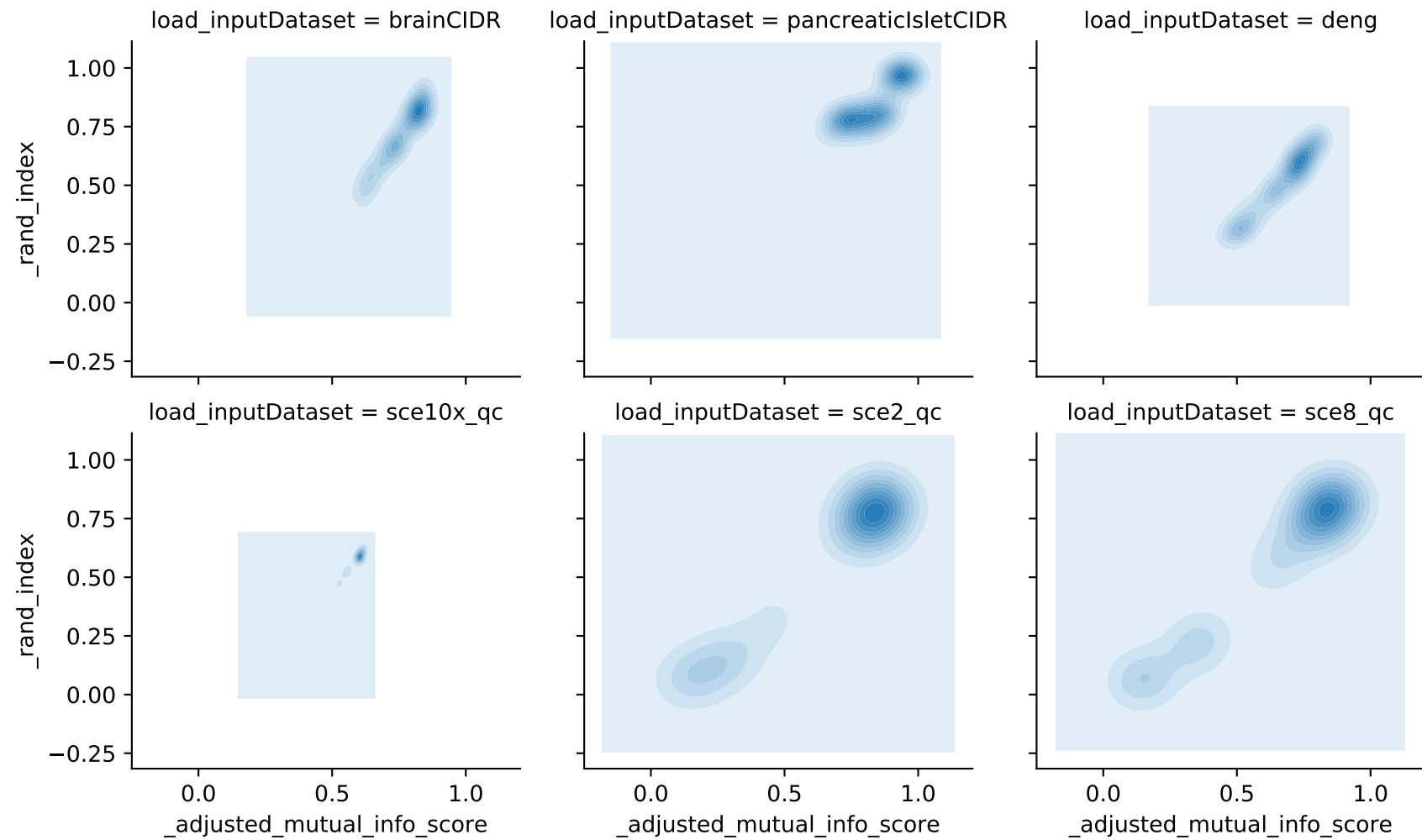
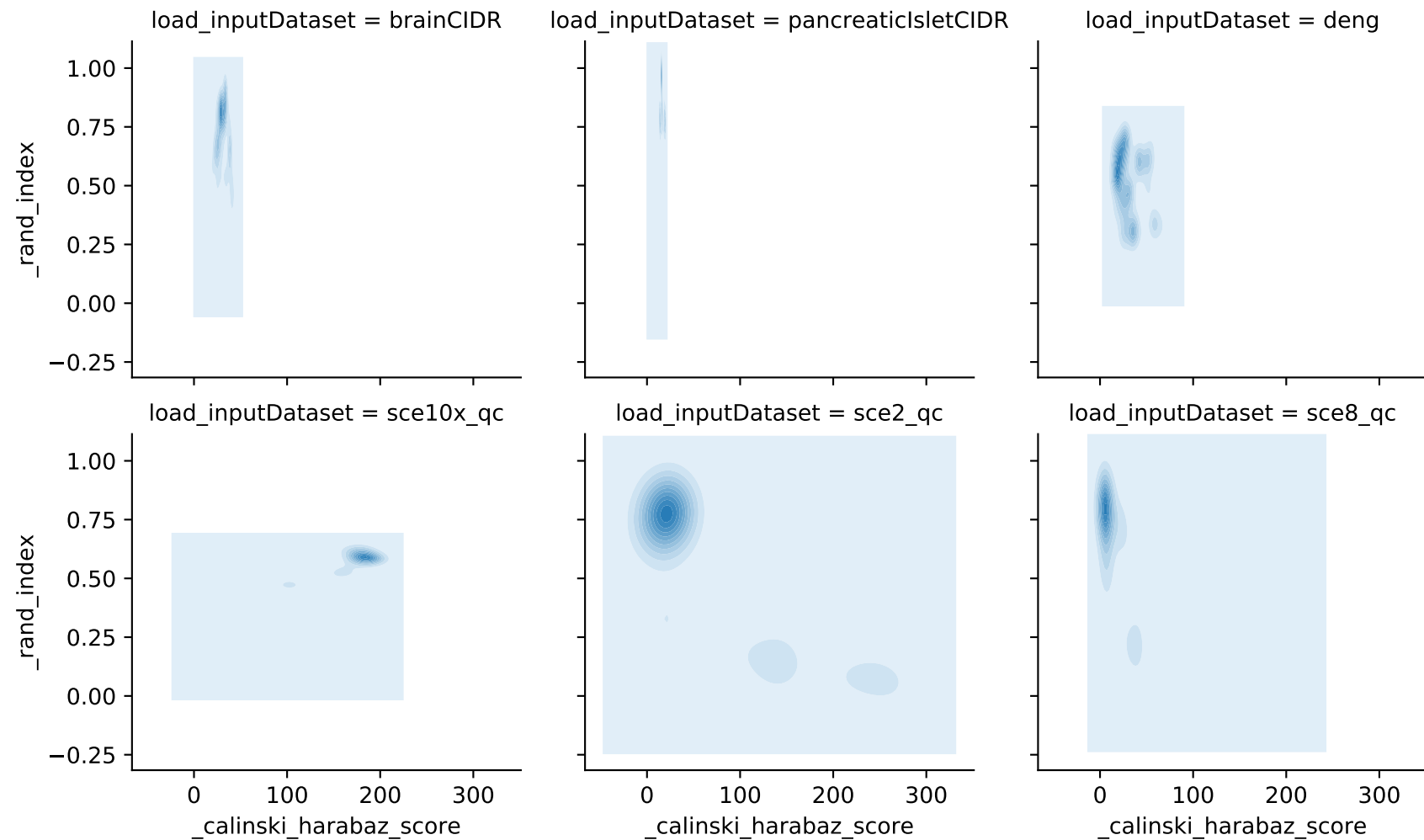


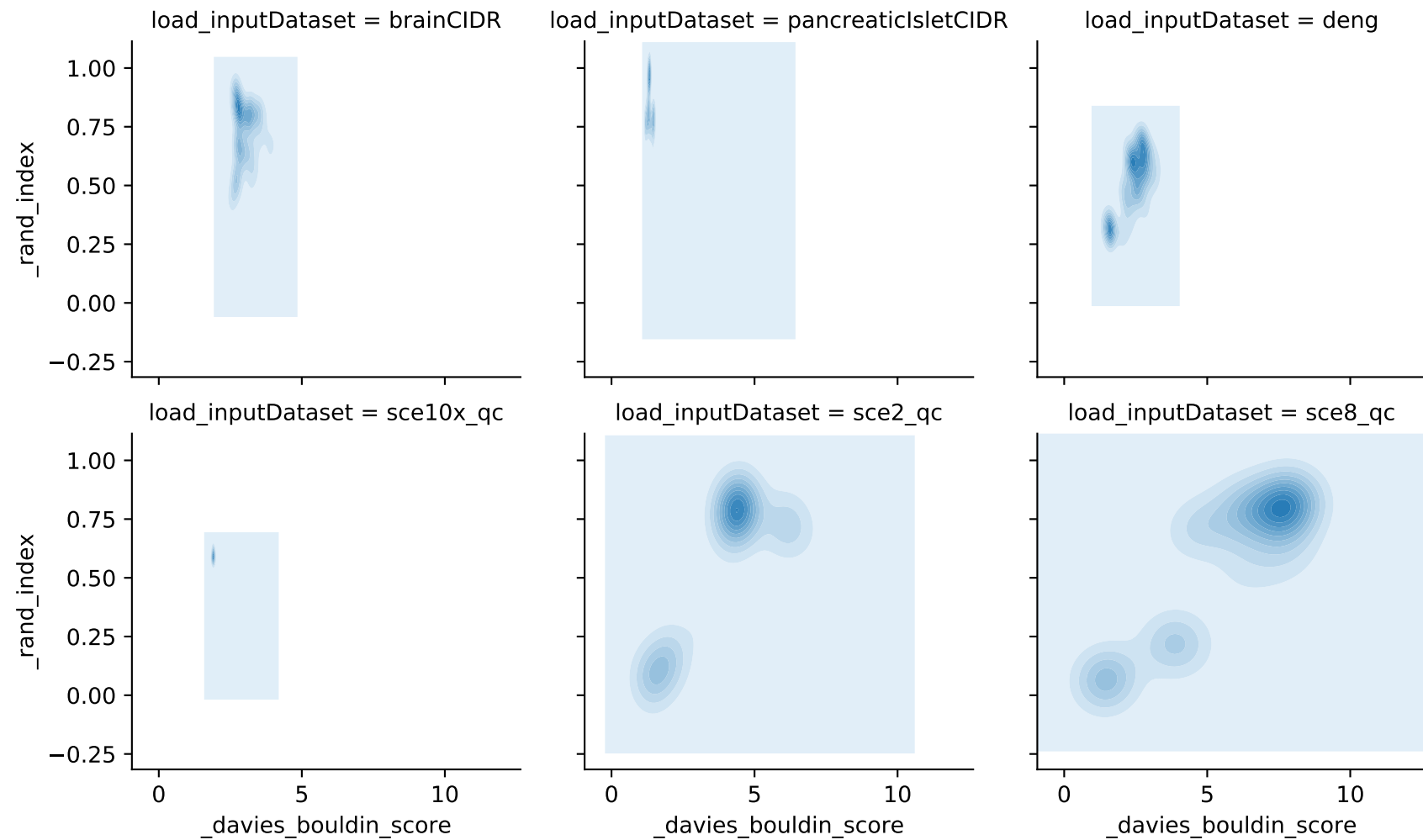
Distribution of _adjusted_mutual_info_score



