Documented targets (N=7)
(Okumura et al., 2008; Myers et al., 2006)
Genome wide scanning
Motifs from orthologues of known targets in other strains (N=30)
Motif’s PWM
Build new PWM
Homology-driven clustering of protein sequences encoded by target genes
Conserved VirR regulon (High confidence)
Strain-specific VirR regulons (Low confidence)