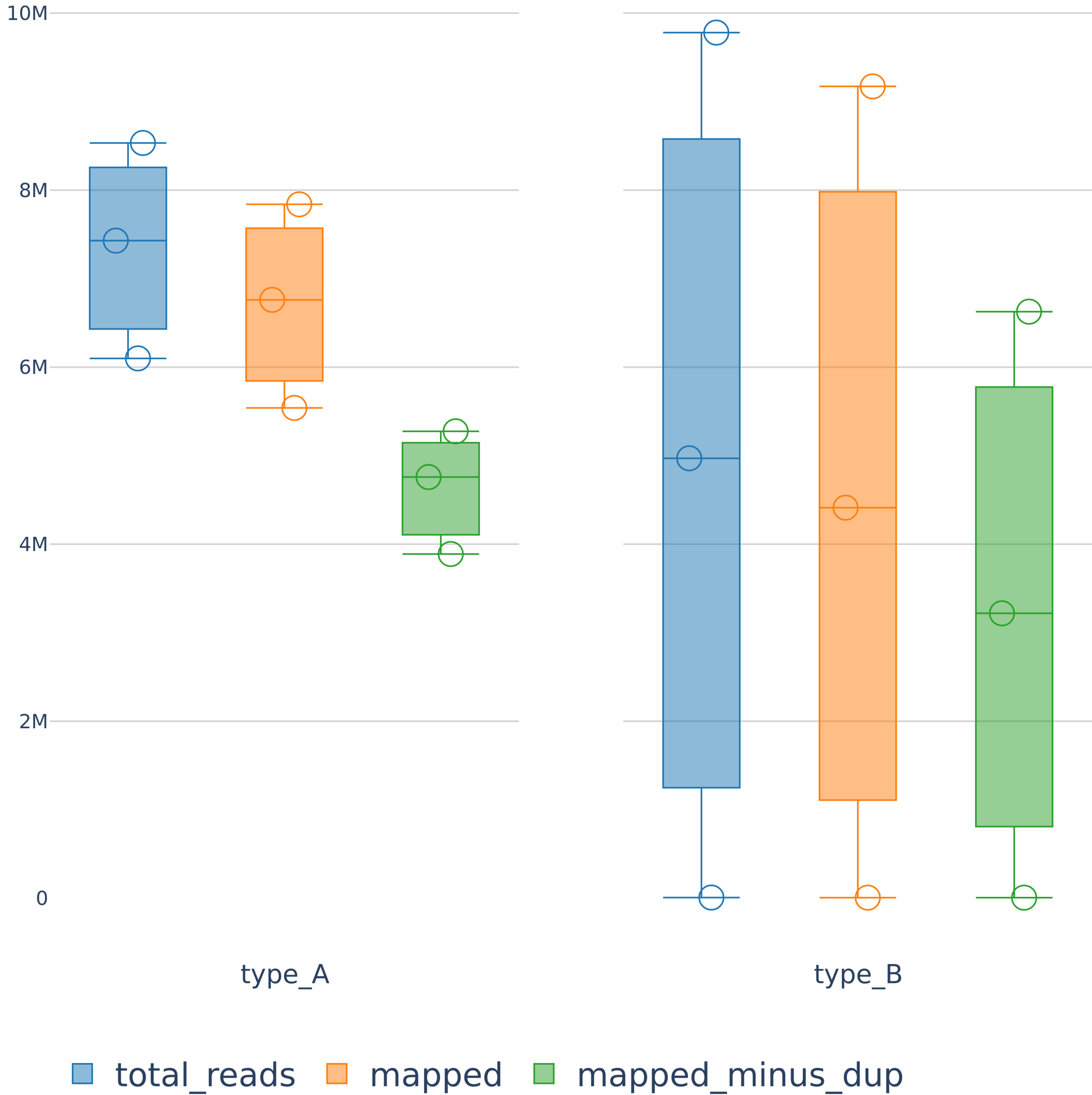
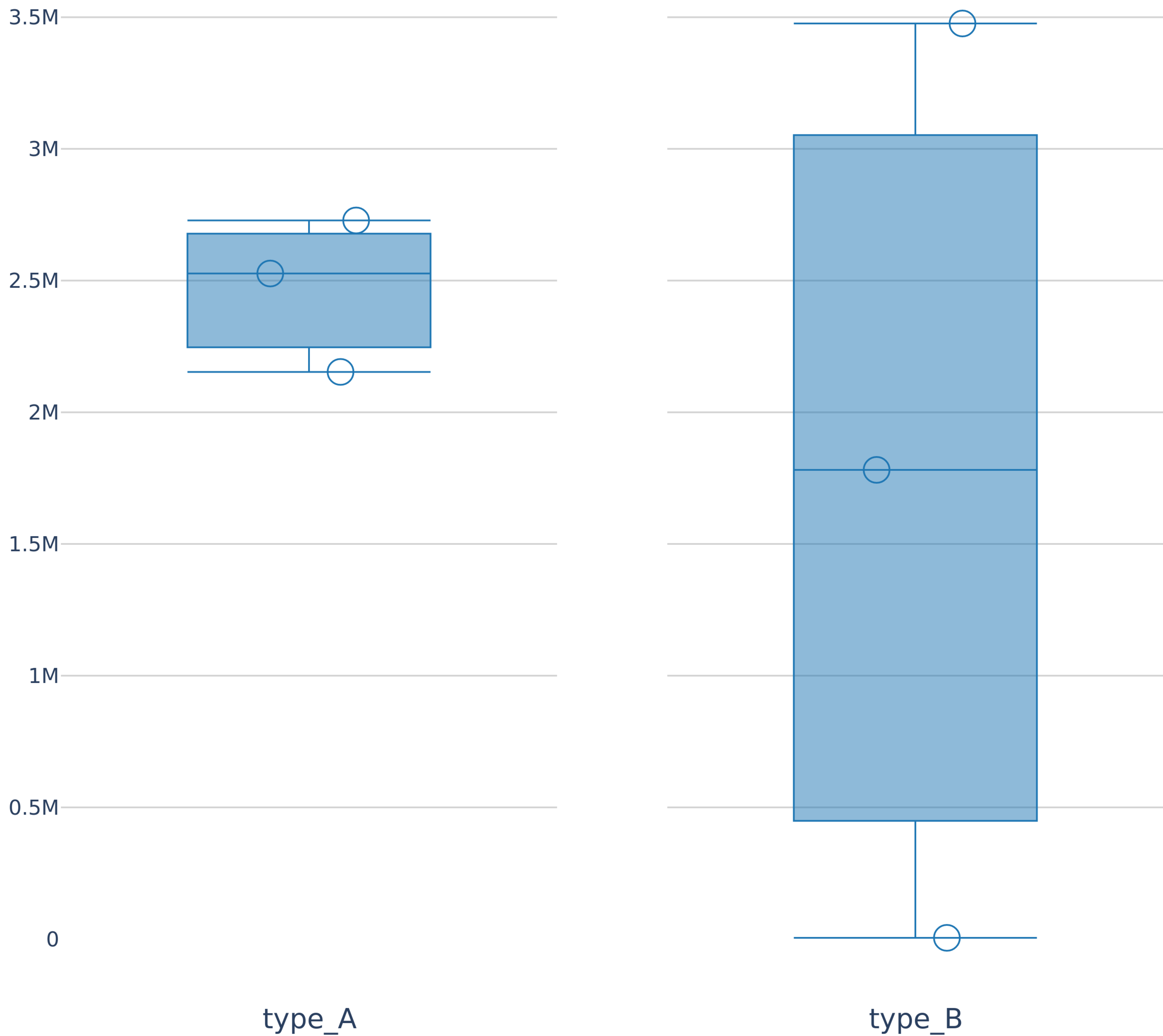


EXAMPLE total_reads,mapped,mapped_minus_dup

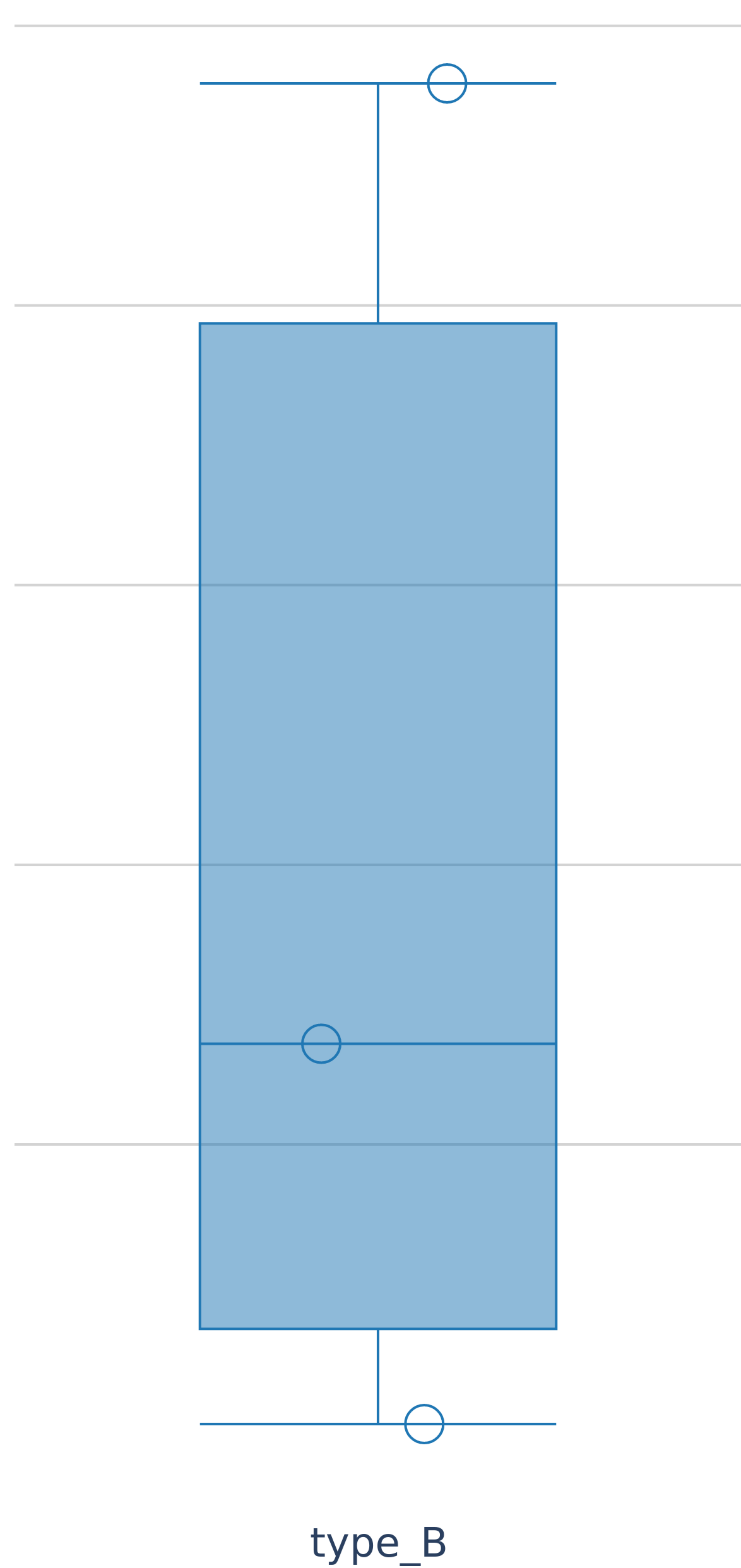
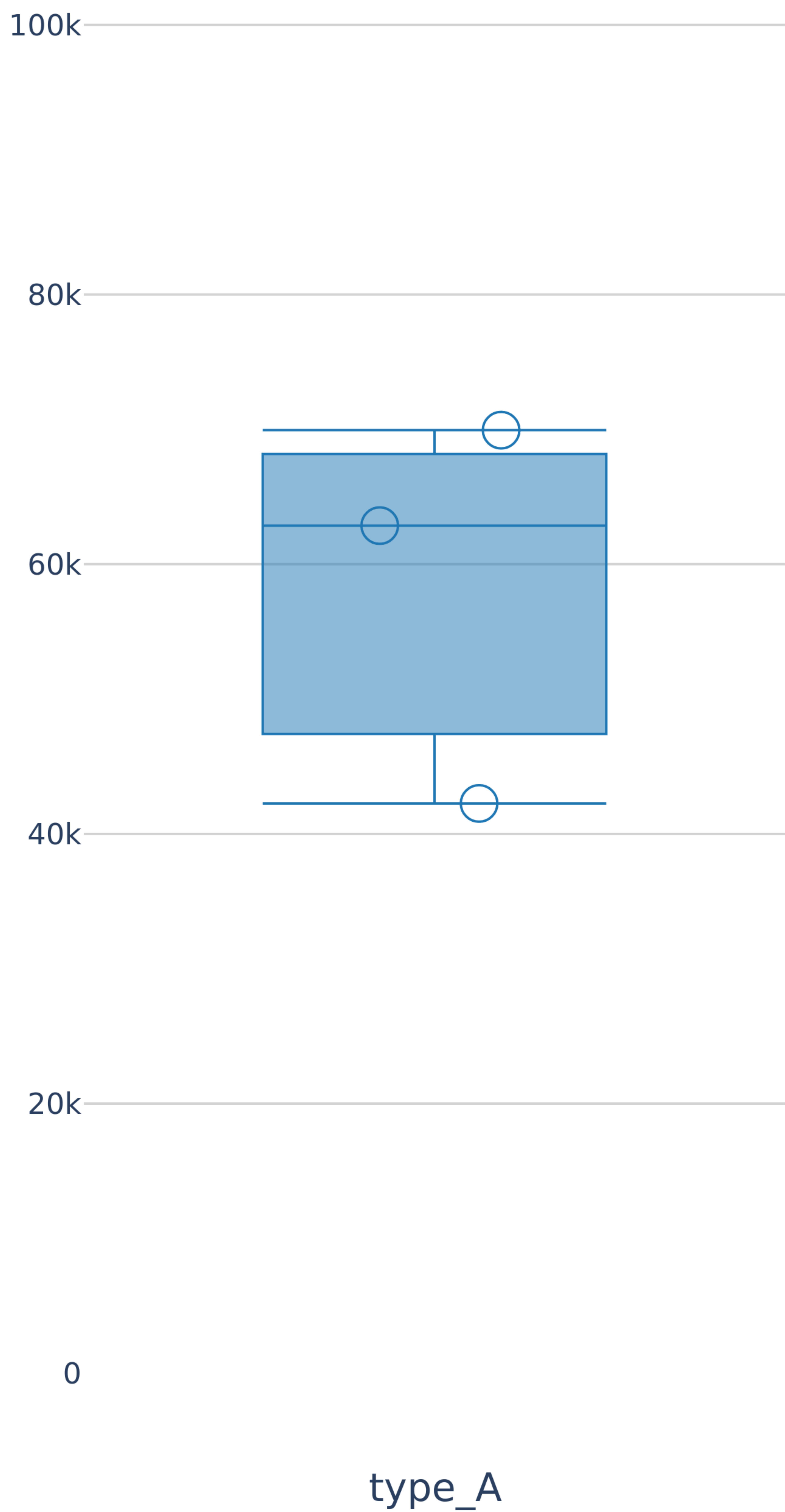


