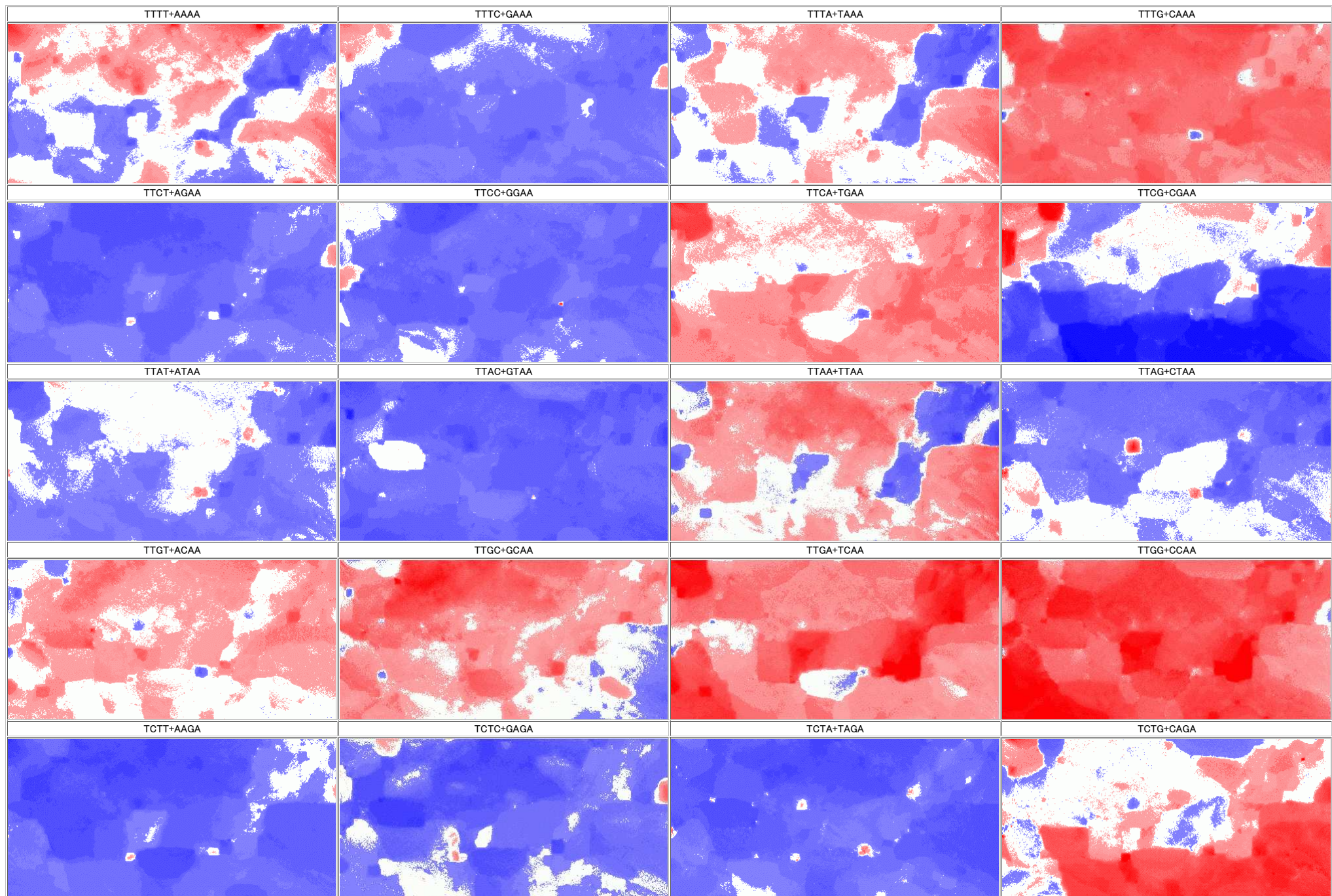
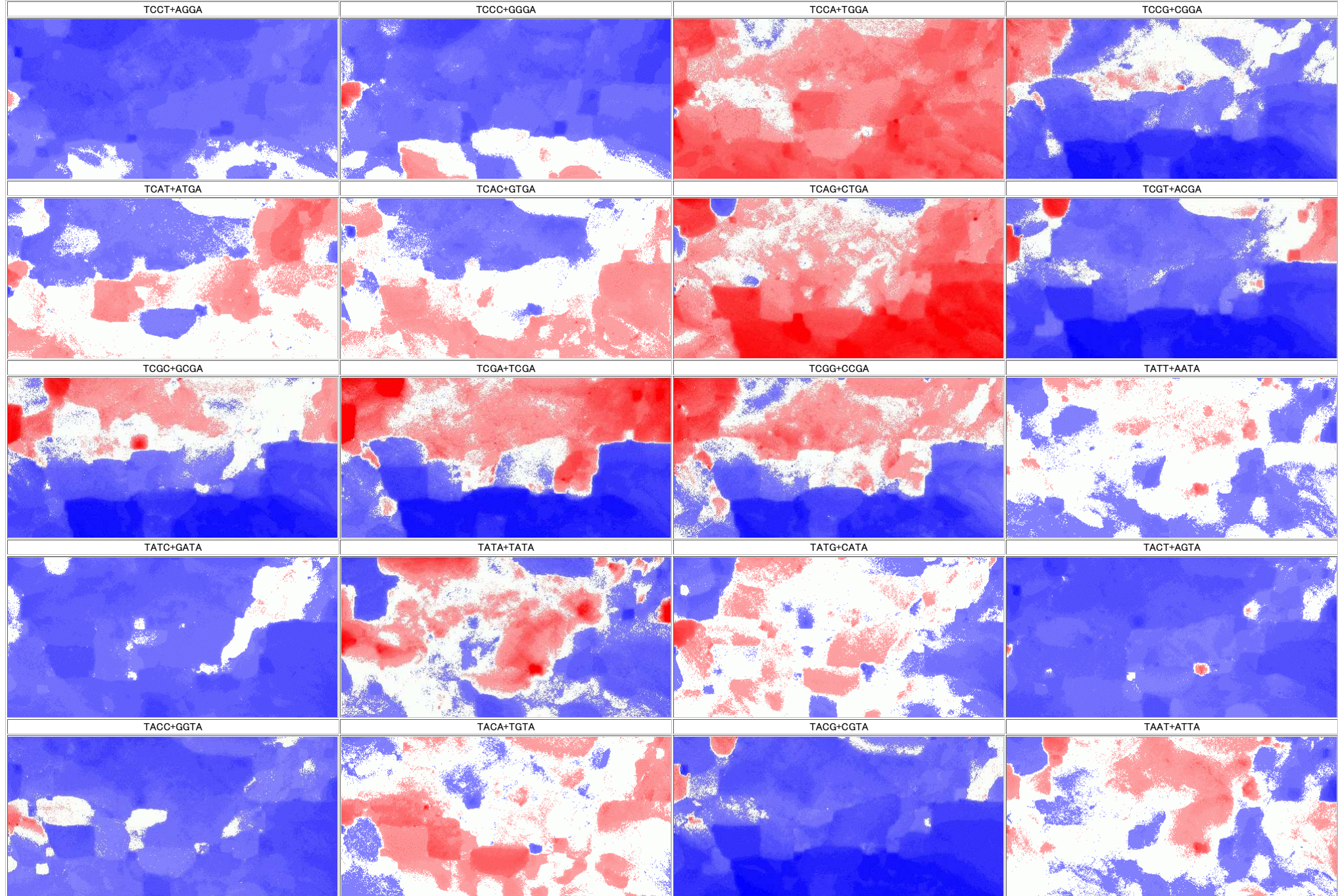
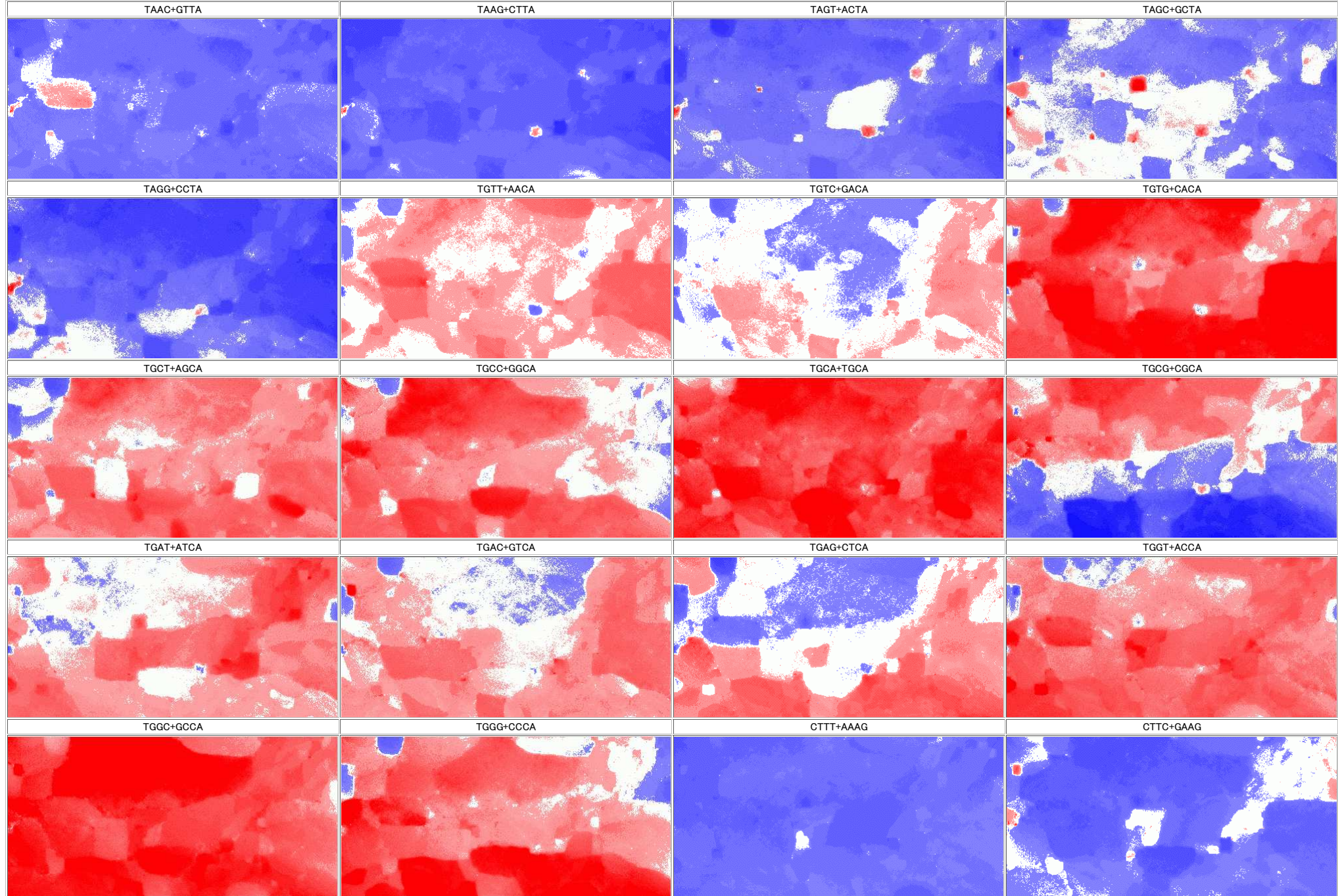


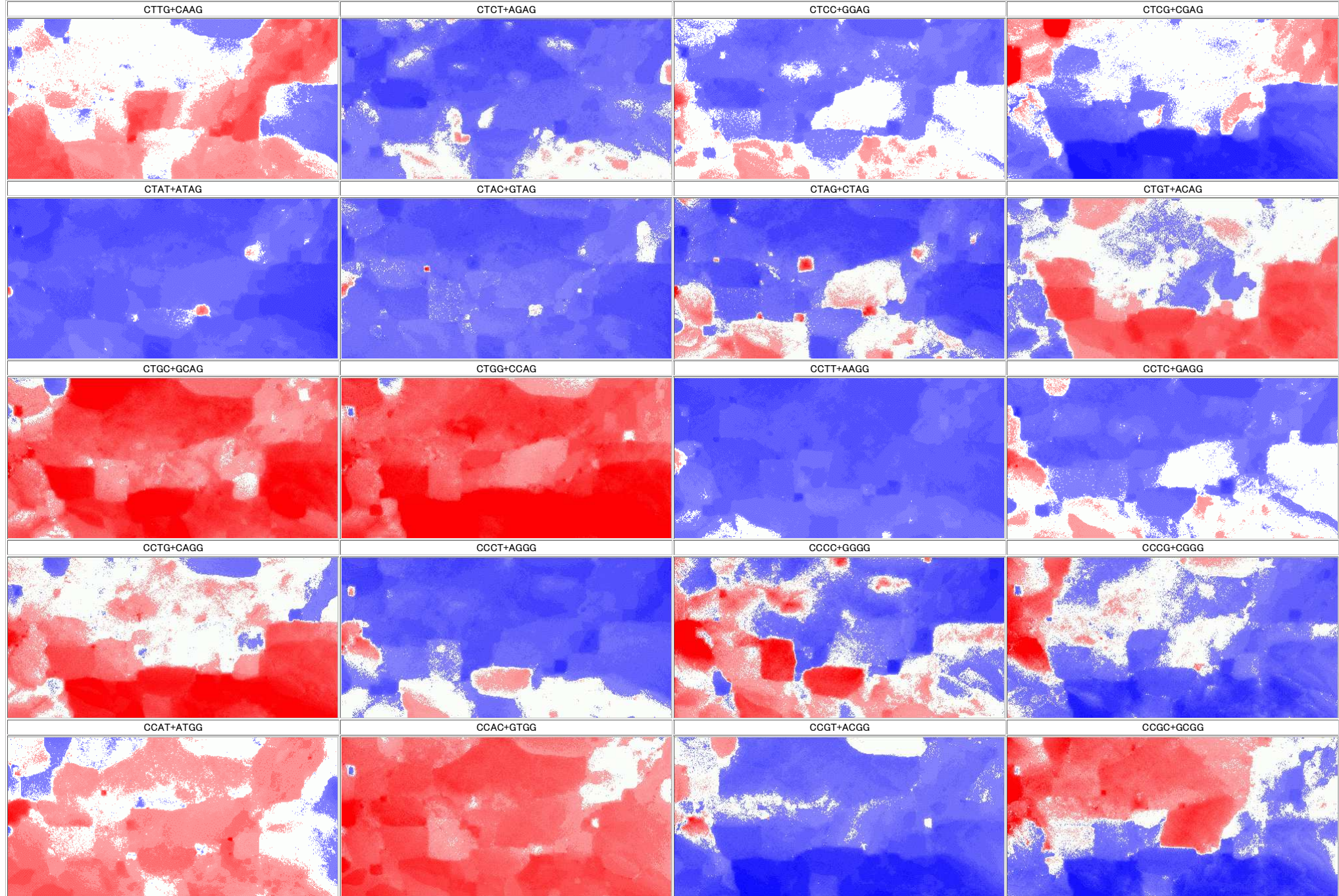
Supplementary Figure S1.

Level of each pair of complimentary tetranucleotide in 100-kb DegeTetra-BLSOM for the overlapping 100-kb sequences of 101 eukaryotic genomes.

