

Appendix for

Convergence of aging- and rejuvenation-related epigenetic alterations on PRC2 targets

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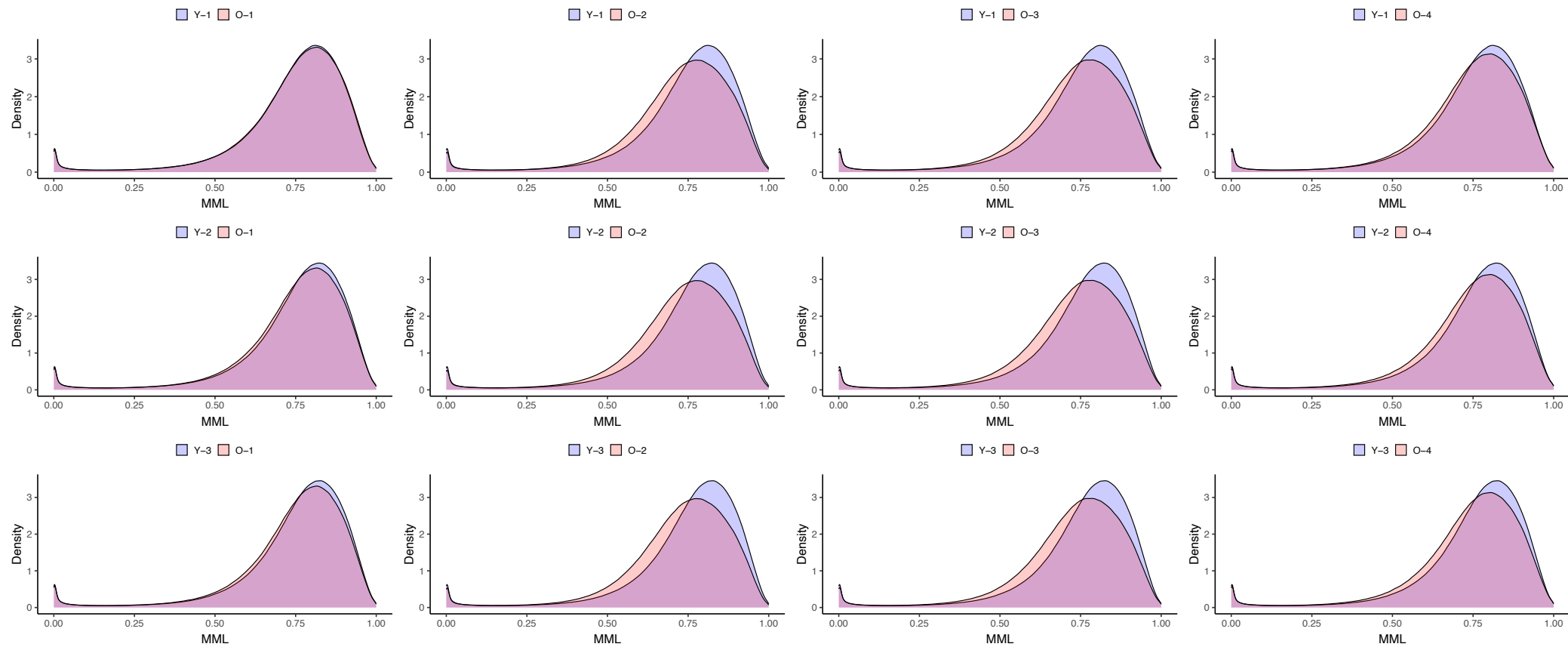
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