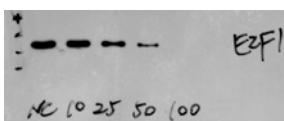
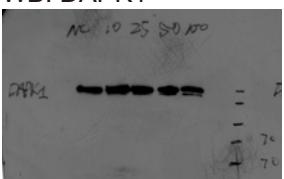


Figure 1b

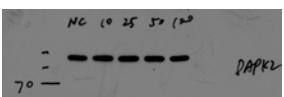
WB: E2F1



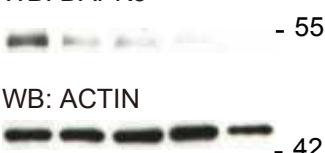
WB: DAPK1



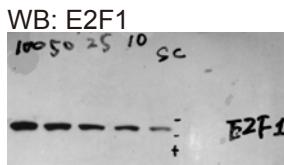
WB: DAPK2



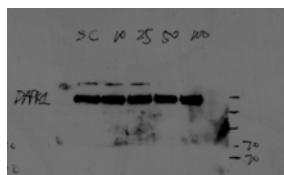
WB: DAPK3



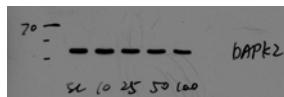
WB: ACTIN

**Figure 1c**

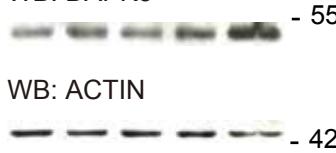
WB: DAPK1



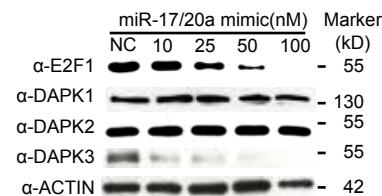
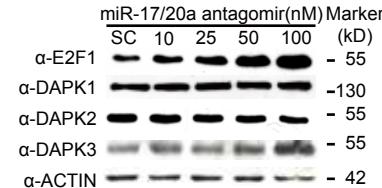
WB: DAPK2



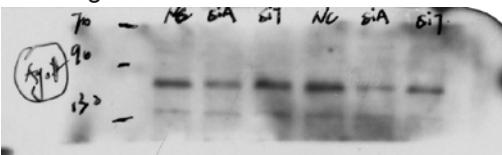
WB: DAPK3



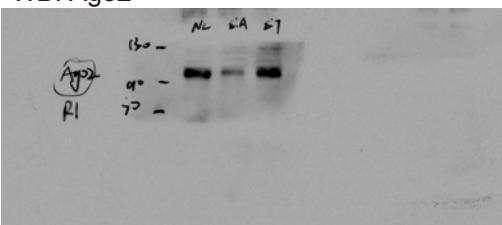
WB: ACTIN

**Figure 1b****Figure 1c****Figure 3a**

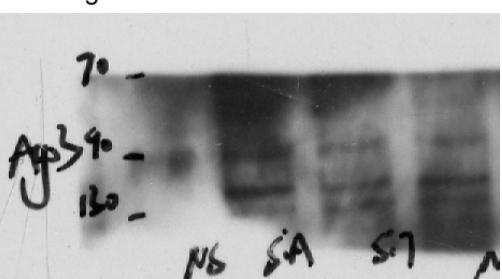
WB: Ago1



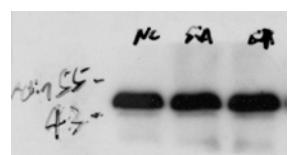
WB: Ago2



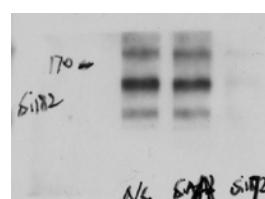
WB: Ago3



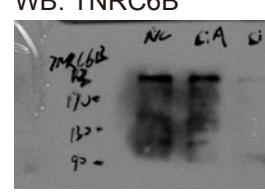
WB: ACTIN



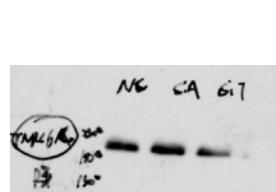
WB: TNRC6A



WB: TNRC6B



WB: TNRC6C



WB: Ago4

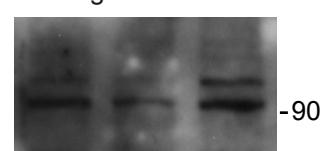
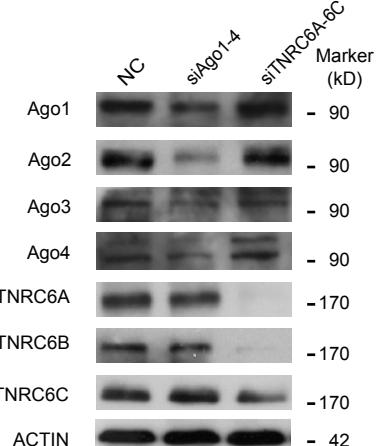
**Figure 3a**

