Epigenetic domains:

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| long-range epigenetic activation regions | LREA |  | [[1](#_ENREF_1)] |
| long-range epigenetic silencing regions | LRES |  | [[2](#_ENREF_2)] |
| Methylation haplotype regions | MHB |  |  |
|  |  |  |  |

1. Bert, S.A., et al., *Regional activation of the cancer genome by long-range epigenetic remodeling.* Cancer Cell, 2013. **23**(1): p. 9-22.

2. Coolen, M.W., et al., *Consolidation of the cancer genome into domains of repressive chromatin by long-range epigenetic silencing (LRES) reduces transcriptional plasticity.* Nat Cell Biol, 2010. **12**(3): p. 235-46.