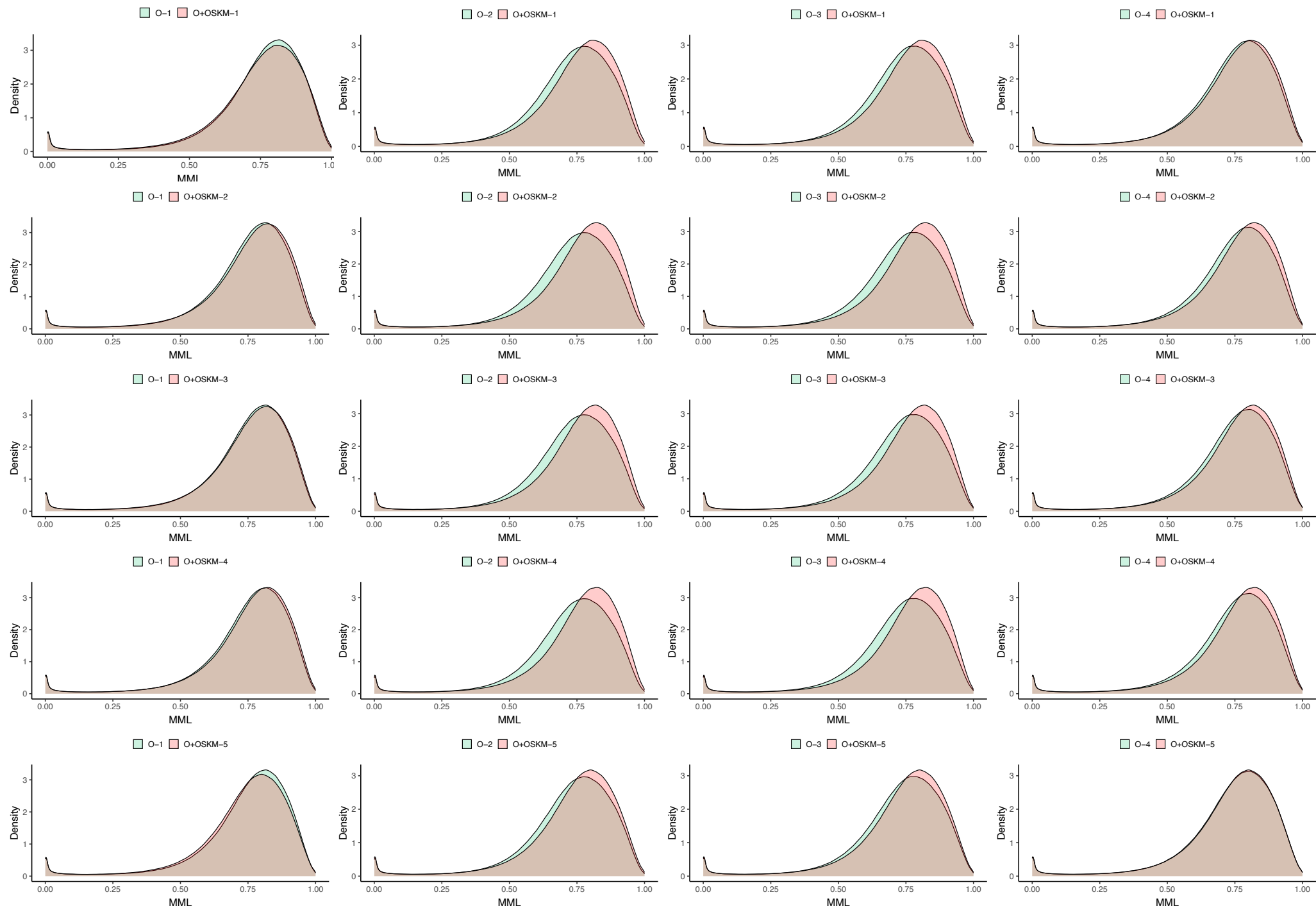
† These authors contributed equally.

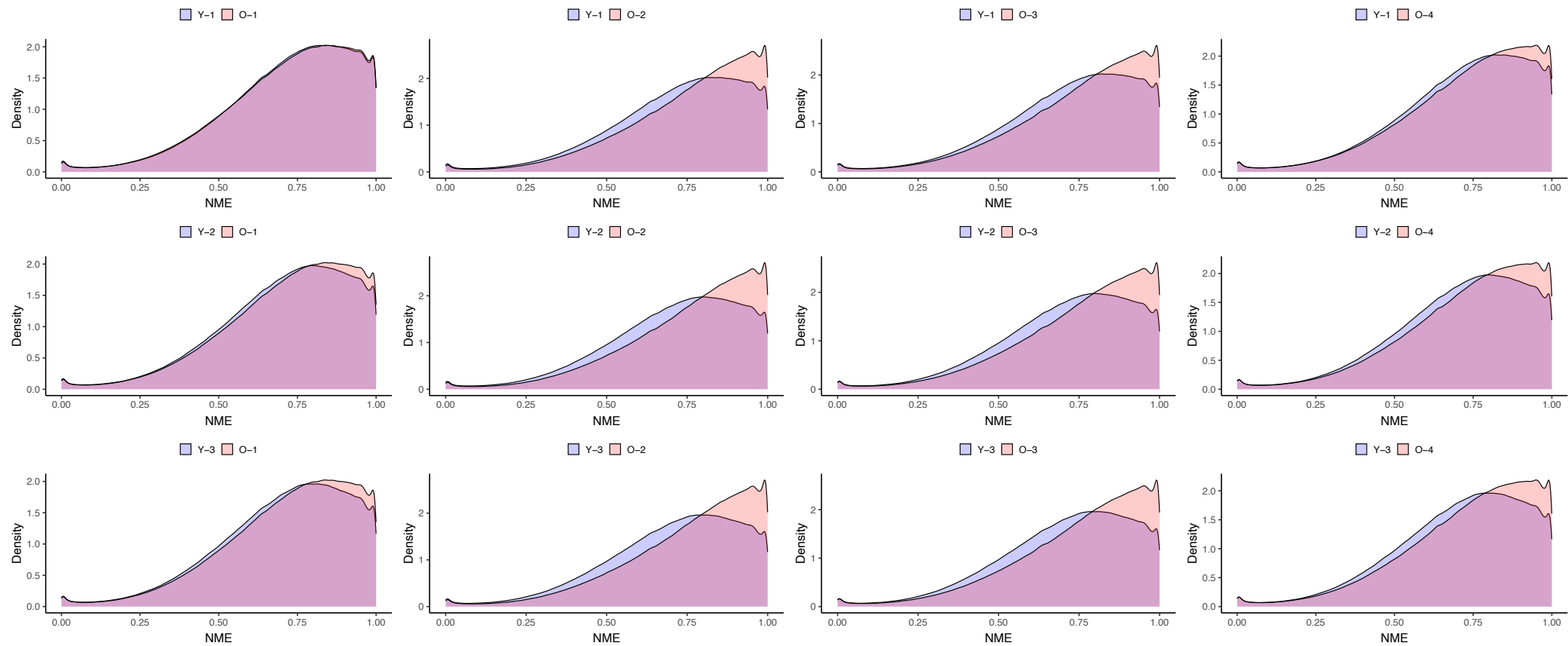
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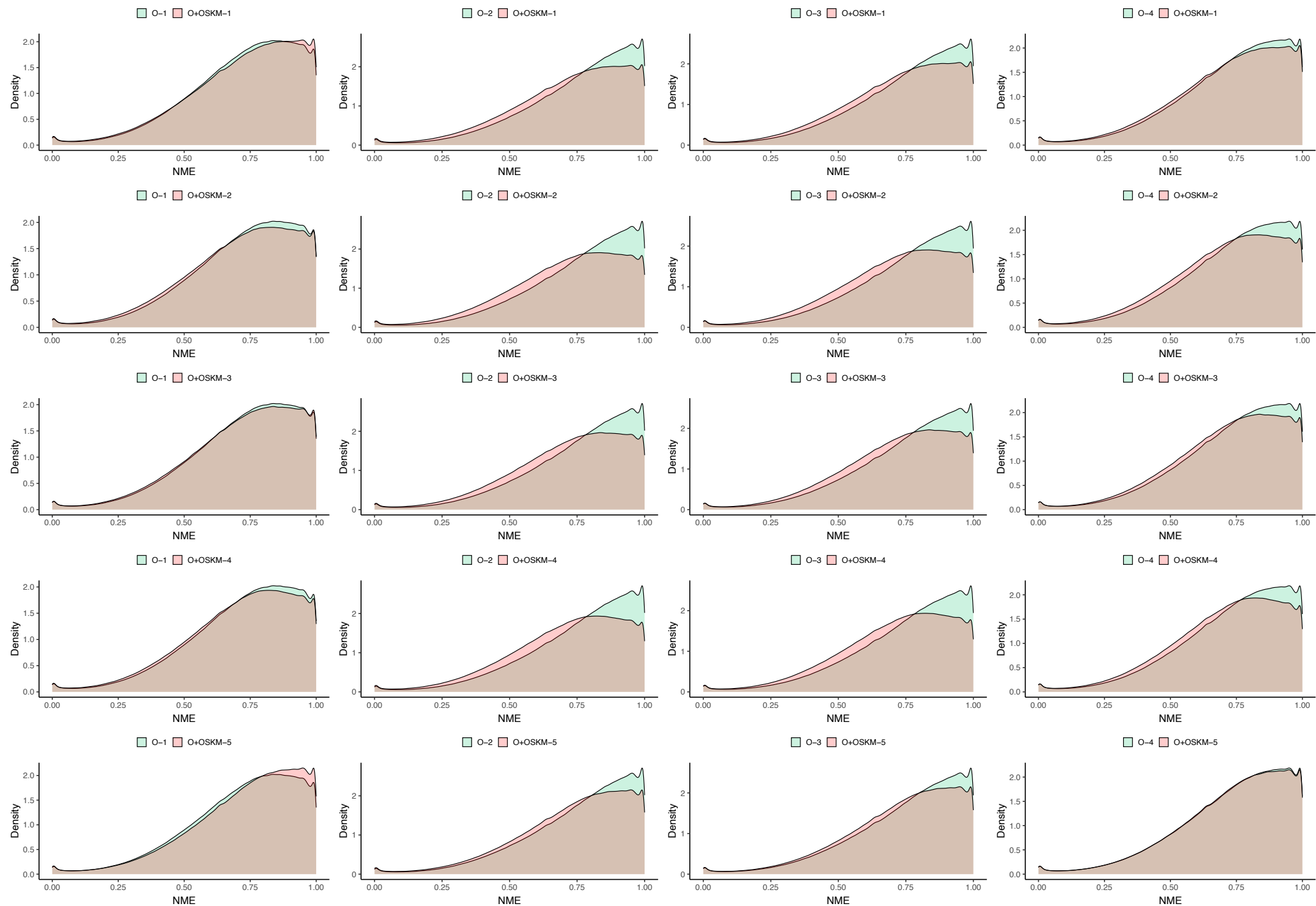
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Appendix Figure S1. Density plots of genome-wide distributions of mean methylation level (MML) and normalized methylation entropy (NME) from pairs of young (Y), old untreated (O), and old treated (O+OSKM) dorsal skin WGBS samples.