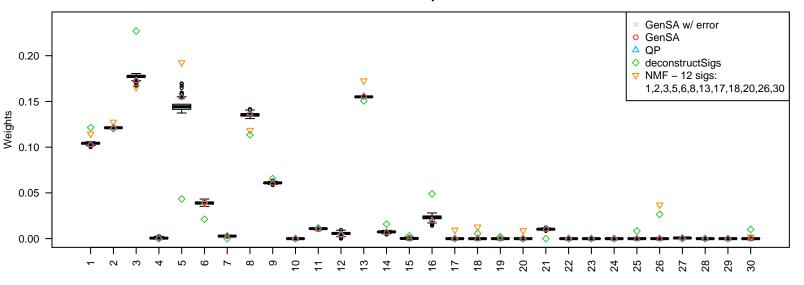
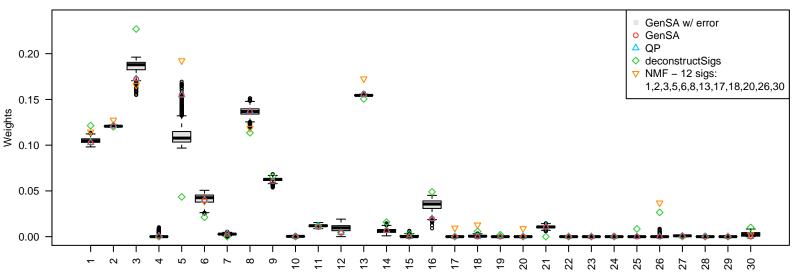
Boxplots based on SA for optimal GenSA error * 1.001 (0.1%) All samples



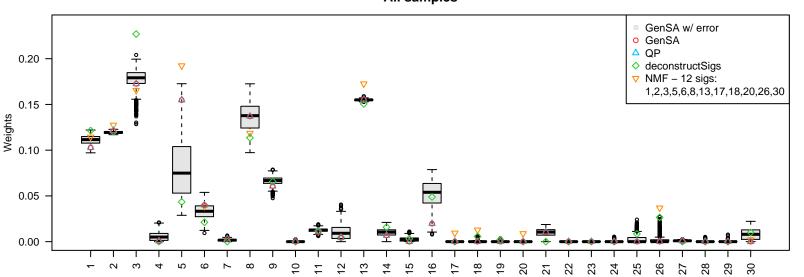
Signatures
GenSA+error(median) 0.00825, GenSA 0.00824, QP 0.00824, deconstructSigs 0.00887, NMF 0.01459

Boxplots based on SA for optimal GenSA error * 1.01 (1%) All samples

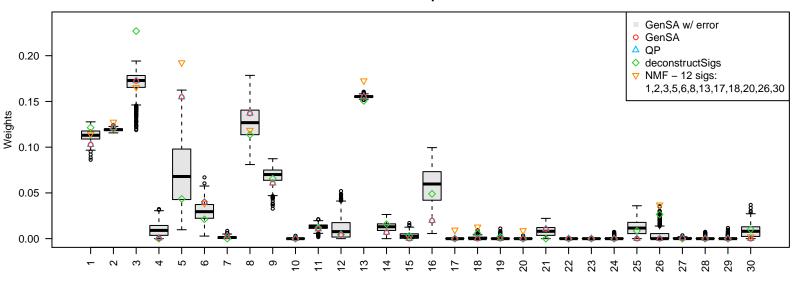


Signatures
GenSA+error(median) 0.00832, GenSA 0.00824, QP 0.00824, deconstructSigs 0.00887, NMF 0.01459

Boxplots based on SA for optimal GenSA error * 1.05 (5%) All samples



Boxplots based on SA for optimal GenSA error * 1.1 (10%) All samples



Signatures
GenSA+error(median) 0.00899, GenSA 0.00824, QP 0.00824, deconstructSigs 0.00887, NMF 0.01459