Best K-means clustering plot 0 cluster 103 **54** 105 45 + -2 **-**PC1

PCA based on SNPs present in candidate structural variant 0.1 Sa21_3 So8_12 So8_6 Sa8_12 Sa6_7 So10_9 So21_20_30 So8_10 So21_11 So10_30 So10_14 So21_8 So10_26 So21_26 So8_14 Sa21_12 Sa8_10 **Groups** So10_16⁷ hom1 Sa21_5 So10_19 hz hom2 So10_25 So10_5 So10 11