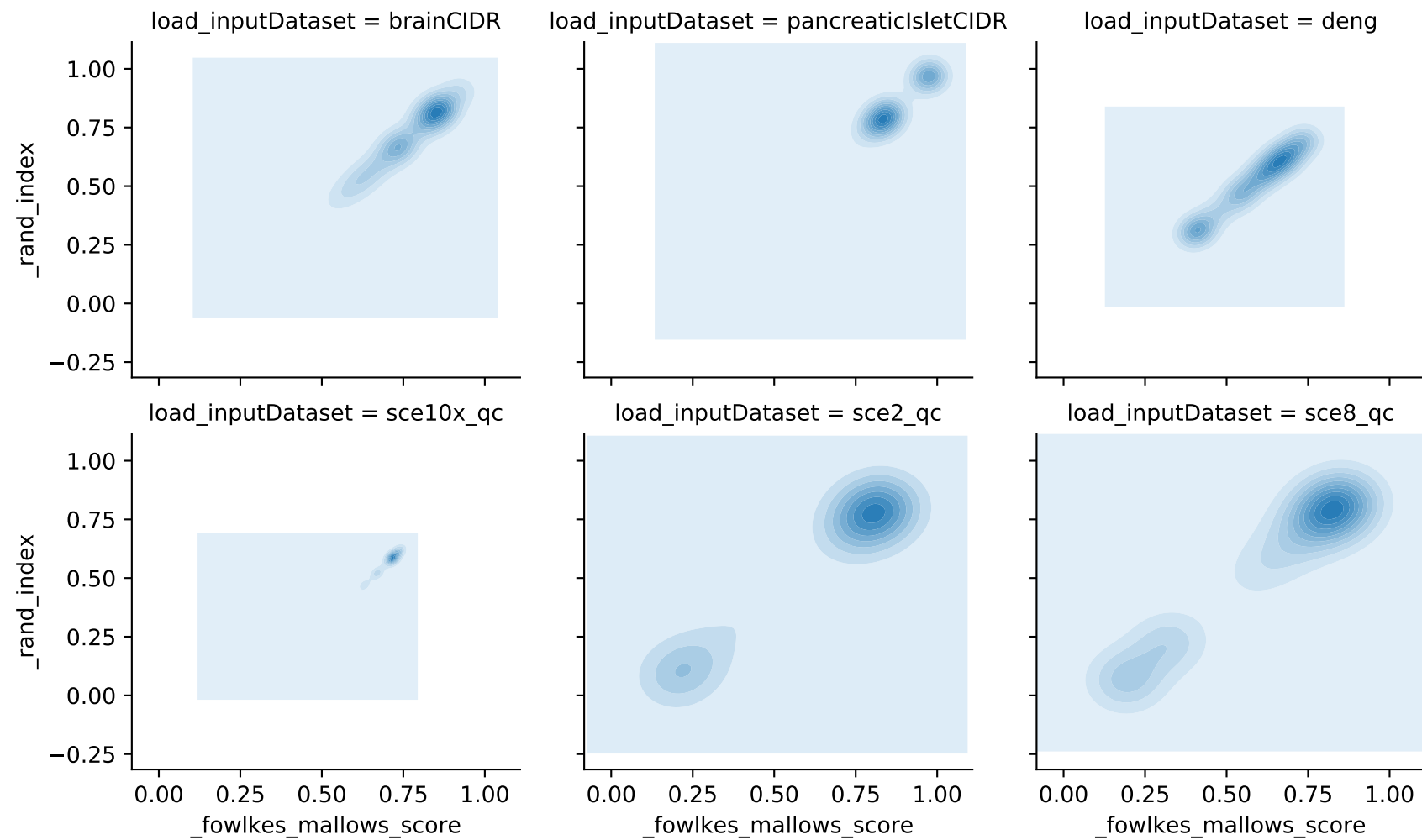
Distribution of _calinski_harabaz_score



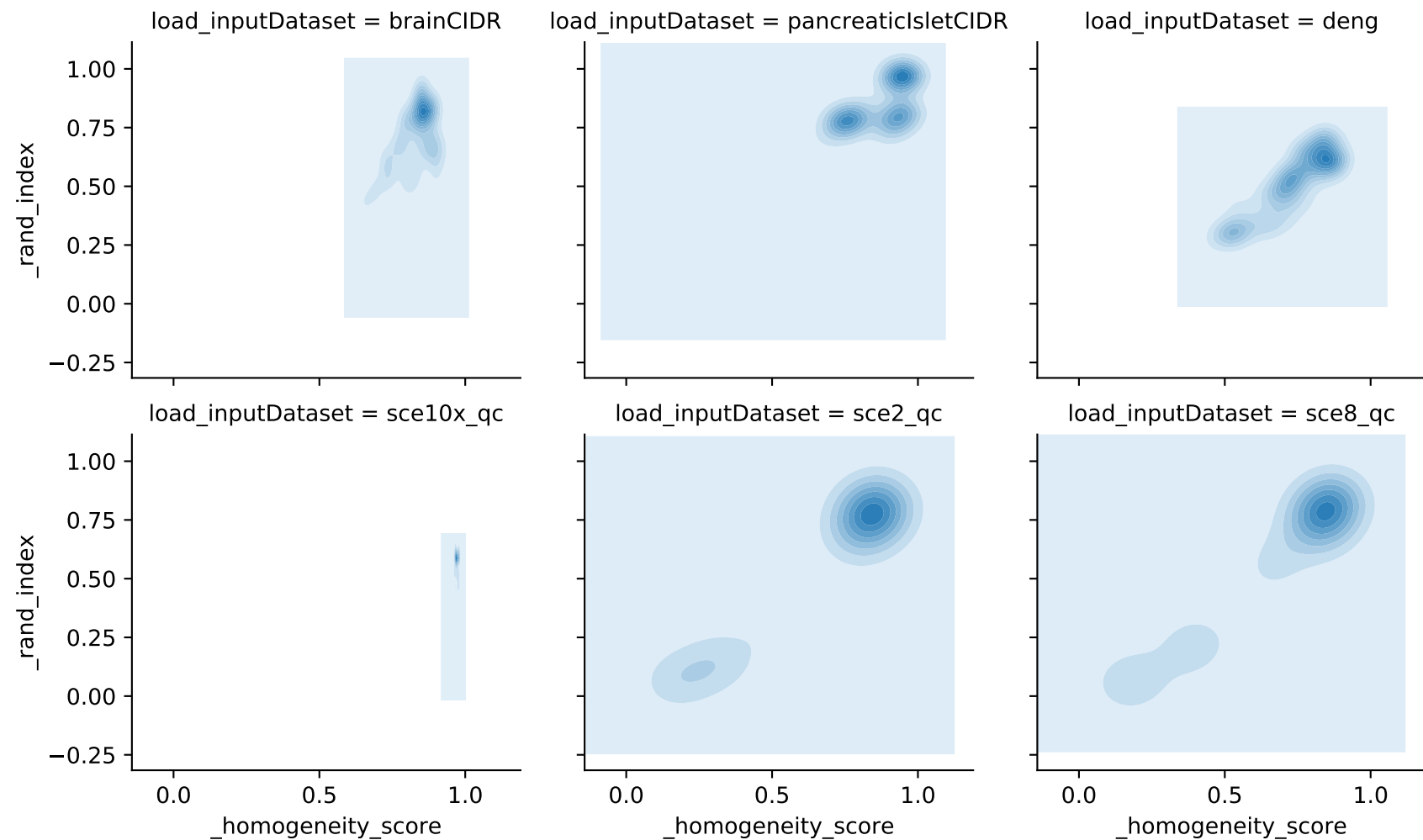
Distribution of _davies_bouldin_score



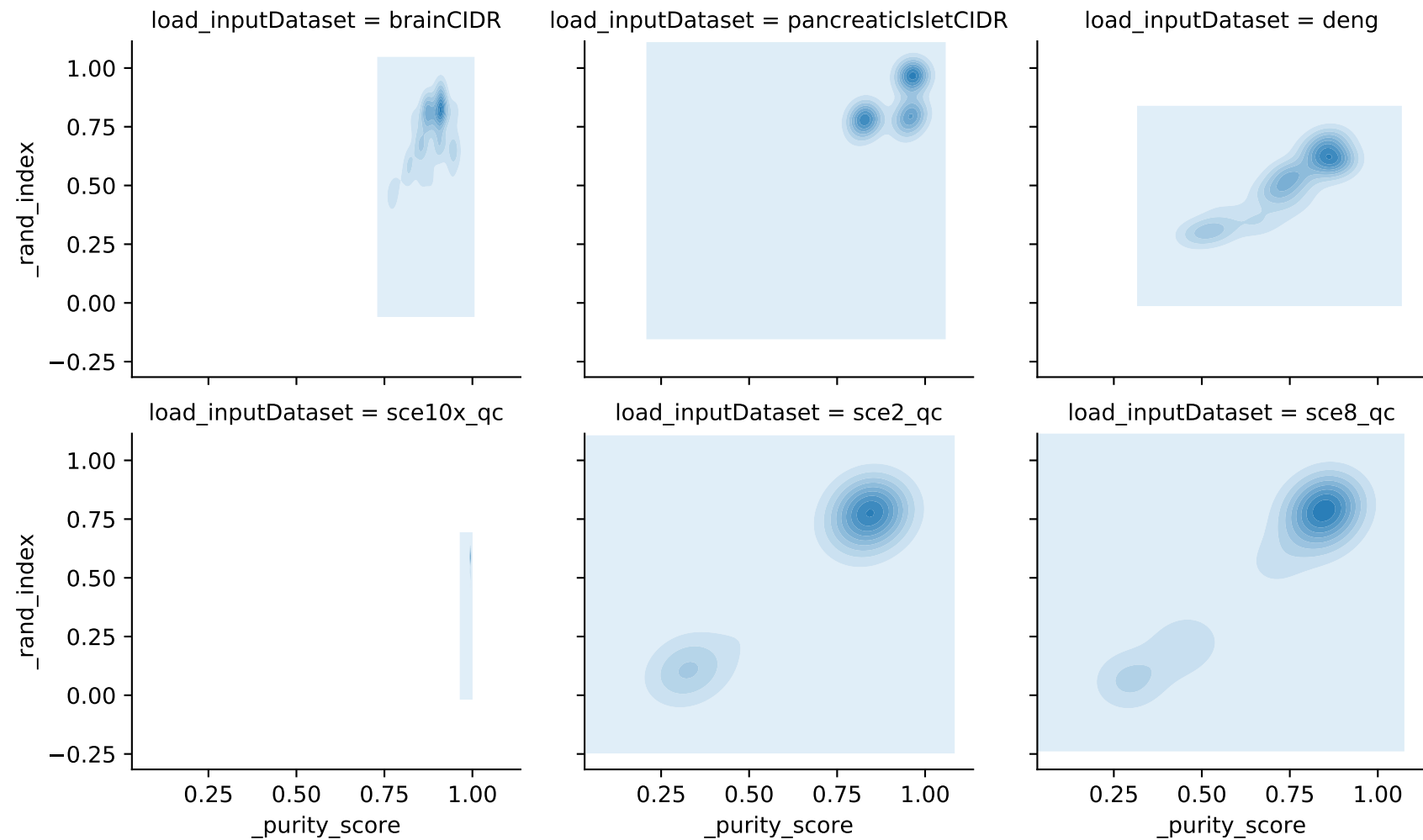
Distribution of _fowlkes_mallows_score



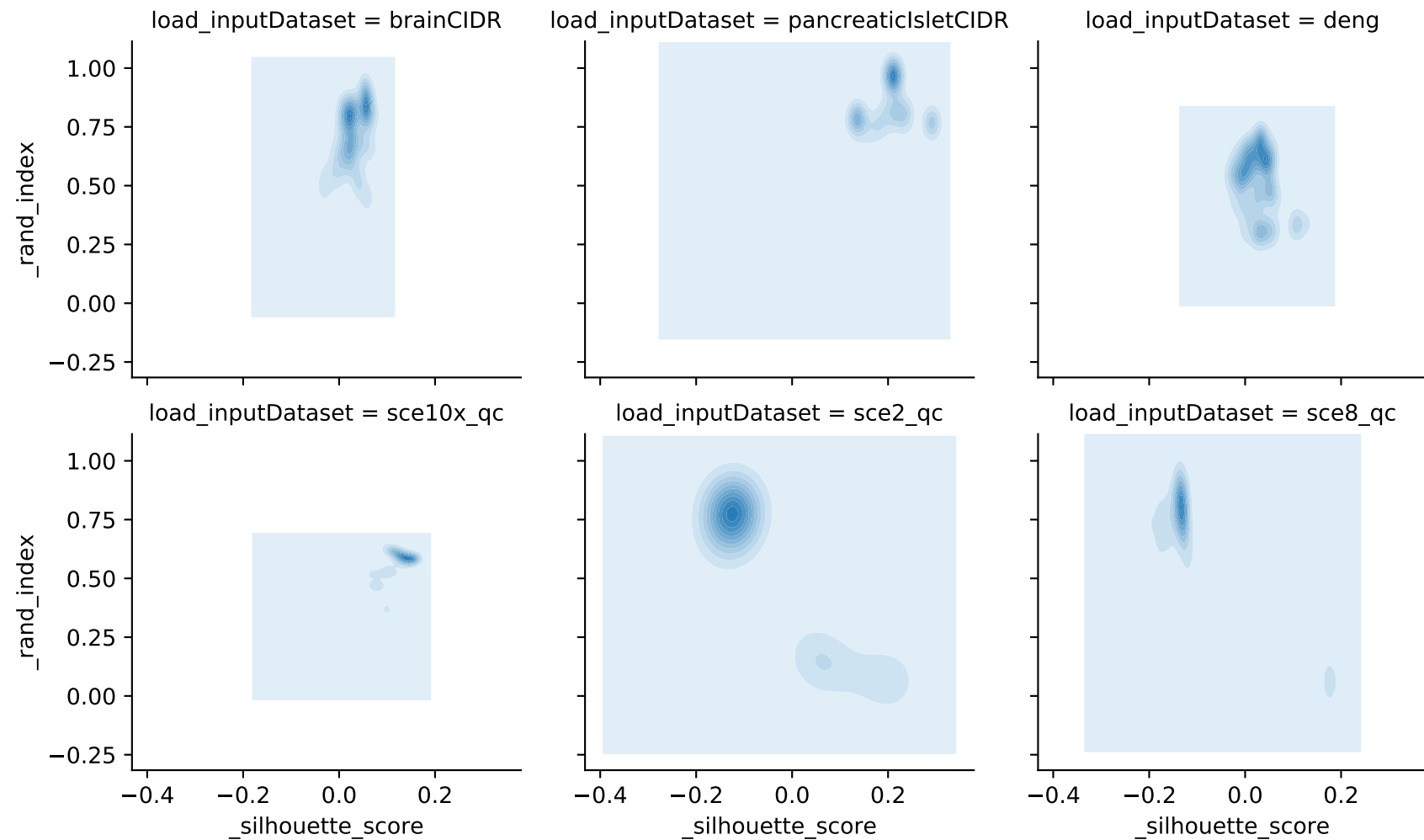
Distribution of _homogeneity_score



