

Supplementary information, Table S1. Mass-spectrum ID results of the Exo10 sample.

Only proteins with score larger than 30 are listed. Relative abundance was calculated by averaging PSMs (peptide-spectrum matches) by molecular weight. Rrp44 is used as normalization reference.

Description	Score	# Unique Peptides	# PSMs	MW [kDa]	Relative abundance
Exosome complex exonuclease RRP44	1373.73	74	484	113.6	1.00
Exosome complex component RRP4	825.27	30	273	39.4	1.63
Exosome complex component SKI6	594.30	18	201	27.5	1.71
Exosome complex component RRP45	454.76	24	149	33.9	1.03
Exosome complex component RRP43	436.50	27	184	44.0	0.98
Exosome complex component MTR3	430.26	14	137	27.6	1.17
Exosome complex component RRP4	402.21	19	134	26.5	1.19
Exosome complex component CSL4	383.13	20	146	31.6	1.09
Exosome complex component RRP46	179.68	11	74	24.4	0.71
Exosome complex component RRP42	146.27	11	70	29.0	0.57
Superkiller protein 7	29.17	9	11	84.7	0.03