Figure 3d

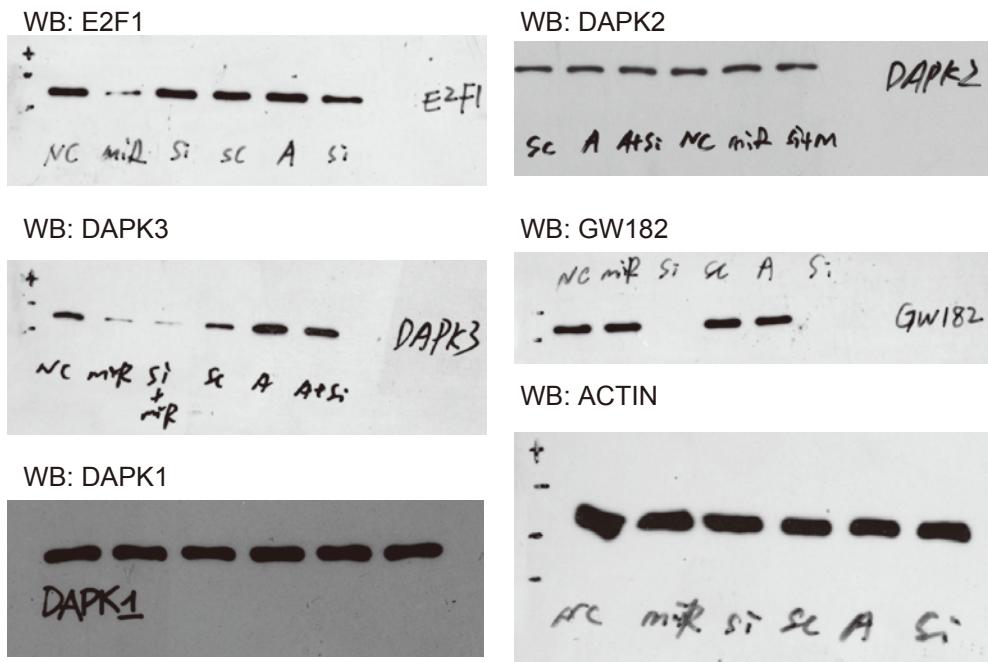


Figure 3d

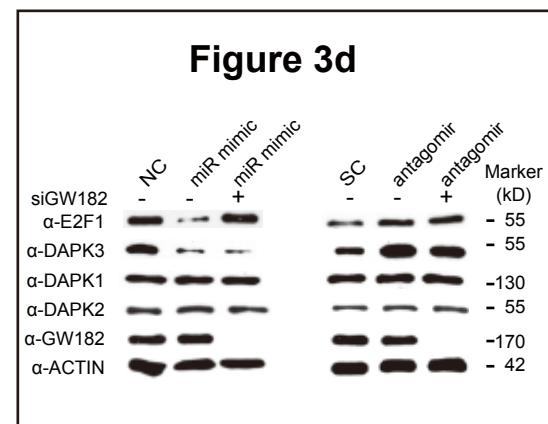
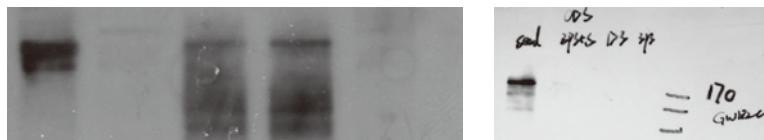


