

Table S3:

disiRNA-6	
ds6(+f1)	GGT GTT GYA TTC TGG GAA AGY TTG
ds6(+f2)	GGA TTT TGA GGG YAG YAG AYG GAG
ds6(+r2)	CTT AAC CCT TTT CTC TTA RAC CTR CRT C
ds6(+r1)	CAA CCC CAC CTA CCT CTT CAT C
disiRNA-29	
d29(+f1)	GTA YGG AGT ATT GGY TGG YTG TG
d29(+f2)	GGT TGG YYG GTY GTT GTT GTA AG
d29(+r1)	CTC RAT ARC CTRACC CCT TTC TCC
d29(+r2)	CAT CAC CAC CAC ATC CRT CRA C
disiRNA-47	
d47(-)f1	AAC TTT TCR ATC CRC TCR ATC CCC T
d47(-)r1	GGT ATT GYG ATA TGG GYA GAG TTG G
d47(-)f2	CTT RCC CTC CCT CCC CTT TCC AC
d47(-)r2	GTY YAG TTG AGG GTA TAG TGT TGG AAG
control primers: am locus	
am(+f1)	GTA GGT TTA TAA GGG TAY GTY TGA GAG A
am(+f2)	GTG TTG TTT GGG AGG AYG AYA AYG
am(+r2)	CCT TAC CAA TAA RRA CAC CCT CRA AAC
am(+r1)	CTC AAC CTC CTC CTT RRA AAC CTCA
control primers: al-1 gene	
al-1(+f1)	GTG YAG AGA GAA GGA GYG AYG ATA AG
al-1(+f2)	GAA GGT YGA GAT YGA AAA GTG GGA AG
al-1(+r2)	CTT RAC RCC CAT CCT CTC TCC AA
al-1(+r1)	CAC CTC RCC RTT CTC AAR CTR RAC AC
control primers: ζ-η	
ζ-η (+)f1	GAA GAA AGG AAG AAA AYA AAG AGA AAY YGA
ζ-η (+)f2	GTT ATT GTA TAT YGA TAG GGA GAG AYY GG
ζ-η (+)r1	CTA TCT CRC CCA ART CRA ARC CTC
ζ-η (+)r2	CCC TCC RCT AAC CCT TAT TAC TAT TAC