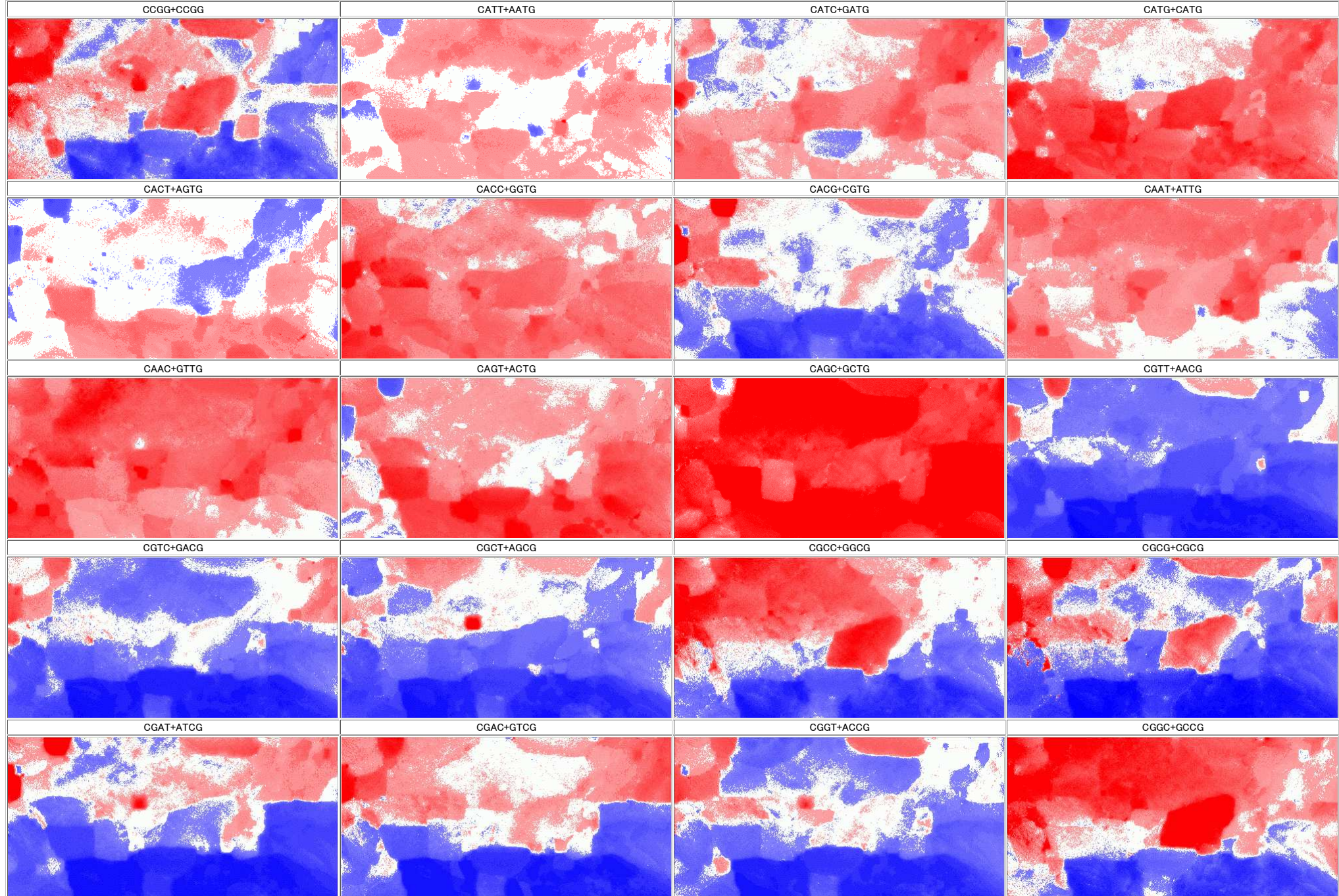


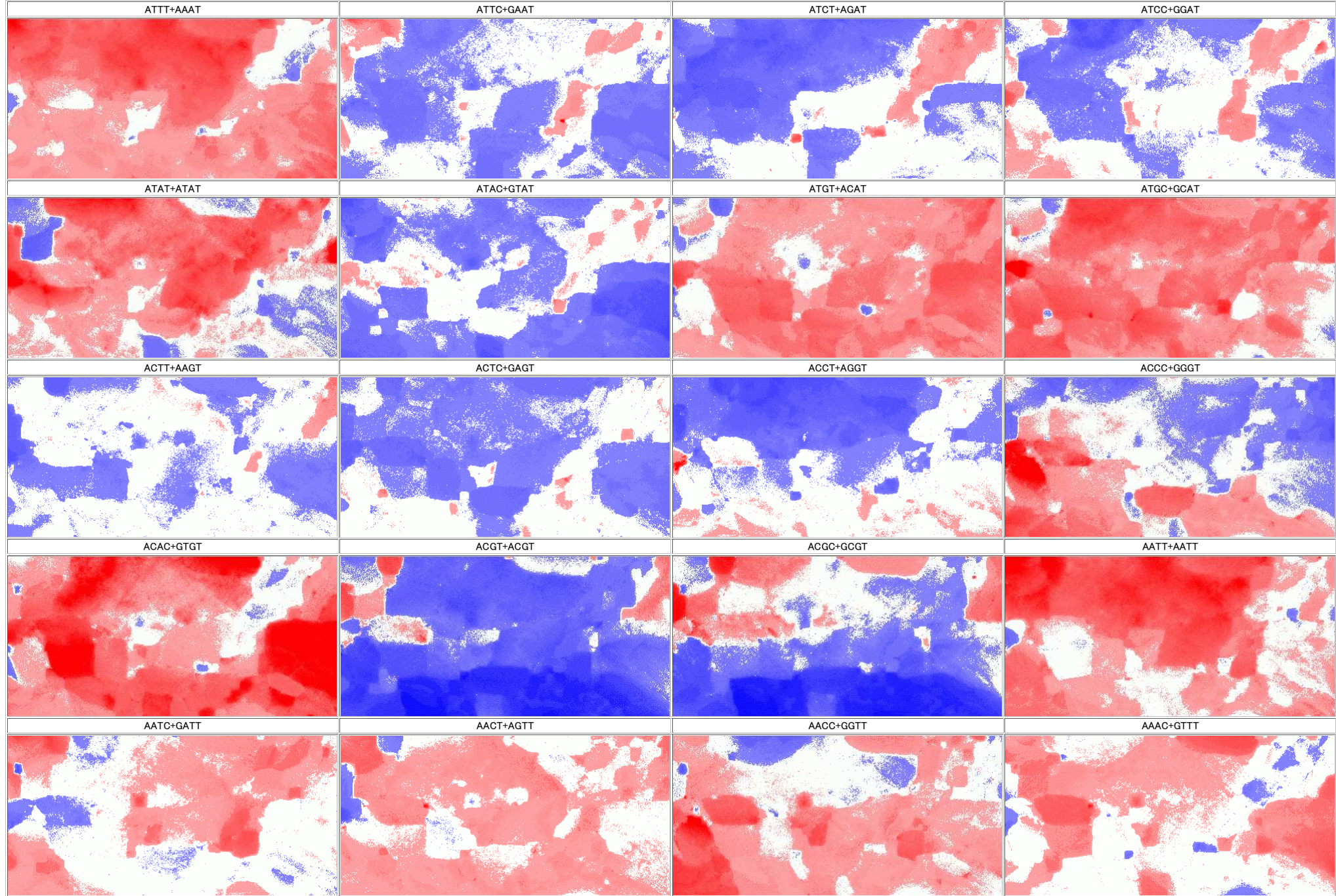


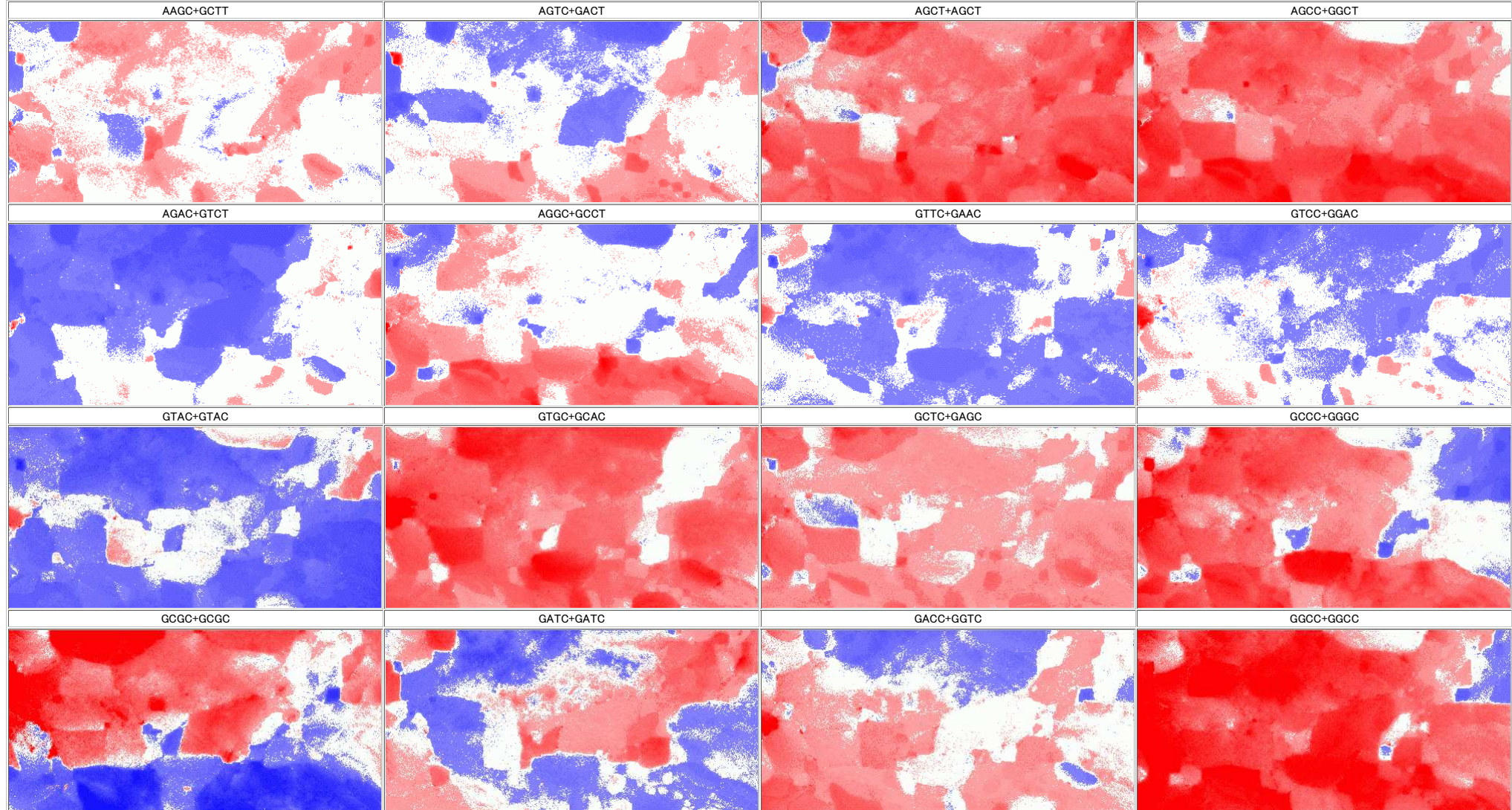




Supplementary Figure S1

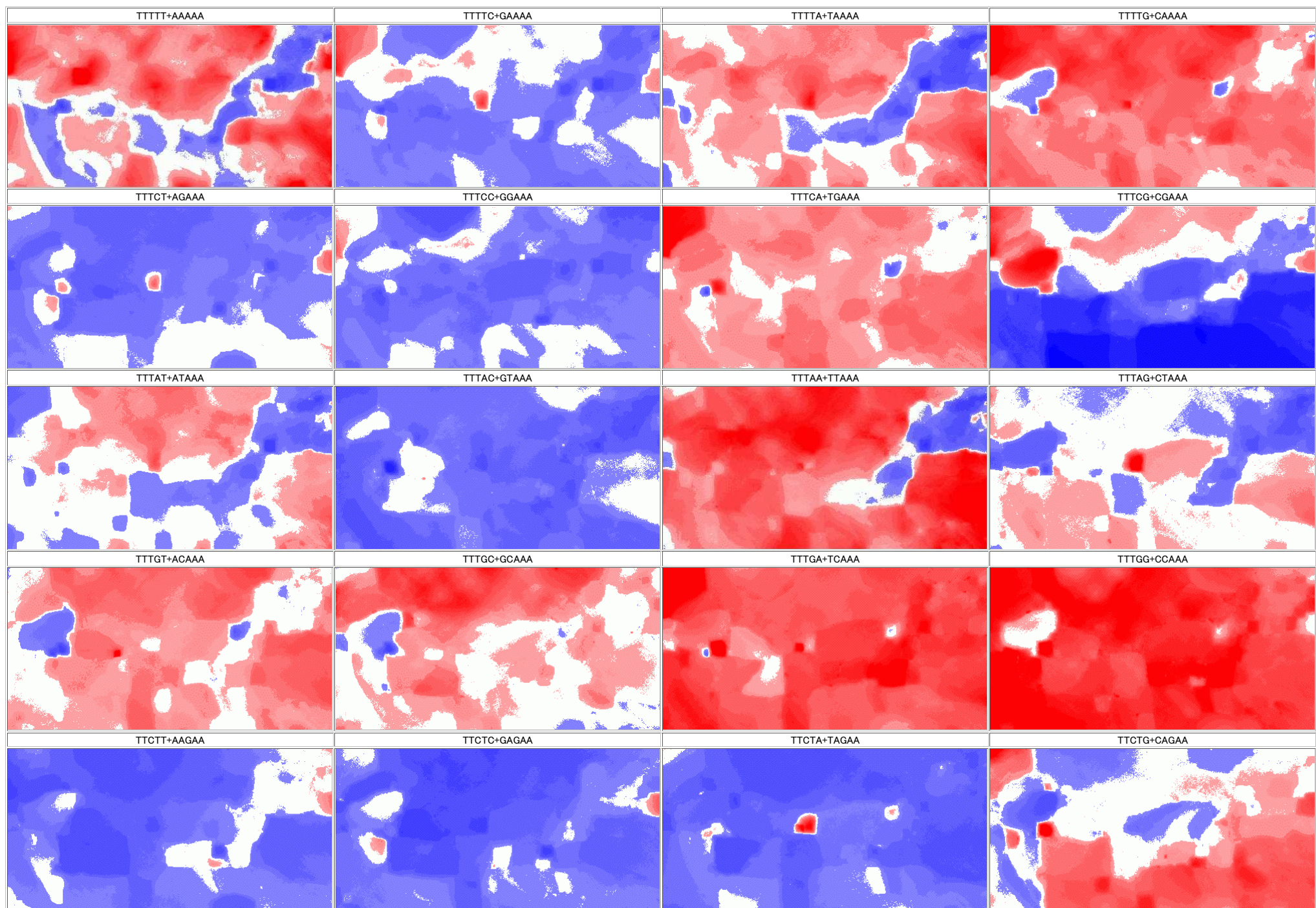


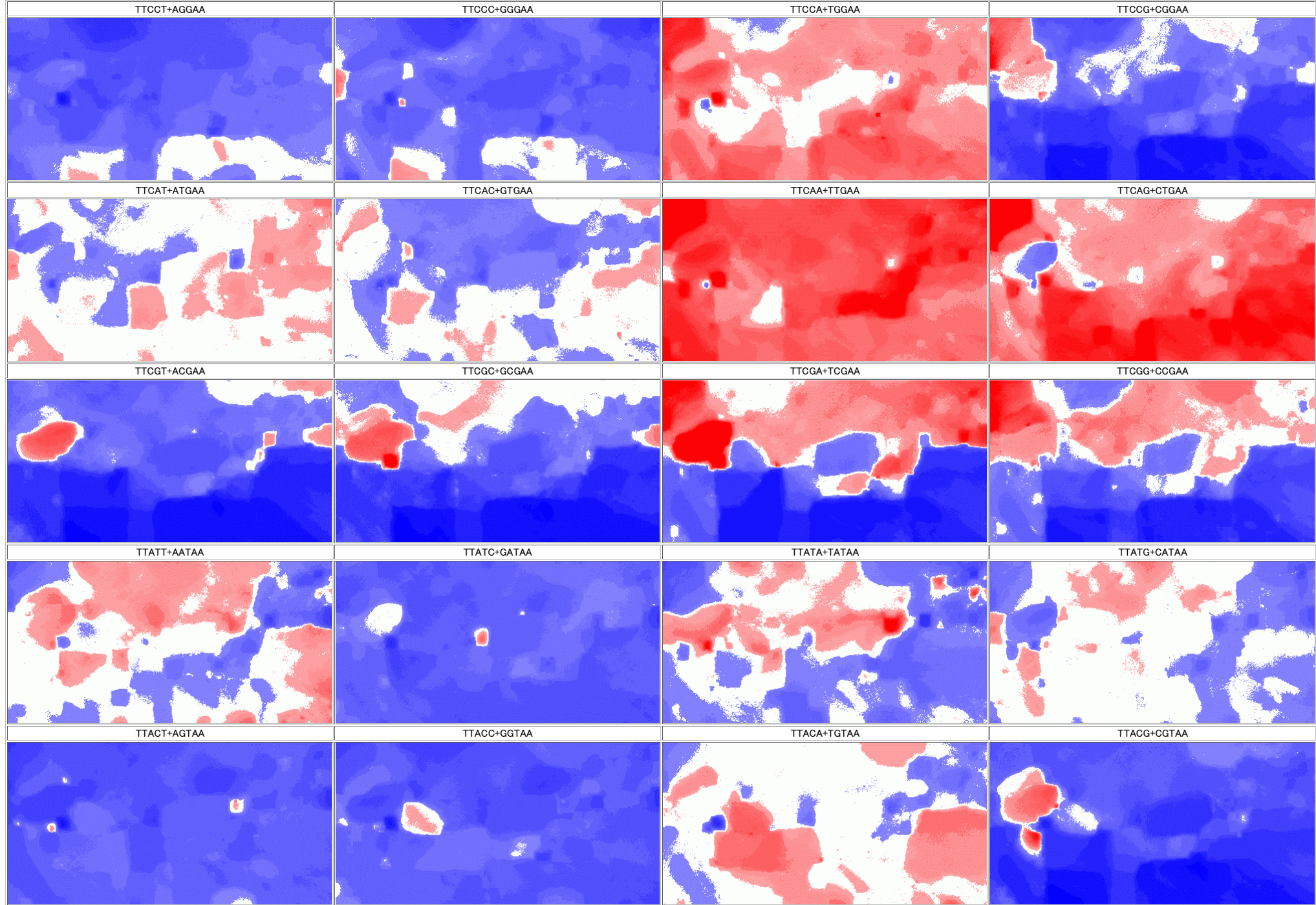