EXAMPLE cpg_count



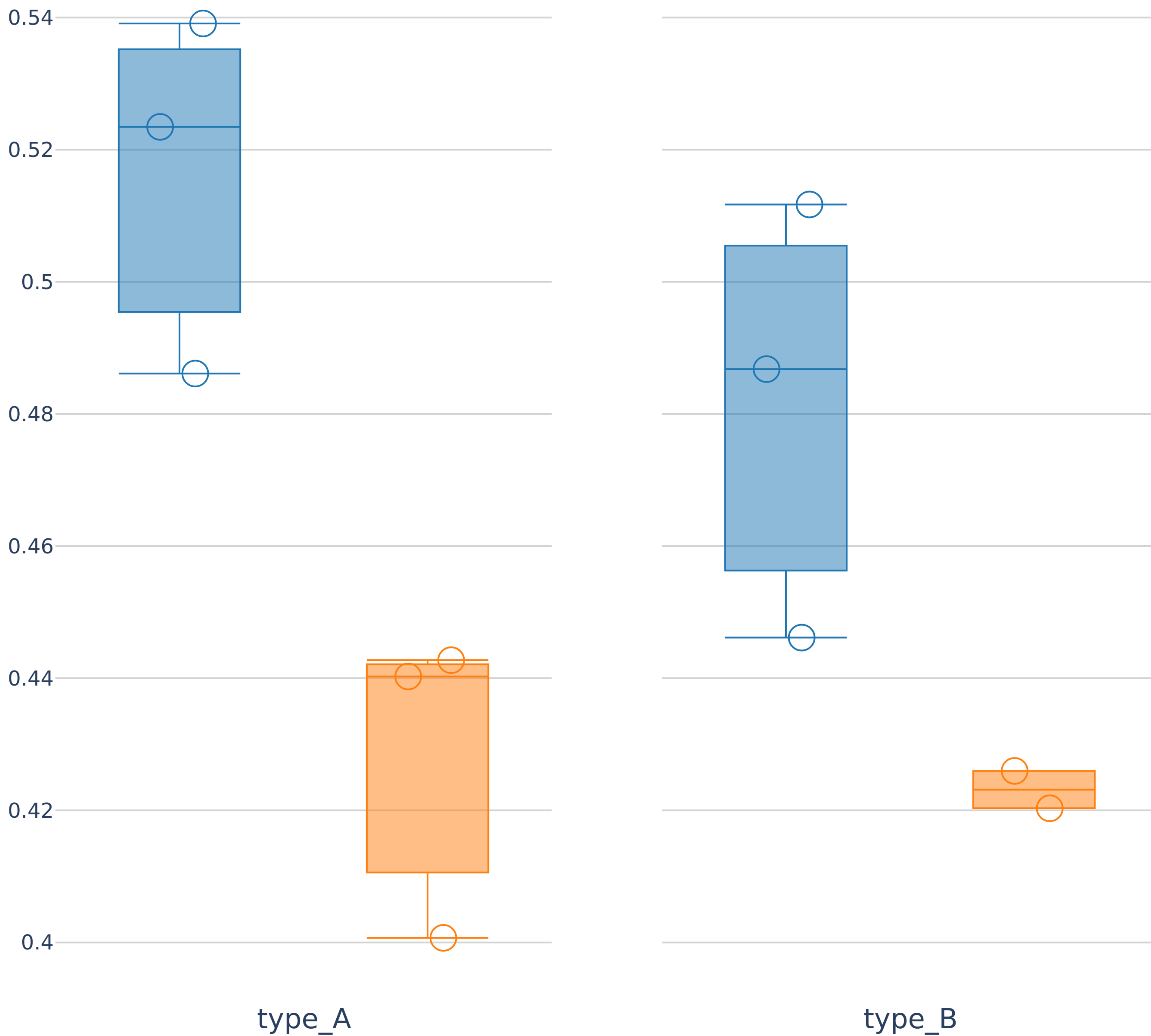
■ cpg_count

EXAMPLE cpg_count_cov3



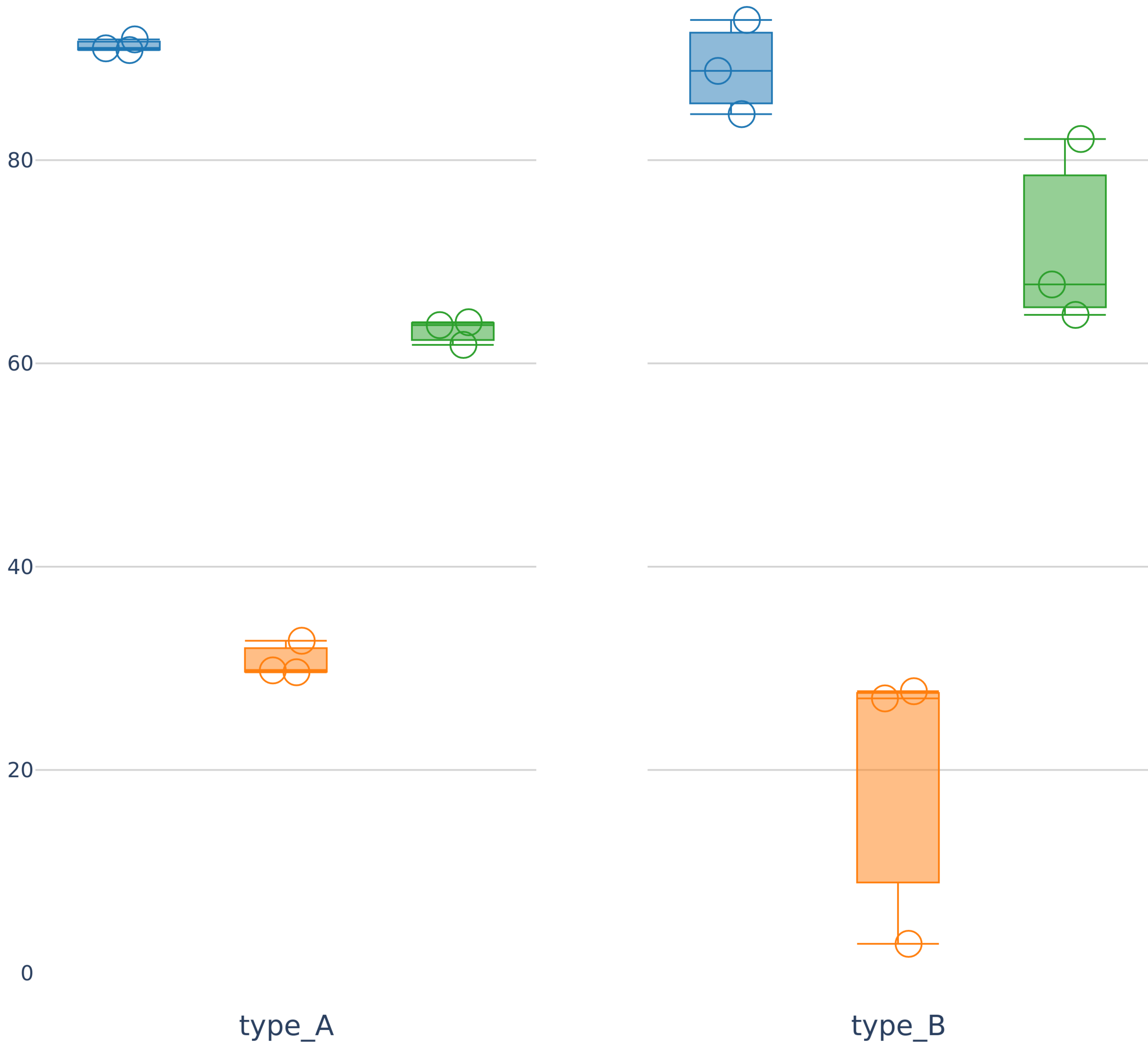
■ cpg_count_cov3

EXAMPLE average_meth,average_meth_cov3



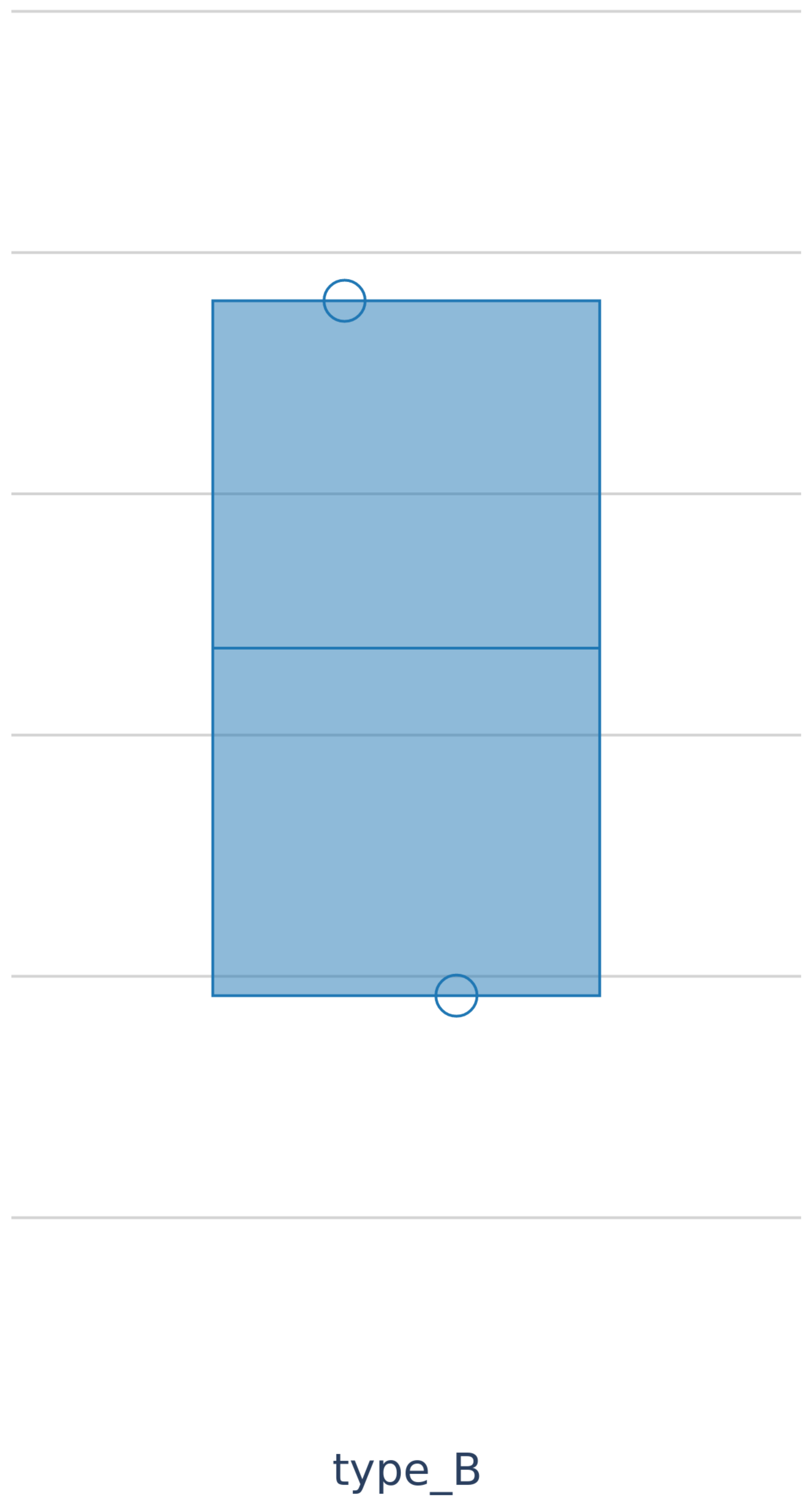
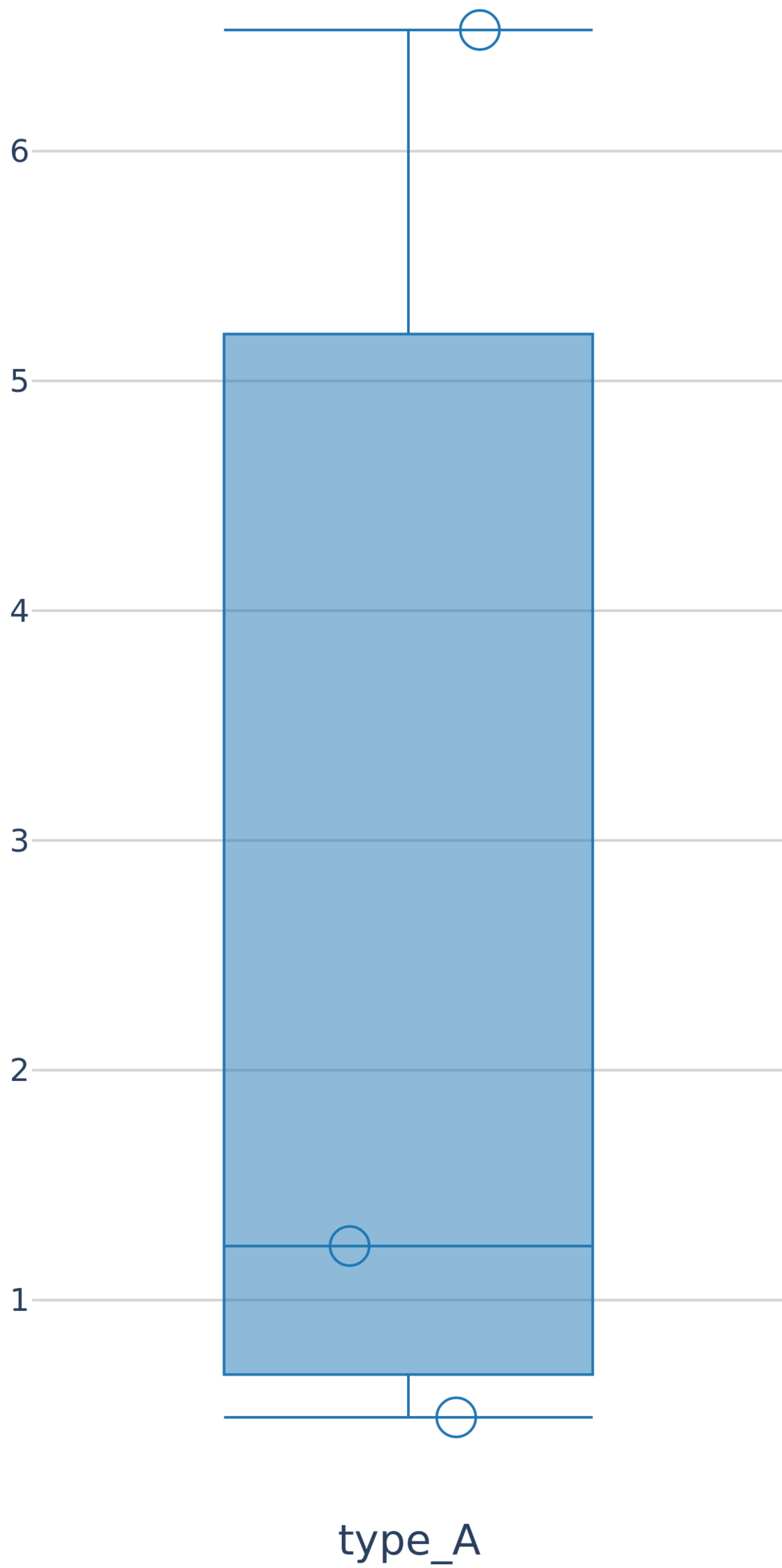
■ average_meth ■ average_meth_cov3

EXAMPLE mapped%,dup_rate%,mapped_minus_dup%



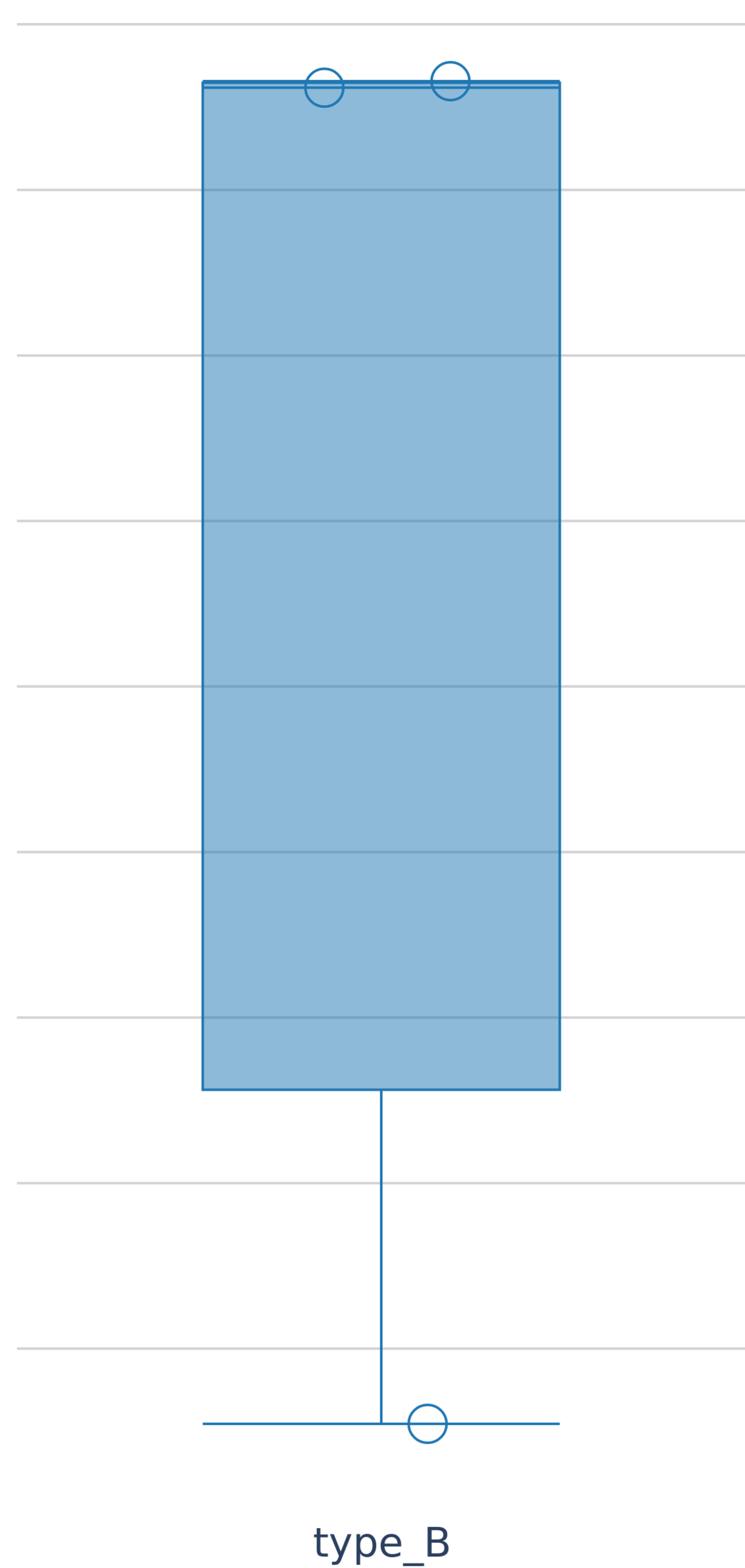
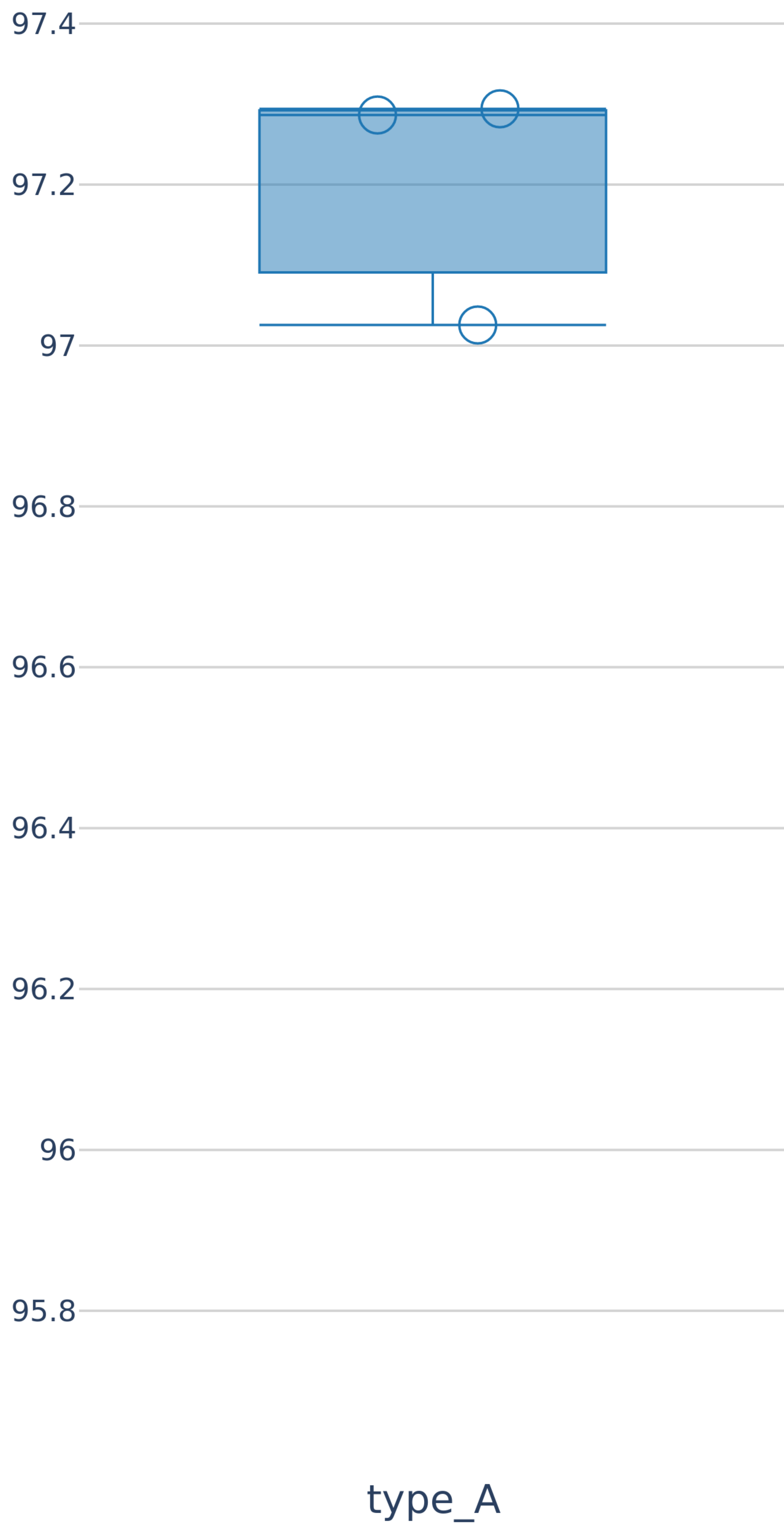
■ mapped% ■ dup_rate% ■ mapped_minus_dup%