Figure 4b

Renilla capture

WB: GW182

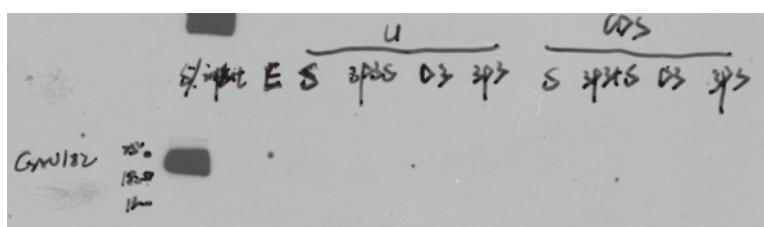


WB: Ago2



Firefly capture

WB: GW182



WB: Ago2

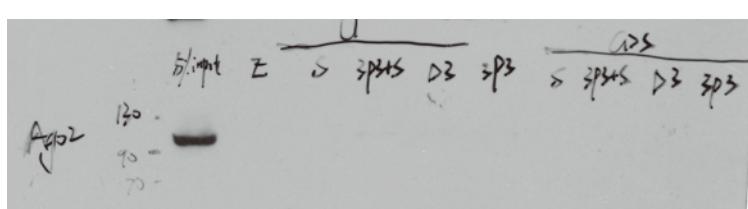


Figure 4b

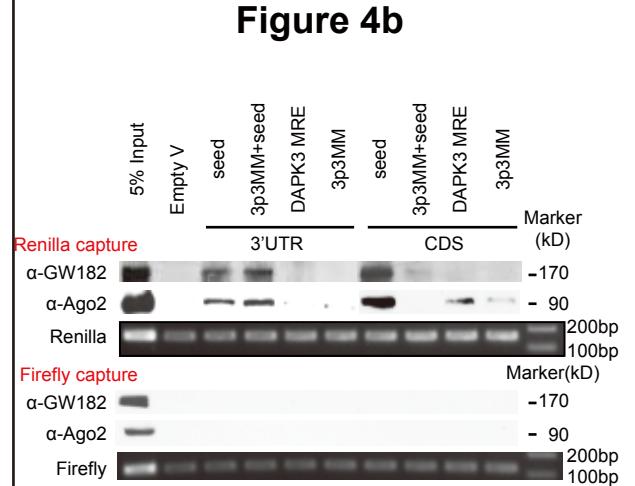


Figure 5b

WB: Flag

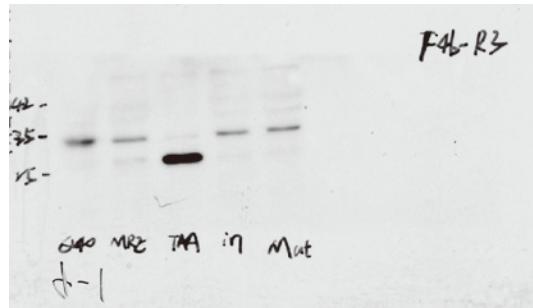


Figure 5b

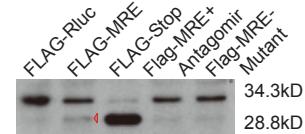
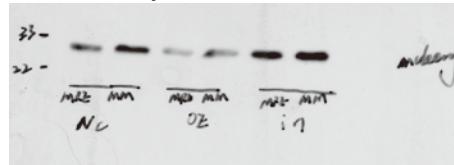


Figure 5c

WB: Mcherry



WB: GFP

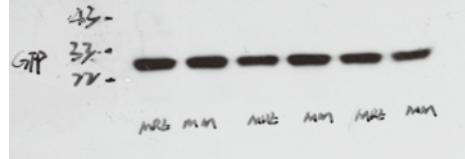
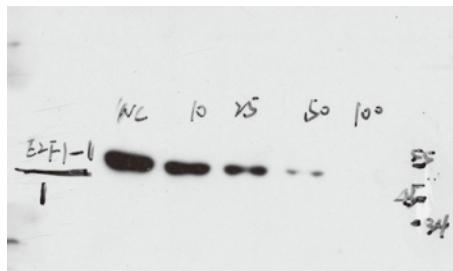