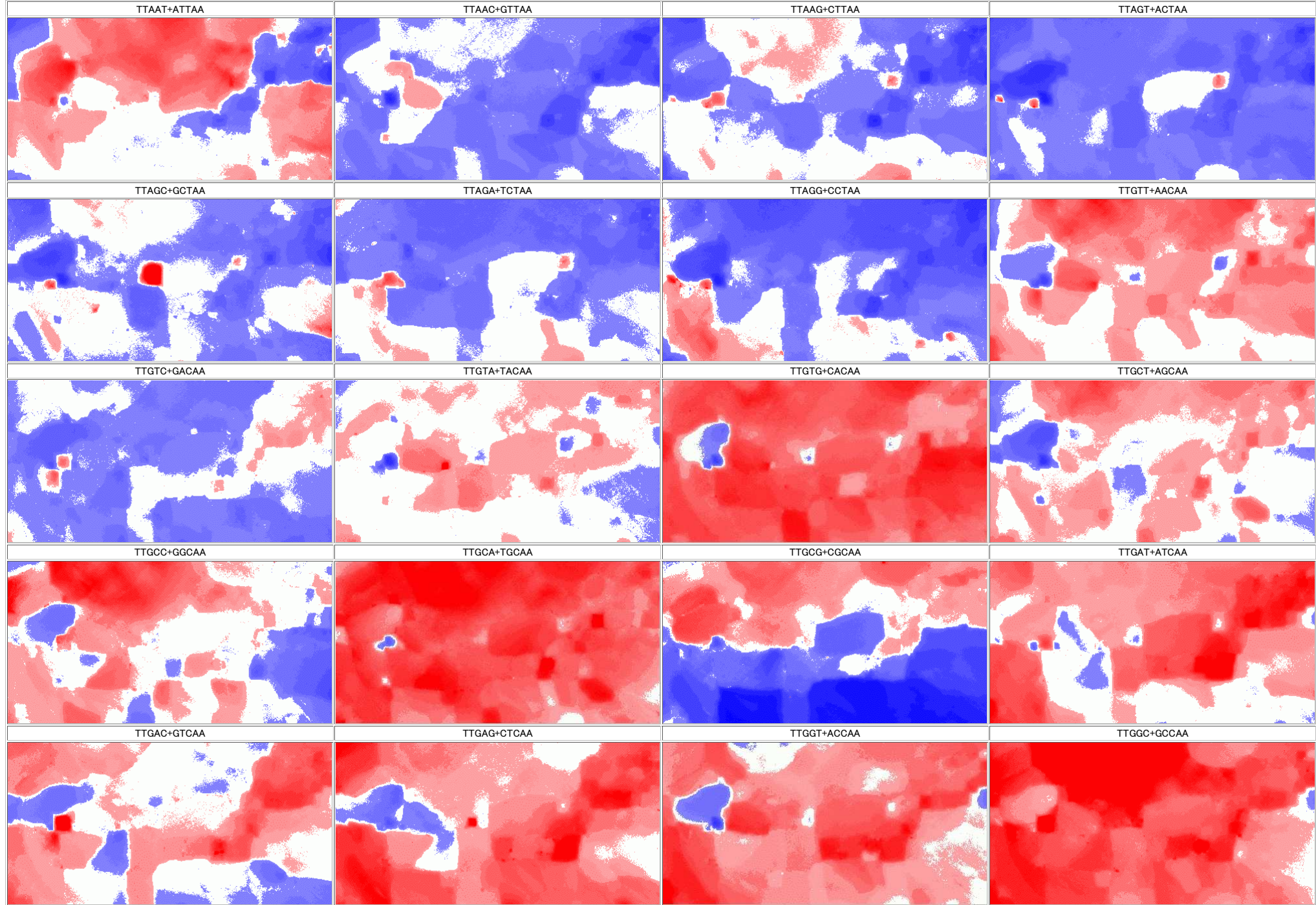


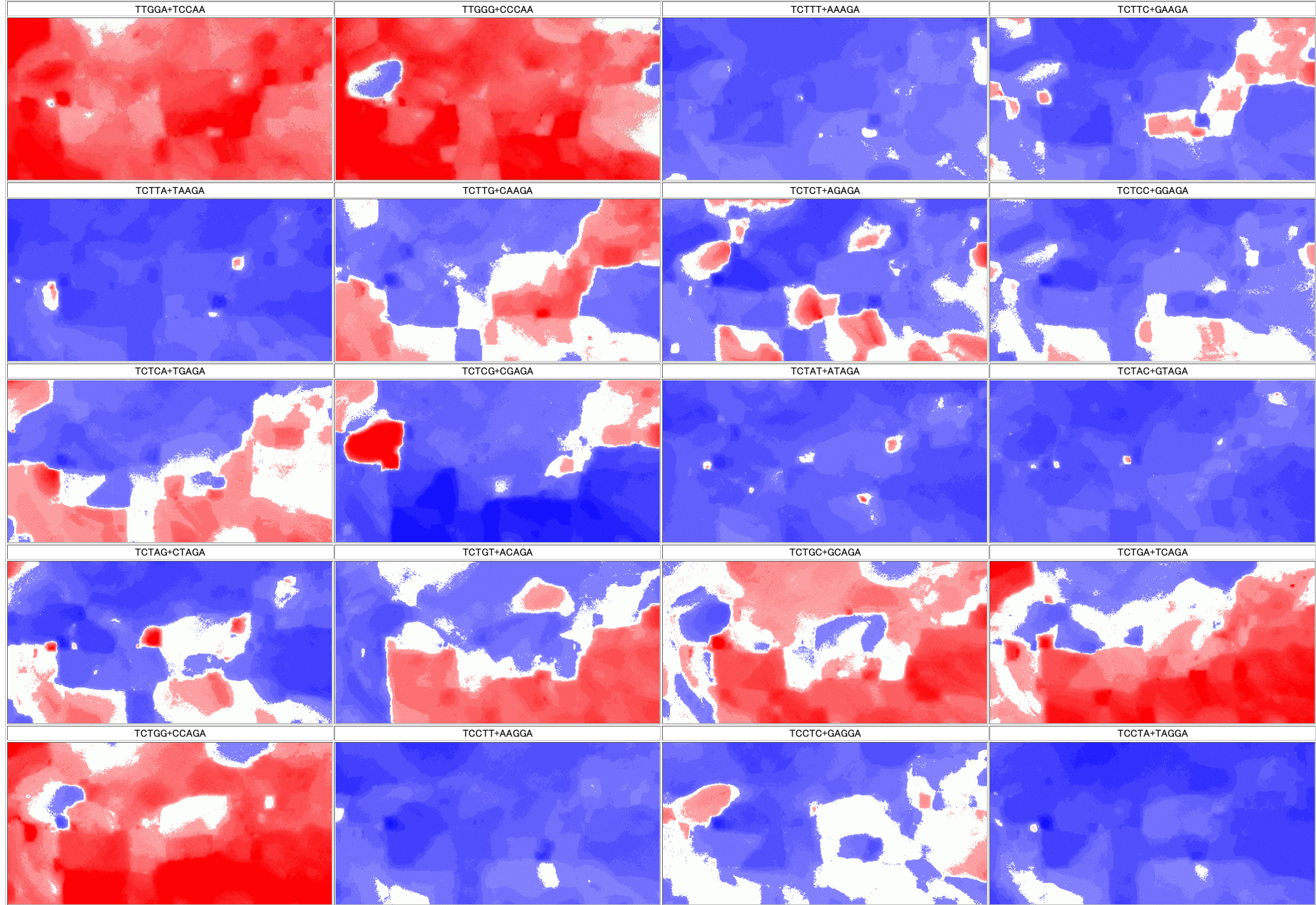
Supplementary Figure S2.

Level of each pair of complimentary pentanucleotide in 100-kb DegePenta-BLSOM for the overlapping 100-kb sequences of 101 eukaryotic genomes.

