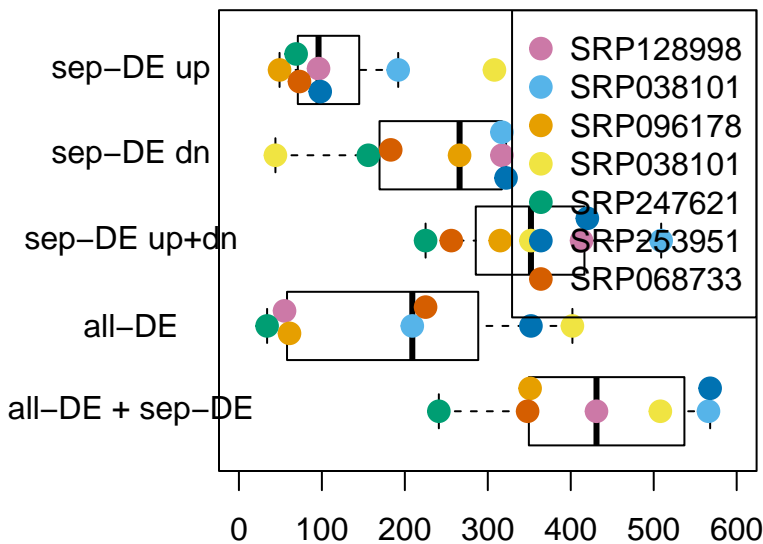
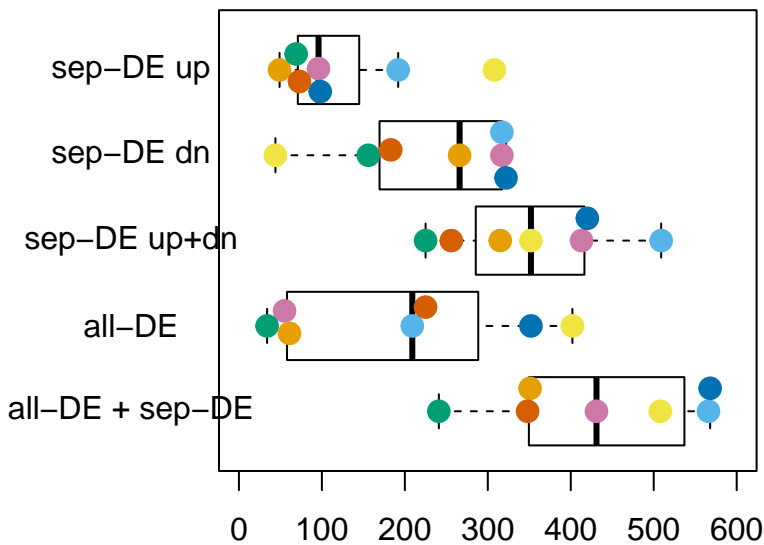


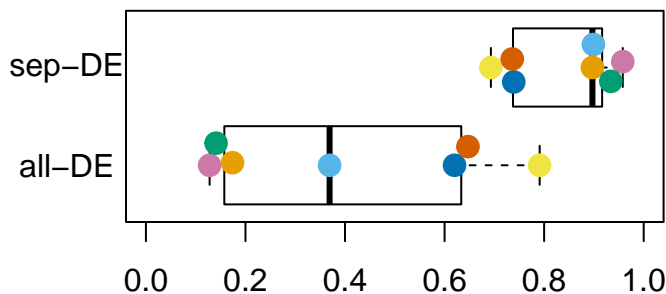
no. gene sets identified



no. gene sets identified



proportion of gene sets identified



no. gene sets unique to each approach