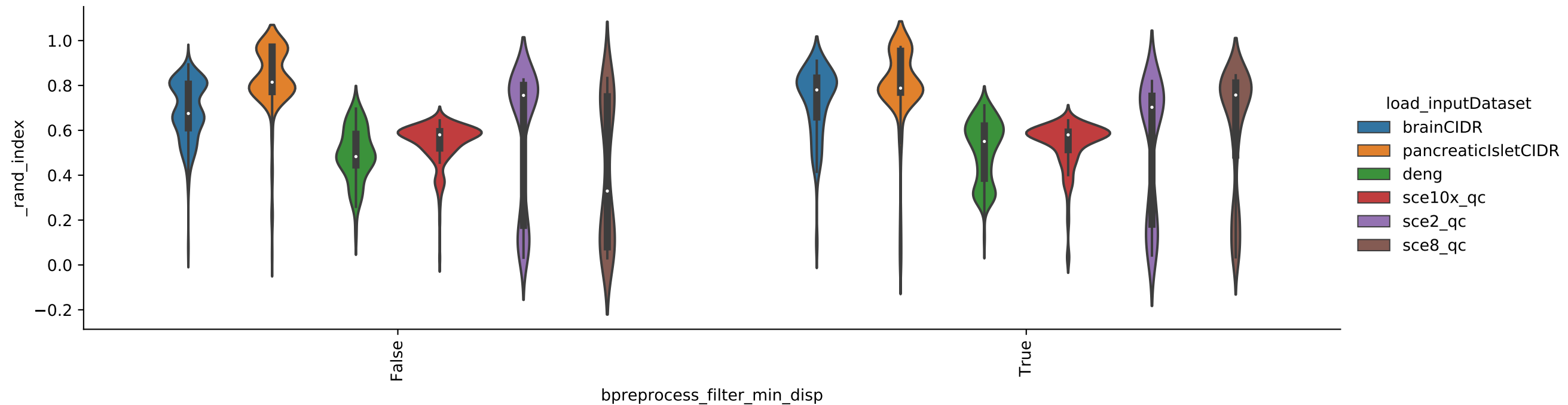
Distribution of _purity_score



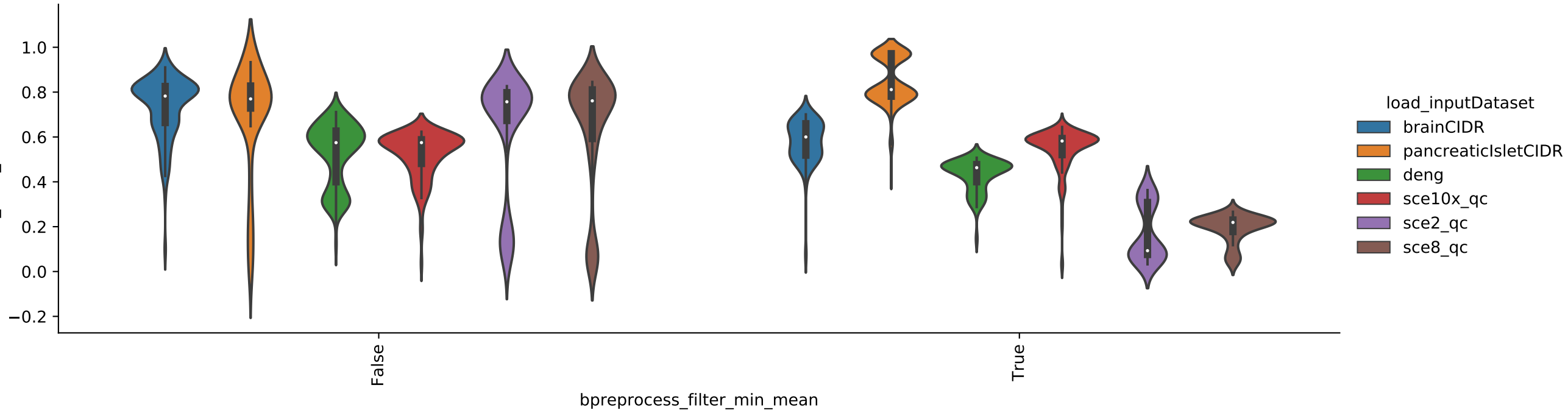
Distribution of _silhouette_score



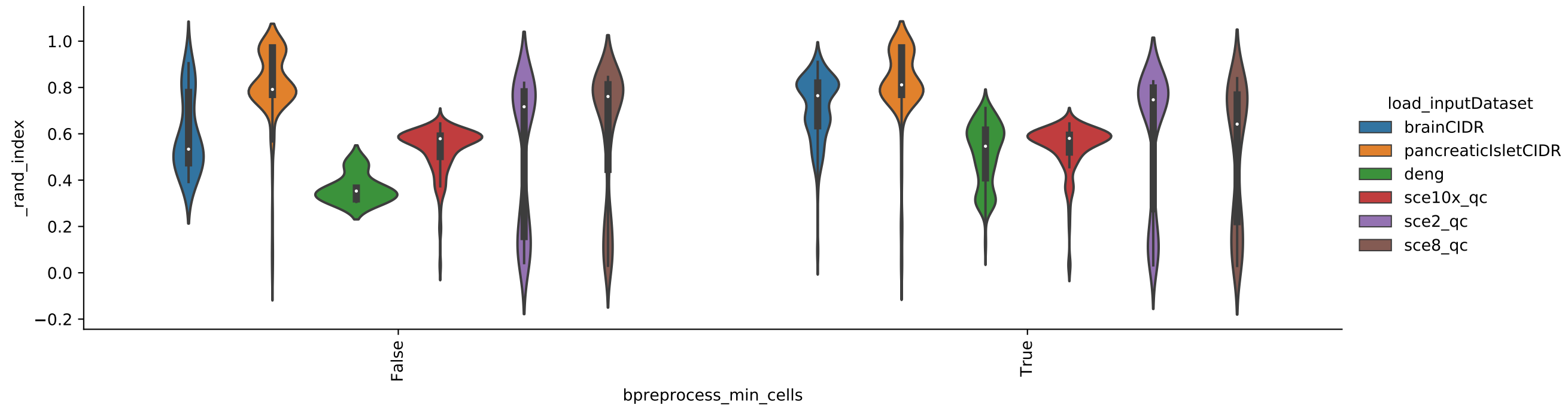
Distribution of bpreprocess_filter_min_disp



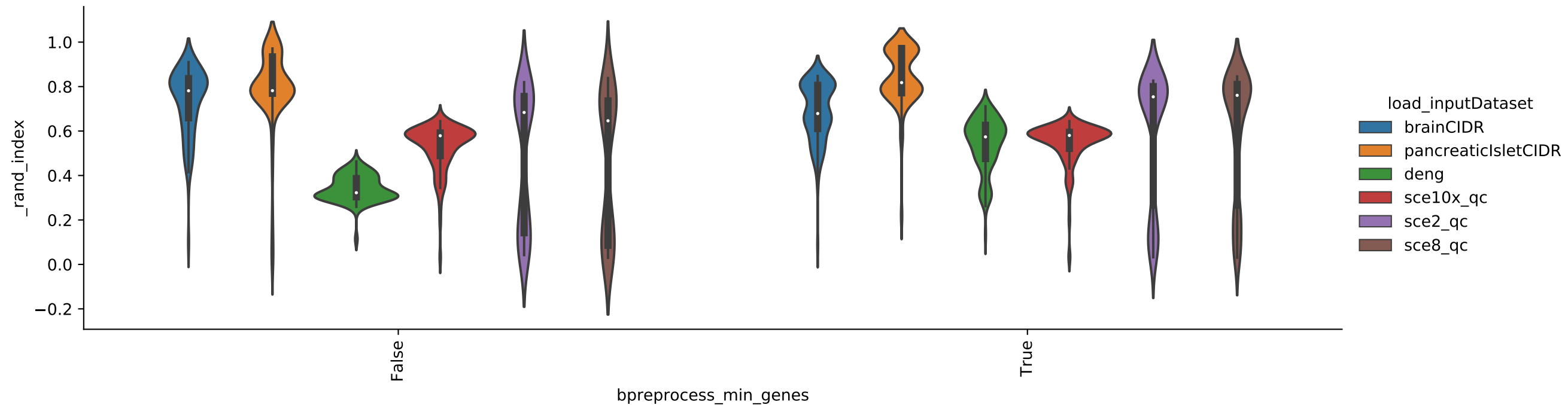
Distribution of bpreprocess_filter_min_mean



