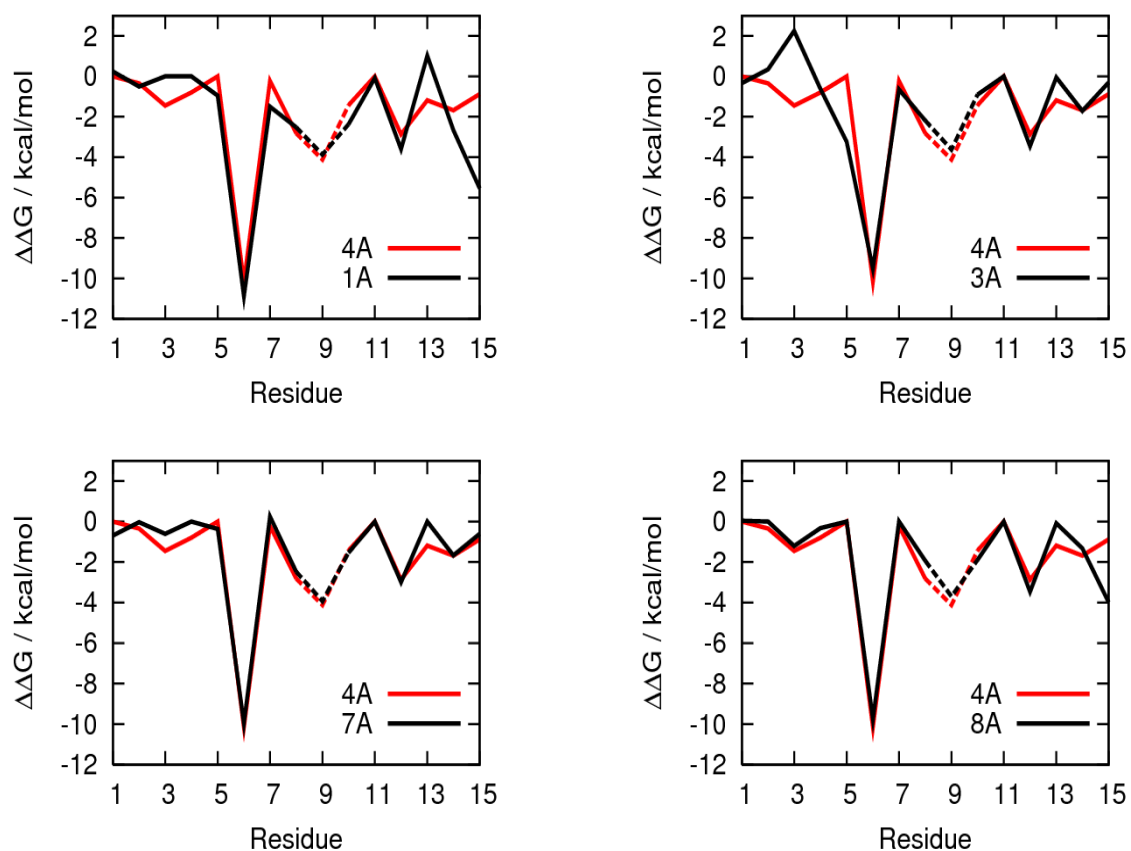




*Figure S3. Computational alanine scans of the BRC4A-like repeats. All show very similar profiles close to the FxTA binding hotspot (residues 06-09) and binding affinity is instead determined by the overall charge of each repeat.*