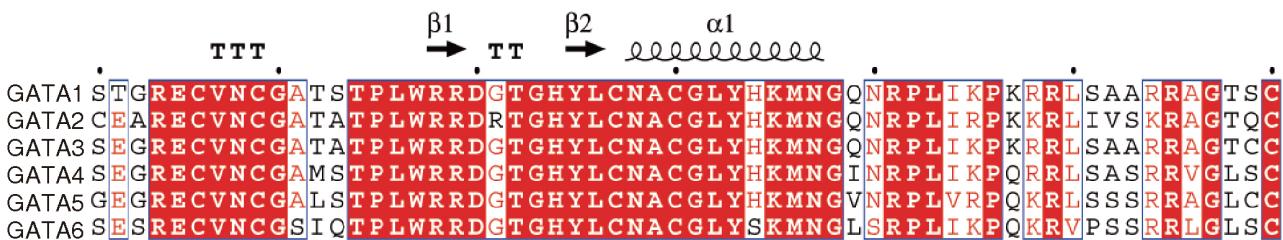


A



B

GATA1-WT	PLPPCEARECVNCGATATPLWRRDRTGHYLCNACGLYHKMNGQNRPLIRPKKRLIVSKRAGTQCTNCQTTTTLWRRNASGDPVCNACGLY
GATA1-N-Mut	PLPPCEARECVNCGATATPLWARDRTGHYLCNACGLYHKMNGQNRPLIRPKKRLIVSKRAGTQCTNCQTTTTLWRRNASGDPVCNACGLY
GATA1-C-Mut	PLPPCEARECVNCGATATPLWRRDRTGHYLCNACGLYHKMNGQNRPLIRPKKRLIVSKRAGTQCTNCQTTTTLWRRNASGDPVCNACGLY
GATA1-D-Mut	PLPPCEARECVNCGATATPLWARDRTGHYLCNACGLYHKMNGQNRPLIRPKKRLIVSKRAGTQCTNCQTTTTLWRRNASGDPVCNACGLY
GATA2-WT	SEGRECVCNCGATATPLWRRDGTGHYLCNACGLYHKMNGQNRPLIKPKRRLSAARRAGTCANCQTTTTLWRRNANGDPVCNACGLYYKLH
GATA2-N-Mut	SEGRECVCNCGATATPLWARDGTGHYLCNACGLYHKMNGQNRPLIKPKRRLSAARRAGTCANCQTTTTLWRRNANGDPVCNACGLYYKLH
GATA2-C-Mut	SEGRECVCNCGATATPLWRRDGTGHYLCNACGLYHKMNGQNRPLIKPKRRLSAARRAGTCANCQTTTTLWARNANGDPVCNACGLYYKLH
GATA2-D-Mut	SEGRECVCNCGATATPLWARDGTGHYLCNACGLYHKMNGQNRPLIKPKRRLSAARRAGTCANCQTTTTLWARNANGDPVCNACGLYYKLH
GATA3-WT	GRECVNCGATSTPLWRRDGTGHYLCNACGLYHKMNGQNRPLIKPKRRLSAARRAGTCANCQTTTTLWRRNANGDPVCNACGLYYKLH
GATA3-N-Mut	GRECVNCGATSTPLWARDGTGHYLCNACGLYHKMNGQNRPLIKPKRRLSAARRAGTCANCQTTTTLWRRNANGDPVCNACGLYYKLH
GATA3-C-Mut	GRECVNCGATSTPLWRRDGTGHYLCNACGLYHKMNGQNRPLIKPKRRLSAARRAGTCANCQTTTTLWARNANGDPVCNACGLYYKLH
GATA3-D-Mut	GRECVNCGATSTPLWARDGTGHYLCNACGLYHKMNGQNRPLIKPKRRLSAARRAGTCANCQTTTTLWARNANGDPVCNACGLYYKLH
GATA4-WT	FSEGRECVCNCGAMSTPLWRRDGTGHYLCNACGLYHKMNGINRPLIKPKRRLSASRRVGLSCANCQTTTTLWRRNAEGERPVCNACGLYMK
GATA4-N-Mut	FSEGRECVCNCGAMSTPLWARDGTGHYLCNACGLYHKMNGINRPLIKPKRRLSASRRVGLSCANCQTTTTLWRRNAEGERPVCNACGLYMK
GATA4-C-Mut	FSEGRECVCNCGAMSTPLWRRDGTGHYLCNACGLYHKMNGINRPLIKPKRRLSASRRVGLSCANCQTTTTLWARNAEGERPVCNACGLYMK
GATA4-D-Mut	FSEGRECVCNCGAMSTPLWARDGTGHYLCNACGLYHKMNGINRPLIKPKRRLSASRRVGLSCANCQTTTTLWARNAEGERPVCNACGLYMK
GATA5-WT	EGRECVNCGALSTPLWRRDGTGHYLCNACGLYHKMNGVRPLVRPQKRRLSSSRAGLCCTNCHTTNTLWRRNSEGERPVCNACGLYMK
GATA5-N-Mut	EGRECVNCGALSTPLWARDGTGHYLCNACGLYHKMNGVRPLVRPQKRRLSSSRAGLCCTNCHTTNTLWRRNSEGERPVCNACGLYMK
GATA5-C-Mut	EGRECVNCGALSTPLWRRDGTGHYLCNACGLYHKMNGVRPLVRPQKRRLSSSRAGLCCTNCHTTNTLWARNSEGERPVCNACGLYMK
GATA5-D-Mut	EGRECVNCGALSTPLWARDGTGHYLCNACGLYHKMNGVRPLVRPQKRRLSSSRAGLCCTNCHTTNTLWARNSEGERPVCNACGLYMK
GATA6-WT	RECVCNCGSIQTPLWRRDGTGHYLCNACGLYSKMNGLSRPLIKPKRVPSSRRGLSCANCHTTTTLWRRNAAEGERPVCNACGLYMKLHG
GATA6-N-Mut	RECVCNCGSIQTPLWARDGTGHYLCNACGLYSKMNGLSRPLIKPKRVPSSRRGLSCANCHTTTTLWRRNAAEGERPVCNACGLYMKLHG
GATA6-C-Mut	RECVCNCGSIQTPLWRRDGTGHYLCNACGLYSKMNGLSRPLIKPKRVPSSRRGLSCANCHTTTTLWARNAAEGERPVCNACGLYMKLHG
GATA6-D-Mut	RECVCNCGSIQTPLWARDGTGHYLCNACGLYSKMNGLSRPLIKPKRVPSSRRGLSCANCHTTTTLWARNAAEGERPVCNACGLYMKLHG

**Supplementary Information, Figure S4 Strategies for mutation of GATA family members.** (A) A schematic diagram illustrating two conserved zinc-finger domains of GATA family (red). The secondary structure of GATA3 is depicted in this figure. (B) Mutation strategy of GATA family members. A diagram depicting detailed protein sequences of the conserved zinc-finger domains of GATA family members. Mutated sites are highlighted in green.