

			Duration (in seconds)				Memory (in MB)		
		Numer of reads (in millions)	ABRIDGE	BAM	CRAM		ABRIDGE	BAM	CRAM
RNA-Seq	Single ended	24	9	11	2		44.44	3.5	2.82
		50	9	12	1		36.14	6.05	2.84
		76	18	22	1		58.65	9.28	2.62
		98	24	32	3		86.17	14.11	2.77
		128	31	40	3		101.09	15.4	2.79
	Paired ended	24	48	53	6		125.96	18.62	2.85
		50	23	25	2		65.47	4.6	2.64
		76	30	37	4		89.61	5.02	2.95
		98	35	48	4		109.32	7.37	3
		128	53	58	4		126.96	8.1	3.01
DNA-Seq	Single ended	47	44	22	2		48.04	2.91	2.66
		90	46	21	1		42.86	2.93	2.88
		136	59	40	2		64.24	4.85	2.96
		177	118	60	2		86.12	2.89	3.07
		245	125	75	5		105.39	3.23	3.14
	Paired ended	47	150	105	5		132.4	3.14	3.26
		90	89	39	2		77.77	3.09	2.84
		136	110	64	4		114.85	3.02	2.85
		177	139	81	5		144.95	3.01	3
		245	198	109	6		195.78	3.05	2.99