

S3 Table. Sample information of RNA used in the RNA-Seq cohort.**Sample Info**

Well	RIN ^c	Vol. [ul]	Conc. [ng/ul]	Sample Description
A0				
A1	8.8	40	293	CTRL1
B1	8.5	40	265	CTRL2
C1	8.4	40	290	CTRL3
D1	8.6	40	277	CTRL4
E1	8.7	40	269	CTRL5
F1	8.3	40	267	CTRL6
G1	8.5	40	269	CTRL7
H1	8.6	40	262	CTRL8
A2	8.9	40	281	CTRL9
B2	8.4	40	279	CTRL10
C2	8.5	40	256	AAAD1
D2	8.7	40	273	AAAD2
E2	8.8	40	298	AAAD3
F2	8.6	40	257	AAAD4
G2	8.3	40	271	AAAD5
H2	8.5	40	286	AAAD6
A3	8.9	40	287	AAAD7
B3	8.7	40	285	AAAD8
D3	8.4	40	281	AAAD9
E3	8.5	40	257	AAAD10