EXAMPLE T7_conversion



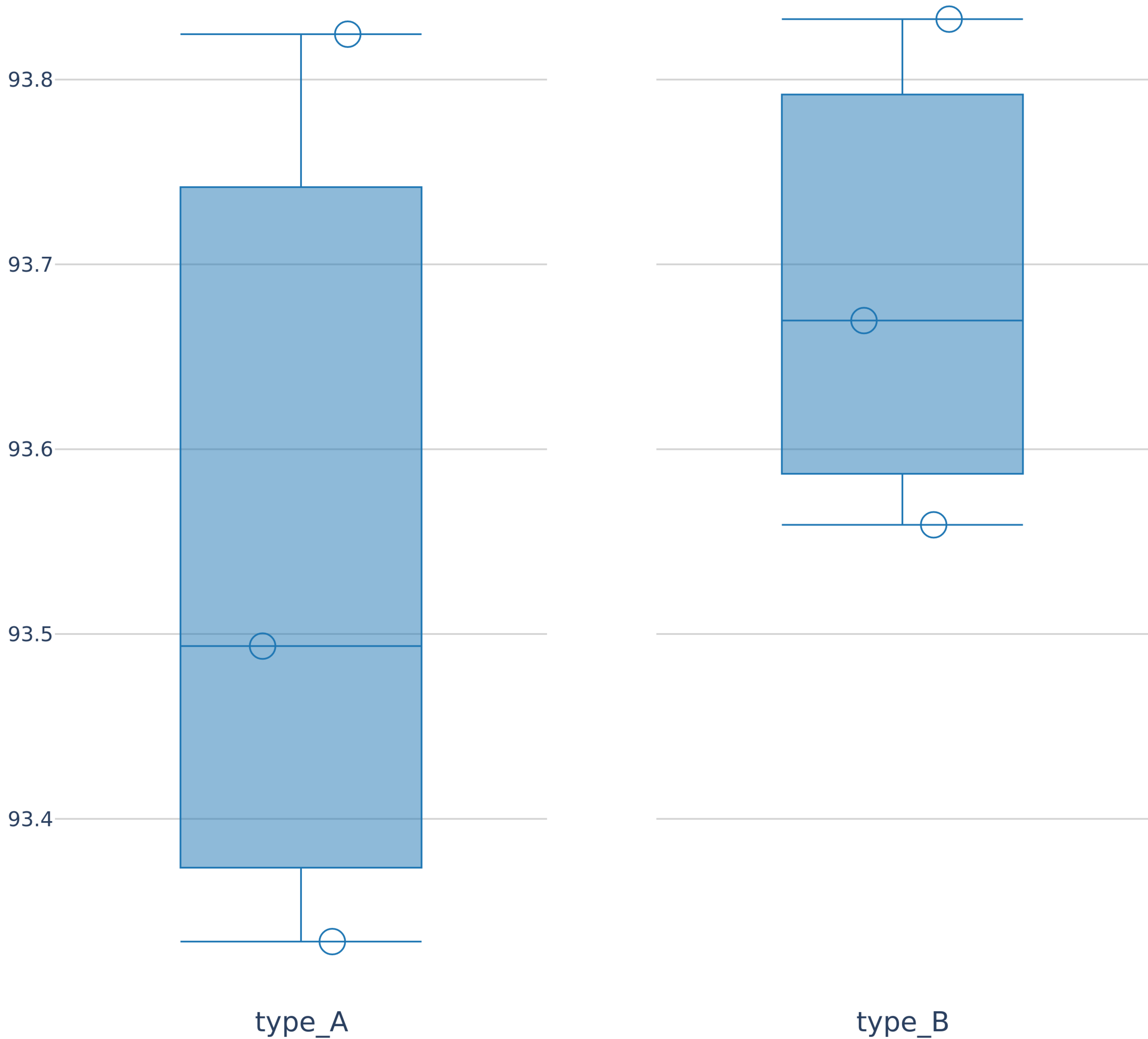
■ T7_conversion

EXAMPLE lambda_conversion



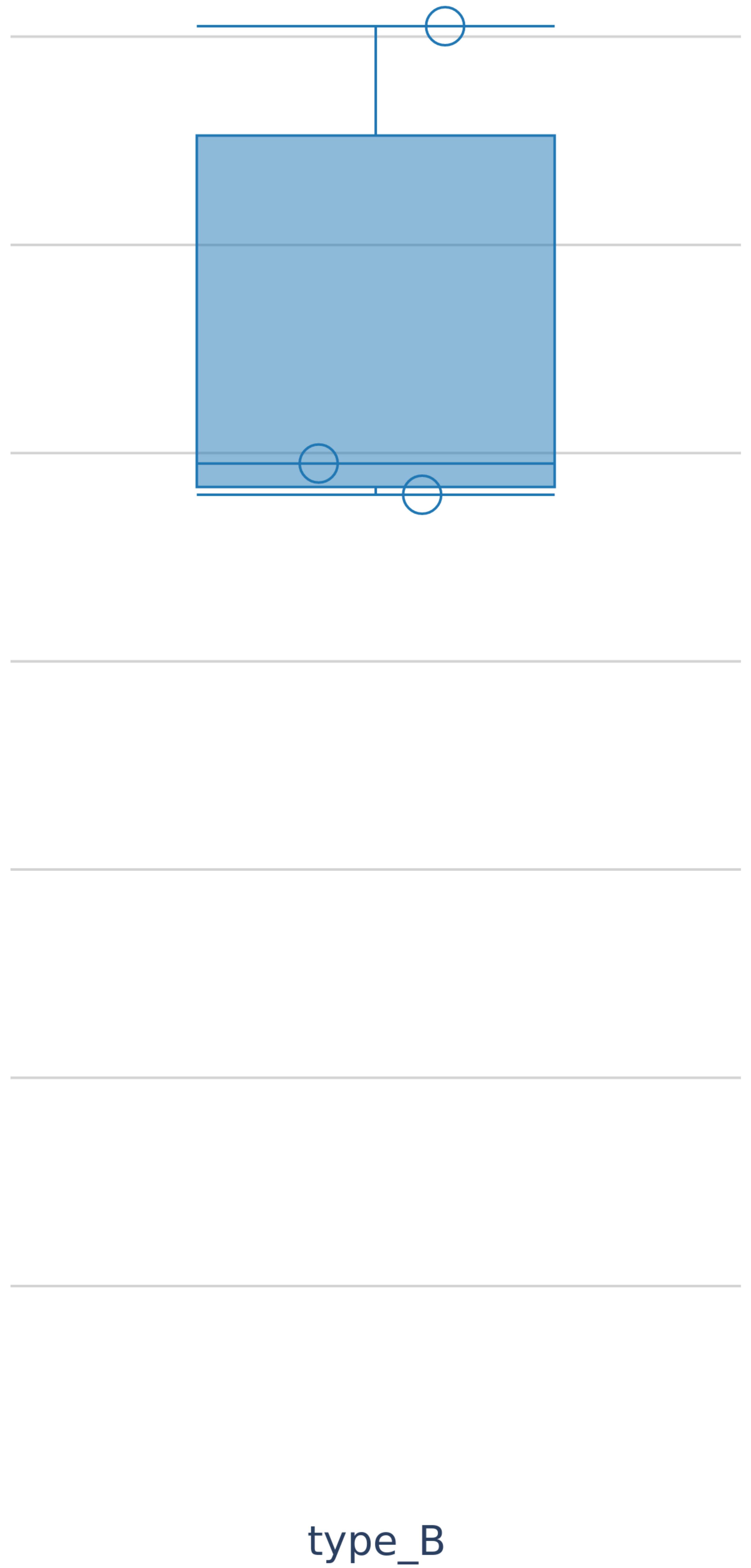
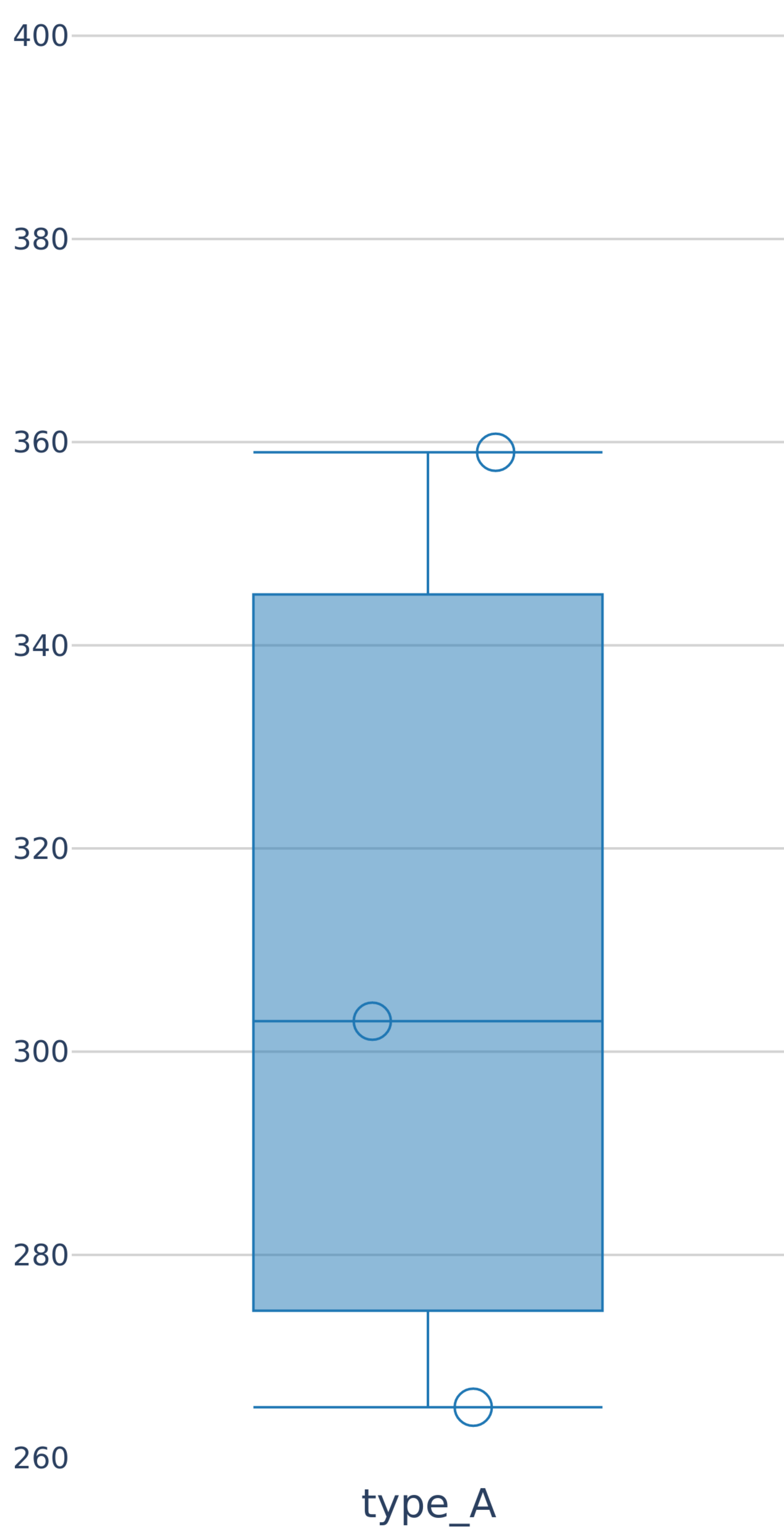
■ lambda_conversion

