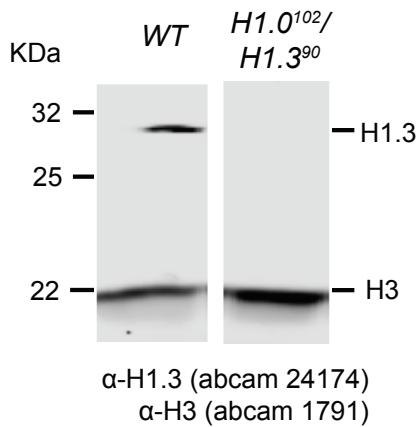


Sanger sequencing

WT 5' GTCTCTCTATATAAACCC...**ATG...TAG**...GTTGTGCGGCCACGGTCTAT-3'
H1.0¹⁰²/H1.3⁹⁰ 5'-GTCTC ----- GGTCTAT-3'
893 bp deletion

Western Blot analysis



Relative mRNA expression

