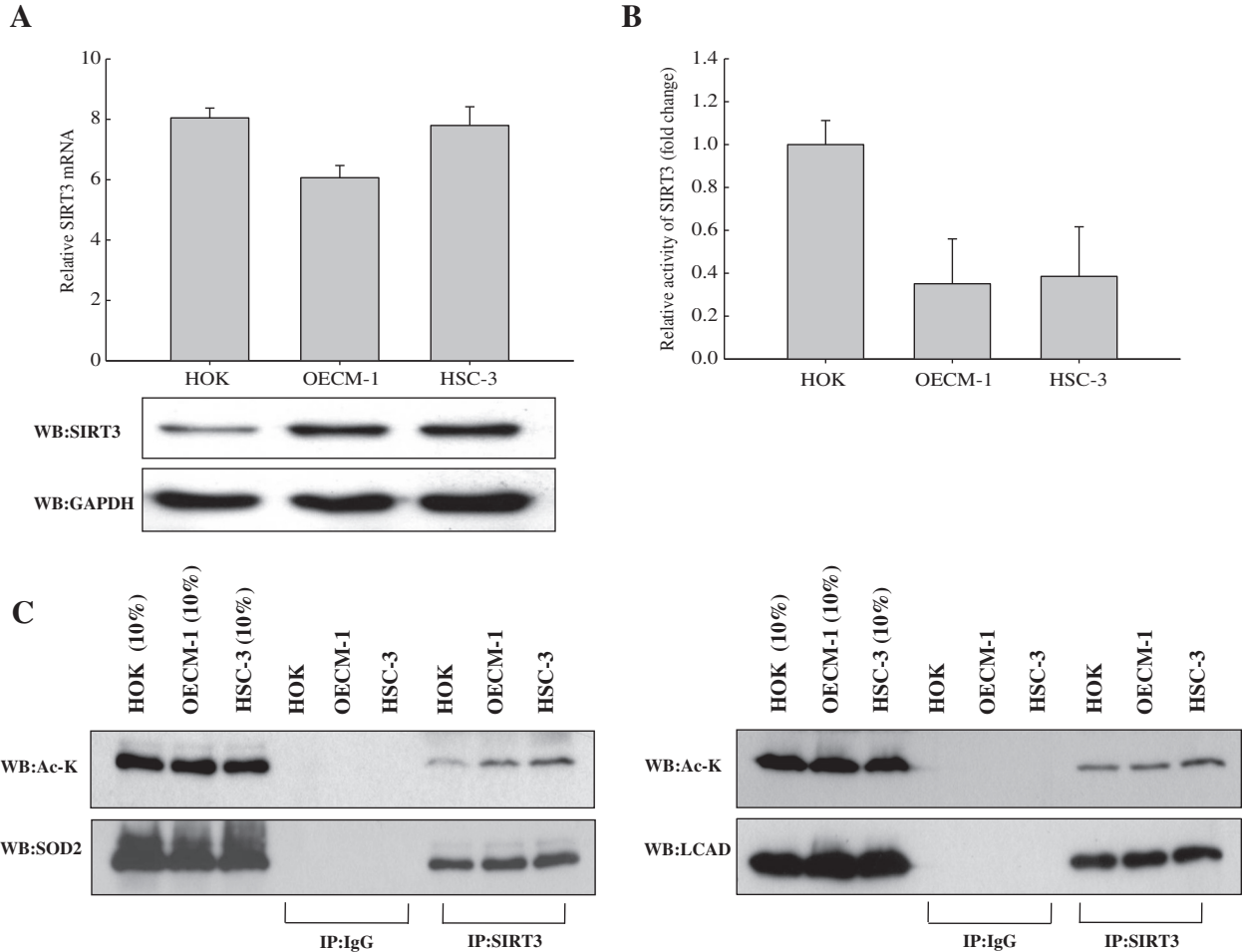
A
Relative SIRT3 mRNA
10
8
6
4
2
0
HOK
OECM-1
HSC-3
WB:SIRT3
WB:GAPDH
B
Relative activity of SIRT3 (fold change)
1.4
1.2
1.0
0.8
0.6
0.4
0.2
0.0
HOK
OECM-1
HSC-3
C
HOK (10%)
OECM-1 (10%)
HSC-3 (10%)
HOK
OECM-1
HSC-3
HOK
OECM-1
HSC-3
WB:Ac-K
WB:SOD2
IP:IgG
IP:SIRT3
HOK (10%)
OECM-1 (10%)
HSC-3 (10%)
HOK
OECM-1
HSC-3
HOK
OECM-1
HSC-3
WB:Ac-K
WB:LCAD
IP:IgG
IP:SIRT3