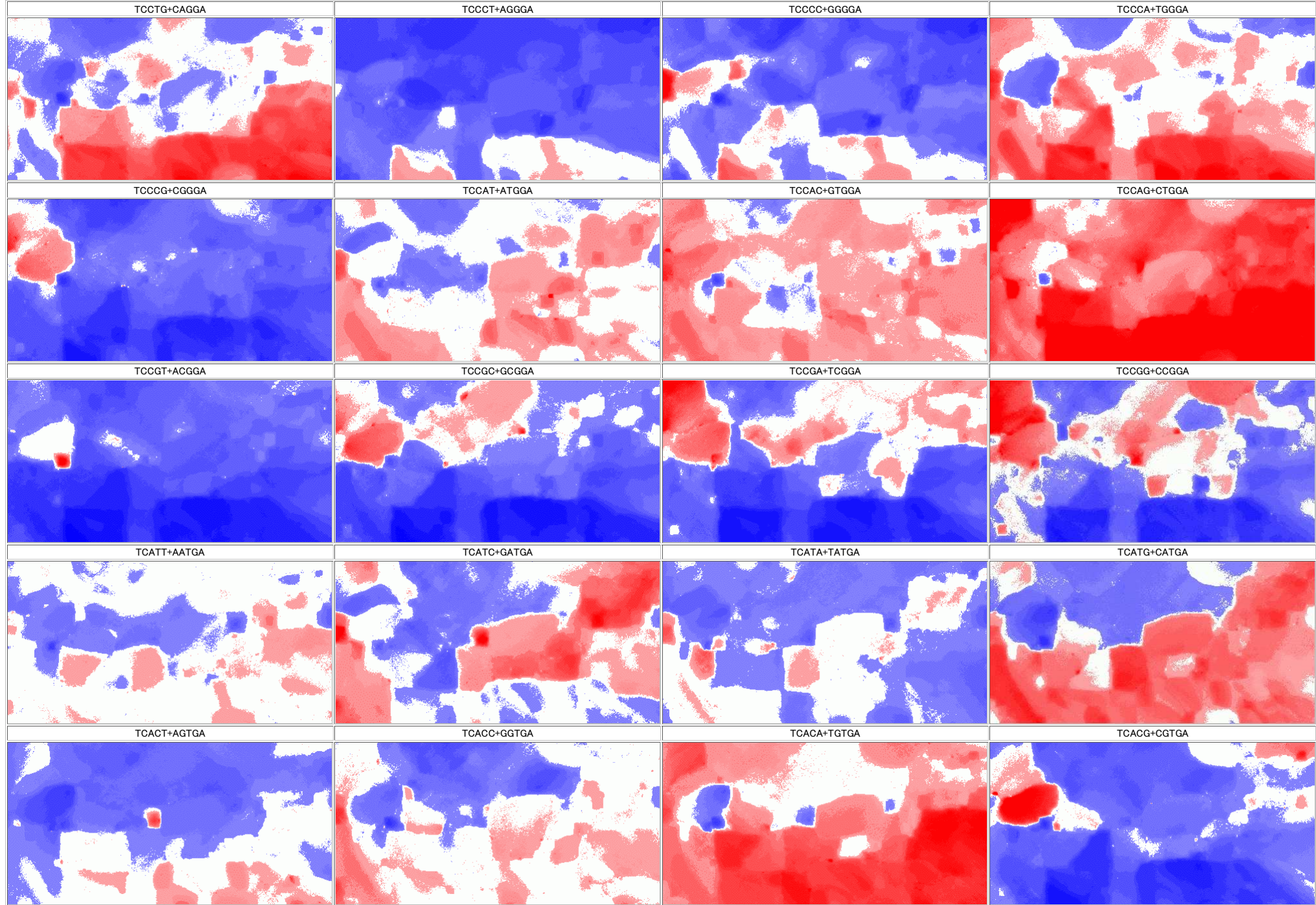


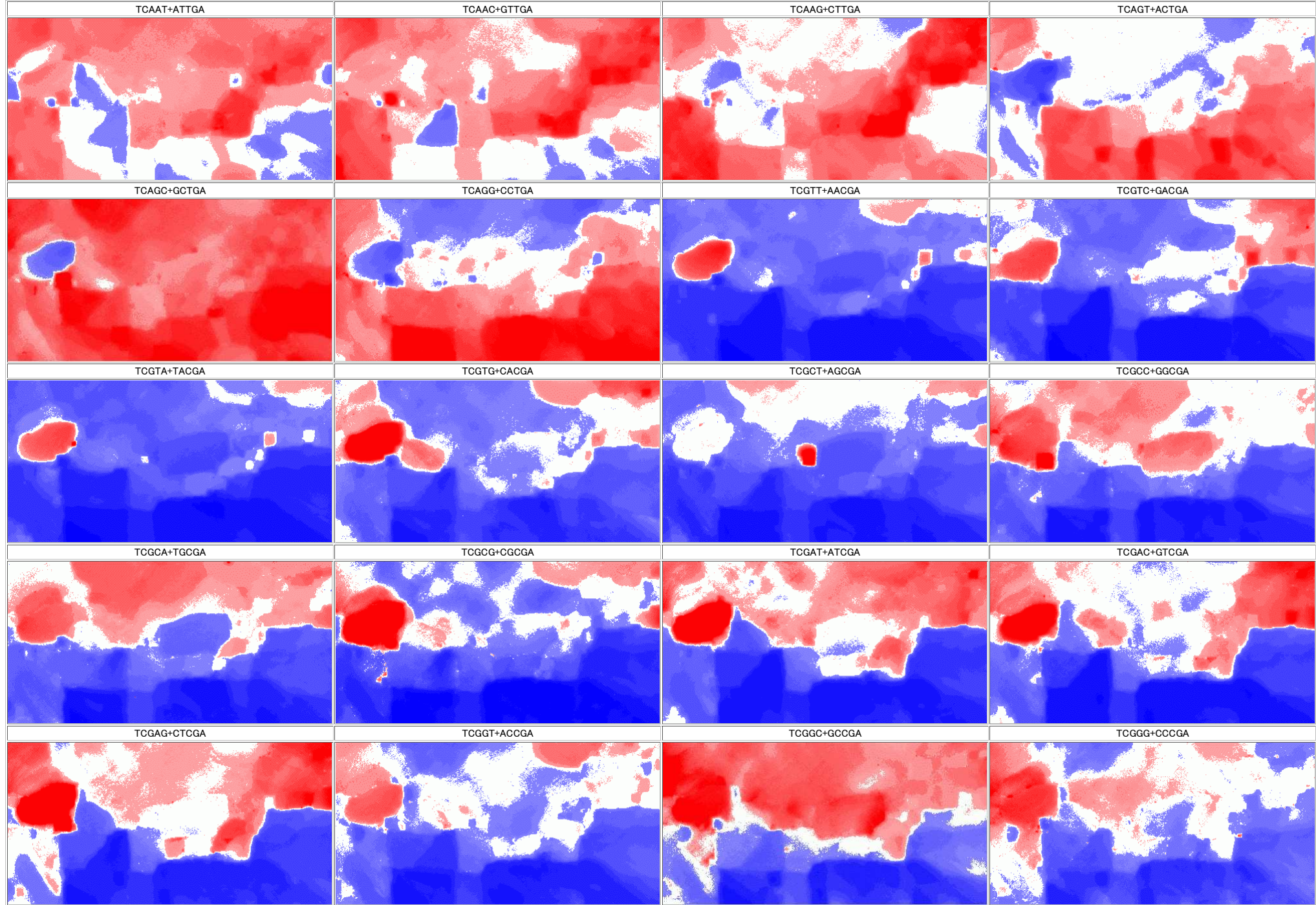


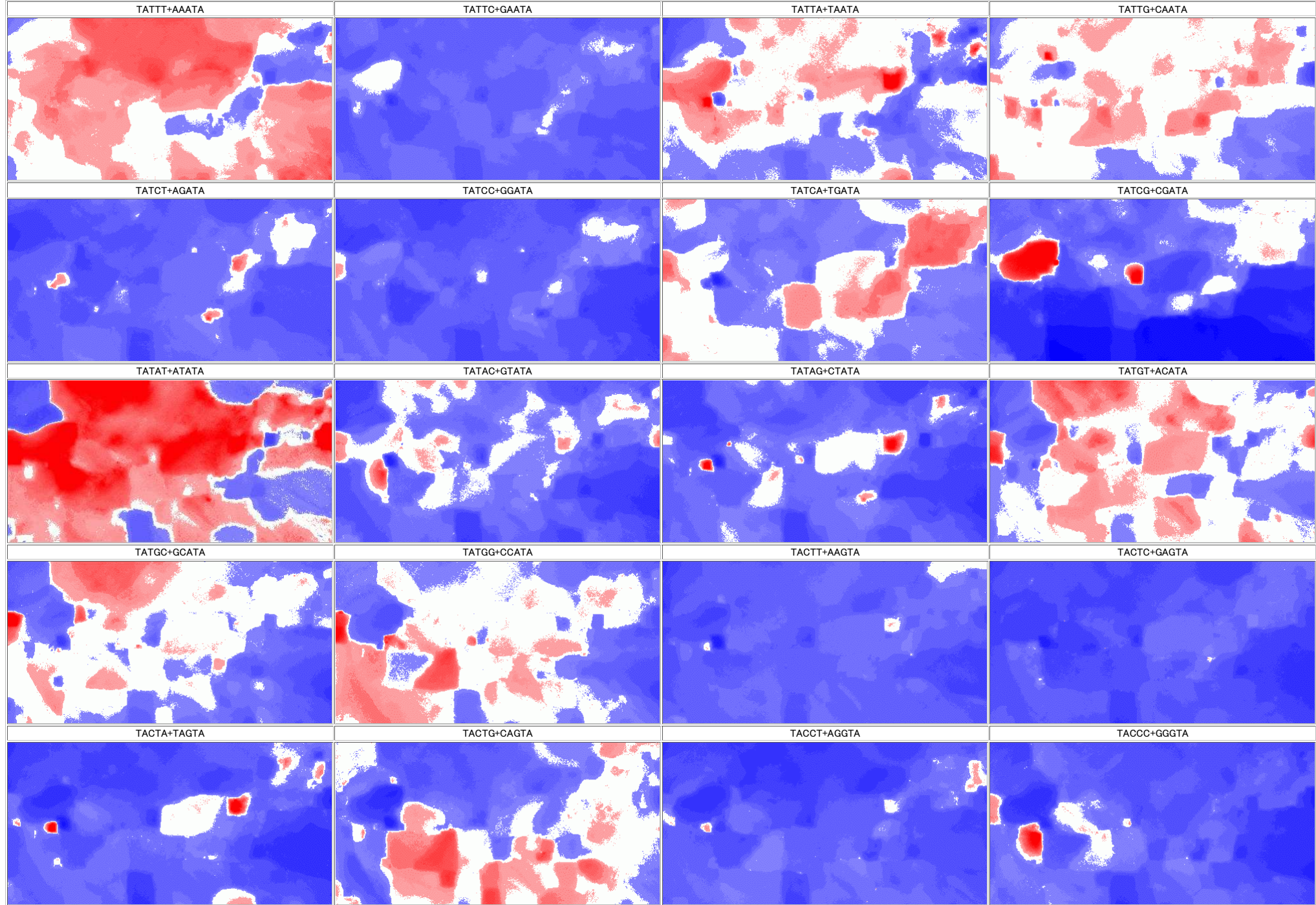


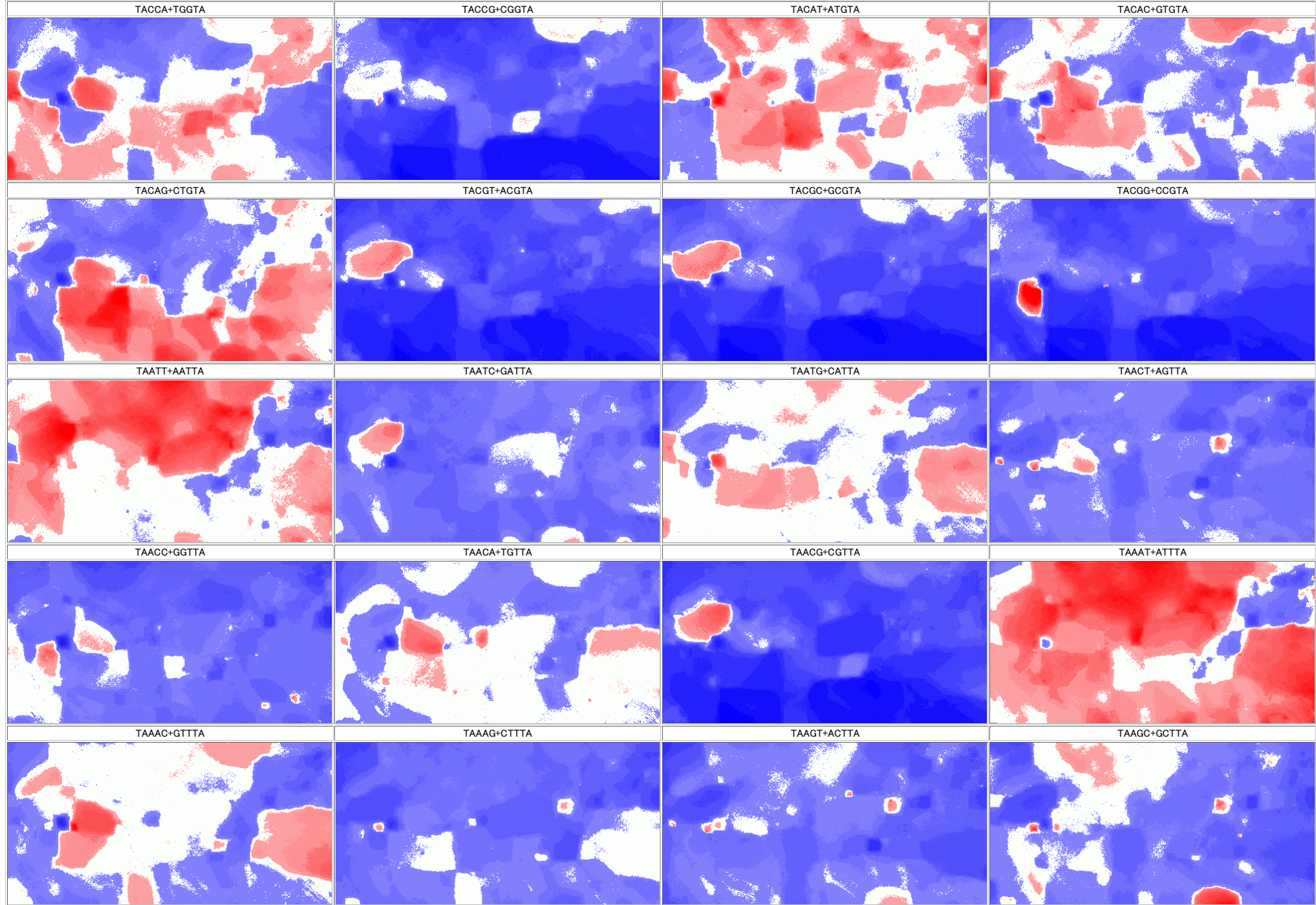


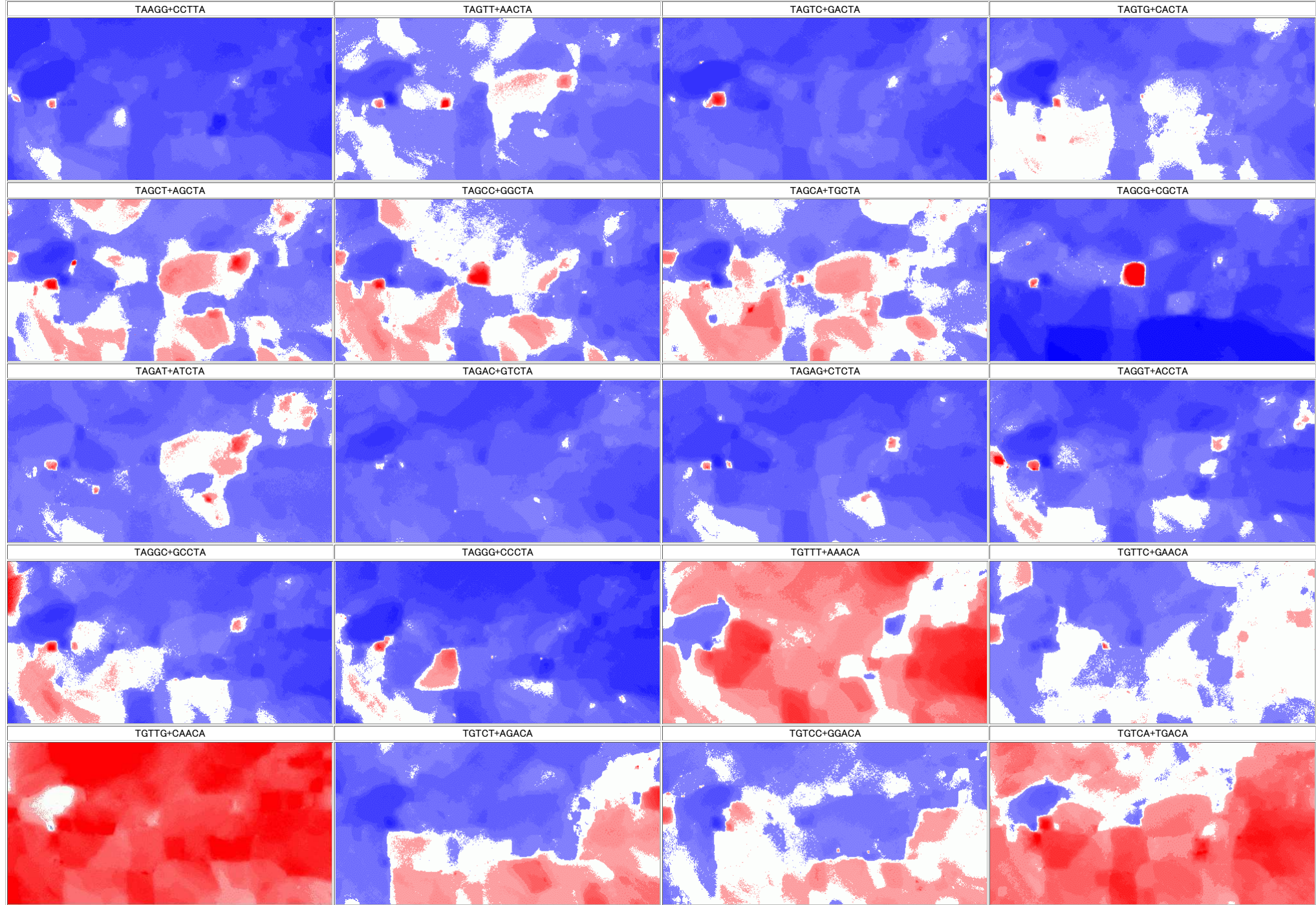
Supplementary Figure S2