EXAMPLE general_conversion

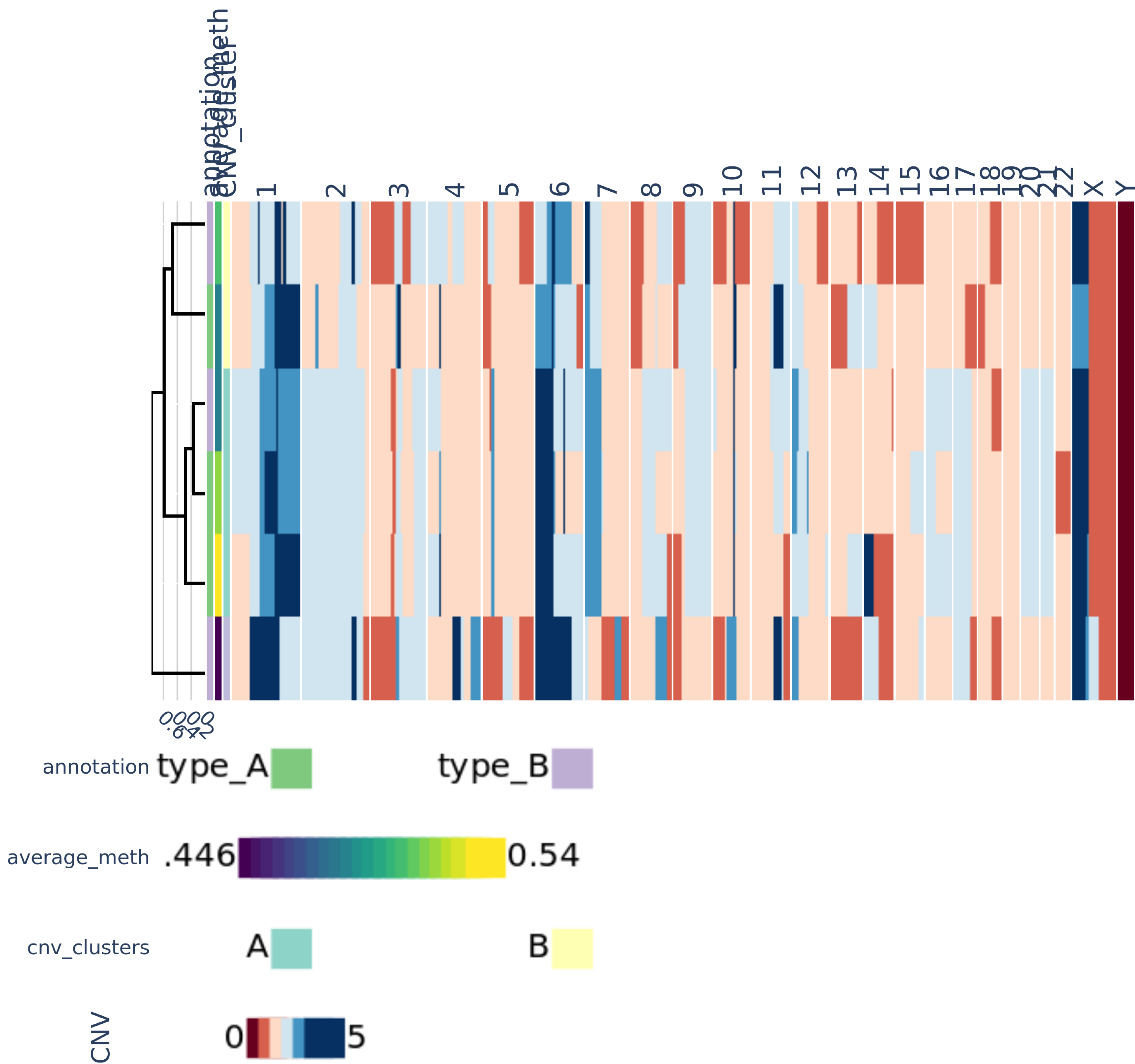


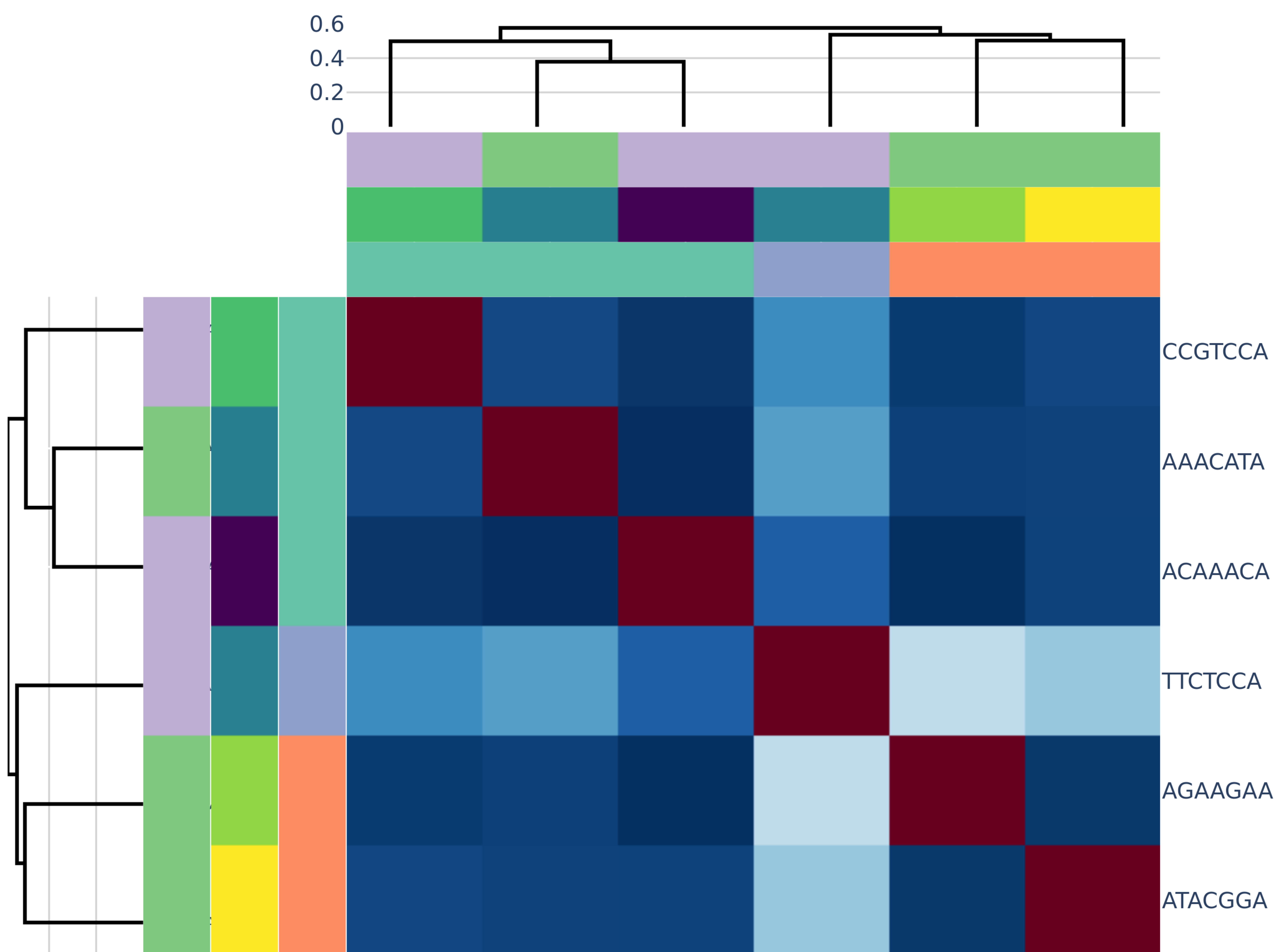
■ general_conversion

EXAMPLE CNV



■ CNV





0.4 0.2 0

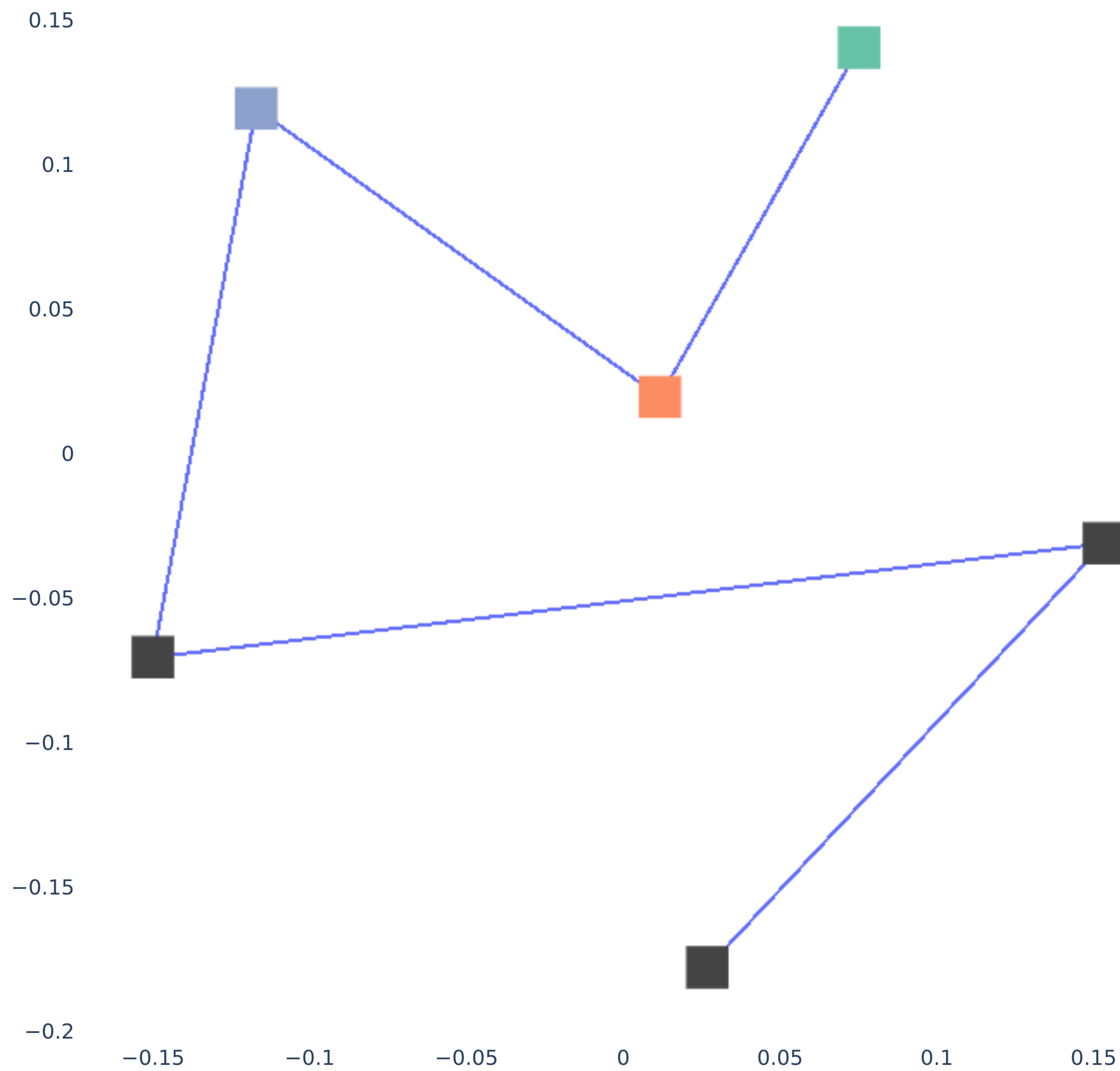
annotation CCGTCCA type A CATA ACAAACA TTCTCCA AGAAGAA type B ATACGGA

