

Additional file 1: Table S1. List of PC4 protein identifications by mass spectrometry.

PMSS*	Seq. cov. (%)	Sample analyzed
65.8	26	1xMS2-purified
48.6	26	3xMS2-purified
13.0	19	3xMS2-purified
16.4	19	negative control sample
72.2	32	gradient, then biotin-purified
39.4	26	gradient, then biotin-purified
22.7	19	gradient, then biotin-purified
37.4	31	gradient, then biotin-purified

*PMSS score is a measure for semiquantitative protein abundance; 1xMS2(3xMS2)-purified sample: fraction eluted after affinity purification of 1xMS2(3x-MS2)-labeled U7 snRNA; biotin-purified: fraction eluted after pull down using antisense biotinylated oligonucleotide; gradient: protein extract was fractionated on 10-50% glycerol gradient and selected fractions were affinity-purified as indicated.