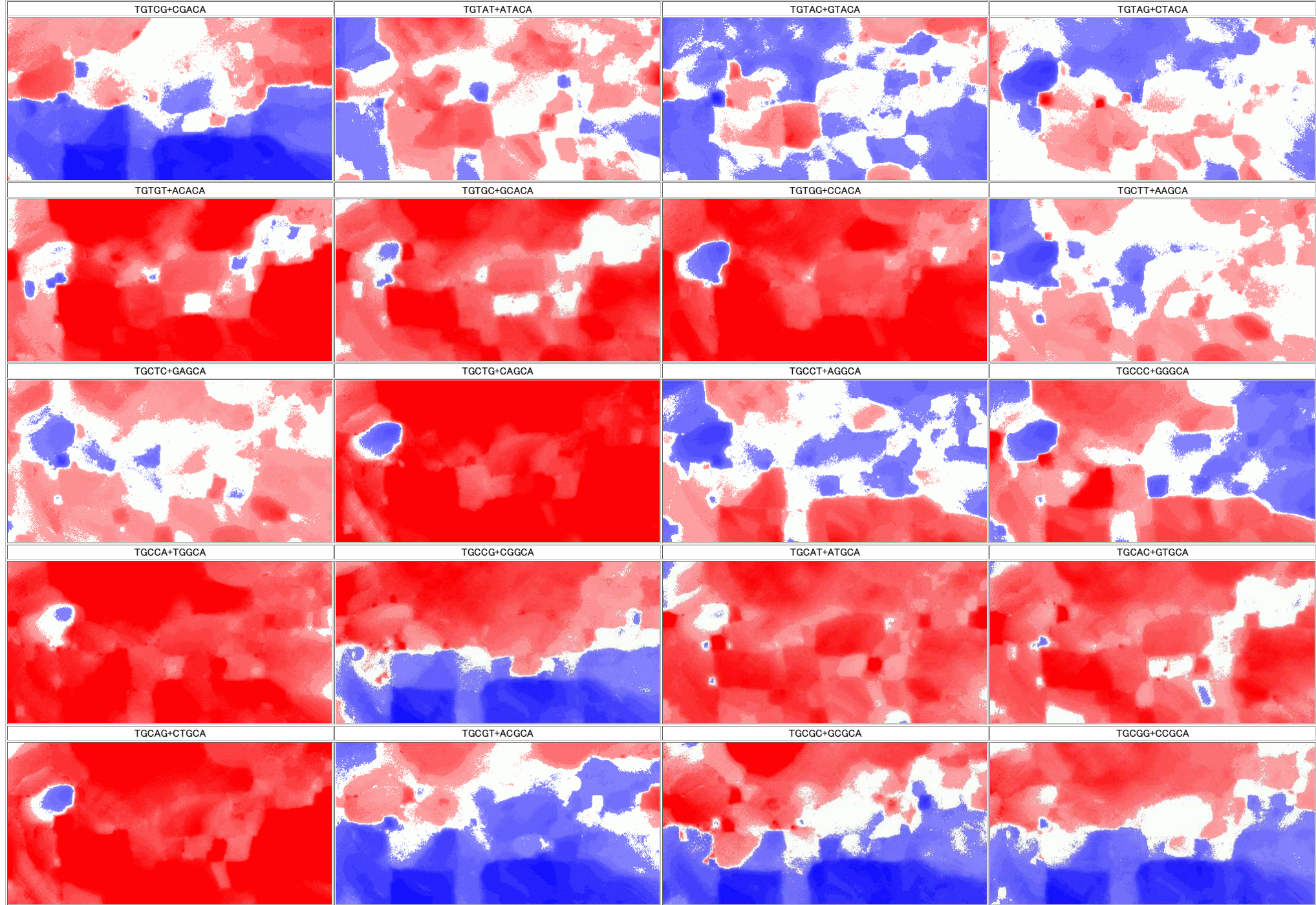


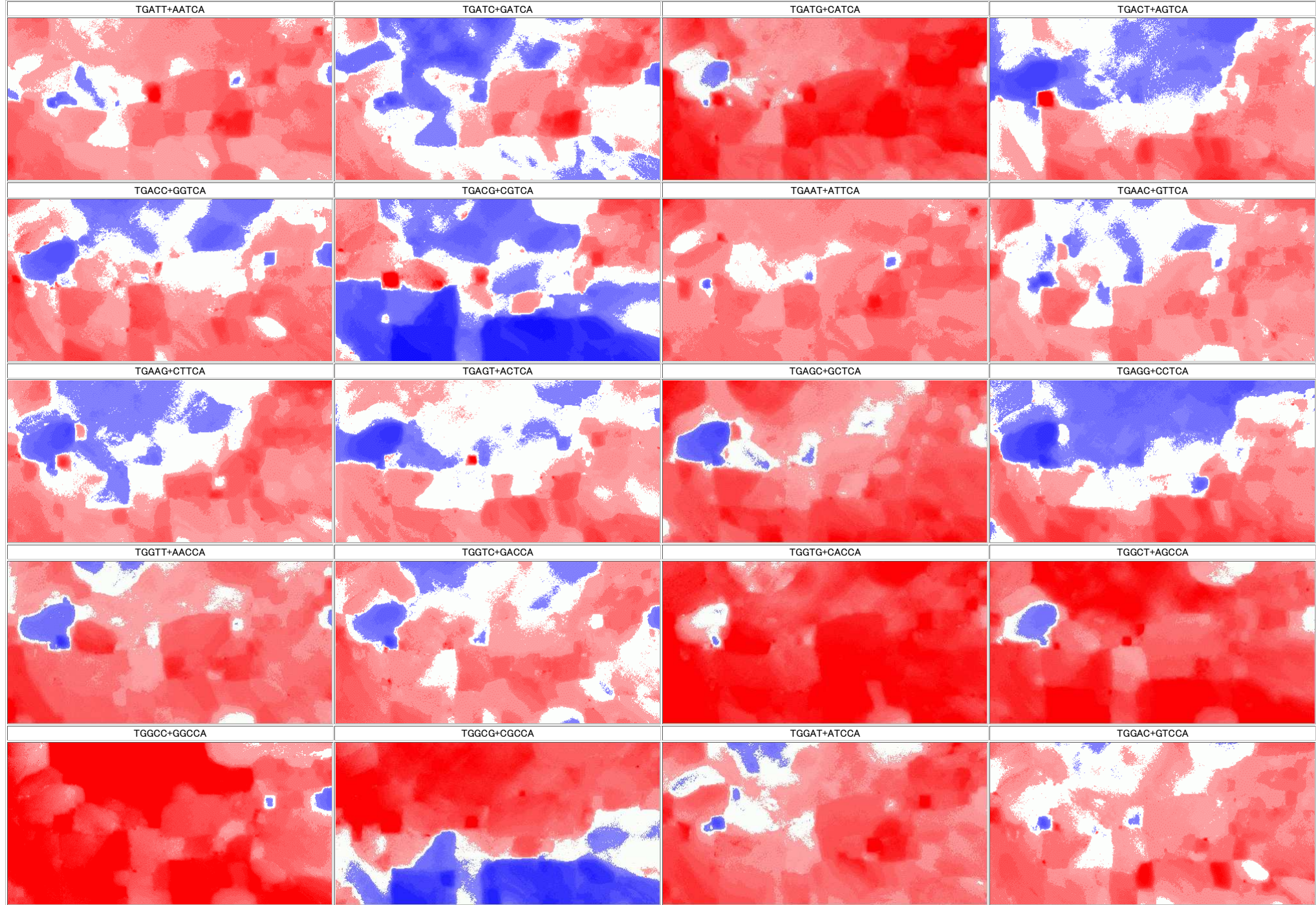


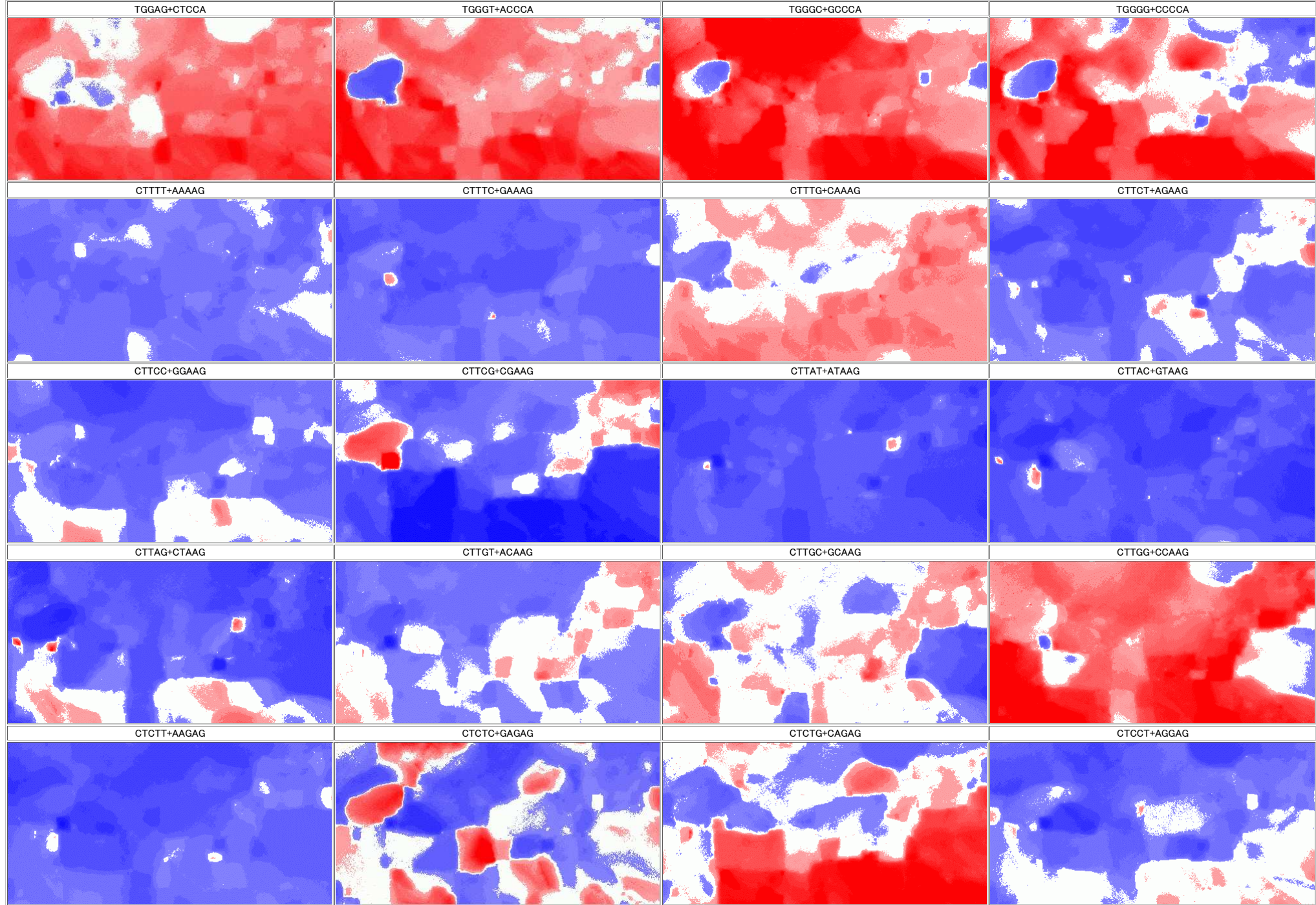


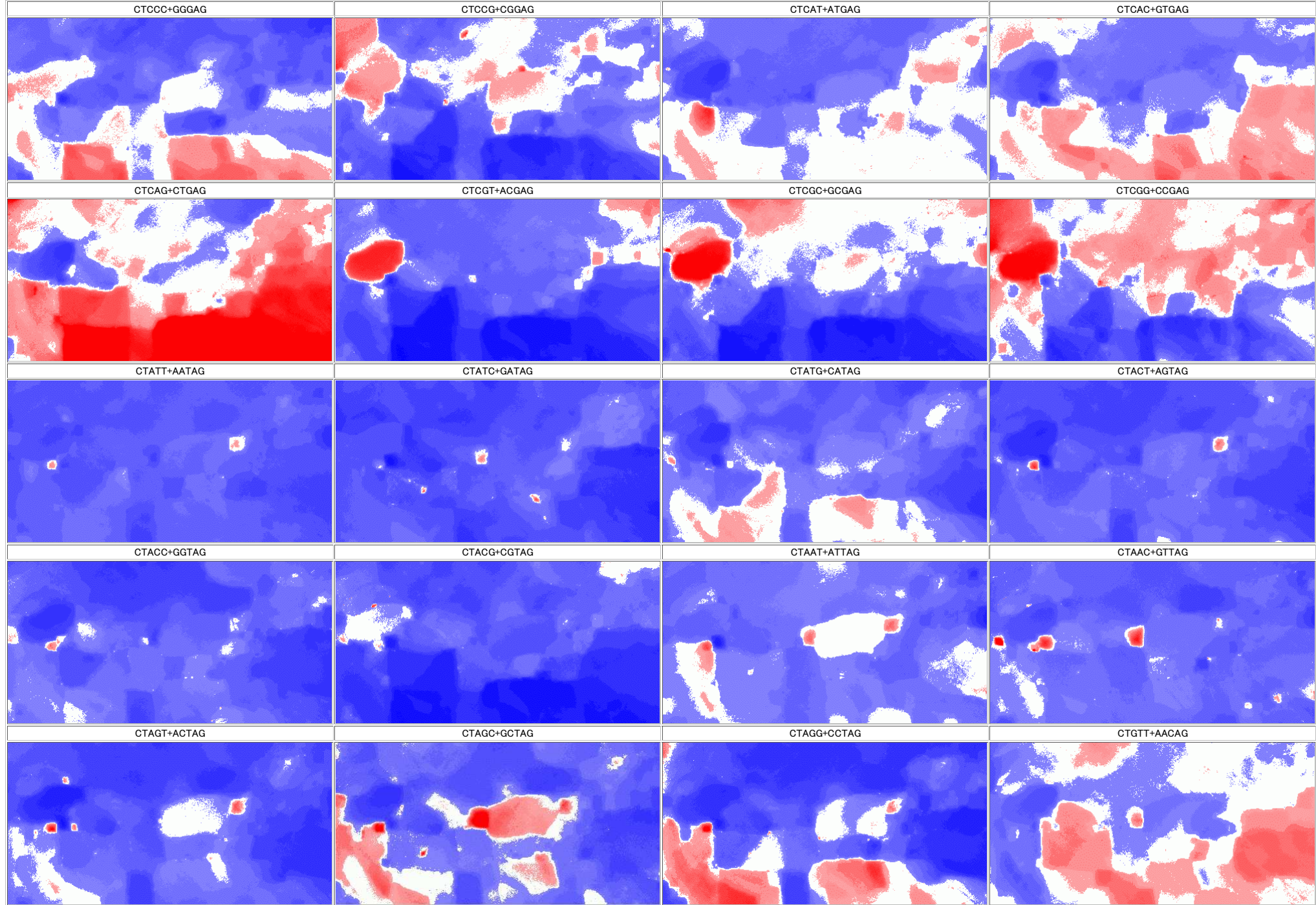


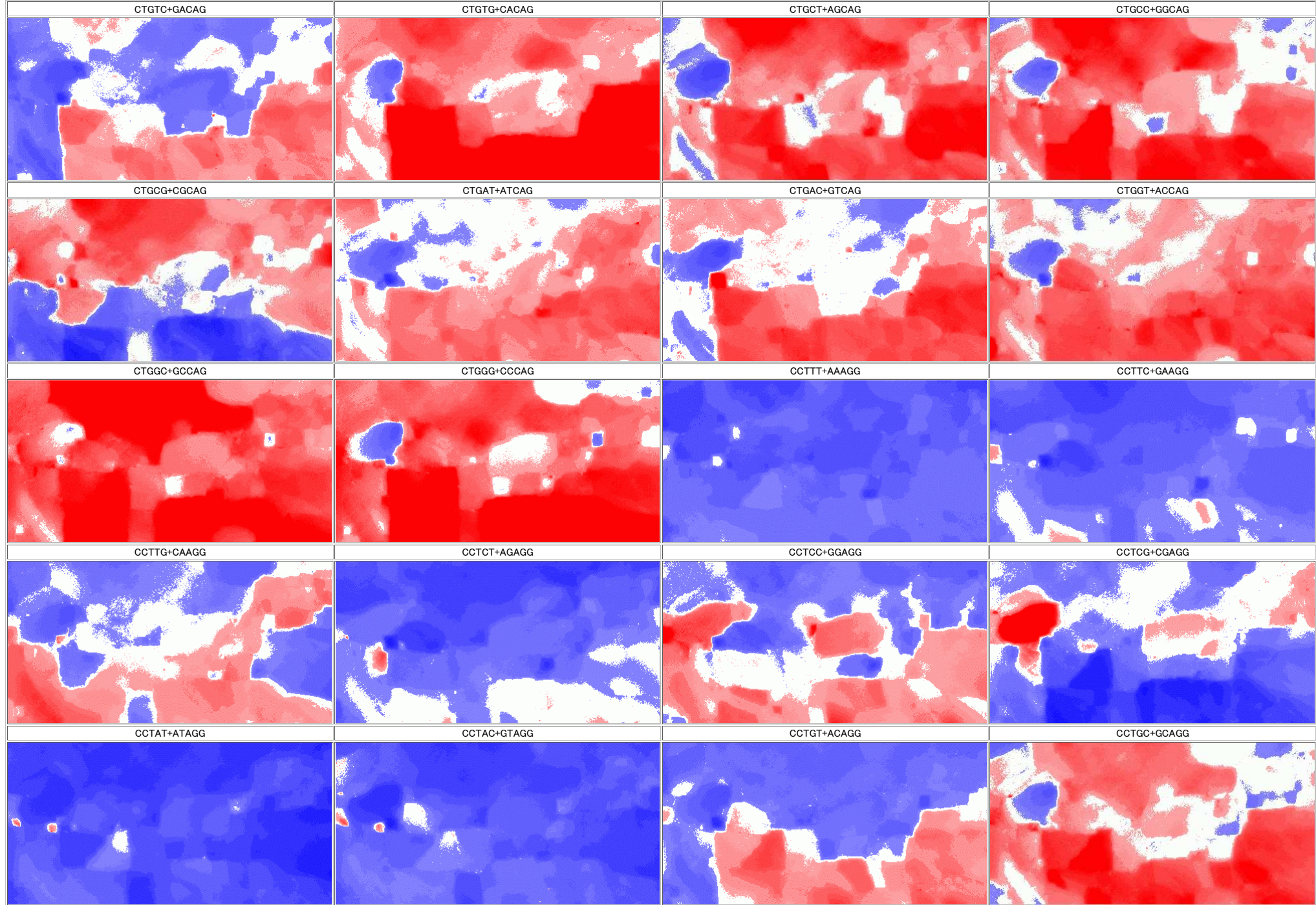