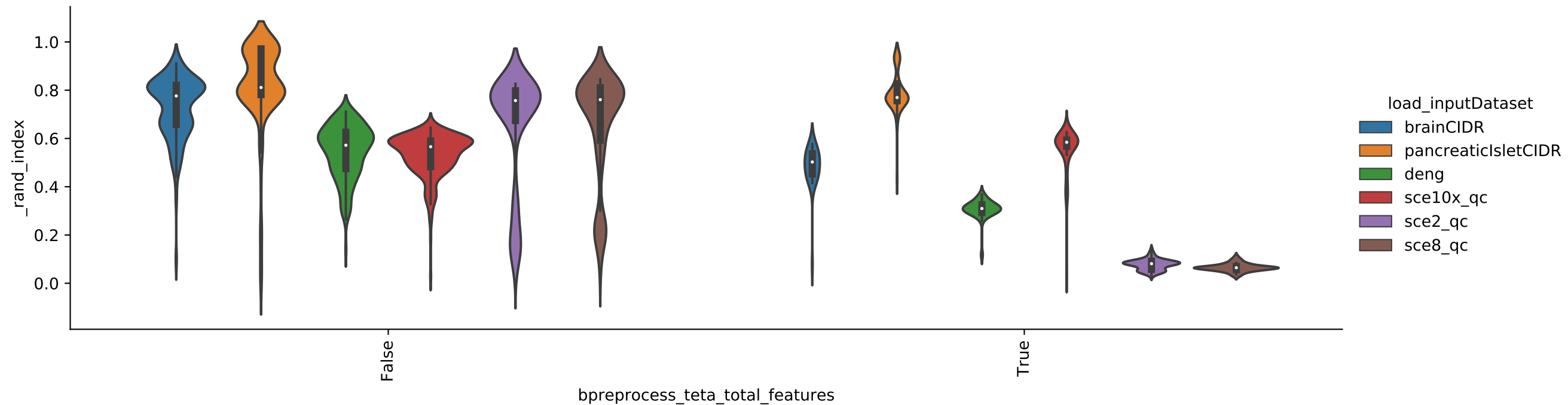
Distribution of bpreprocess_min_cells



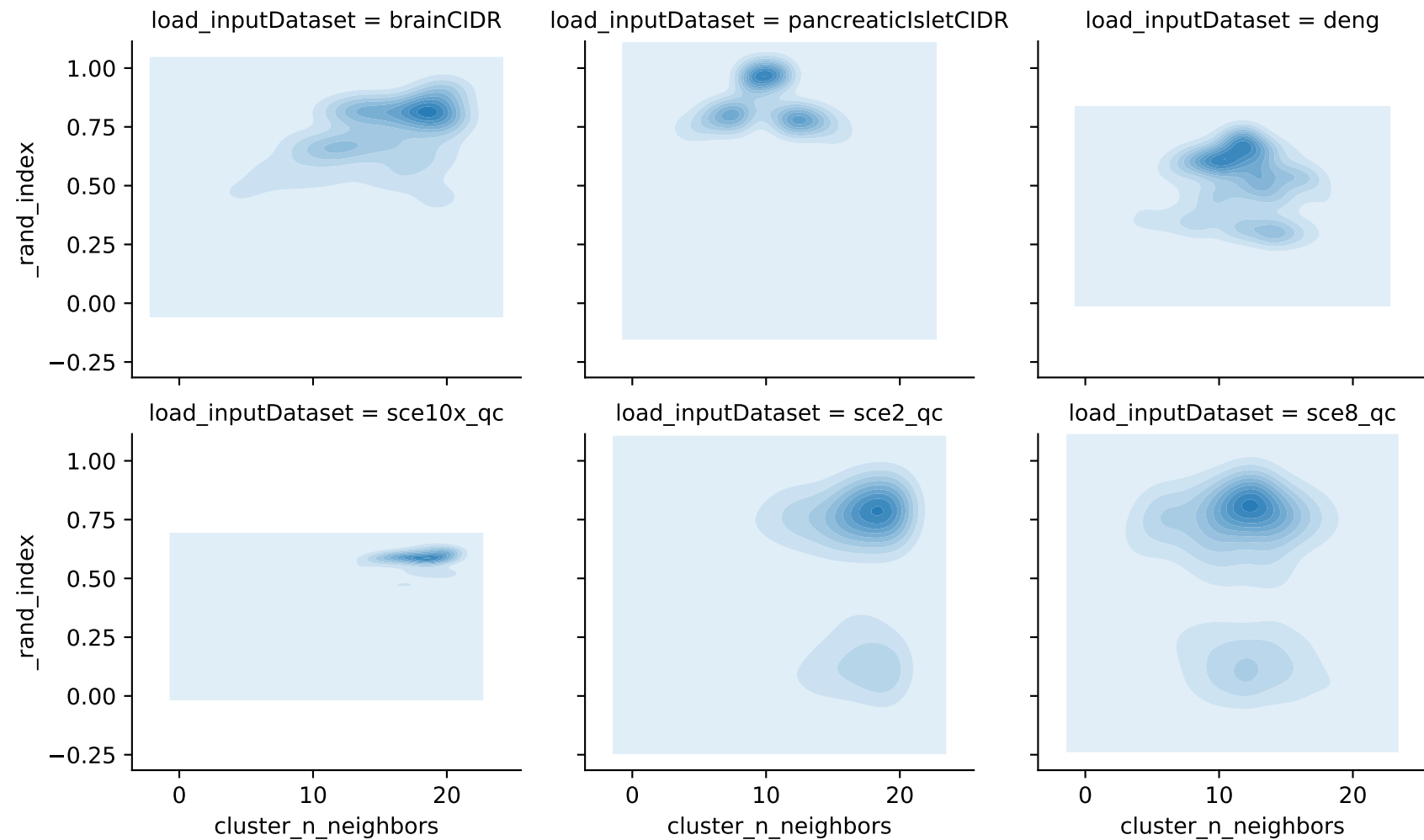
Distribution of bpreprocess_min_genes



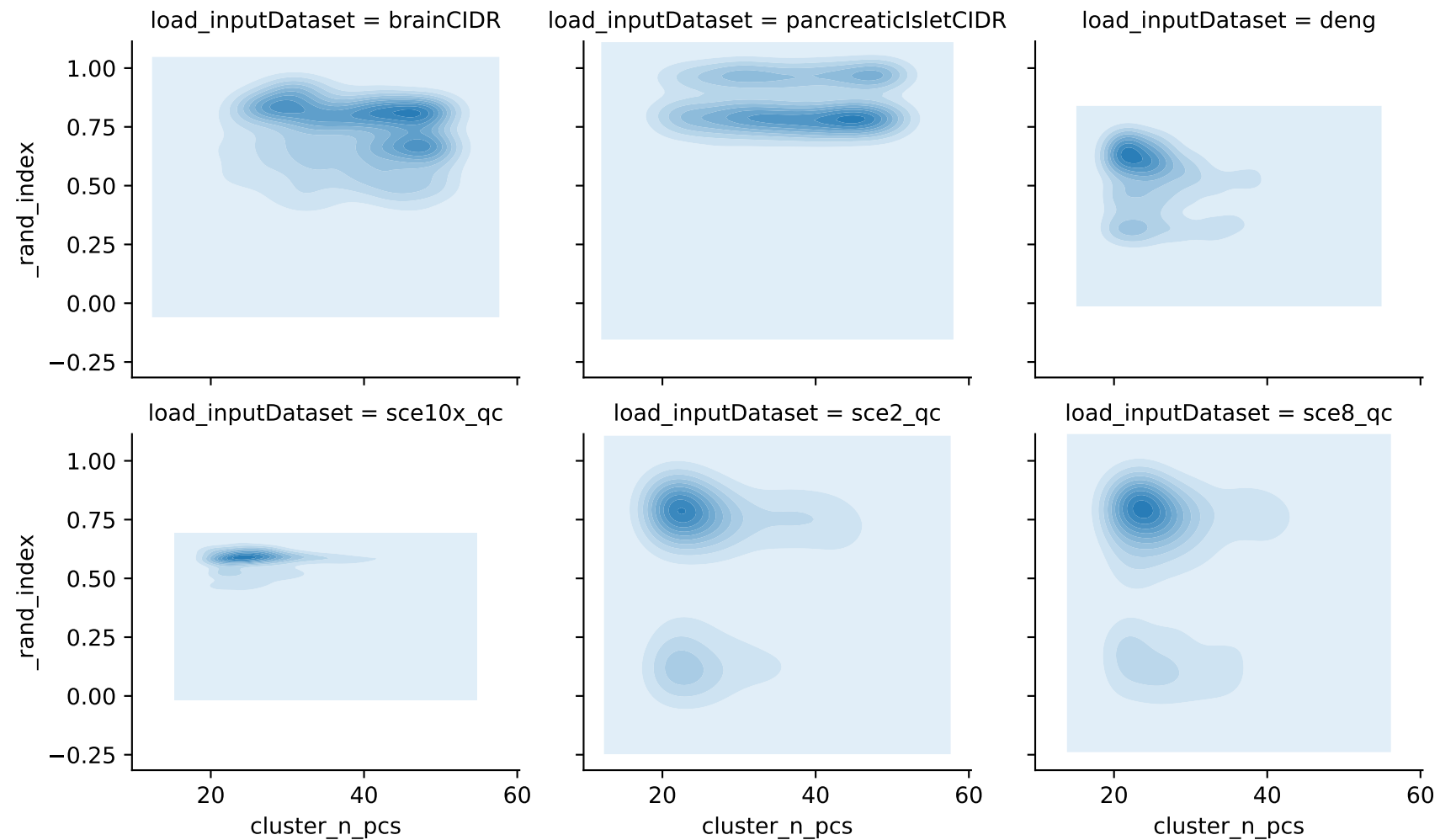
Distribution of bpreprocess_teta_total_features



