Supplemental Figure 1   **shang et al.**

ShRNA Control

Go/G1: 64.47%
S : 23.91%
G2/M : 11.52%

Sh Aurora A

Go/G1: 62.83%
S : 23.21%
G2/M : 13.95%

SK-N-AS
(S-type)

ShRNA Control

Go/G1: 70.86%
S : 24.31%
G2/M : 4.84%

Sh Aurora A

Go/G1 : 53.23%
S : 35.84%
G2/M : 10.95%

SH-SY5Y-Luc
(N-type)

Counts

Relative nuclear DNA content

FL2-A

Supplemental Table 1.   Shang et al.

Table 1. Cox multivariable analysis for progression-free survival

Variables*	Number of Patient (%)	P value	Hazard Ratio	95% CI
Aurora A mRNA				
≤ 3.25	40(64.5%)		1	
>3.25	22(35.5%)	0.006	4.8	1.6 – 14.4
COG Risk Group				
Low-Intermediate	42(65.6%)		1	
High	22(34.4%)	0.03	3.1	1.1 – 8.8

*Final model included only statistically significant variables.