average_meth 0.446 0.54

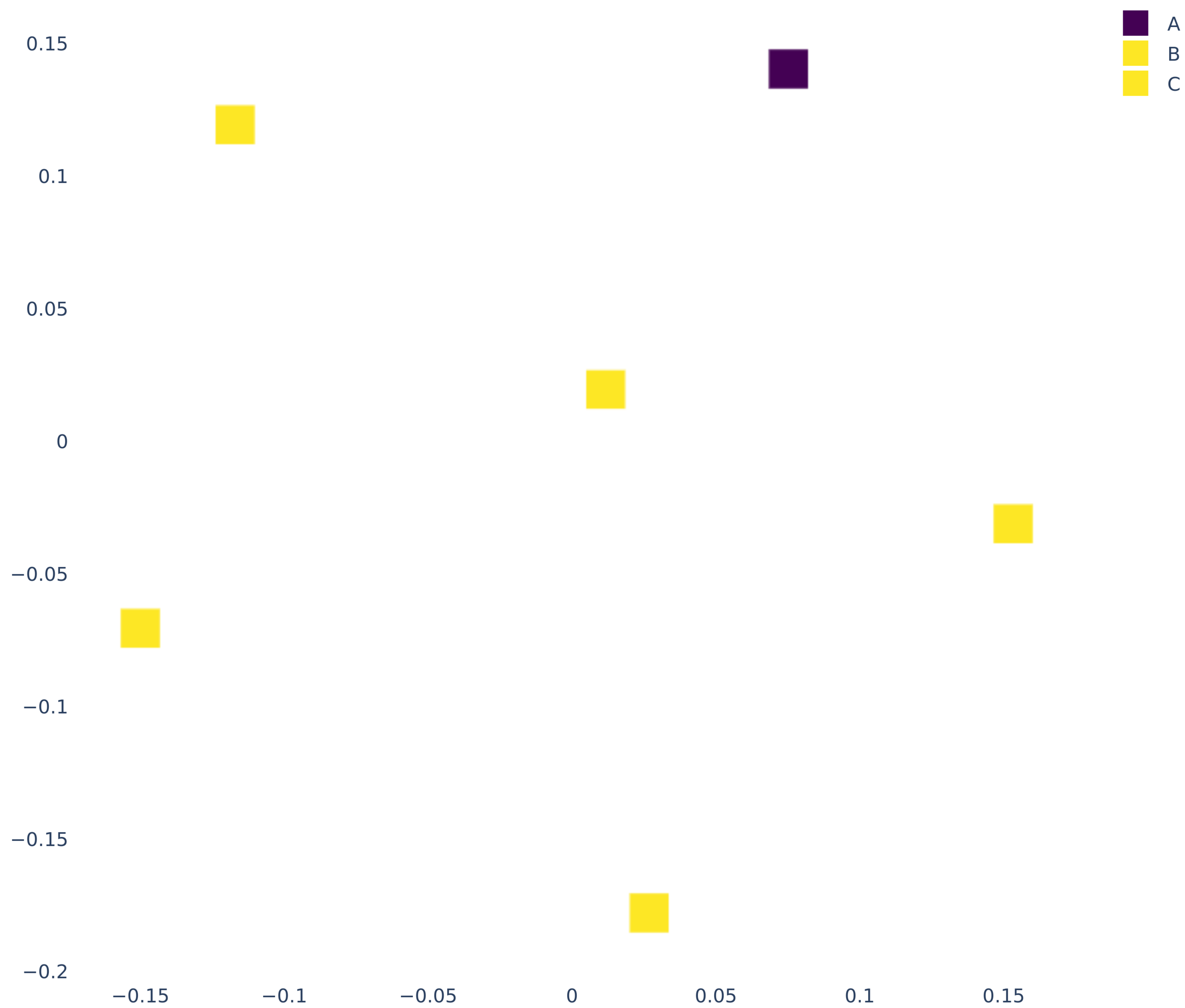
pdclust_clusters A B C

man_dist_scaled 0 0.26

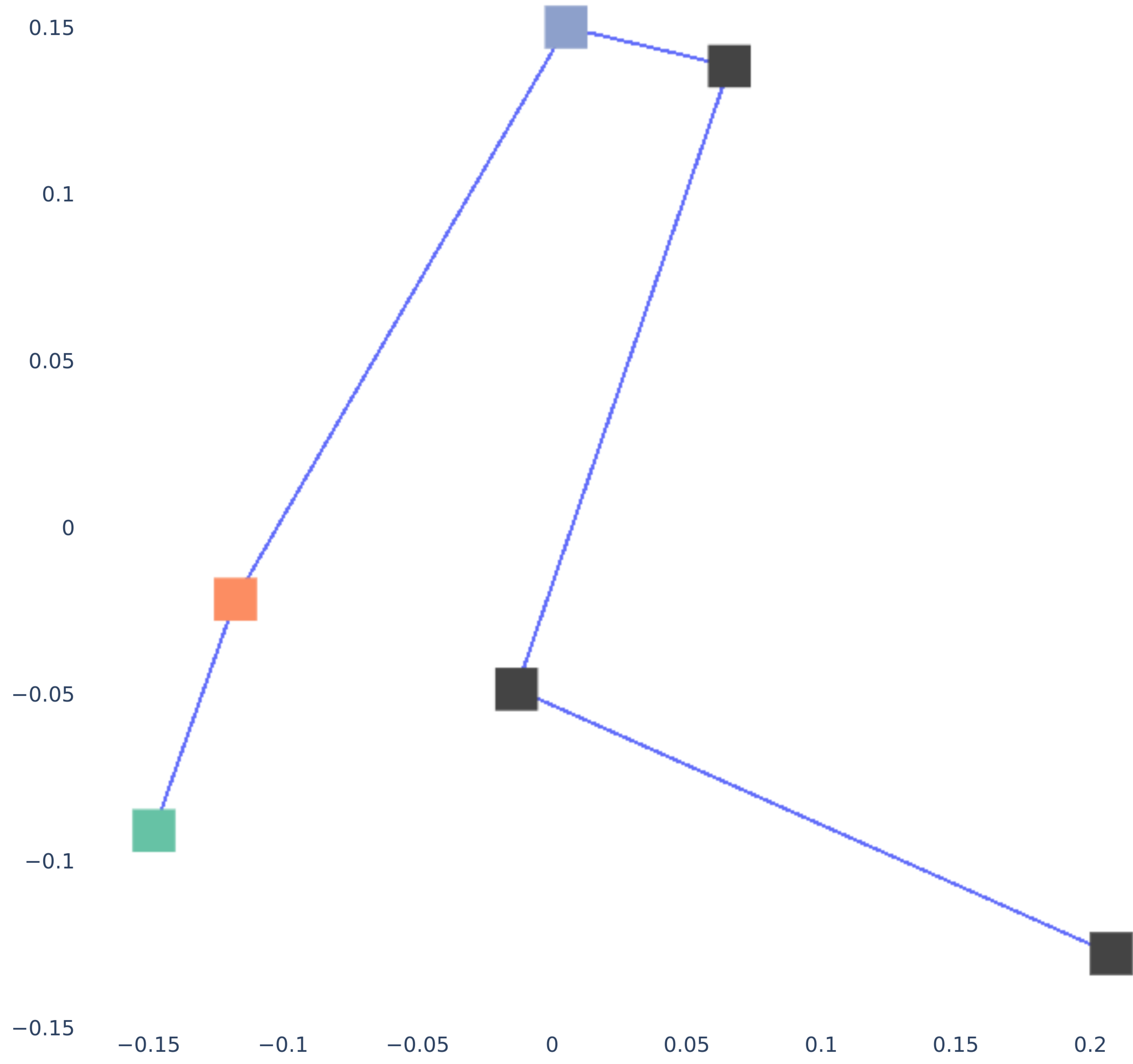
MDS



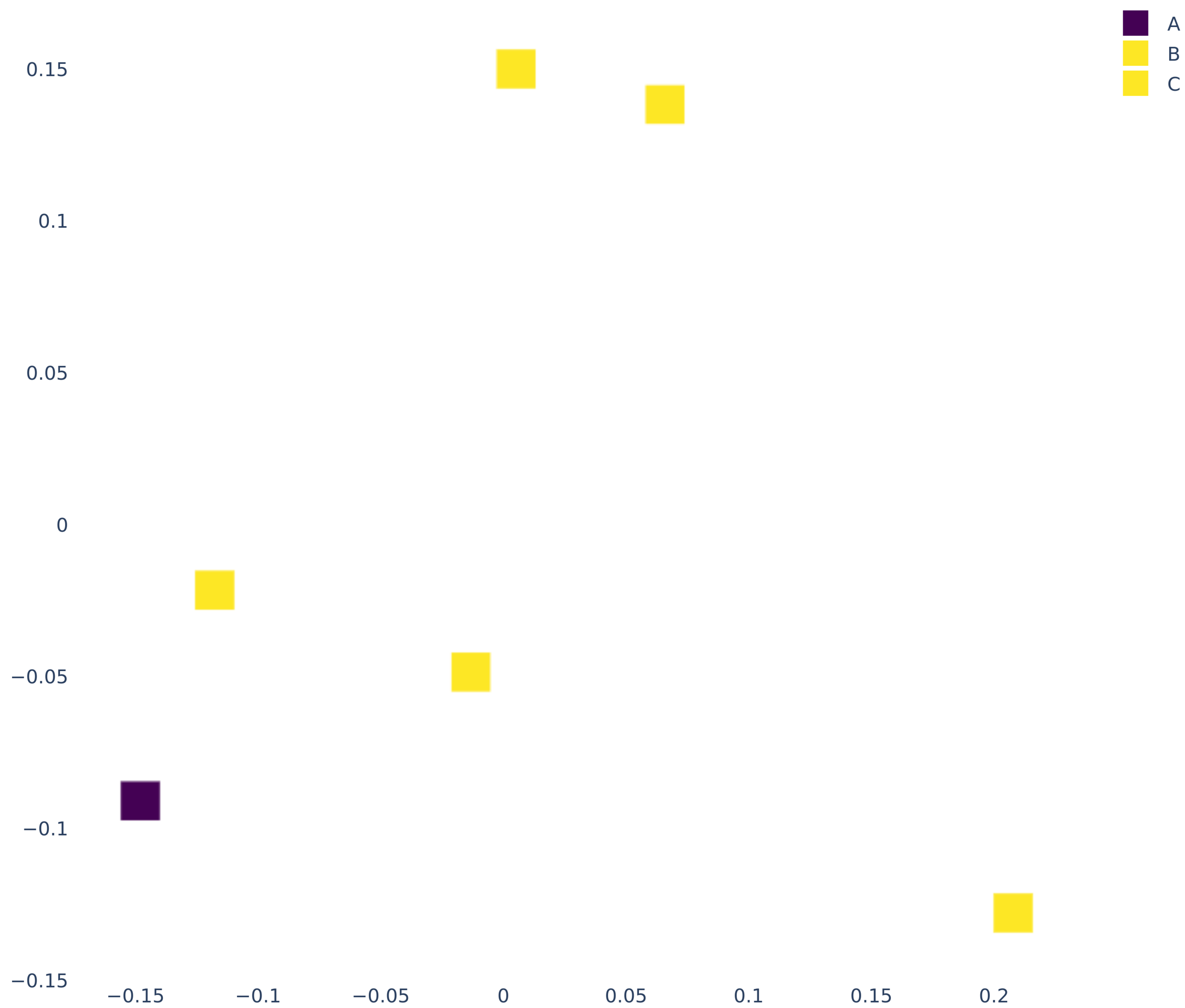
MDS



MDS

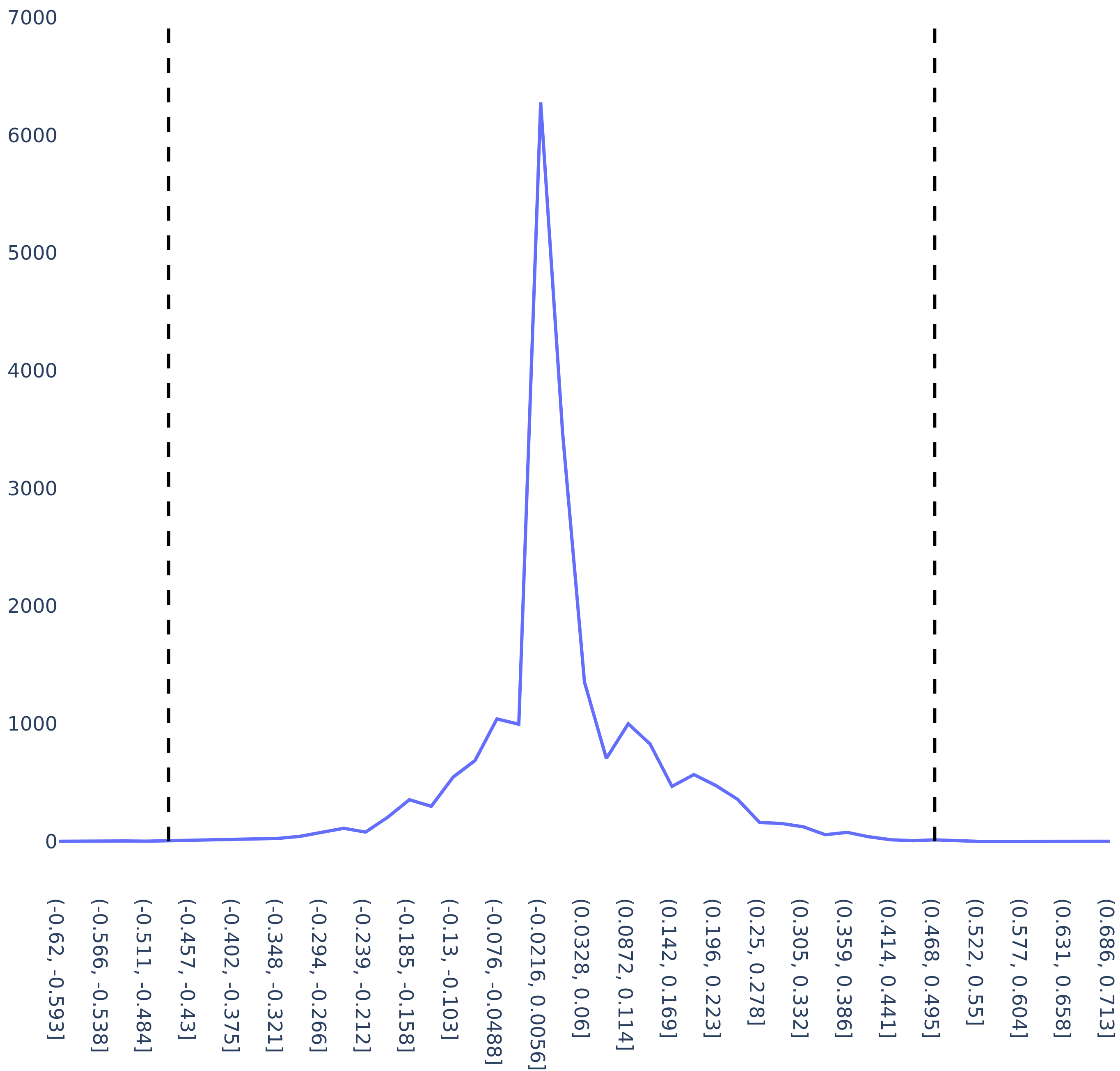


MDS

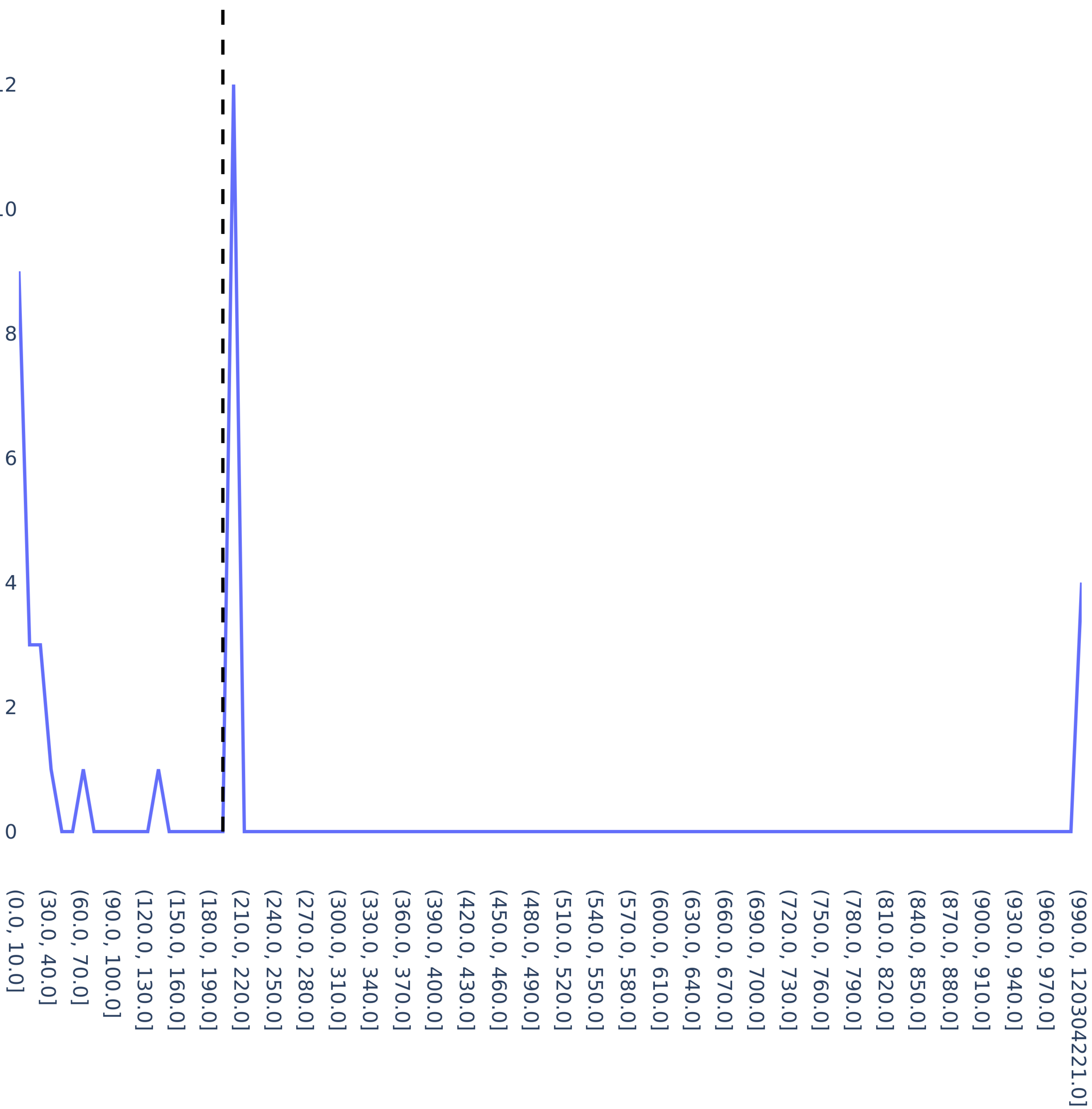


CpG Methylation difference distribution

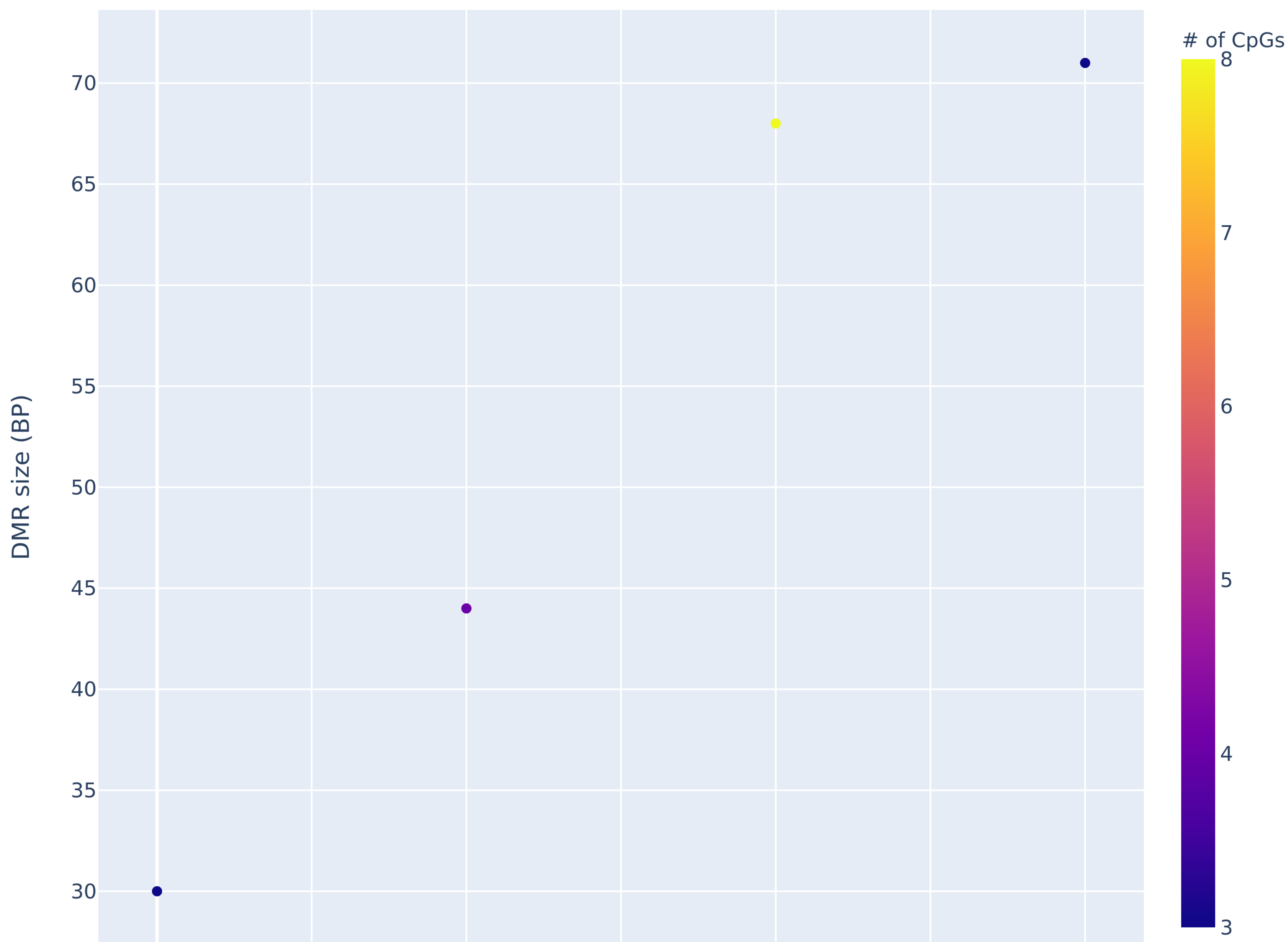
Hyper:16;Hypo:18;Within boundaries:20655