Figure 5c

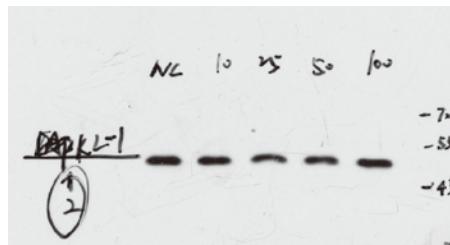


Suppl Fig. 1a

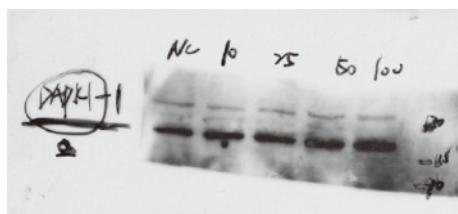
WB: E2F1



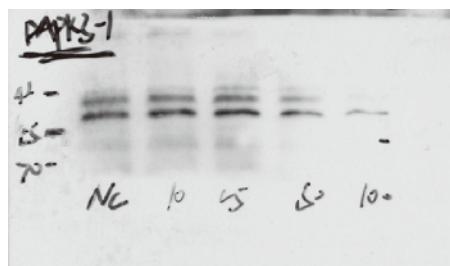
WB: DAPK2



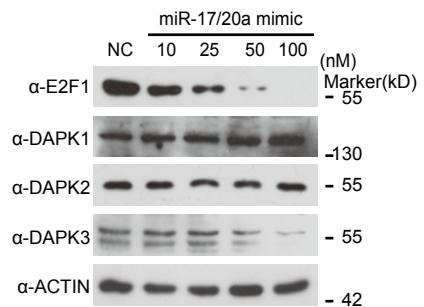
WB: DAPK1



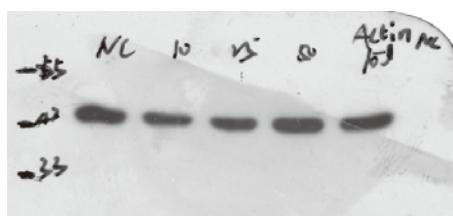
WB: DAPK3



Suppl Fig. 1a

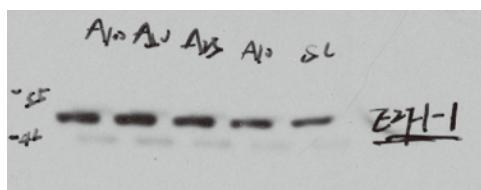


WB: ACTIN

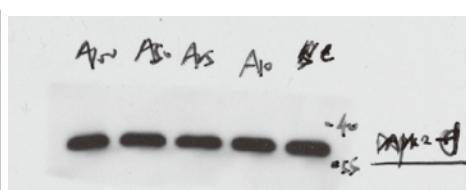


Suppl Fig. 1b

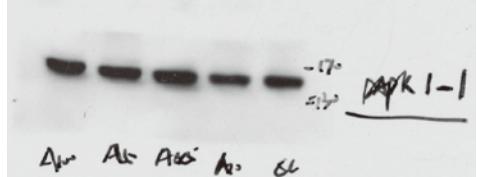
WB: E2F1



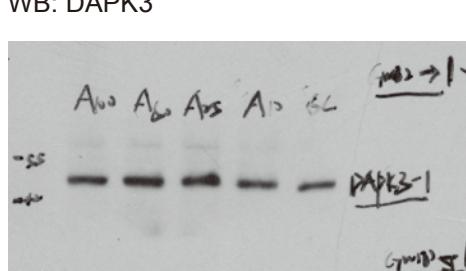
WB: DAPK2



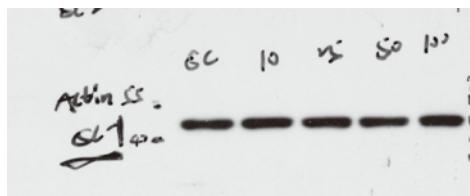
WB: DAPK1



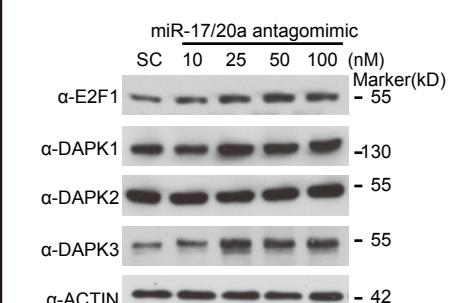
WB: DAPK3



WB: ACTIN

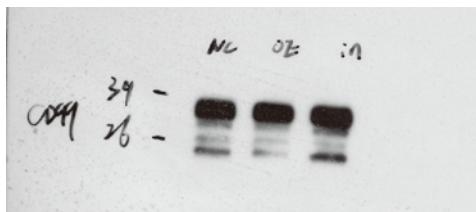


Suppl Fig. 1b

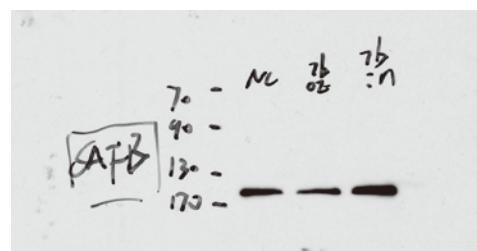


Suppl Fig. 1n

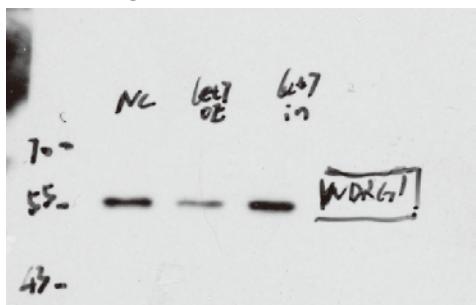
WB: CD99



