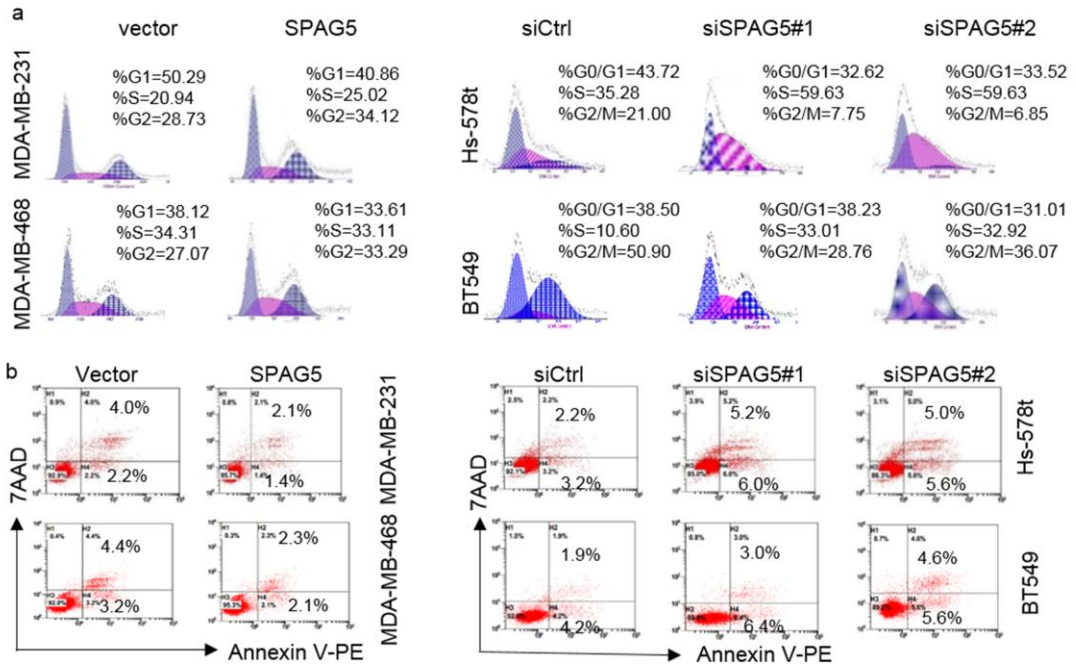
a
vector
SPAG5
siCtrl
siSPAG5#1
siSPAG5#2
MDA-MB-231
%G1=50.29
%S=20.94
%G2=28.73
%G1=40.86
%S=25.02
%G2=34.12
Hs-578t
%G0/G1=43.72
%S=35.28
%G2/M=21.00
%G0/G1=32.62
%S=59.63
%G2/M=7.75
%G0/G1=33.52
%S=59.63
%G2/M=6.85
MDA-MB-468
%G1=38.12
%S=34.31
%G2=27.07
%G1=33.61
%S=33.11
%G2=33.29
BT549
%G0/G1=38.50
%S=10.60
%G2/M=50.90
%G0/G1=38.23
%S=33.01
%G2/M=28.76
%G0/G1=31.01
%S=32.92
%G2/M=36.07
b
Vector
SPAG5
siCtrl
siSPAG5#1
siSPAG5#2
7AAD
4.0%
2.2%
2.1%
1.4%
MDA-MB-231
2.2%
3.2%
5.2%
6.0%
5.0%
5.6%
Hs-578t
4.4%
3.2%
2.3%
2.1%
MDA-MB-468
1.9%
4.2%
3.0%
6.4%
4.6%
5.6%
BT549
Annexin V-PE