Significant CpG distances



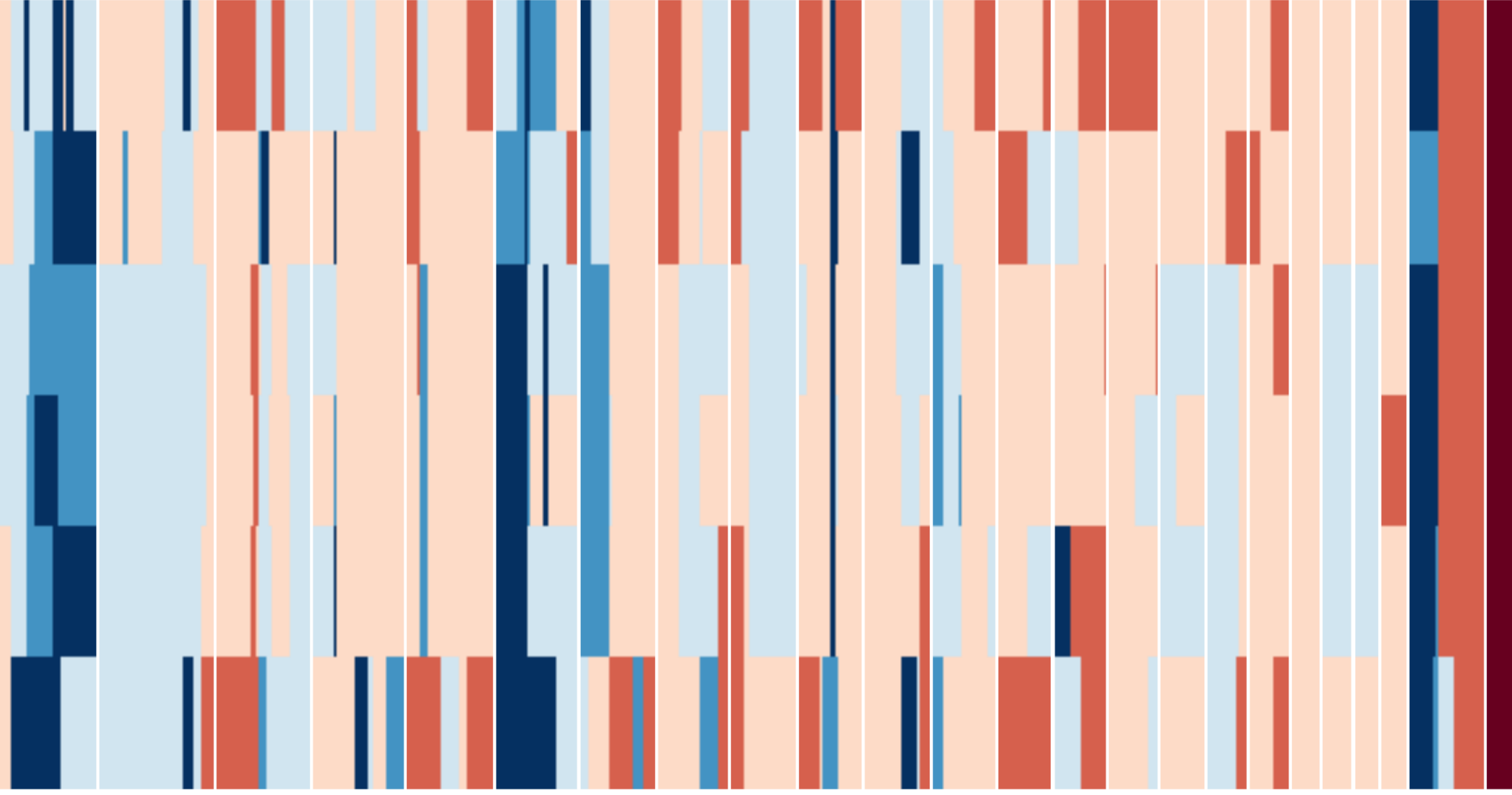
DMR size and # of Sig.CpGs



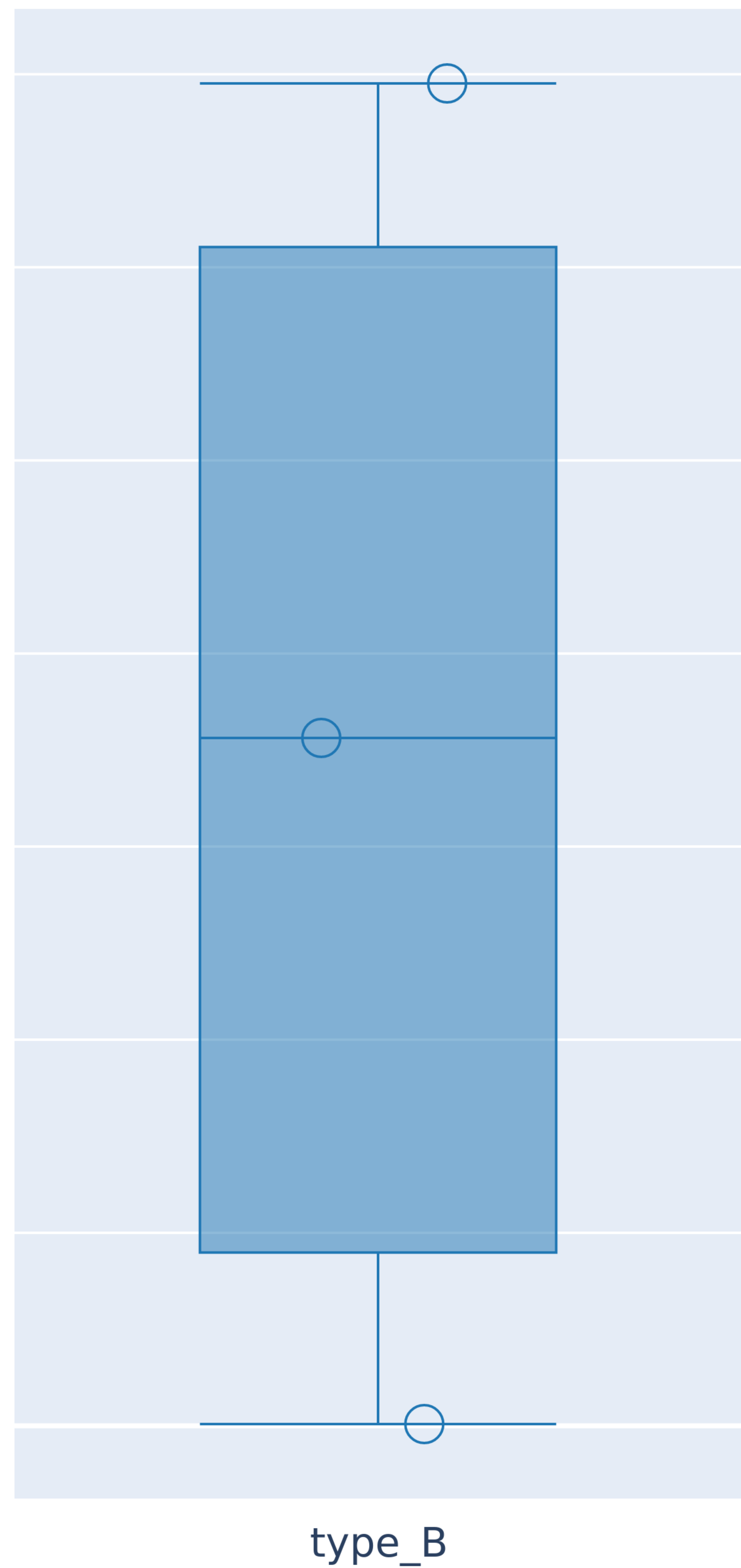
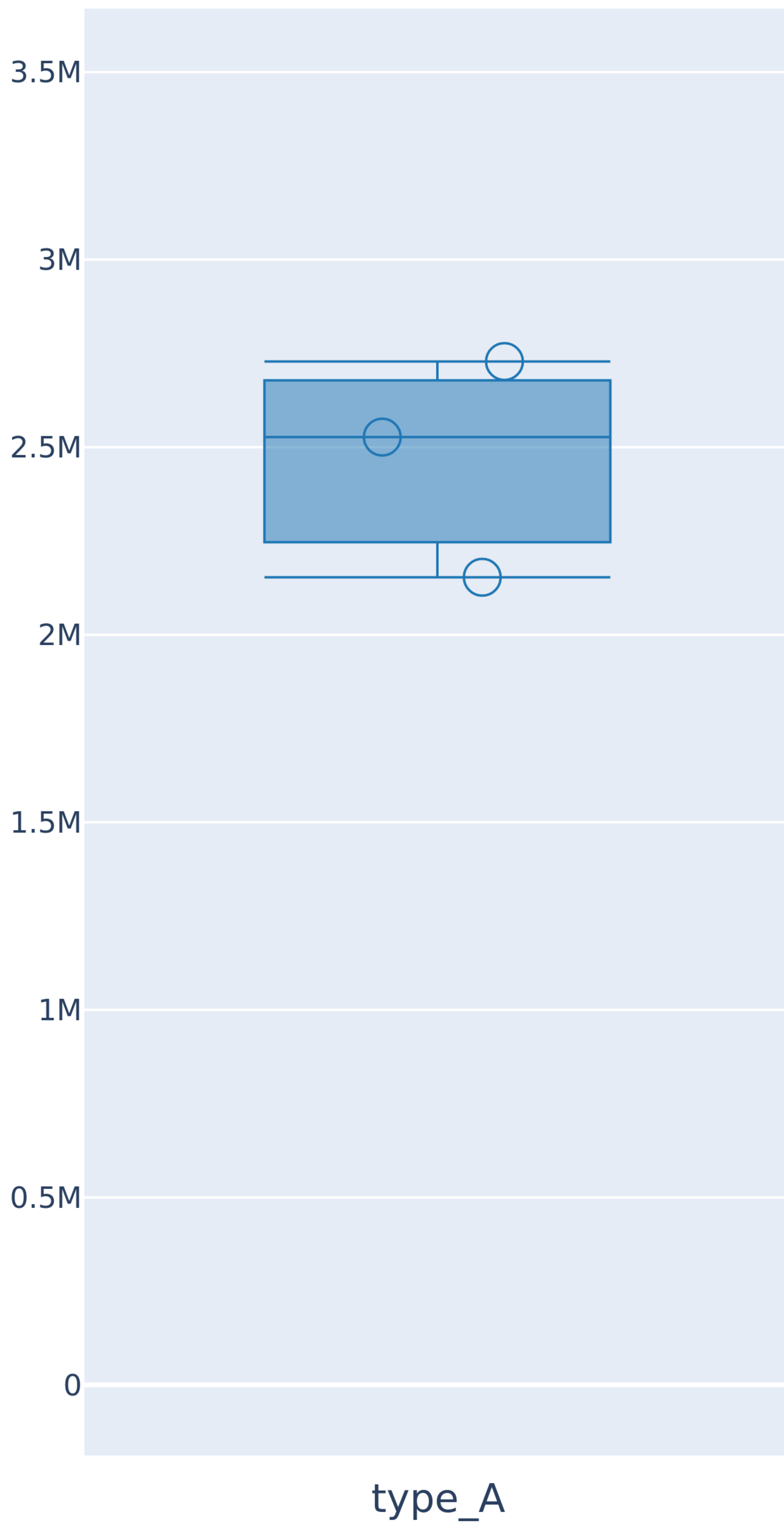
CNVnv_classification_annotation

0000
6420

average_meth

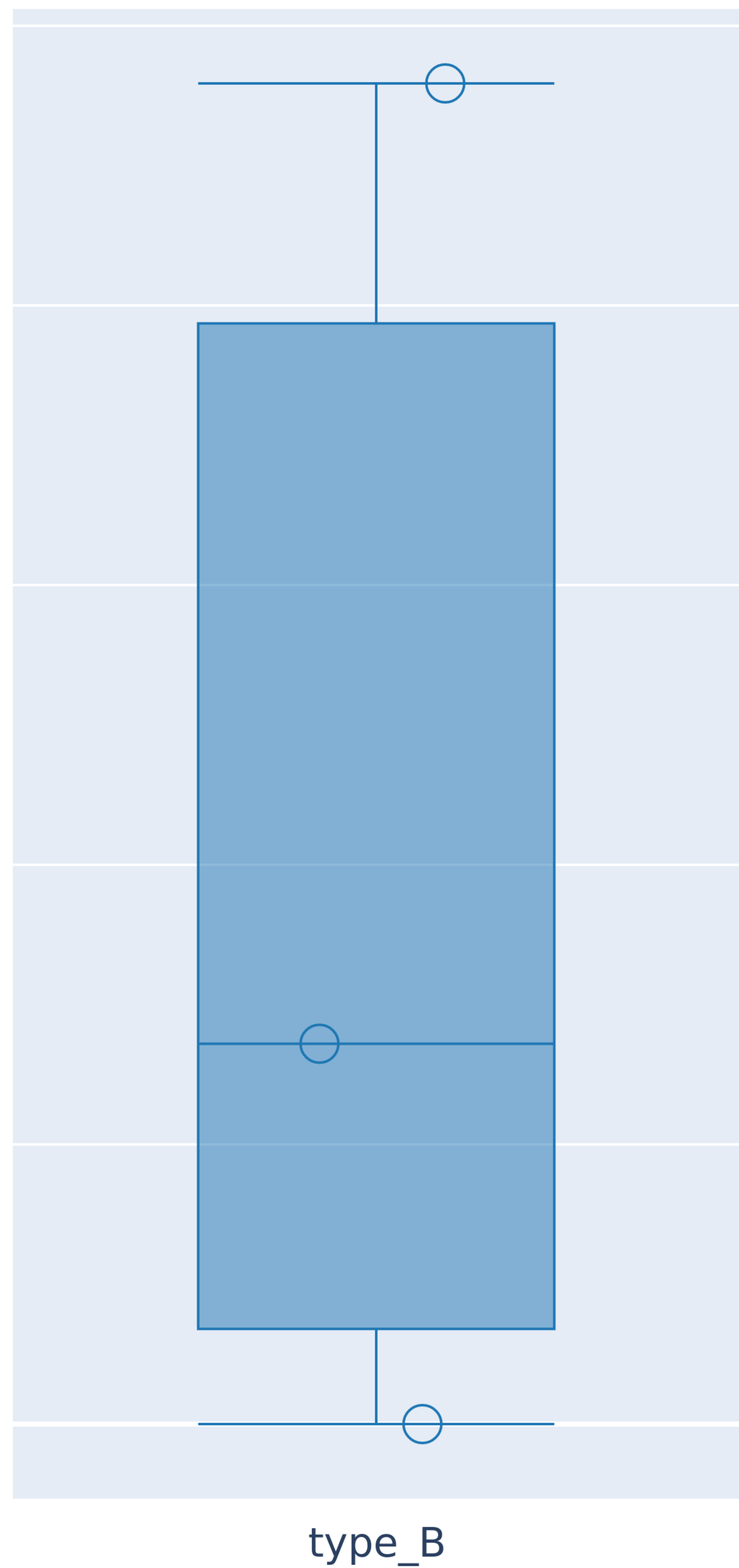
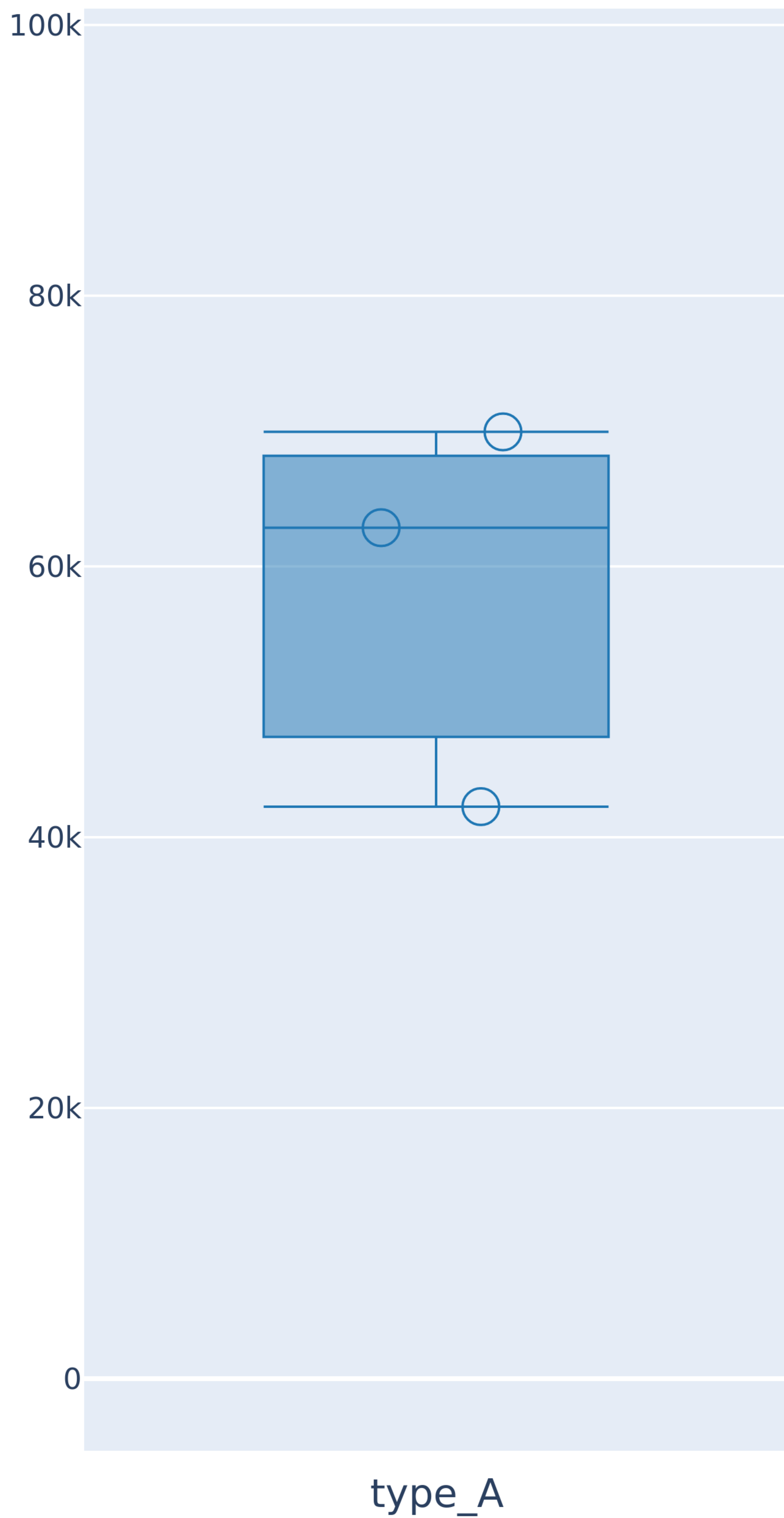


