

Classify each putative Acr-Aca locus and infer targeted CRISPR-Cas subtypes:

- 1. If this Acr-Aca locus has a nearby self-targeting spacer target, it is labeled as "**high confidence**" and inferred to target the CRISPR-Cas locus with the self-targeting spacer
- 2. If this Acr-Aca locus has a self-targeting spacer target in the genome but not nearby, it is labeled as "**medium confidence**" and inferred to target the CRISPR-Cas locus with the self-targeting spacer
- 3. If this Acr-Aca locus has a CRISPR-Cas locus but no self-targeting spacers in the genome, it is labeled as "**low confidence**" and inferred to target the CRISPR-Cas locus