.* 2003, Voskuil *et al.* 2004)



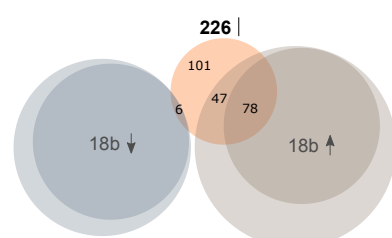
Reactive nitrogen (Ohno *et al.* 2003)



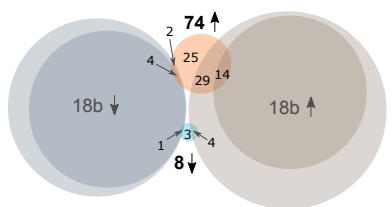
Microaerophilic (NRP1) (Muttucumaru *et al.* 2004)



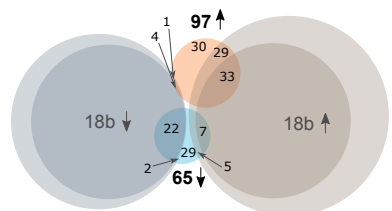
Anaerobic (NRP2) (Muttucumaru *et al.* 2004)



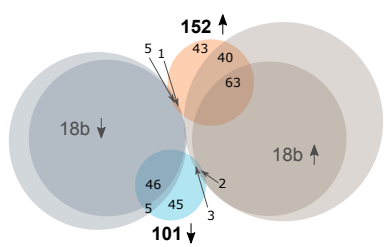
Hypoxia, 7 days (Rustad *et al.* 2008)



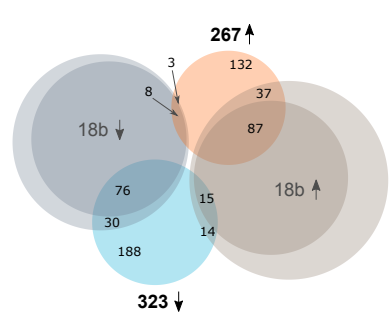
Reduced oxygen (Bacon *et al.* 2004)



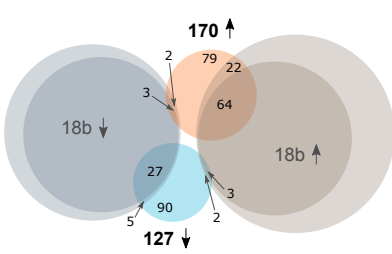
Macrophage (Rhode *et al.* 2007)



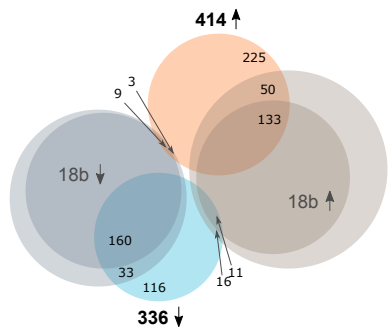
Macrophage (Homolka *et al.* 2010)



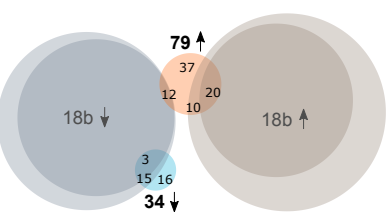
Nutrient depletion (Betts *et al.* 2002)



Nutrient depletion (Hampshire *et al.* 2004)



SDS stress (Manganelli *et al.* 2001)



Phosphostarvation (Rifat *et al.* 2009)