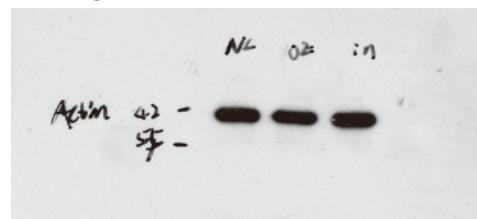
WB: SAFB



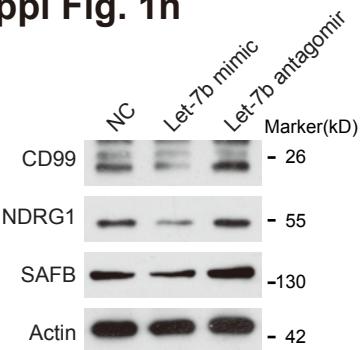
WB: NDRG1



WB: ACTIN

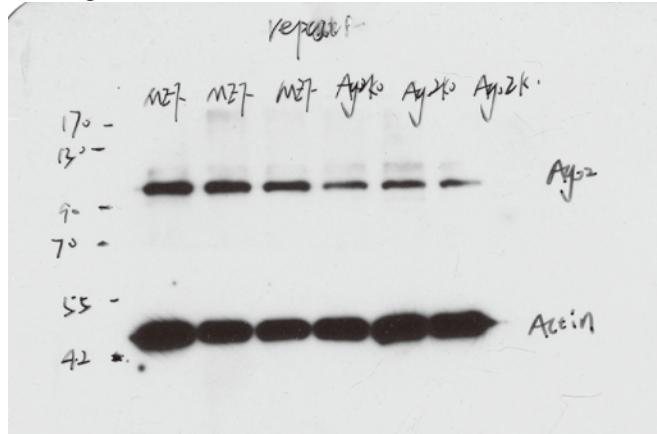


Suppl Fig. 1n



Suppl Fig. 4a

WB: Ago2

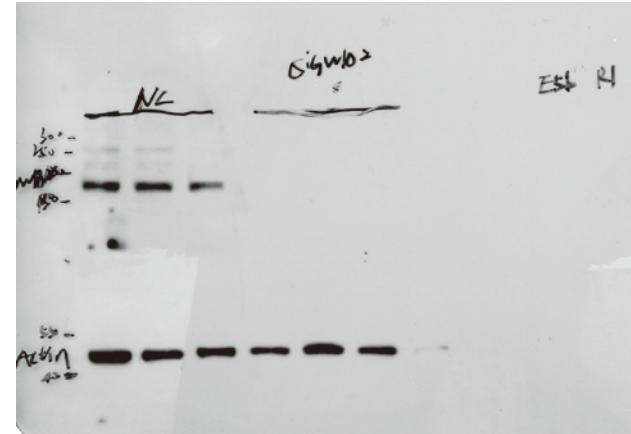


Suppl Fig. 4a

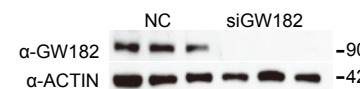


Suppl Fig. 4b

WB: GW182

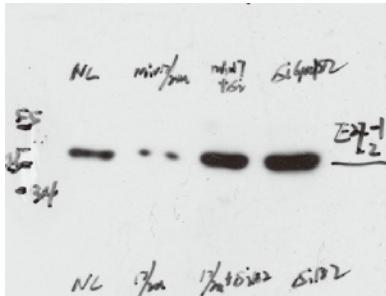


Suppl Fig. 4b

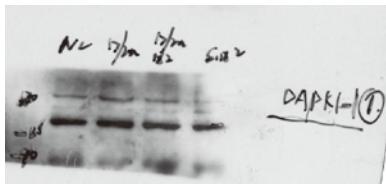


Suppl Fig. 4c left

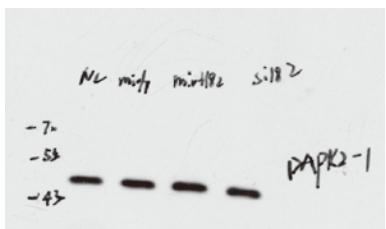
WB: E2F1



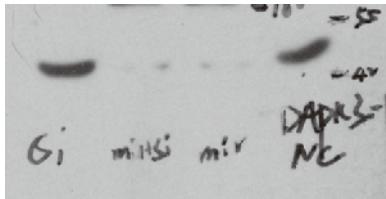
WB: DAPK1



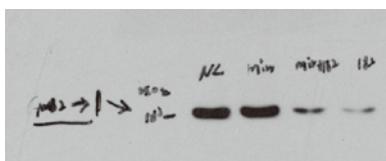
WB: DAPK2



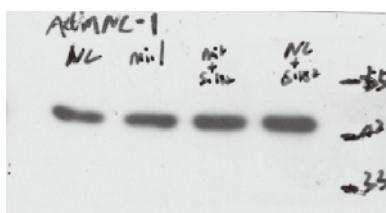
WB: DAPK3



WB: GW182

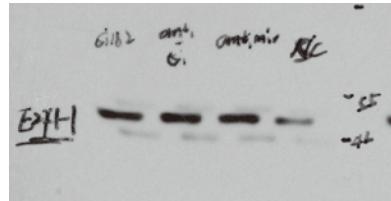


WB: ACTIN

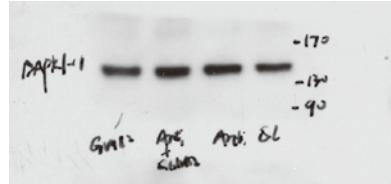


Suppl Fig. 4c right