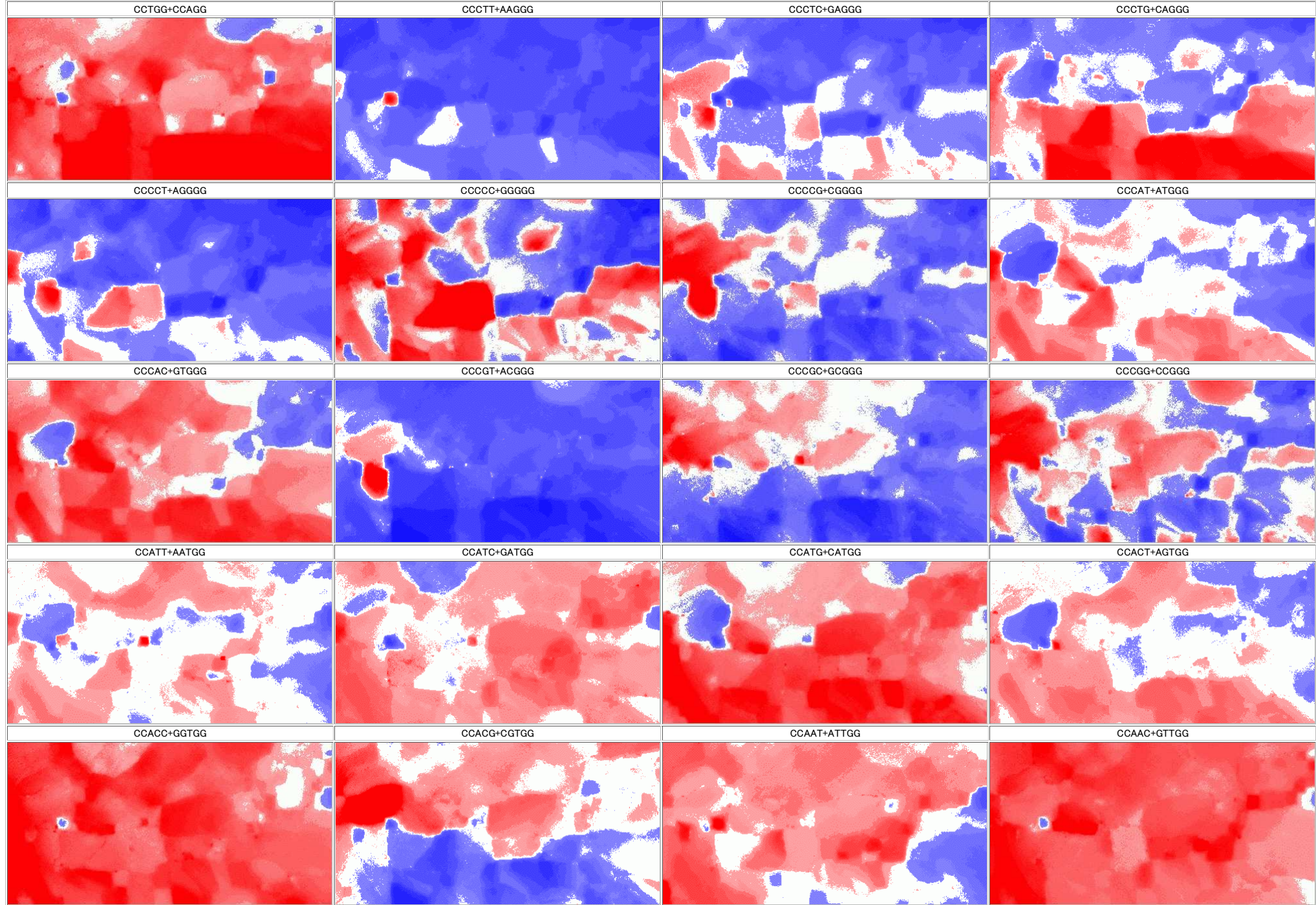


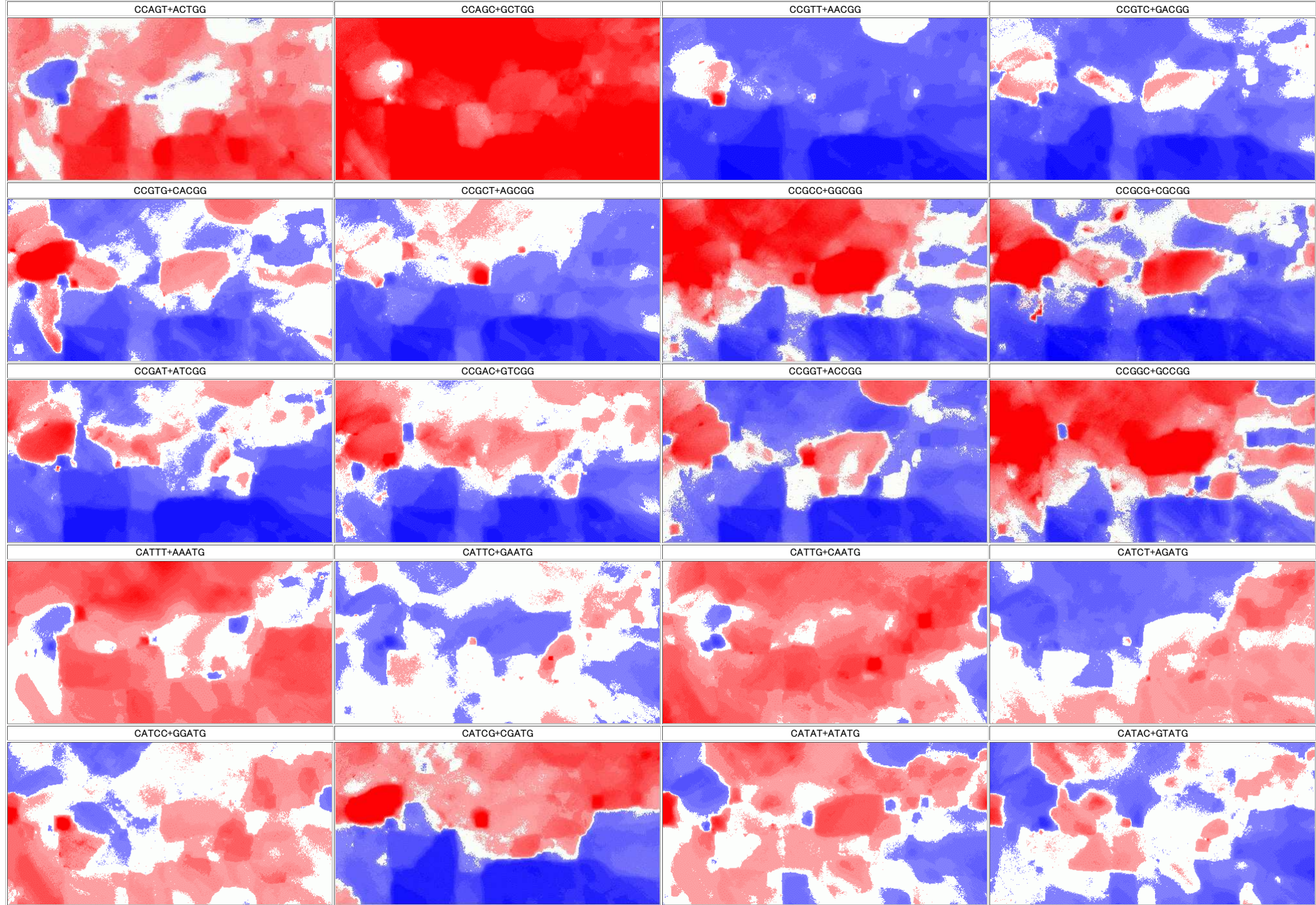


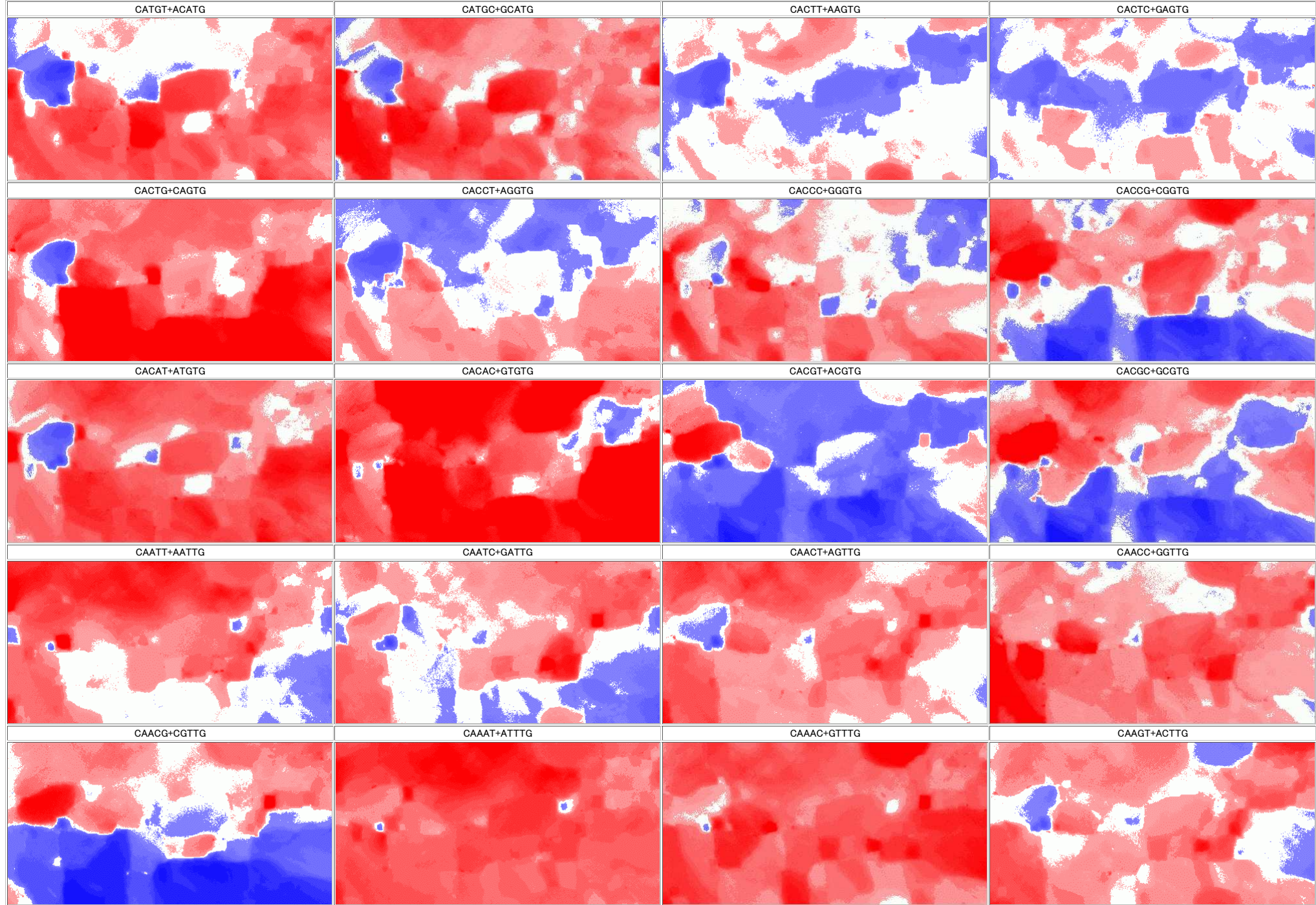


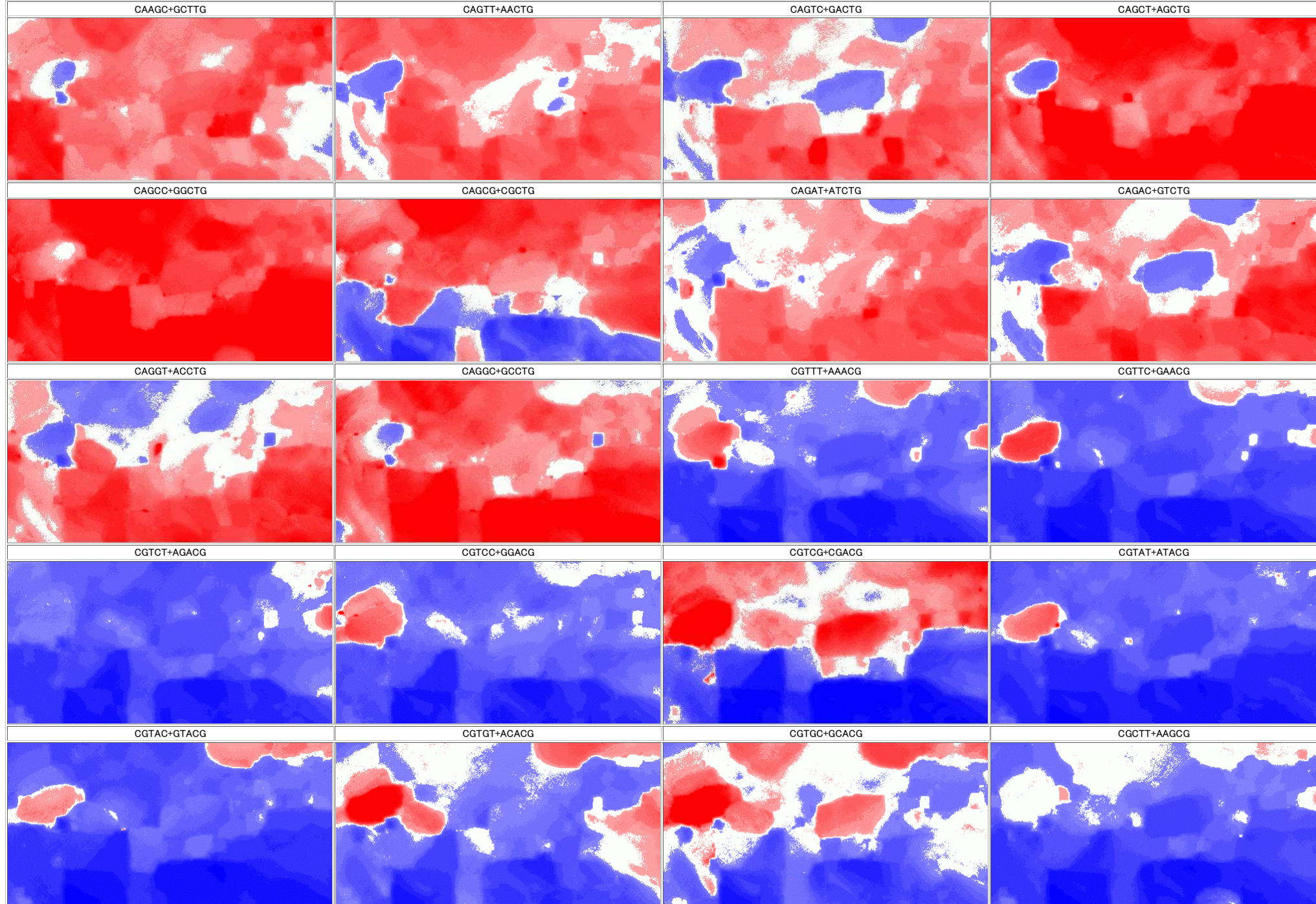


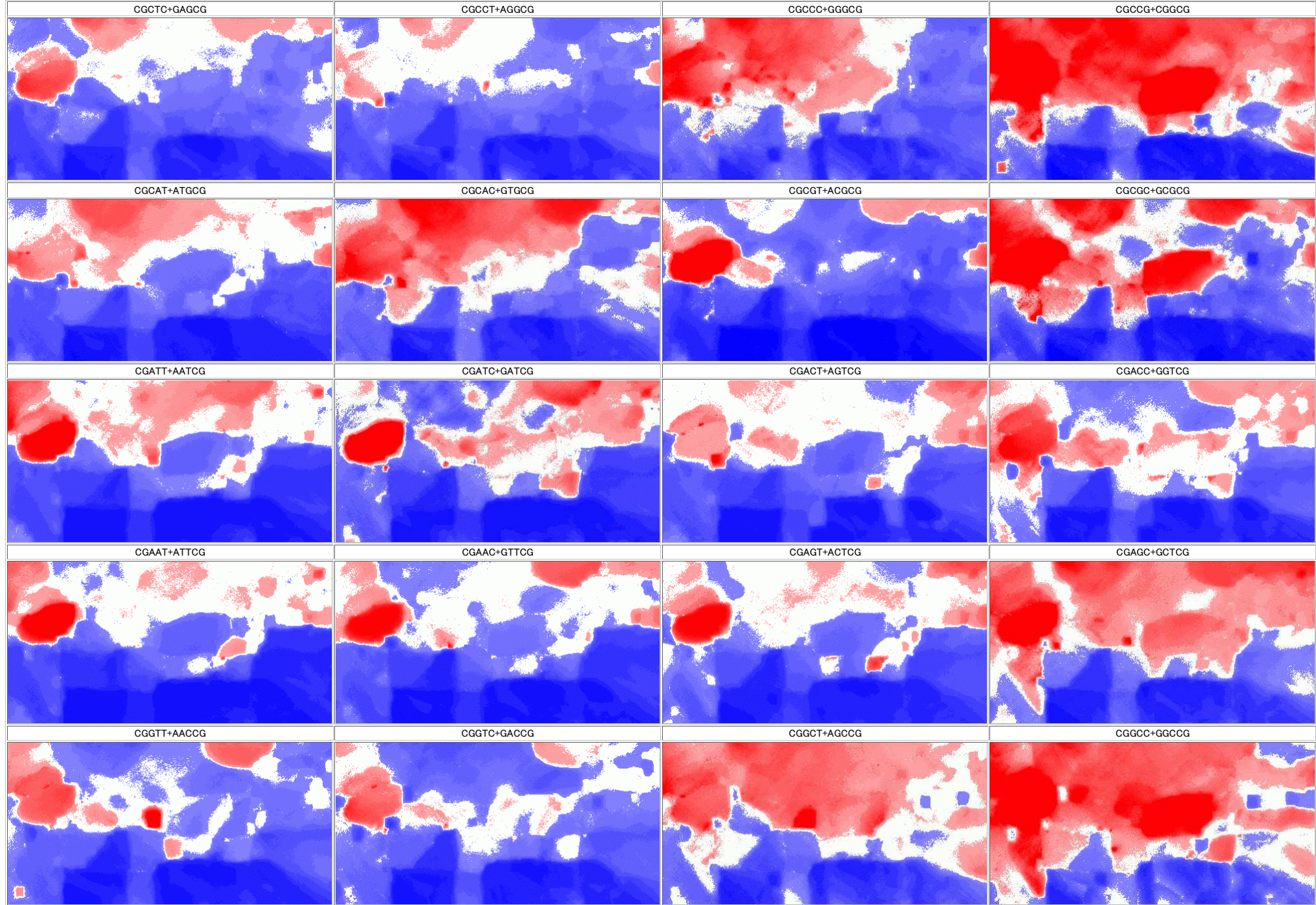


