

sample name	total rna (ng)	RIN	Total read pairs	Total Mapped	Total Mapped, deduplicated	% uniquely mapped	% rRNA after deduplication	DNase treated?	Biological replicate	Final library cDNA yields (ng)
SMARTseq 1	1.508	9.8	31767947	25142260	7935521	79.14348	1.708252	No	1	3.68
SMARTseq 2	1.160	9.5	40416985	25631426	5915352	63.41746	2.315434	Yes	2	3.00
SMARTseq 3	3.600	9.9	37442564	24398942	7394695	65.16365	2.194880	No	3	4.36
SMARTseq 4	1.230	9.8	36977605	23990669	5018616	64.87892	2.223749	Yes	3	3.68
SoLo 1	1.508	9.8	17969902	10483341	3860066	58.33833	11.712460	No	1	59.20
SoLo 2	1.160	9.5	17395388	10075674	3001022	57.92152	15.558520	Yes	2	60.00
SoLo 3	3.600	9.9	21456169	12216428	4393602	56.93667	12.983270	No	3	78.00
SoLo 4	1.230	9.8	15559476	9323351	3010153	59.92073	12.886540	Yes	3	60.60

Table S1: Sample metadata