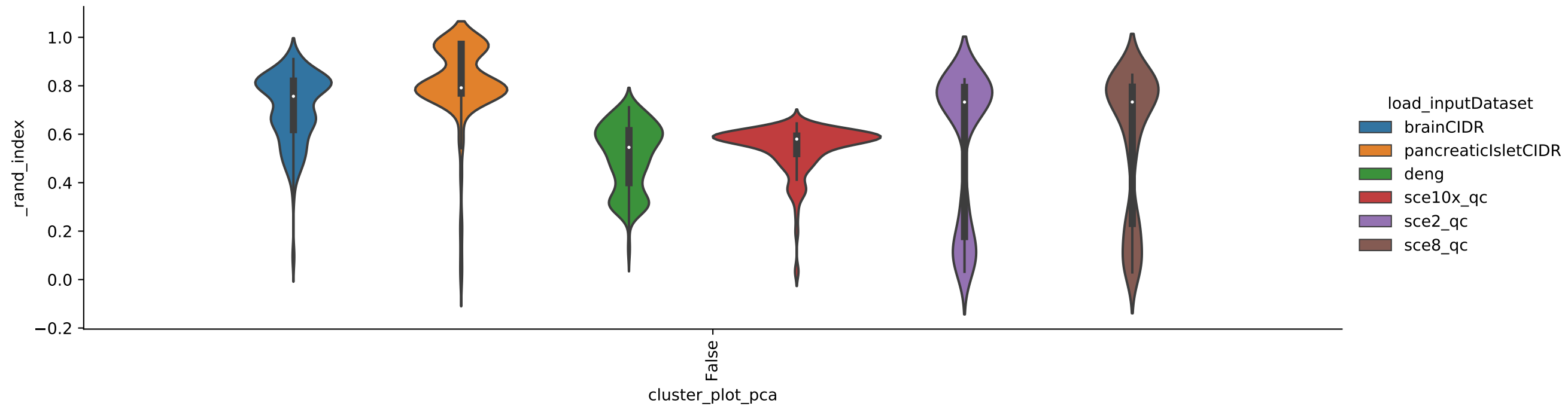
Distribution of cluster_n_neighbors



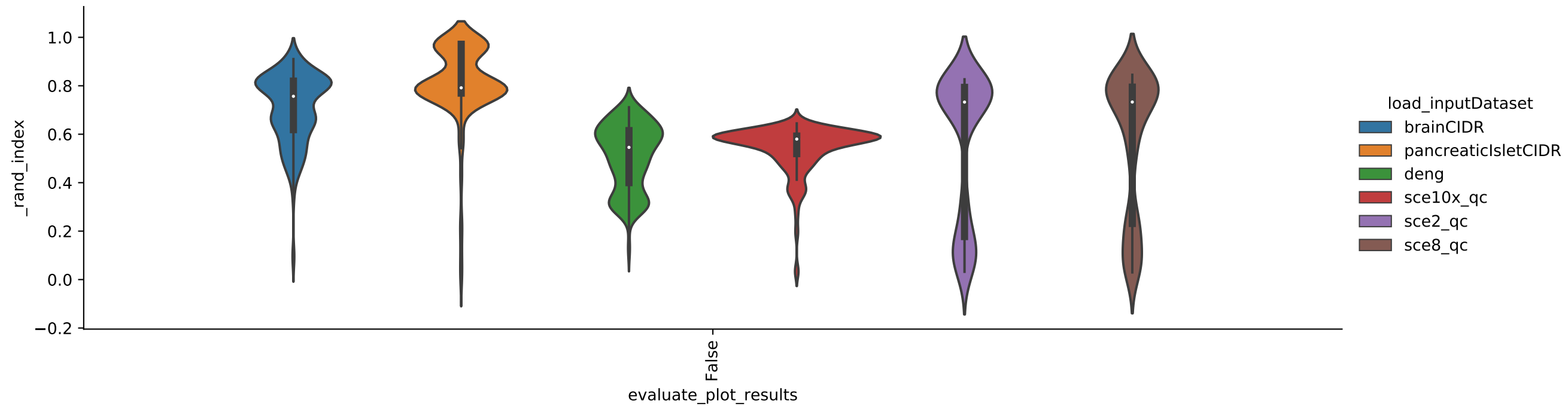
Distribution of cluster_n_pcs



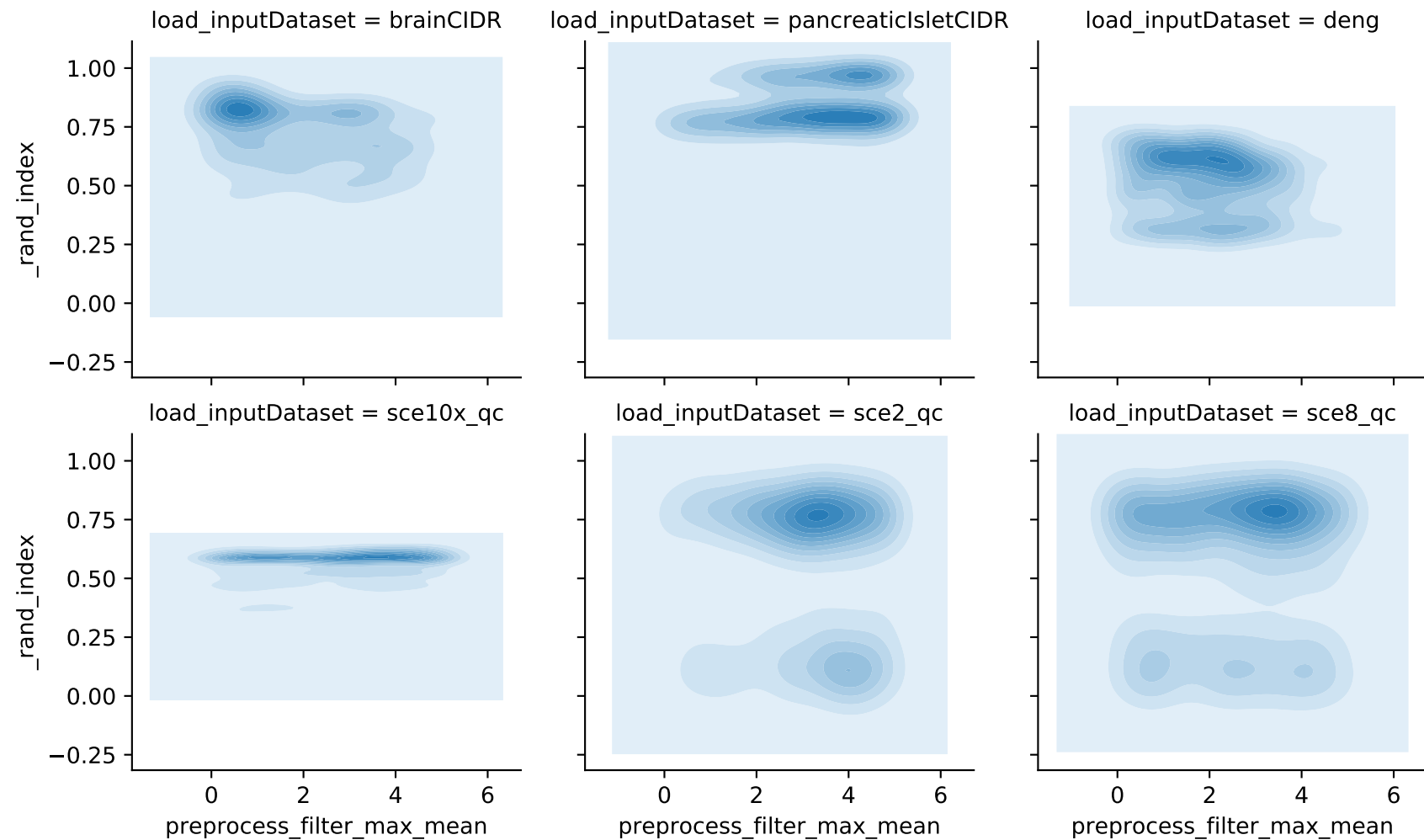