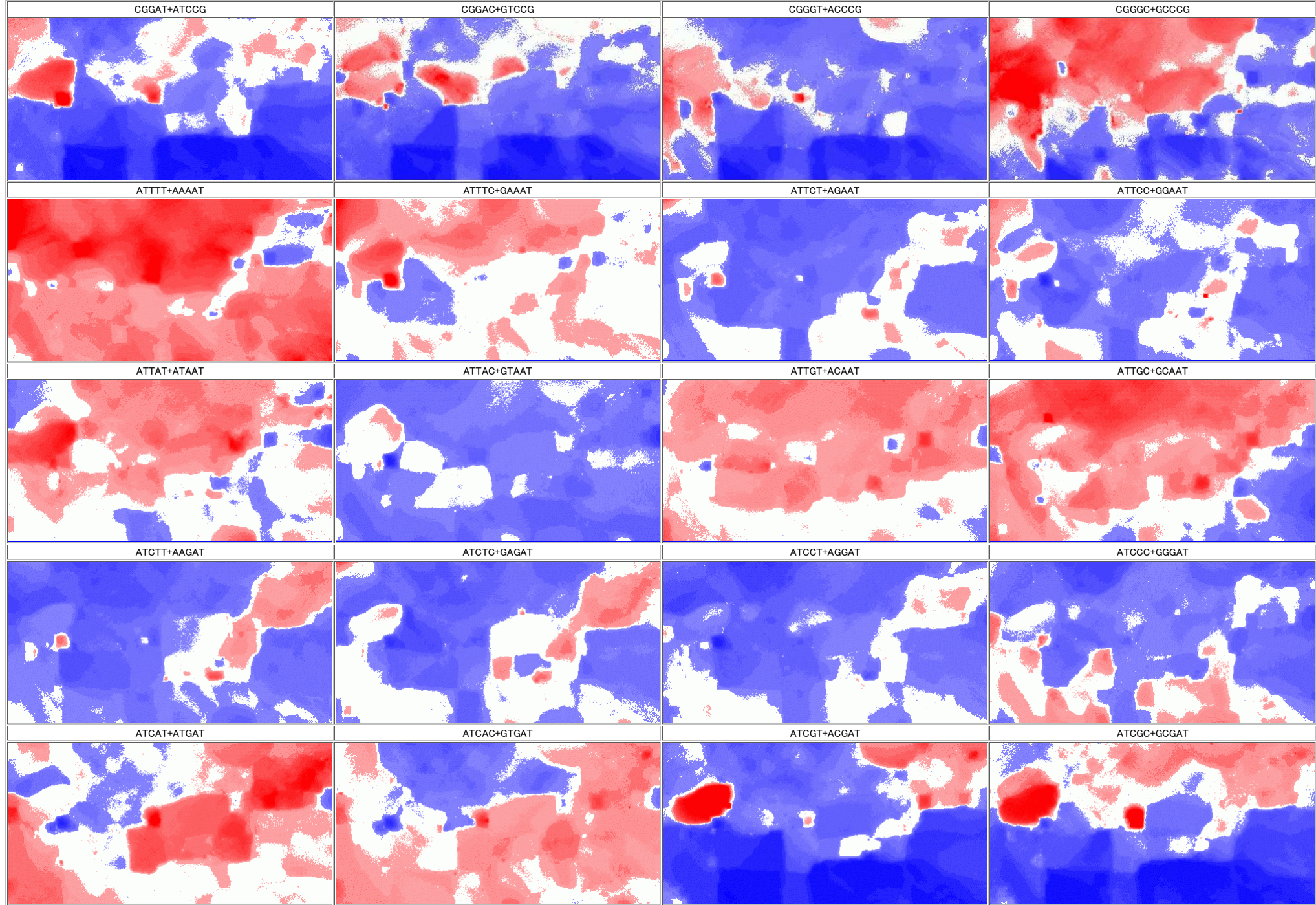


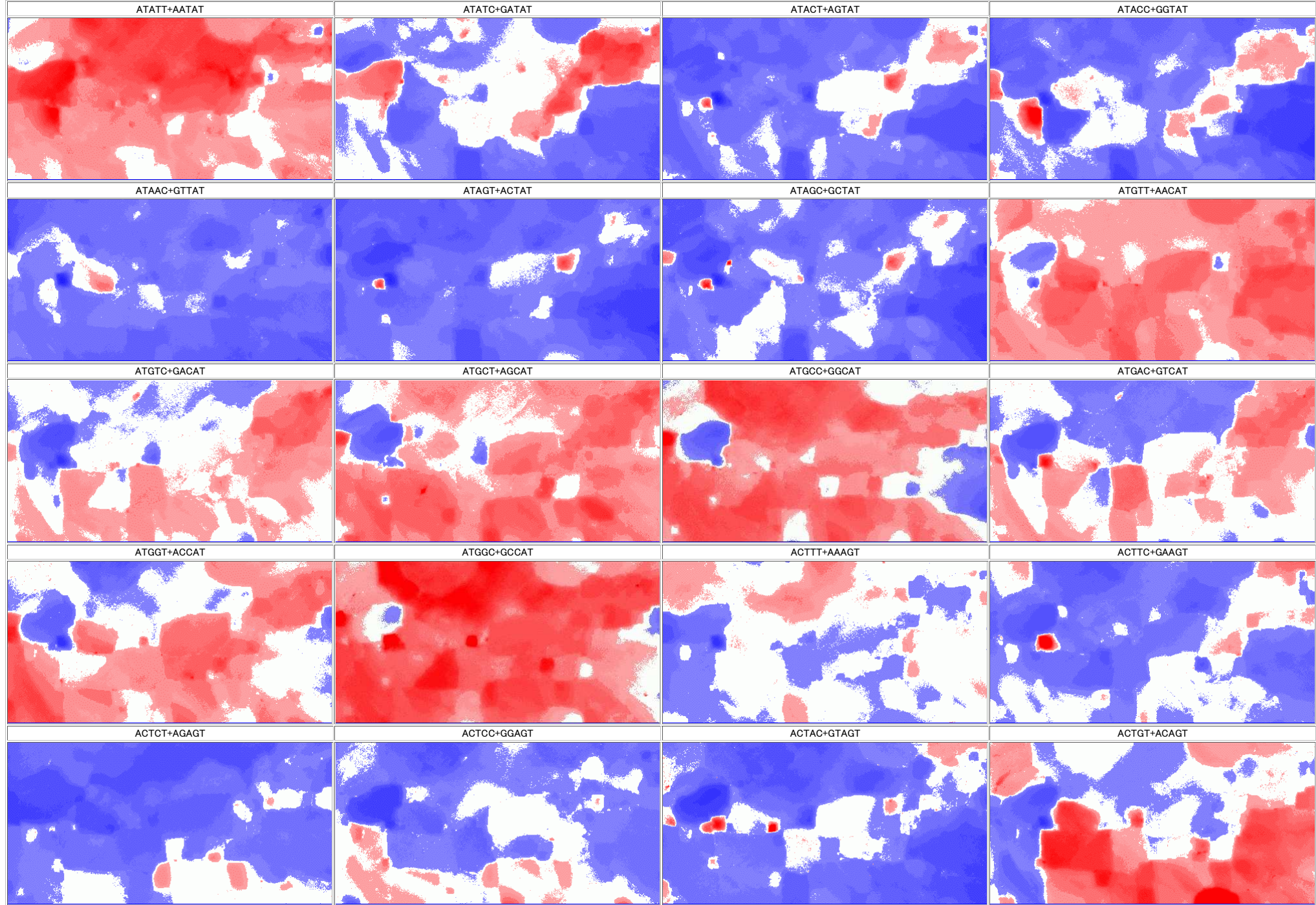


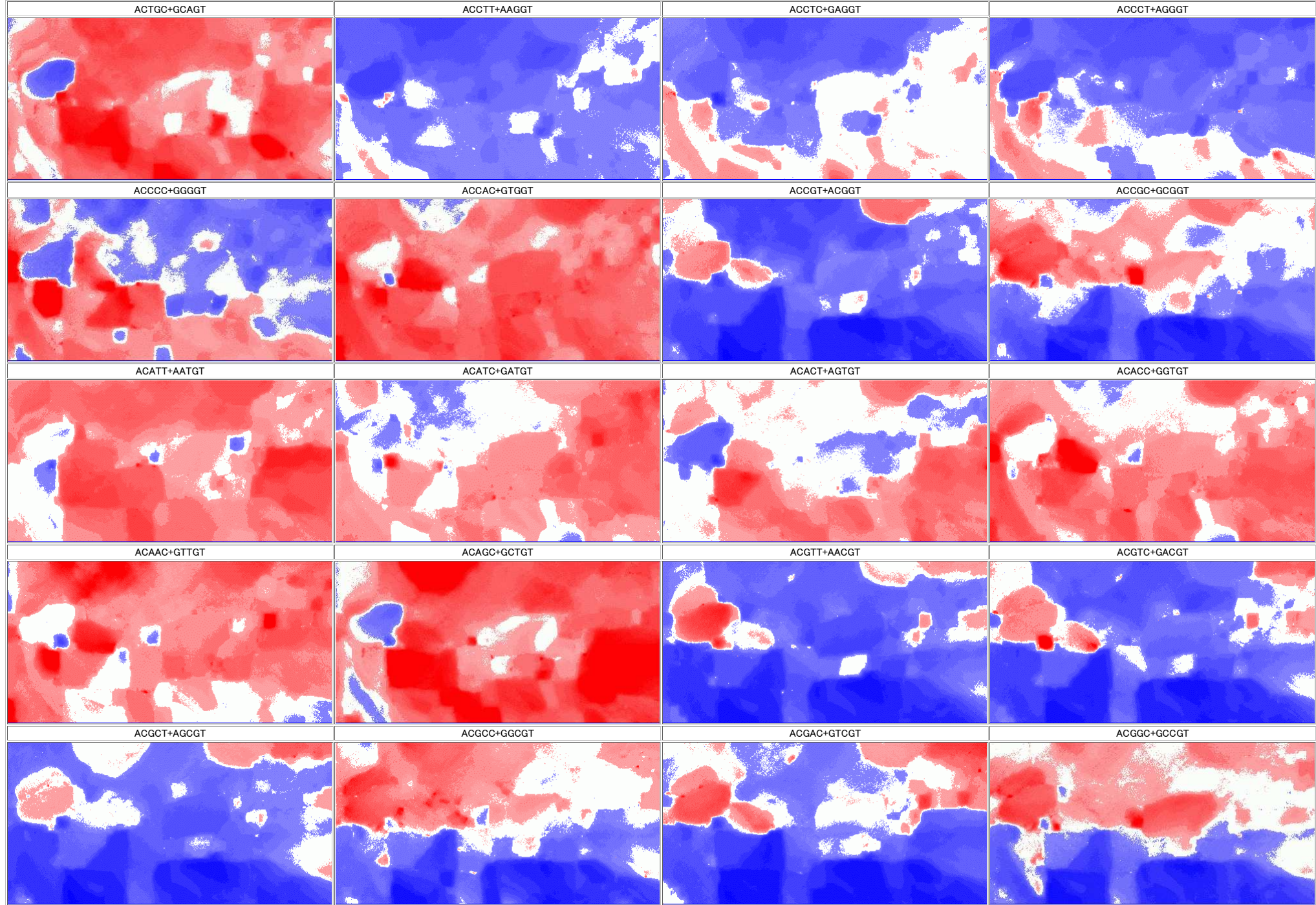


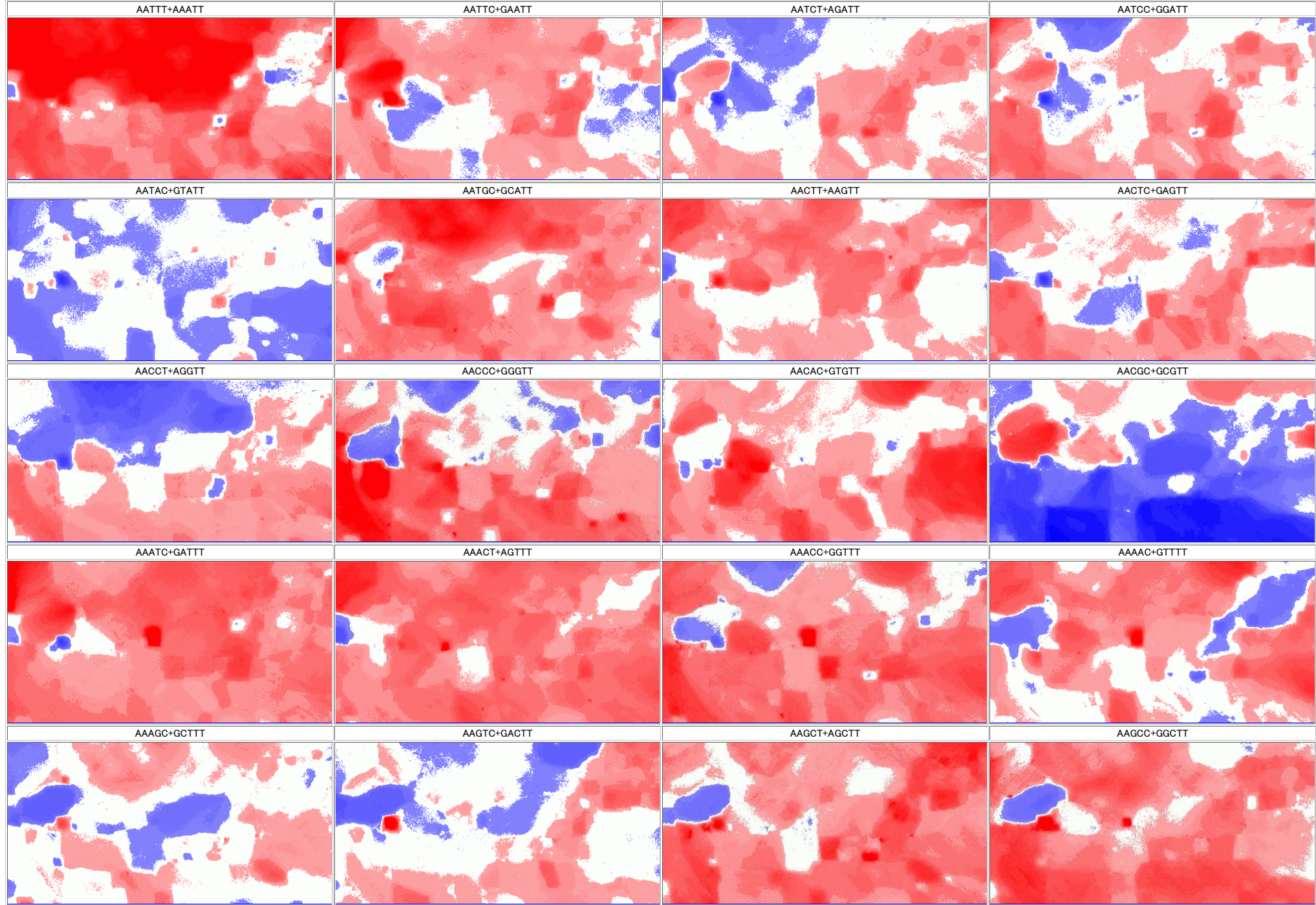


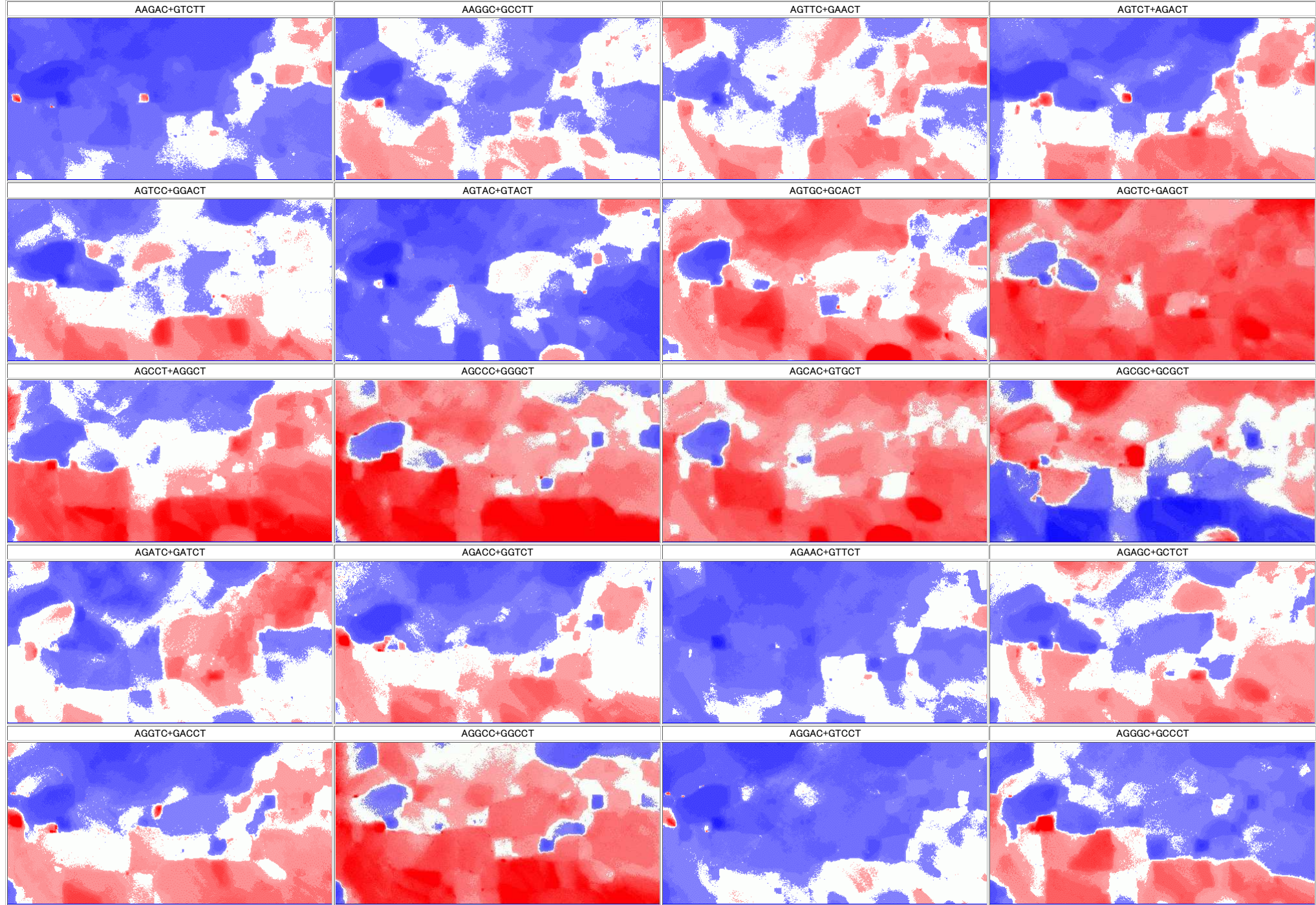


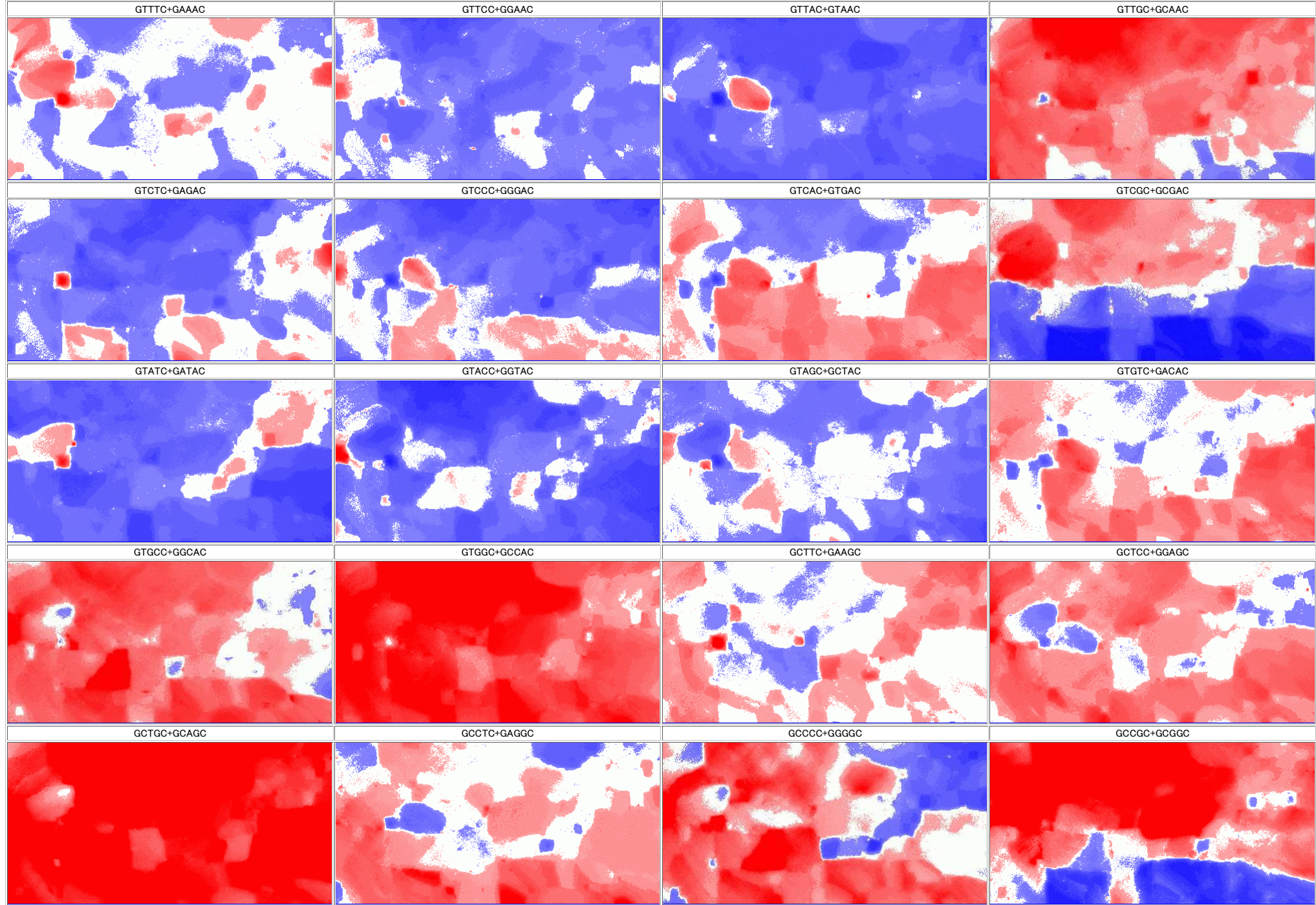


