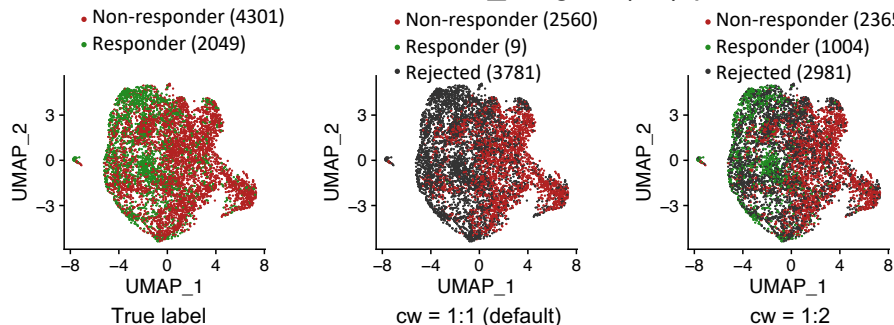
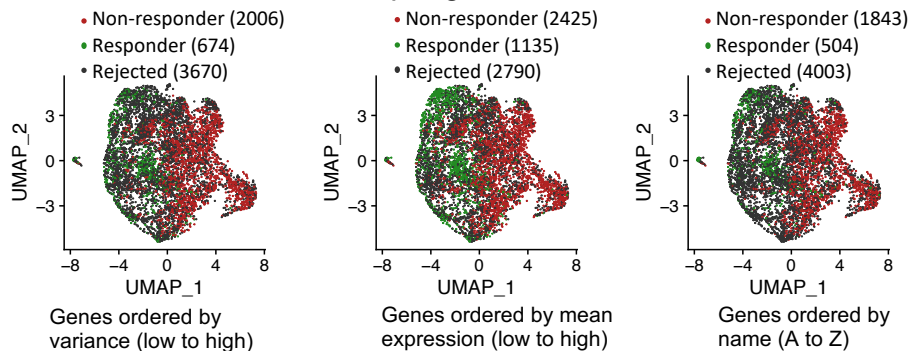


a PENCIL is sensitive to the *class_weights* (cw) parameter setting



b PENCIL is sensitive to input gene order



c PENCIL is sensitive to input cells