EXAMPLEcpg_count



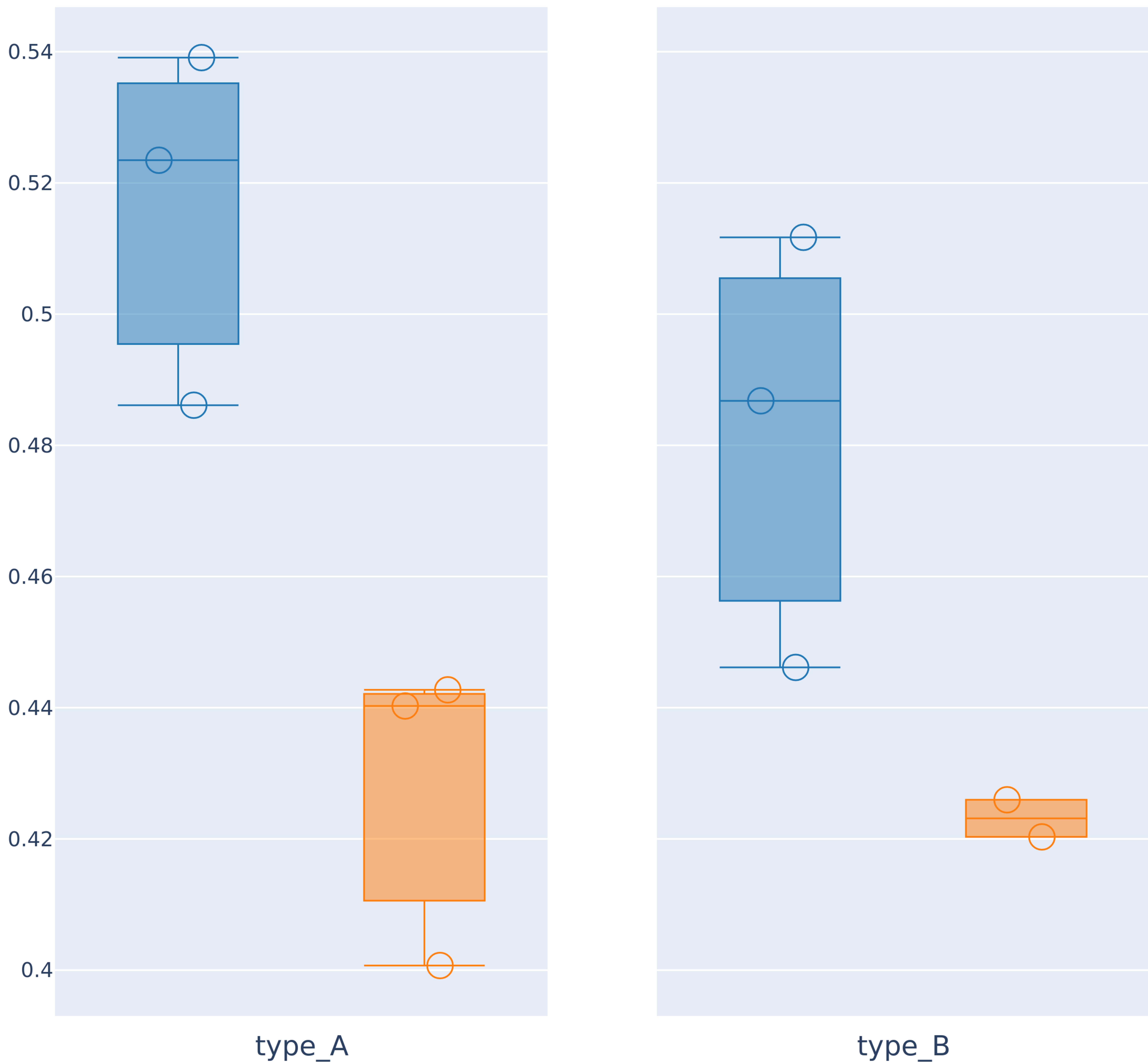
■ cpg_count

EXAMPLEcpg_count_cov3



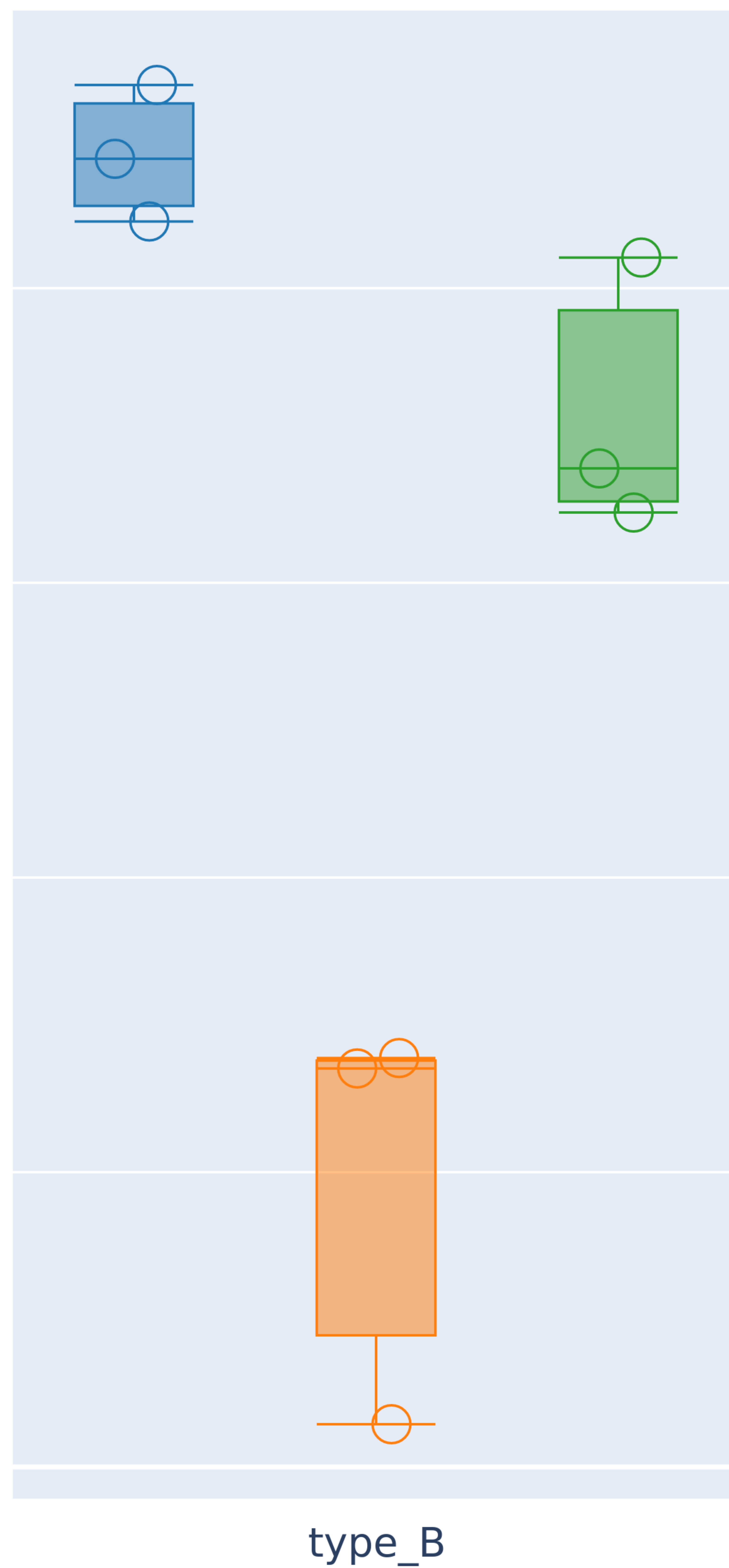
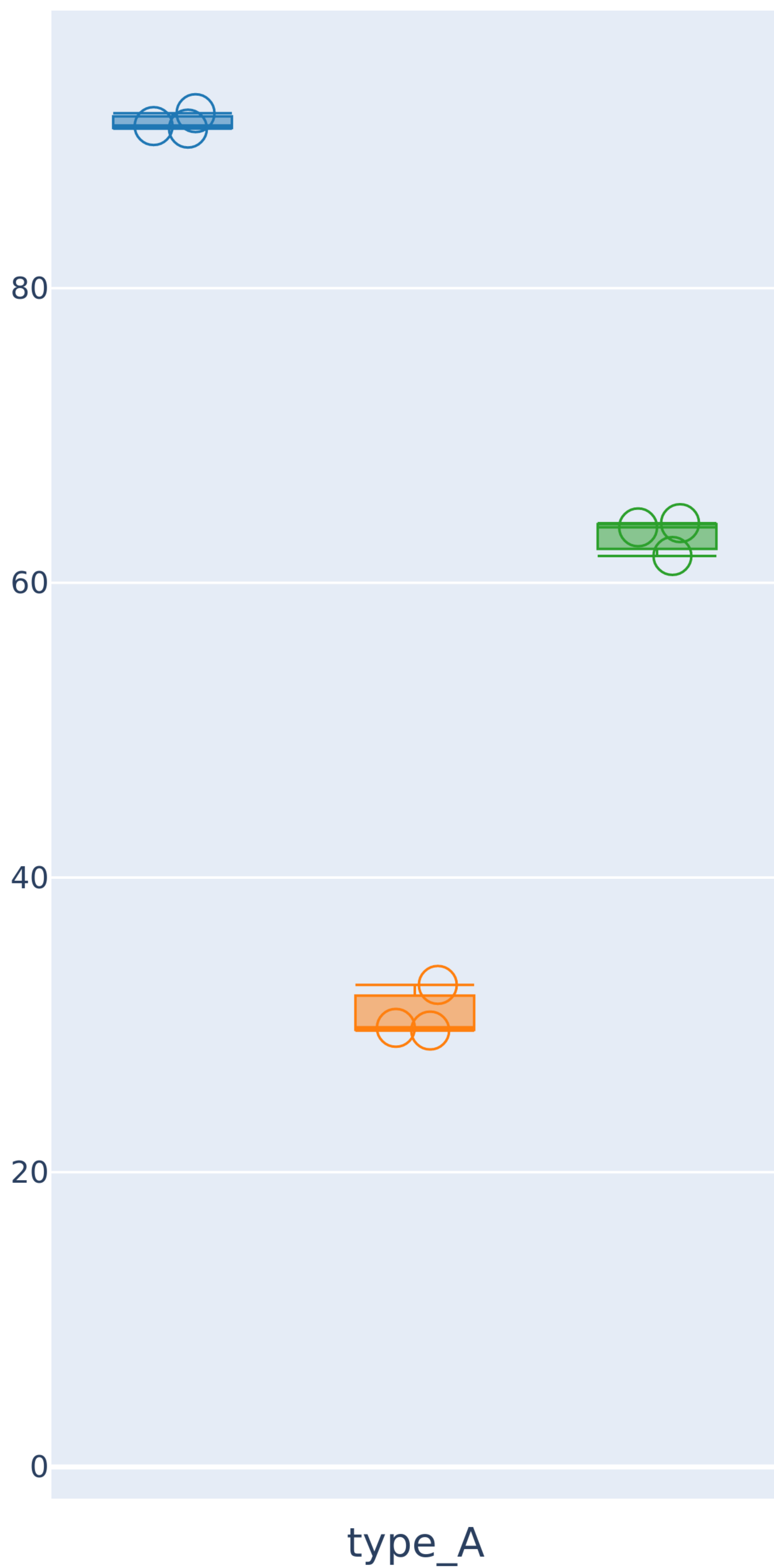
■ cpg_count_cov3

EXAMPLEaverage_meth,average_meth_cov3



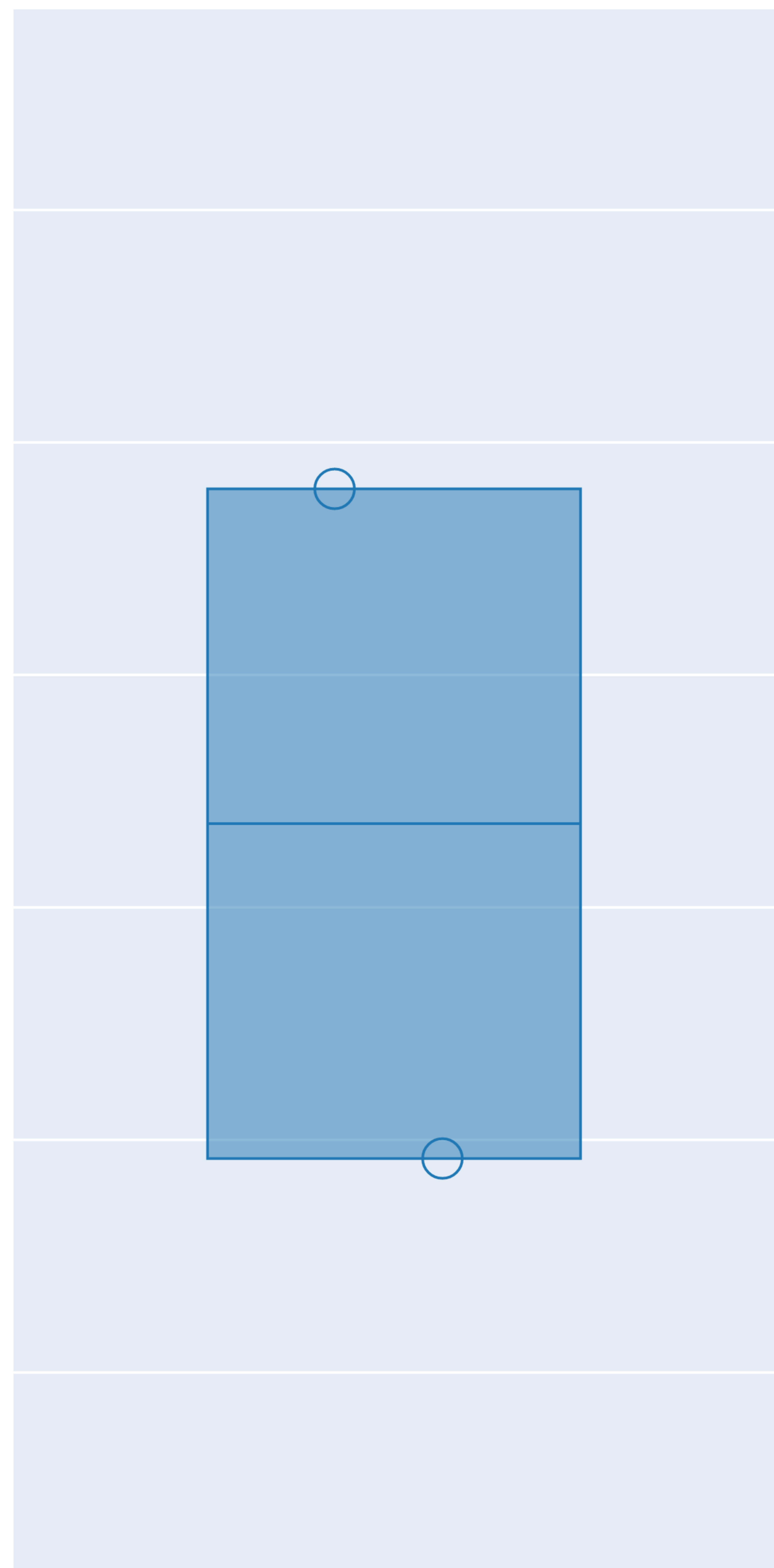
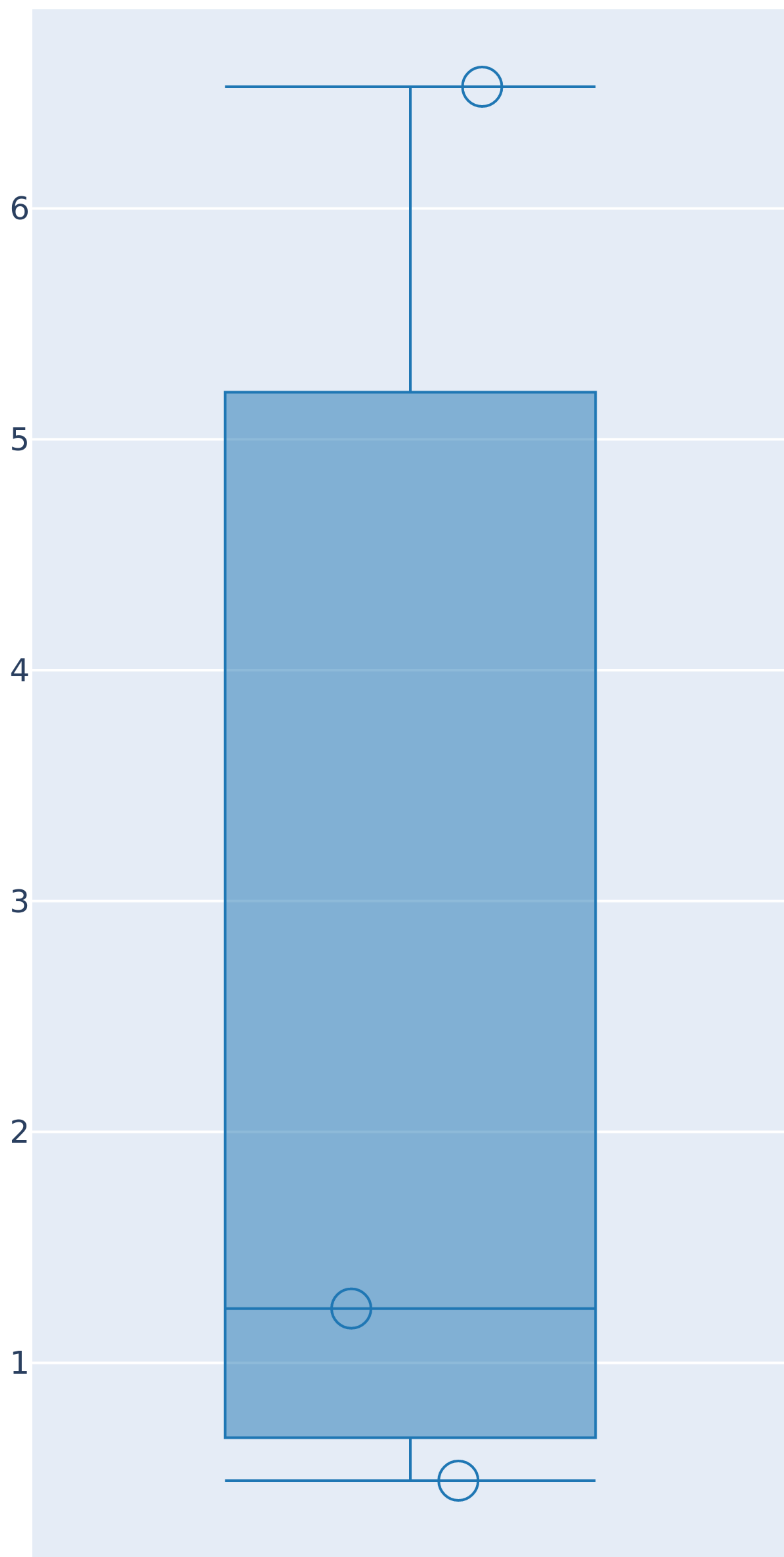
■ average_meth ■ average_meth_cov3

EXAMPLE mapped%,dup_rate%,mapped_minus_dup%



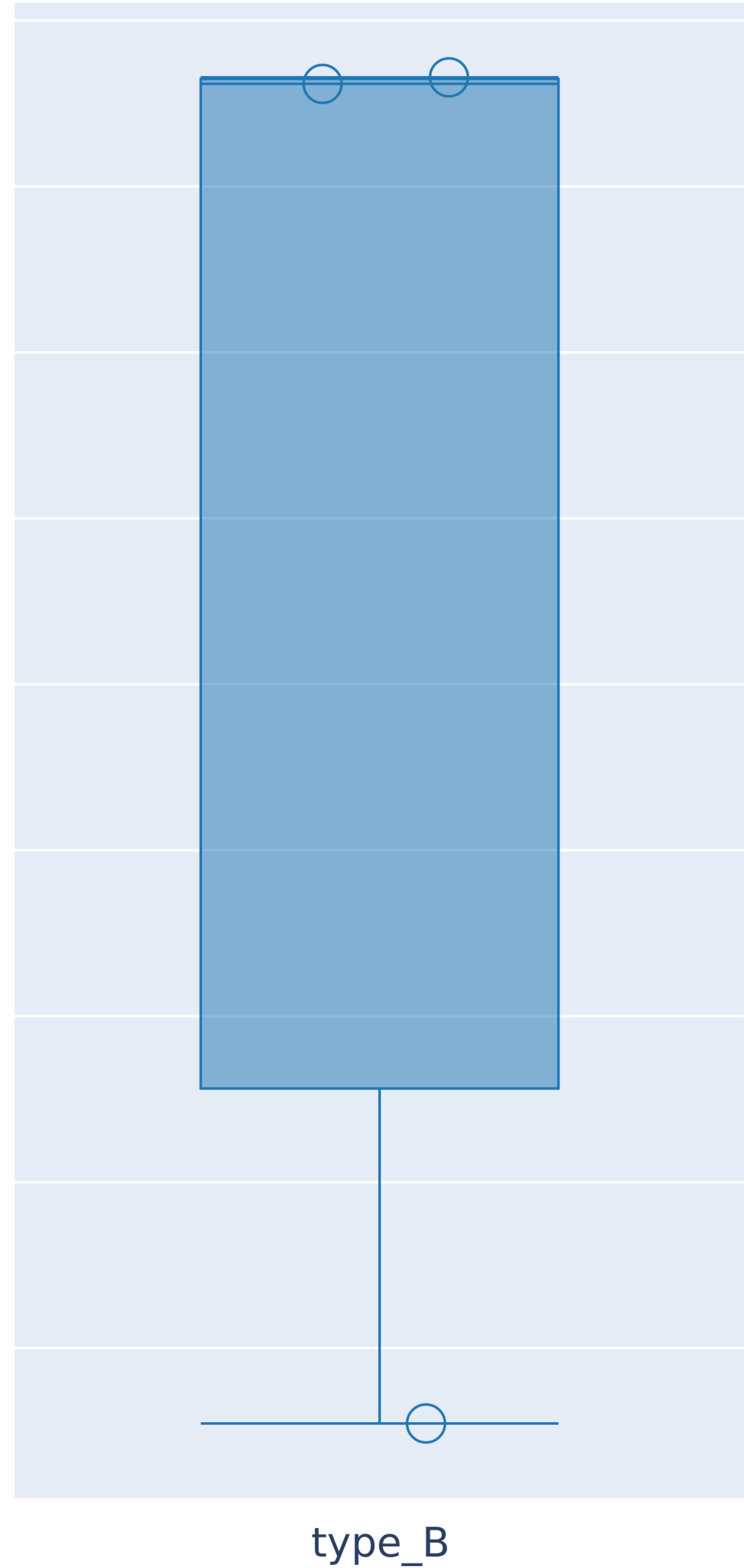
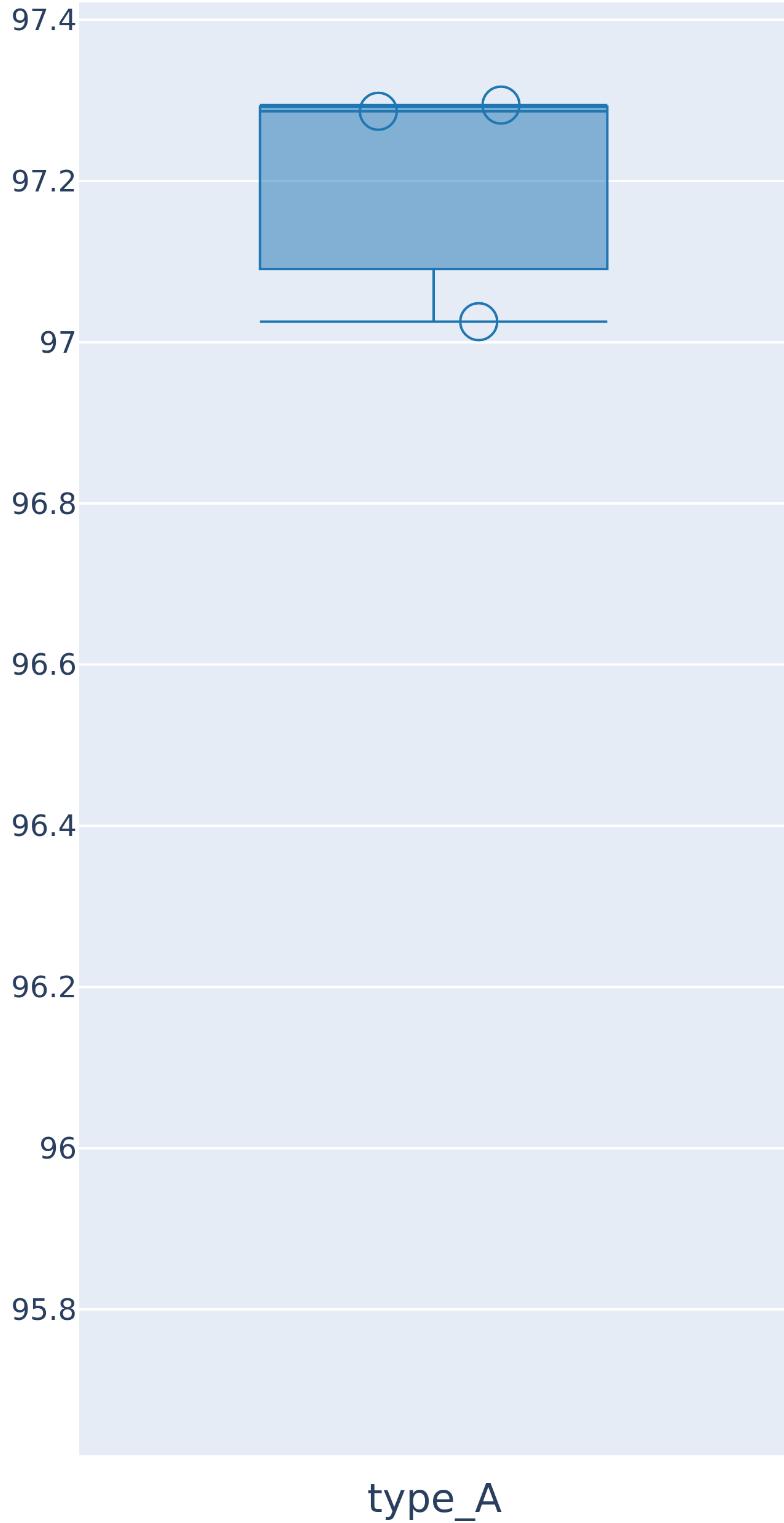
■ mapped% ■ dup_rate% ■ mapped_minus_dup%