Distribution of cluster_plot_pca



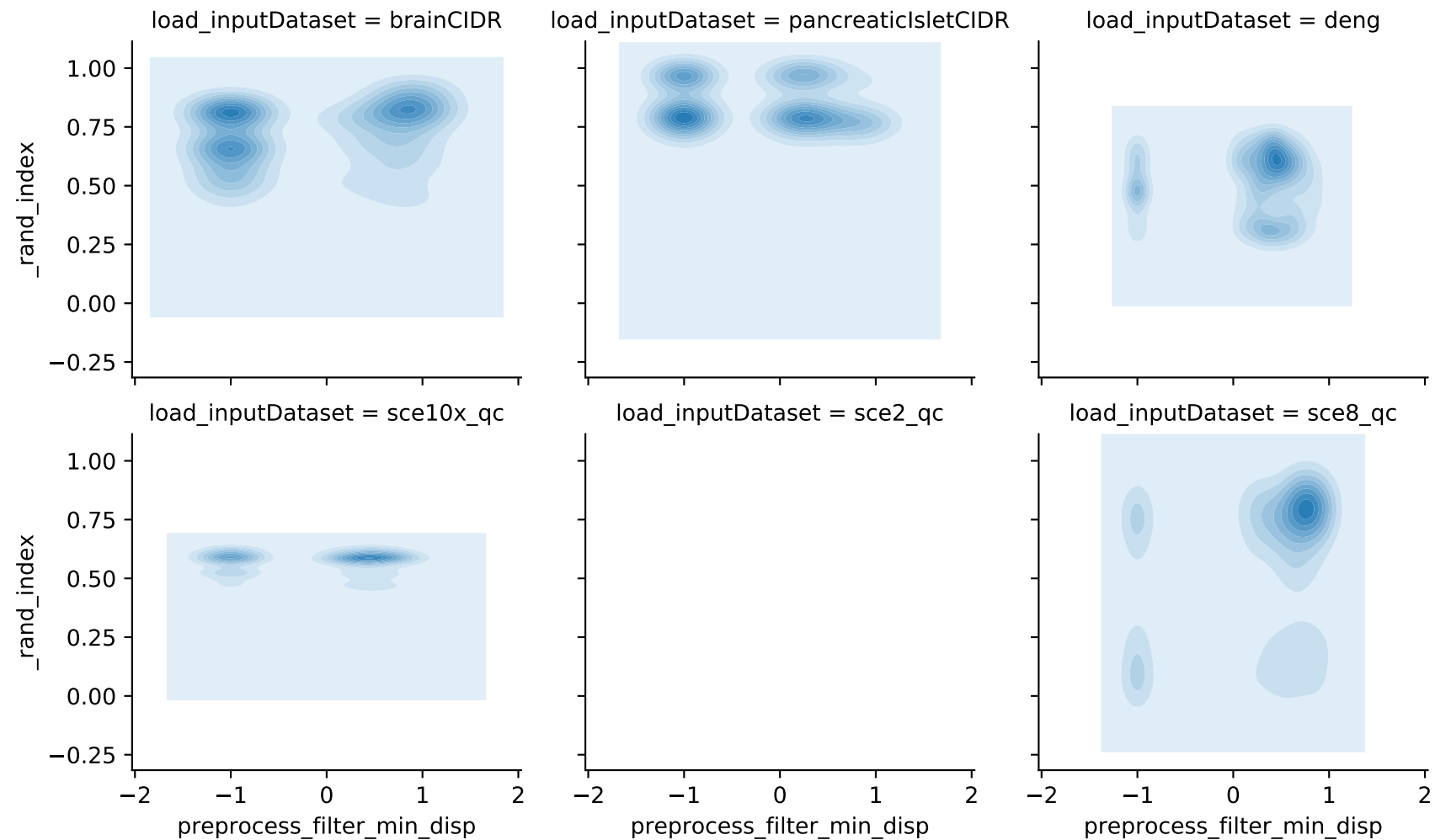
Distribution of evaluate_plot_results



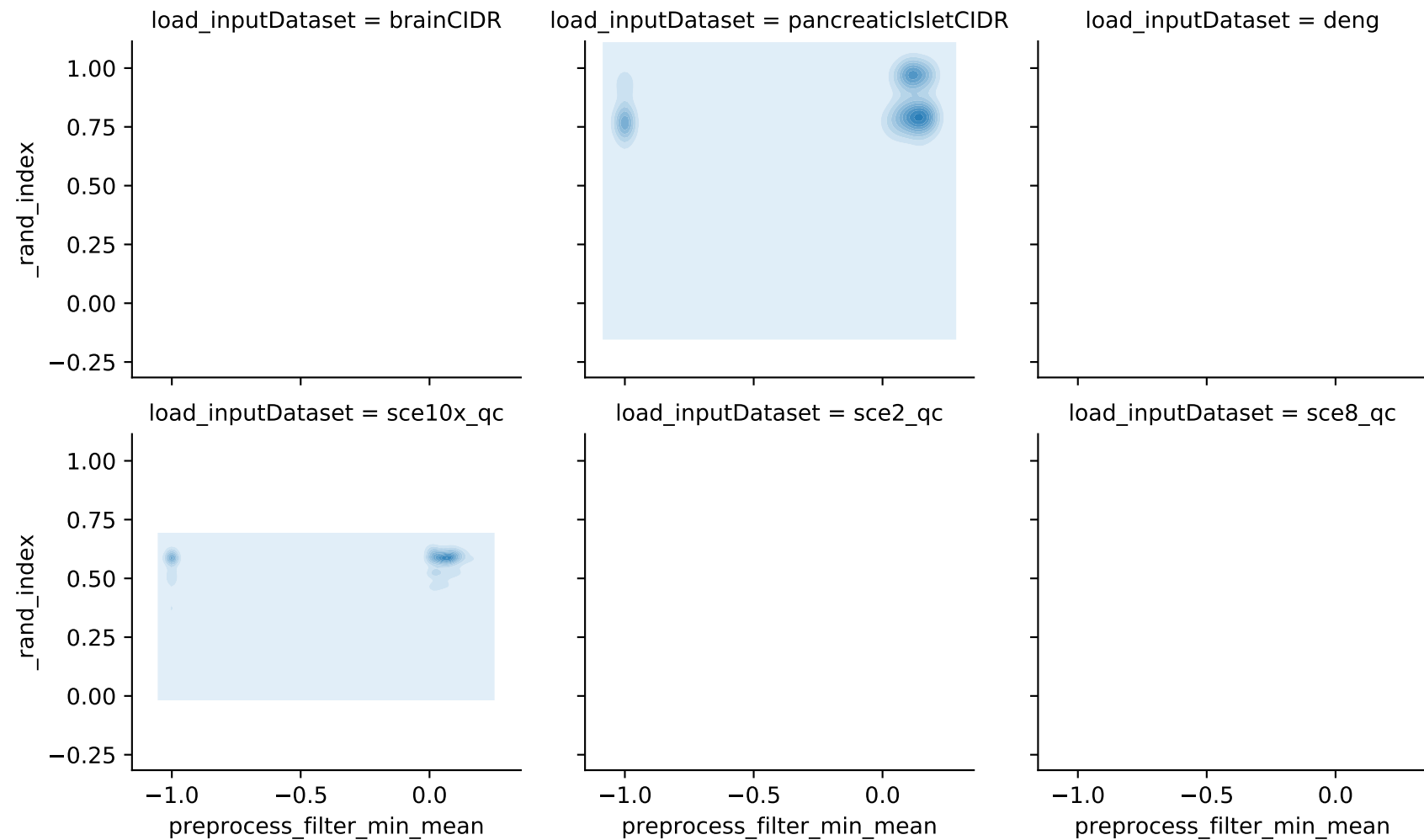
Distribution of preprocess_filter_max_mean



