Cell Cluster Confusion Matrix, ARI:0.92 Precision: 0.88 Recall: 1.00

predicted label \ true label	1	2	3	4	5	6
5			292			
4	188					
2						235
1					443	
0		180		160		
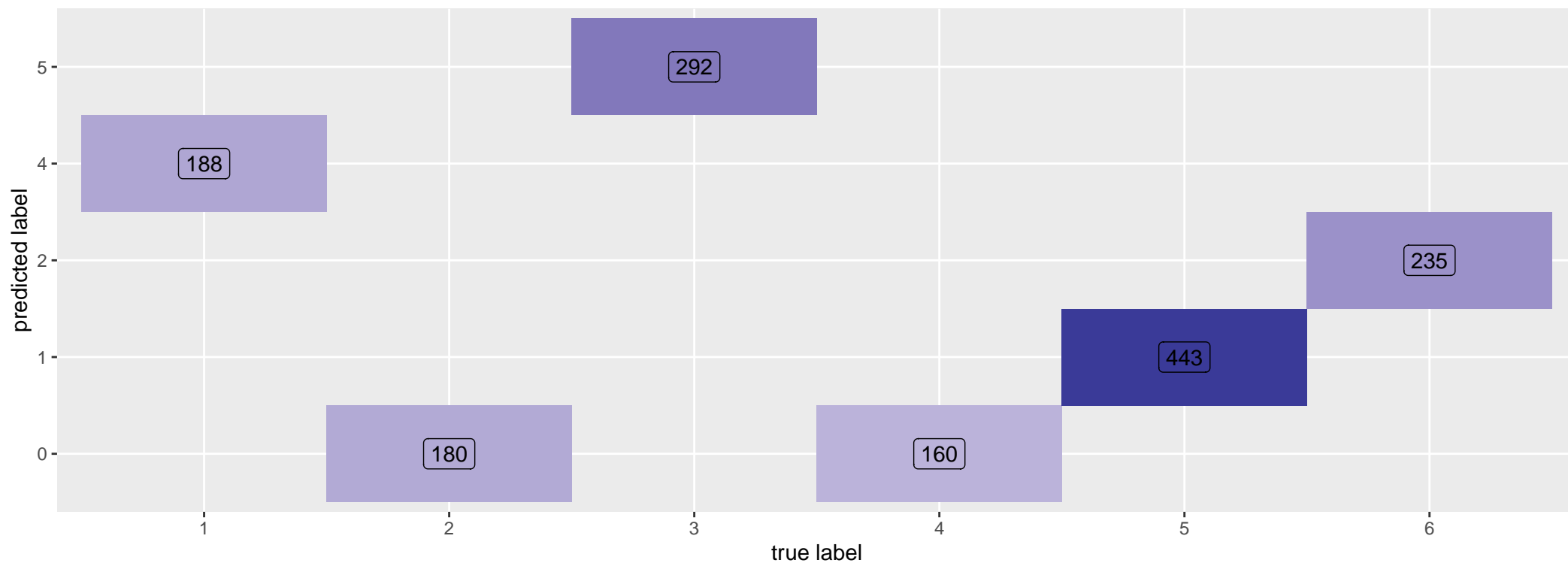
Mutation Cluster Confusion Matrix, ARI:0.37 Precision: 0.42 Recall: 0.62

predicted label \ true label	0	1	2	3	4	5	6
5			2	546	2	2	35
4	1	300	4	2	6	2	18
3			2			6	65
2	7	43	39	55	5	57	431
1	66	2	438	7	1	457	6
0	470	12	656	20	373	21	14