EXAMPLET7_conversion



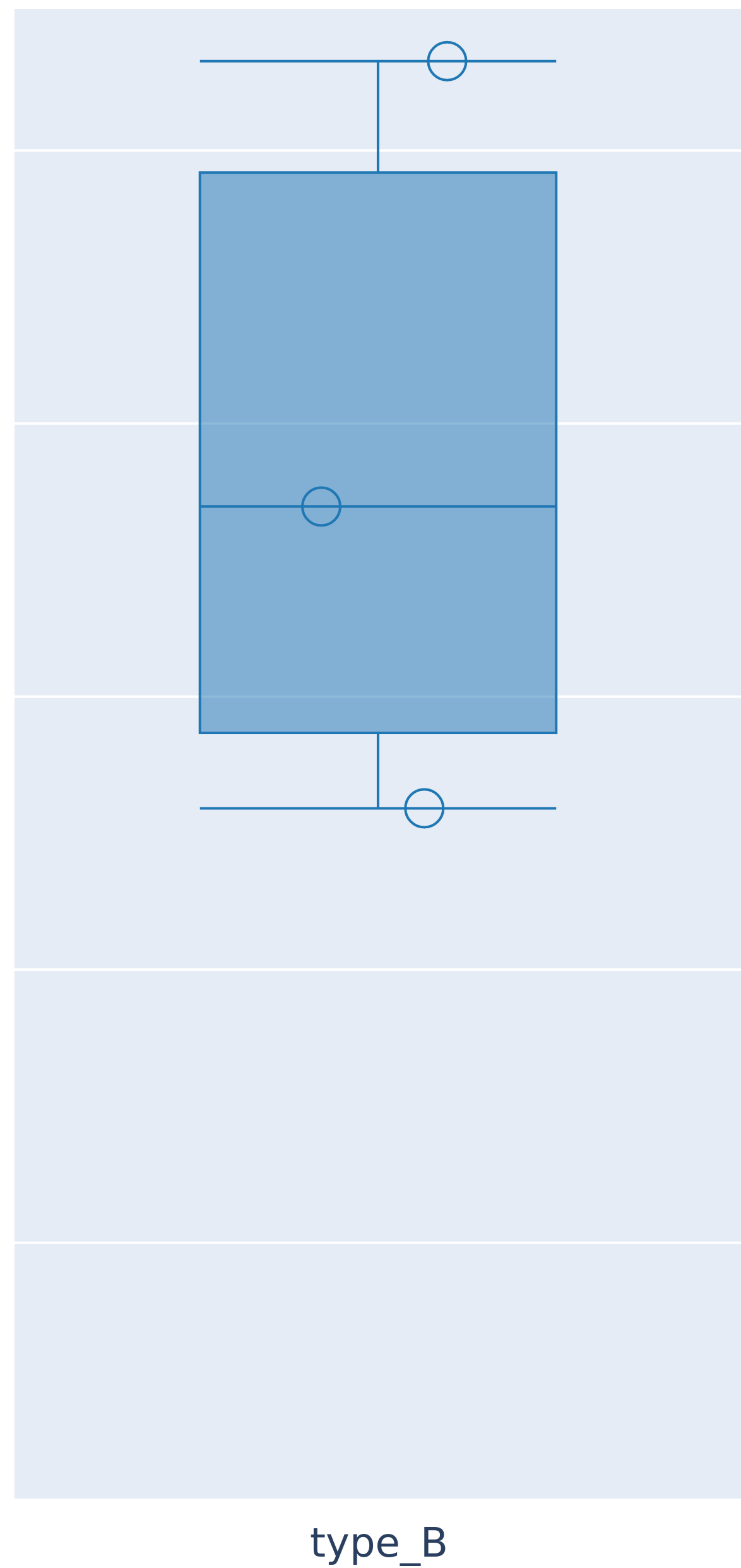
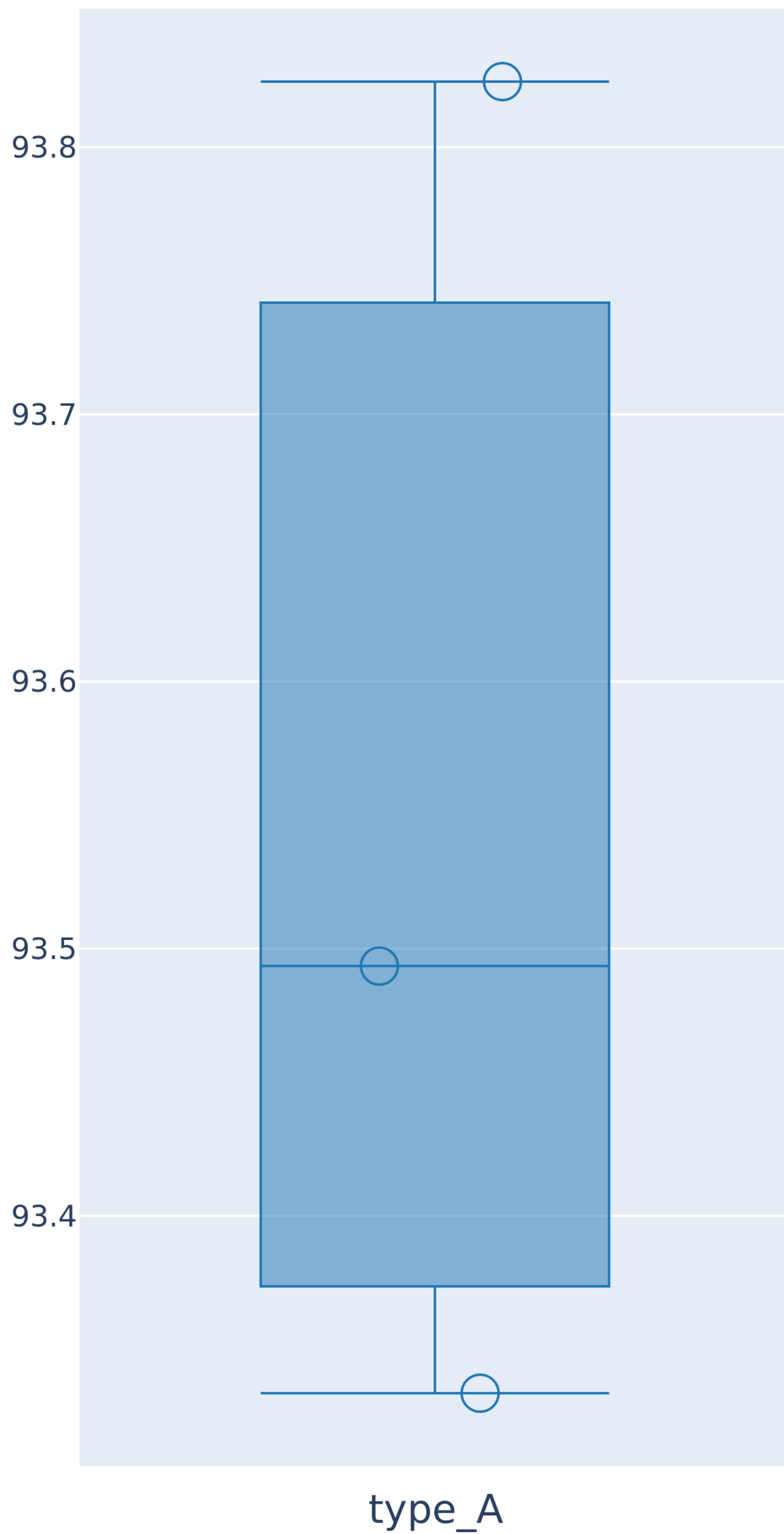
■ T7_conversion

EXAMPLE lambda_conversion



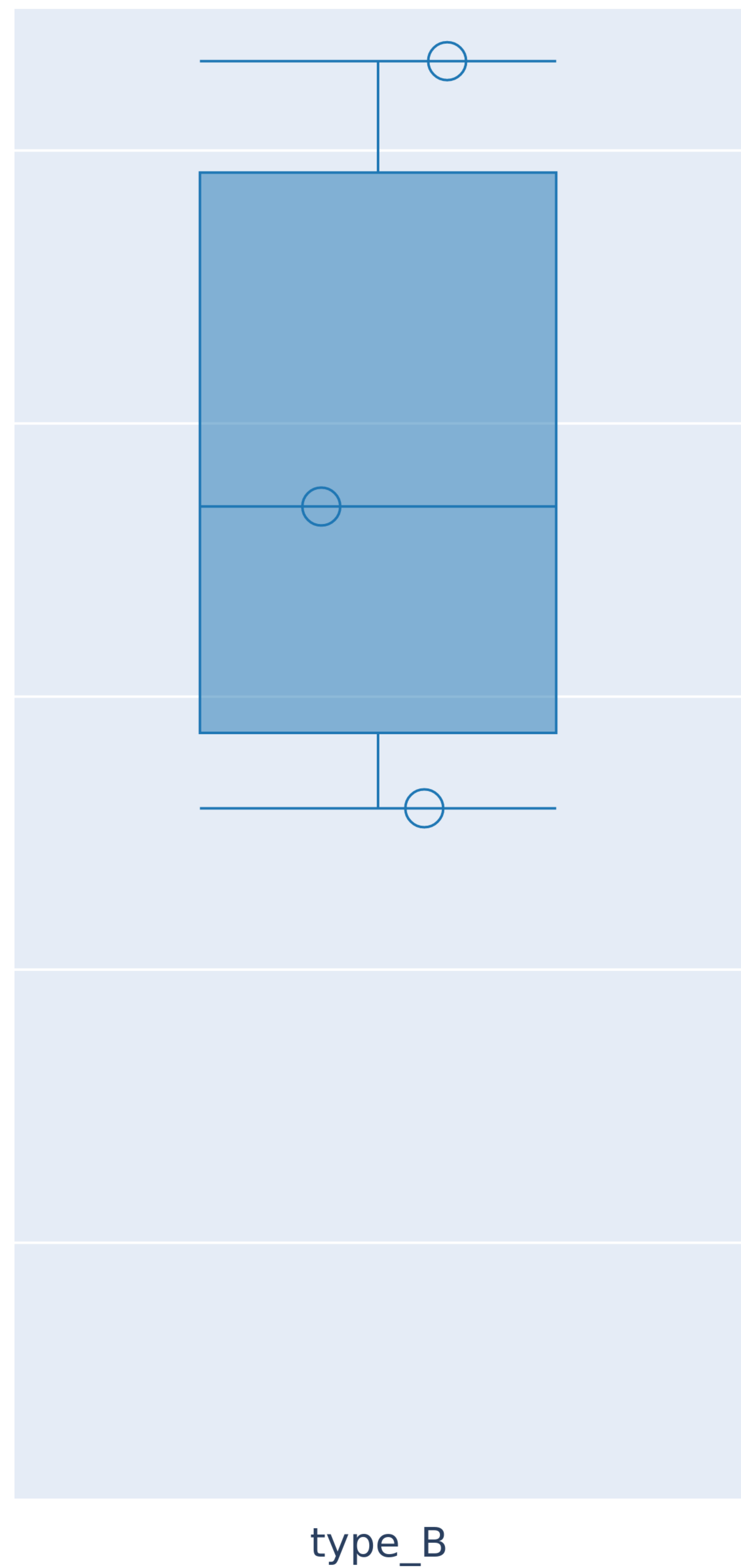
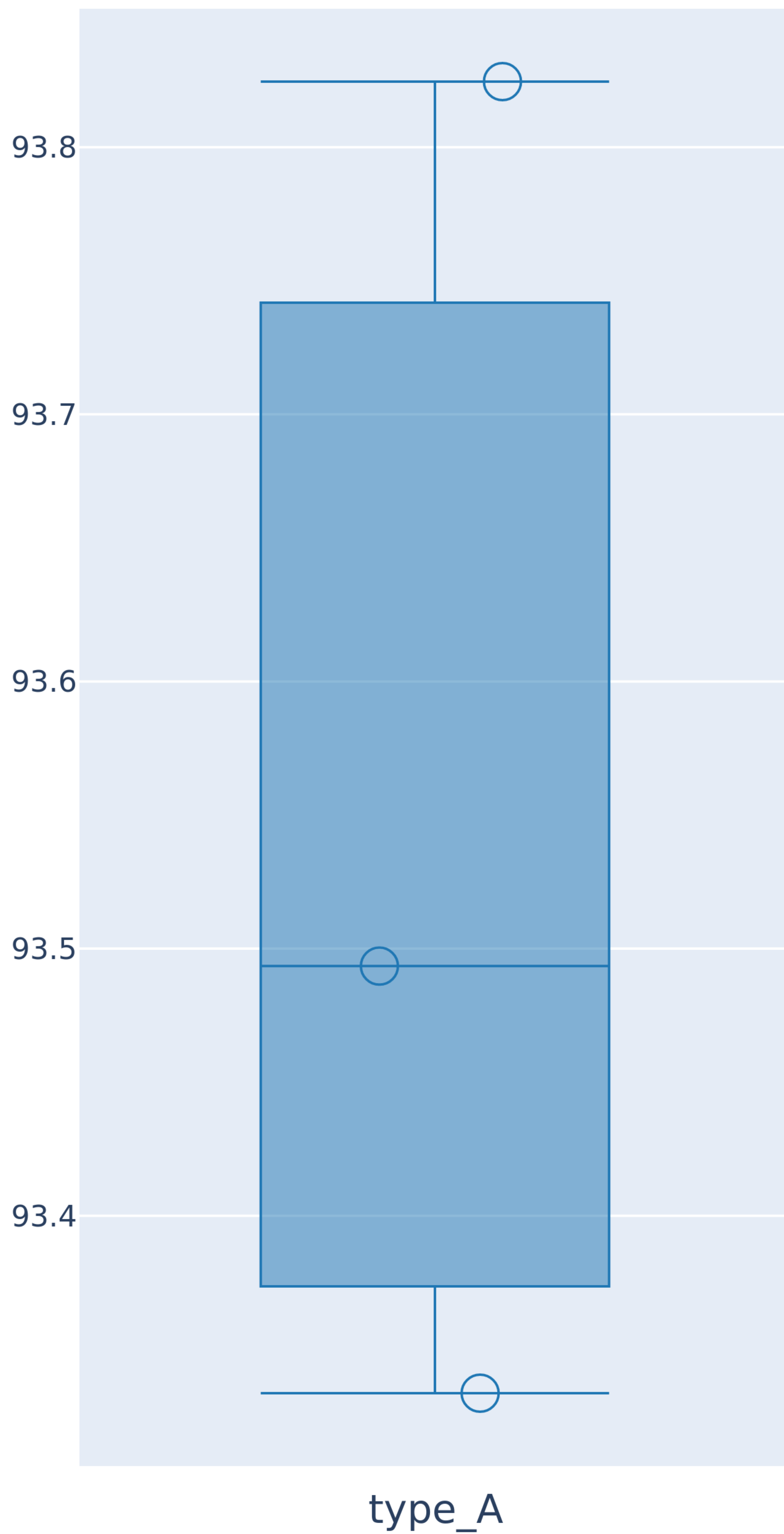
■ lambda_conversion

EXAMPLEgeneral_conversion



■ general_conversion

EXAMPLEgeneral_conversion



■ general_conversion