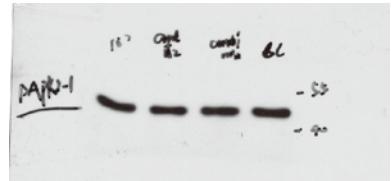
WB: E2F1



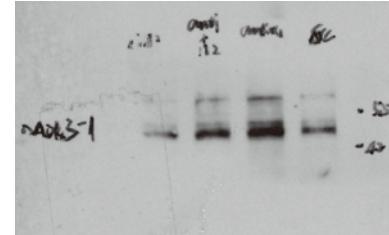
WB: DAPK1



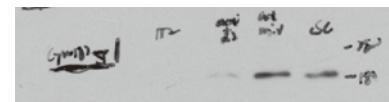
WB: DAPK2



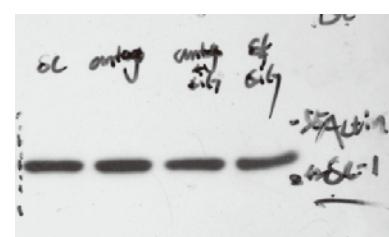
WB: DAPK3



WB: GW182



WB: ACTIN

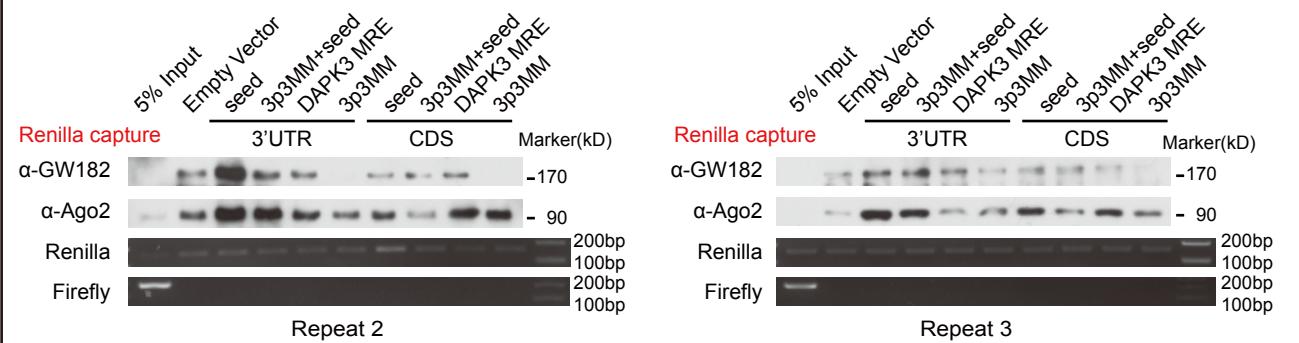


Suppl Fig. 4c

The Western blot shows protein expression levels for α-E2F1, α-DAPK3, α-DAPK1, α-DAPK2, α-GW182, and α-ACTIN across various siRNA and miRNA treatments. The lanes are grouped by treatment: siGW182 (NC, +), miR mimic (NC, +), SC (NC, +), antagonir (NC, +), and Marker (kD). The α-ACTIN row serves as a loading control.

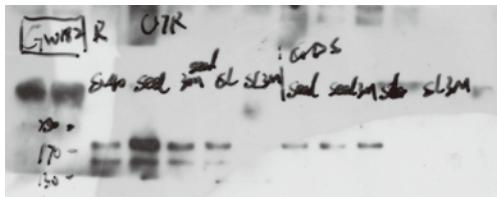
	siGW182	miR mimic	SC	antagonir	Marker (kD)
α-E2F1	-	-	+	+	-
α-DAPK3	-	-	-	-	-
α-DAPK1	-	-	-	-	-
α-DAPK2	-	-	-	-	-
α-GW182	-	-	-	-	-
α-ACTIN	-	-	-	-	-

