



SOM-L	SOM-E	SOM-E	SOM-L	SOM-E
ITSN2 RAB1B TSPYL2 UBB USP9X XPO1 YAP1	ACVR1B	BAZ2B CTNNB1 E124 EPPK1 <sup>a)</sup> GORASP2 NACA PCNX4 SDR16C5 TAPT1 TIMM22 VDAC1 XDH	ABCA13 ASXL2 HK1 ITGA1 VCAN	ABCB6 (P1-ADC1, P1-ADC2) ALMS1 BRAF EGFR (P1-ADC, P5-ADC) <sup>b)</sup> GRIPAP1 KIAA0513 KIF3A NOC3L PHF1 PIGR PSEN2 PTPN13 RBBP8 RP1 SLC25A39 SMAP1

**S10 Fig.** Protein interaction graphs in AIS (A), MIA (B), and ADC (C) in P1. ADC, adenocarcinoma; AIS, adenocarcinoma in situ; MIA, minimally invasive adenocarcinoma; SOM-E, somatic variant overexpressed in the transcriptome; SOM-L, somatic variant lost in the transcriptome. <sup>a)</sup>*CTNNB1* (P1 MIA1, P1 MIA2), *ABCB6* (P1 ADC1, P1 ADC2), <sup>b)</sup>*EGFR* (P1 ADC1, and P5 ADC).