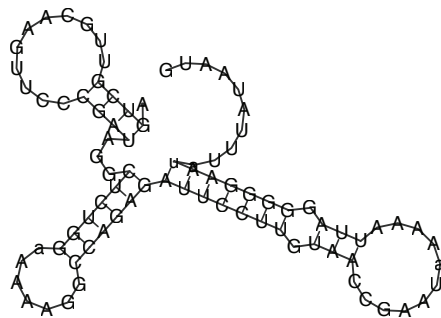
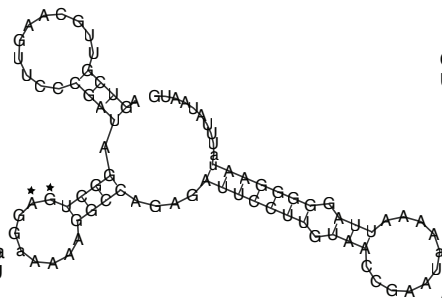


wt



mu1



mu2

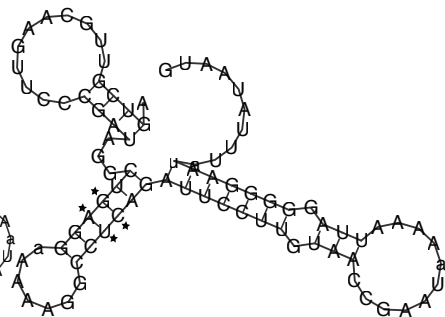


Figure S7. SAM III riboswitch. Secondary structure prediction with RNAfold for three different constructions. wt means the wild type sequence, mu1 means the disrupted mutant and mu2 means the rescue mutation sequence. Stars mark the mutation position.