Suppl Fig. 4d

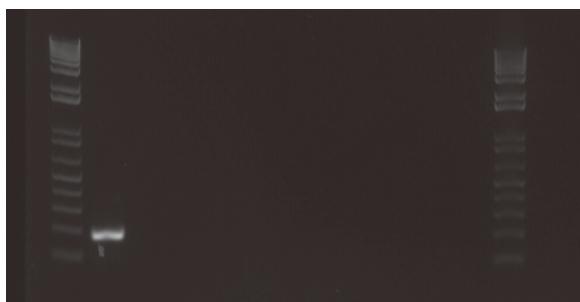
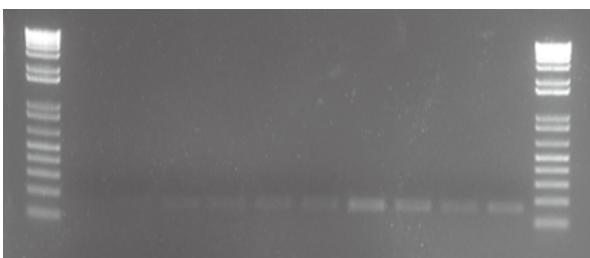
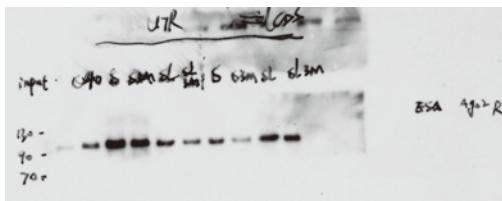


Suppl Fig. 4d left

WB: GW182

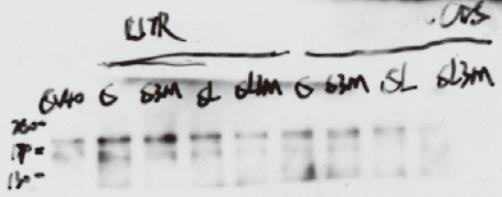


WB: Ago2

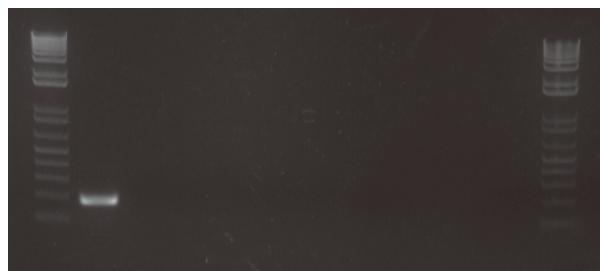
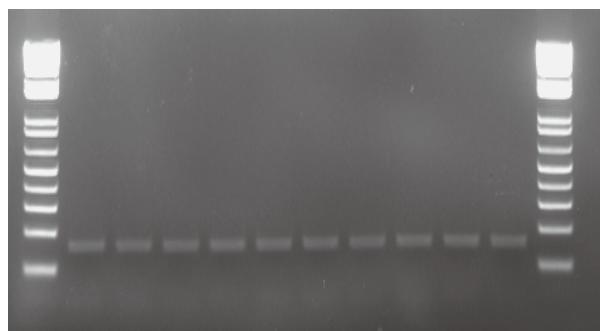
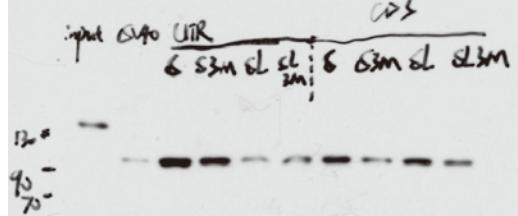