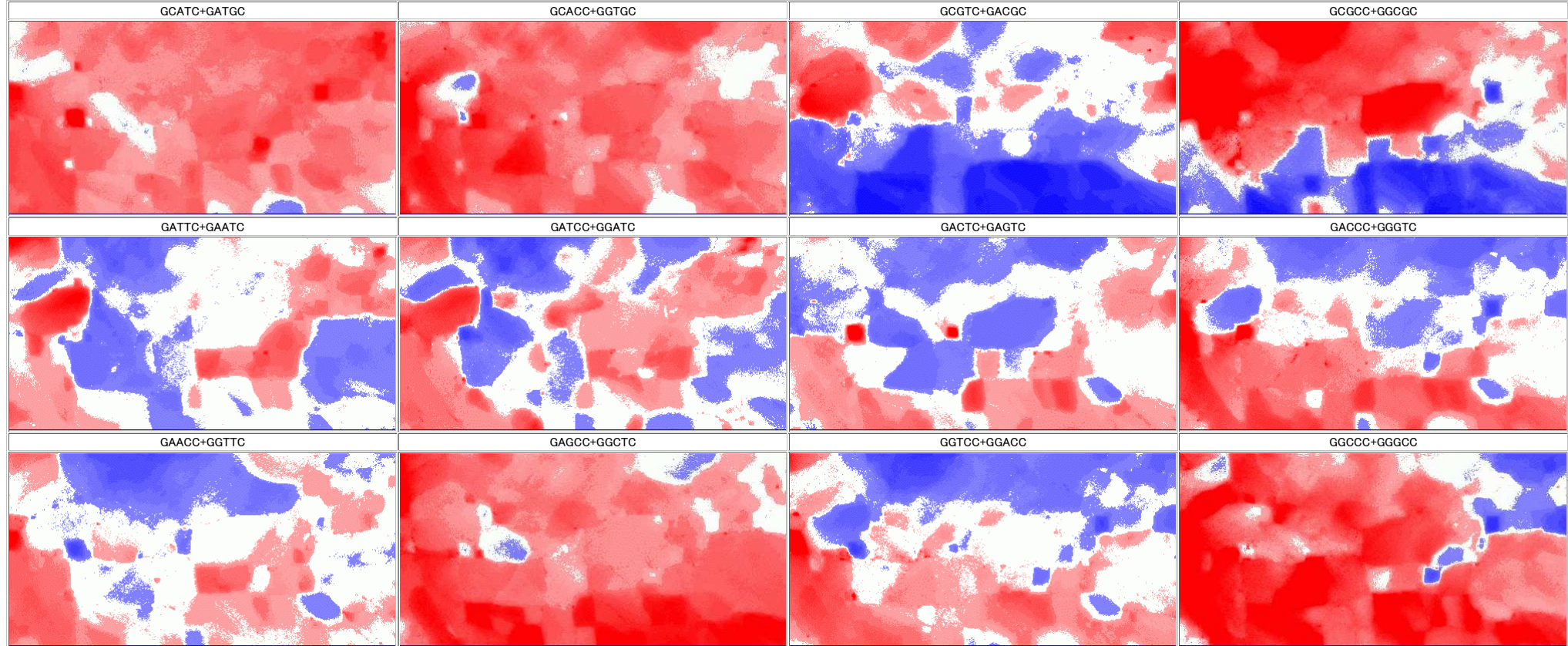












Supplementary Figure S3.

Level of each pair of complimentary pentanucleotide in 100-kb DegePenta-BLSOM for the overlapping 100-kb sequences of 12 *Drosophila* genomes.