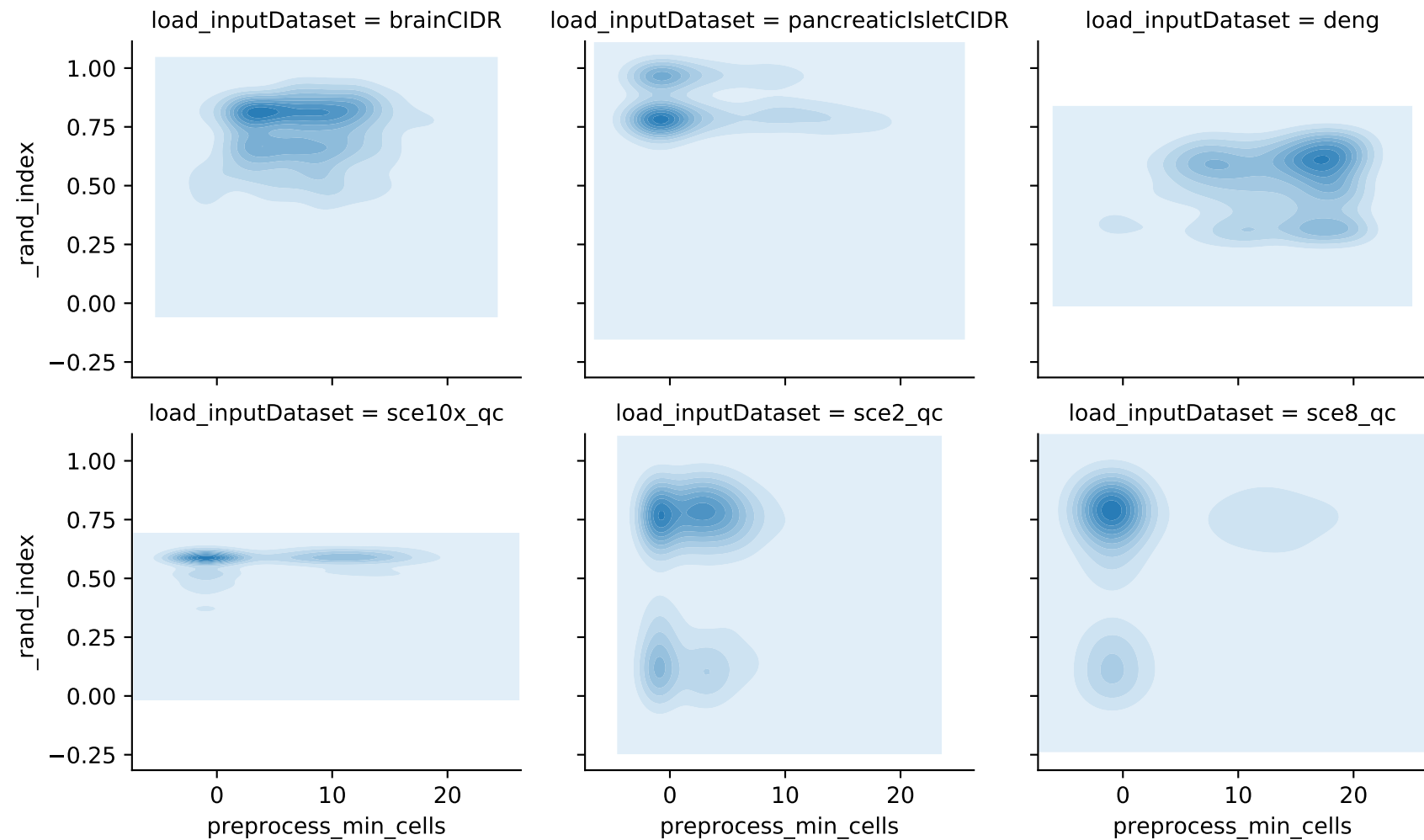
Distribution of preprocess_filter_min_disp



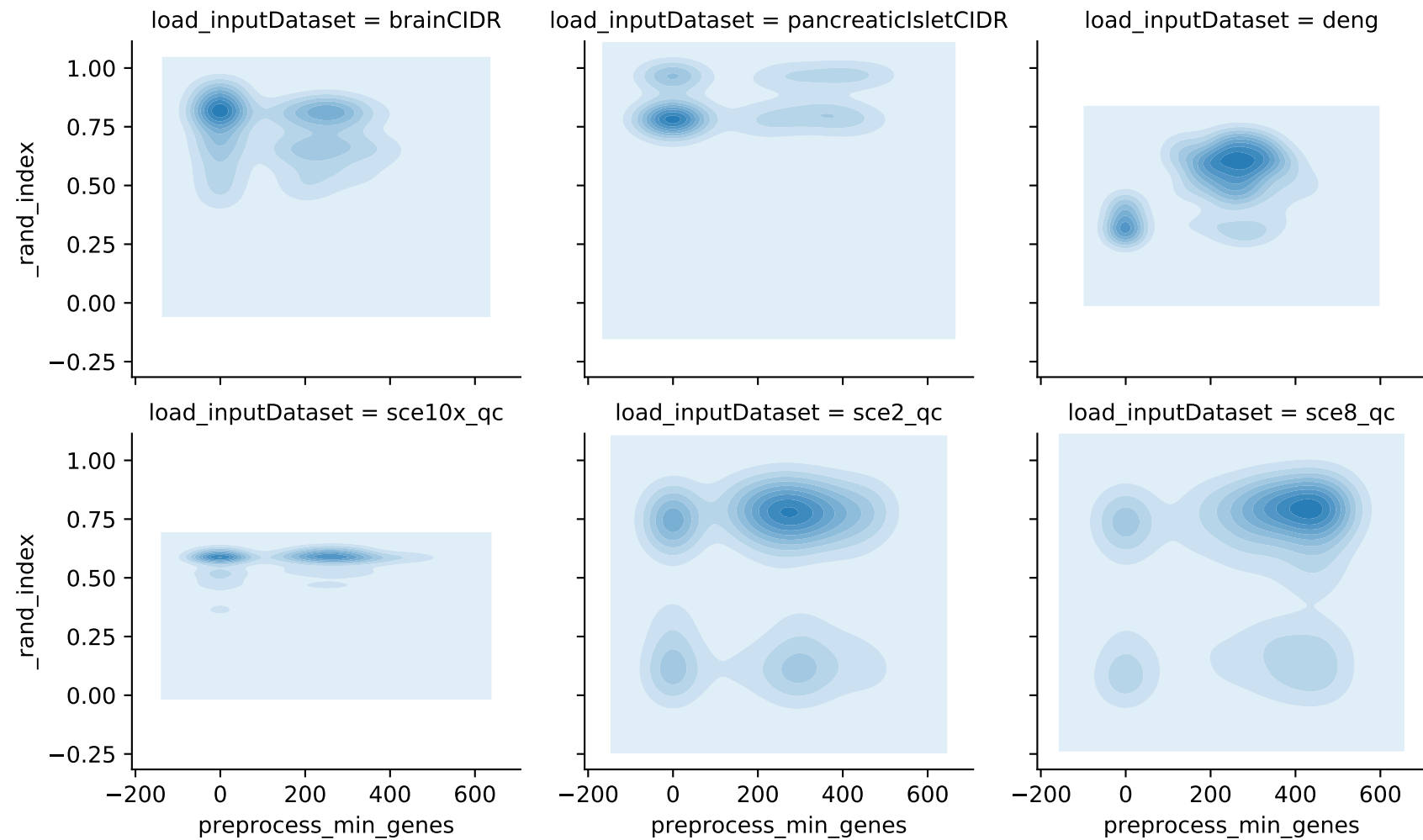
Distribution of preprocess_filter_min_mean