Suppl Fig. 4d right

WB: GW182

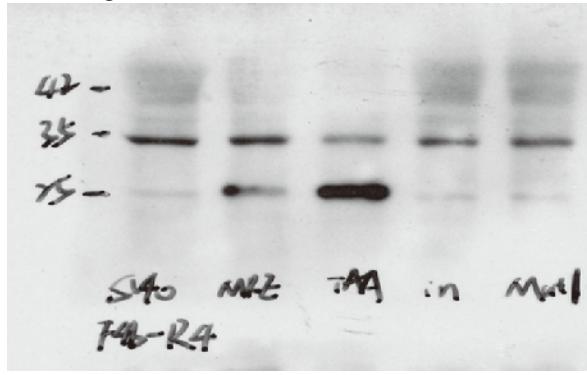


WB: Ago2

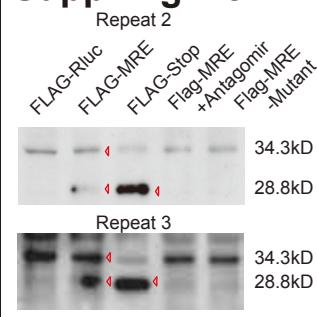


Suppl Fig. 4e up

WB: Flag

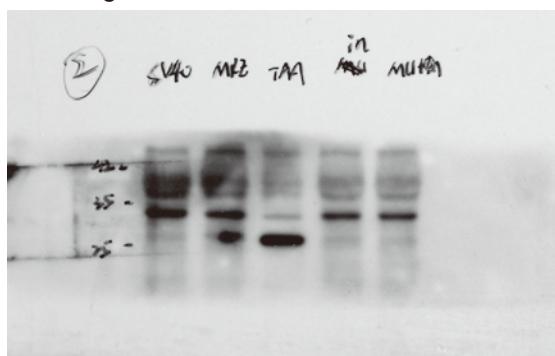


Suppl Fig. 4e

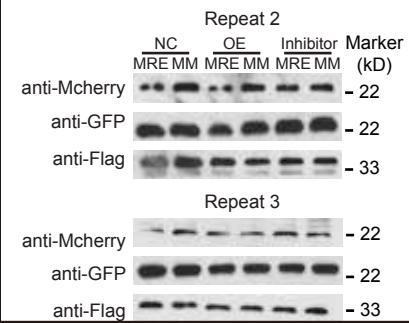


Suppl Fig. 4e down

WB: Flag

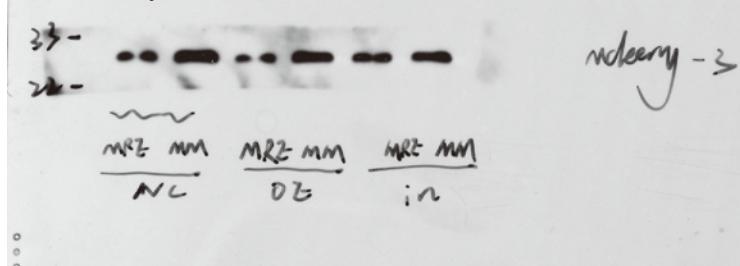


Suppl Fig. 4f



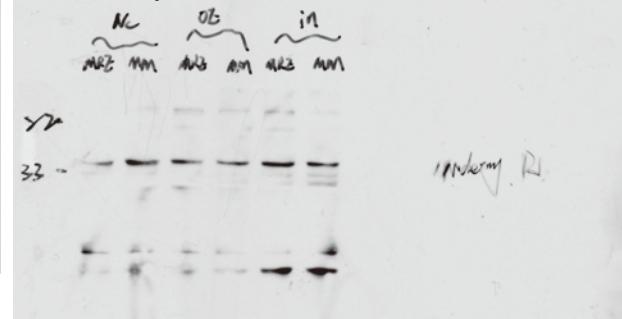
Suppl Fig. 4f up

WB: Mcherry

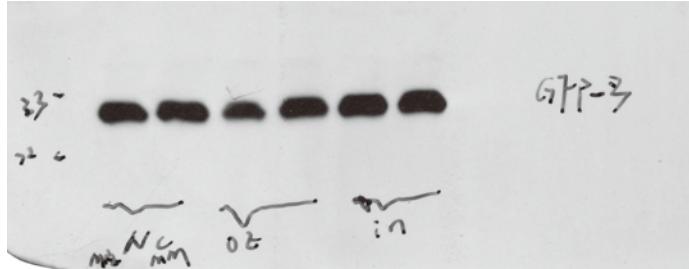


Suppl Fig. 4f down

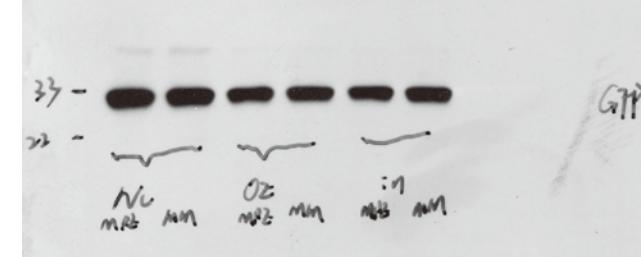
WB: Mcherry



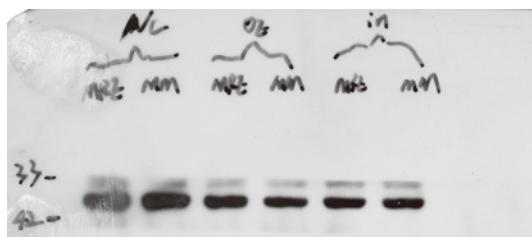
WB: GFP



WB: GFP



WB: Flag



WB: Flag