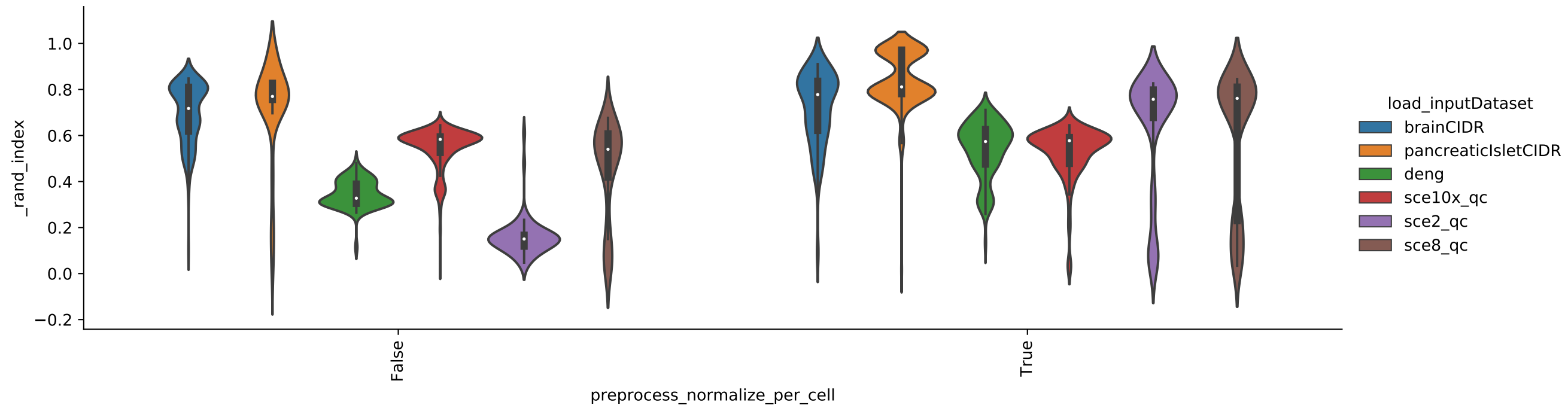
Distribution of preprocess_min_cells



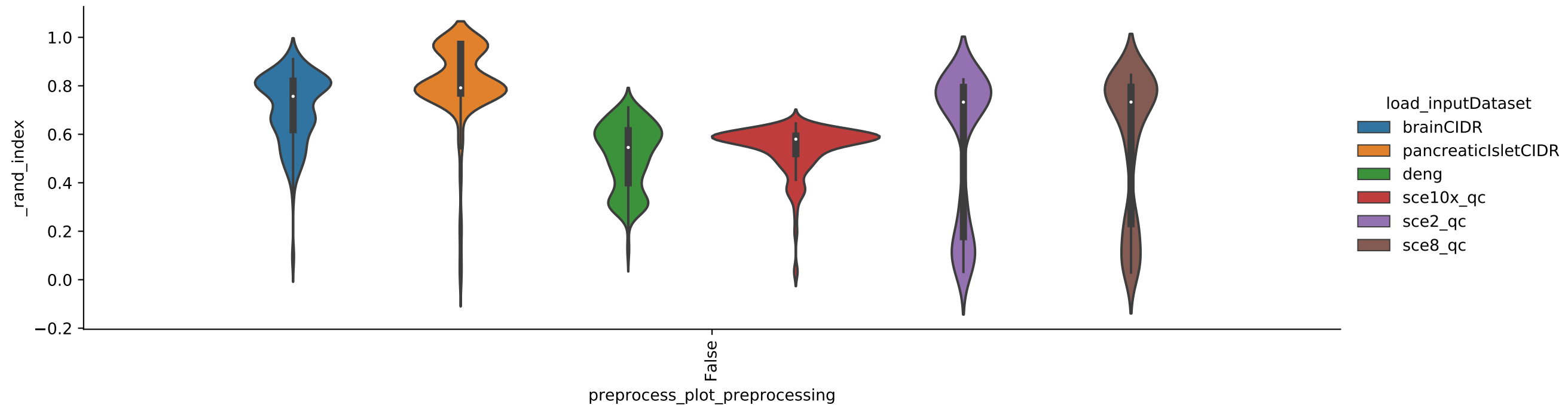
Distribution of preprocess_min_genes



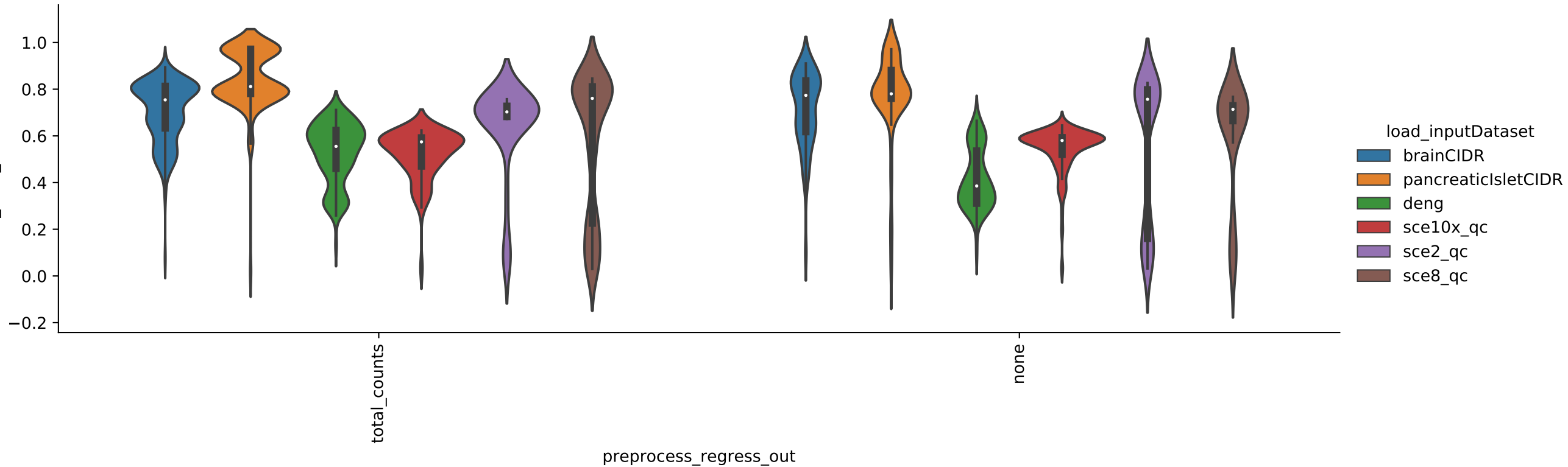
Distribution of preprocess_normalize_per_cell



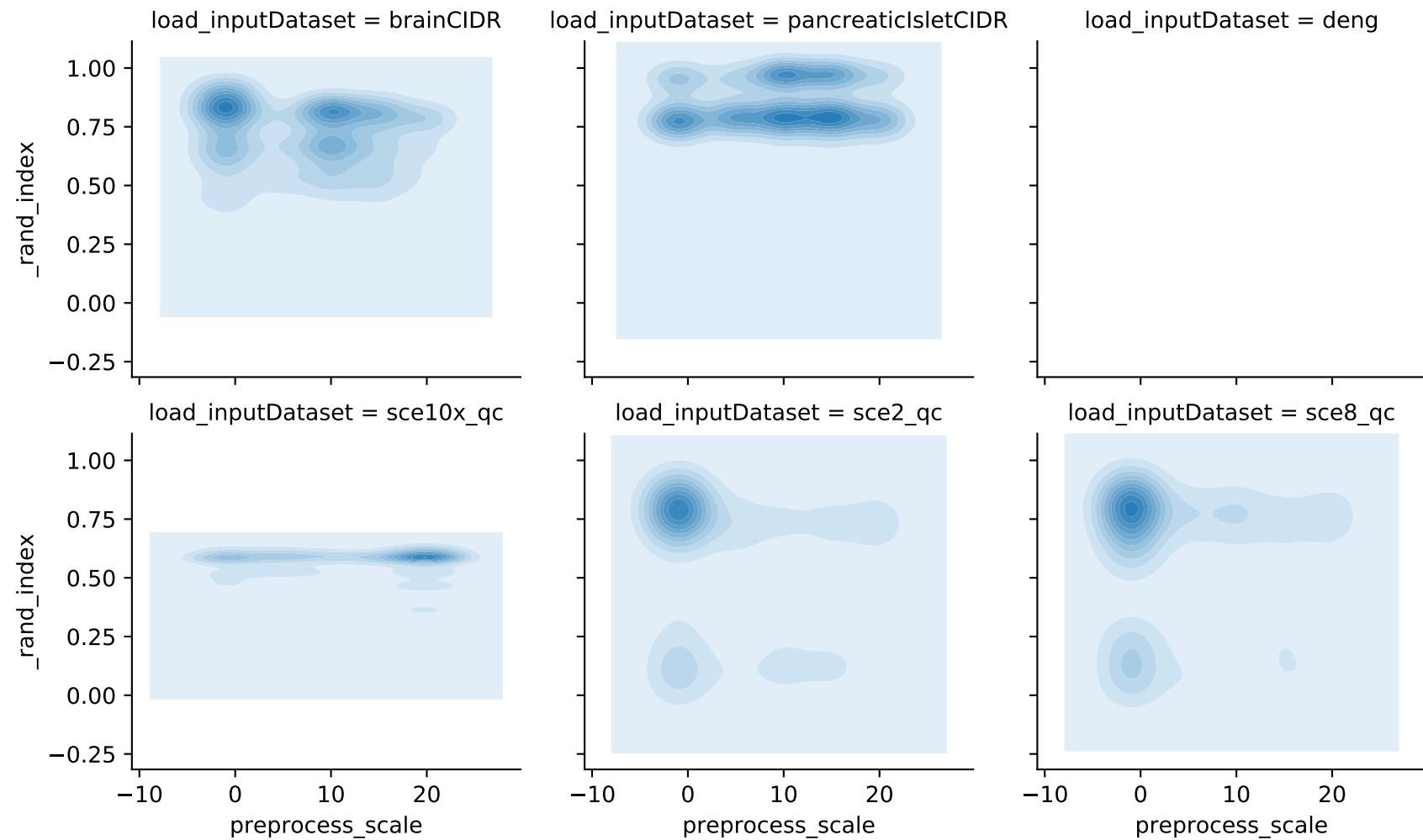
Distribution of preprocess_plot_preprocessing



Distribution of preprocess_regress_out



Distribution of preprocess_scale



Distribution of preprocess_teta_total_features

