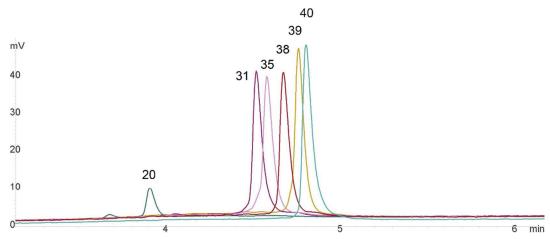
	I RecJ-SSBP experiments, trityl-on						
S1	TTTGTGTCCAAGAATGTTTCCATCTTTAAAAATCAATA						
S2	CCATGATGTATACATTGTGTGAGTTATAGTTGTATTCCAA						
S3	CACCCTCTCCACTGACAGAAAATTTGTGCCCATTAACATC						
S4	ATTTTGAAGTTAACTTTGATTCCATTCTTTTGTTTGTCTG						
S5	GTCTGCTAGTTGAACGCTTCCATCTTCAATGTTGTGTCTA						
S6	GAAGCGTTCAACTAGCAGACCATTATCAACAAAATACTCC						
S7	GGACAGGGCCATCGCCAATTGGAGTATTTTGTTGATAATG						
S8	CTGTCCACACAATCTGCCCTTTCGAAAGATCCCAACGAAA						
S9	CCATGCCATGTGTAATCCCAGCAGCTGTTACAAACTCAAG						
S10	TGGGATTACACATGGCATGGATGAACTATACAAATAGGAG						
	II Failure strand analysis for HPLC, trityl-on / off						
40b	TTCTTGTTGAATTAGATGGTGATGTTAATGGGCACAAATT						
39b	TCTTGTTGAATTAGATGGTGATGTTAATGGGCACAAATT						
38b	CTTGTTGAATTAGATGGTGATGTTAATGGGCACAAATT						
35b	GTTGAATTAGATGGTGATGTTAATGGGCACAAATT						
31b	AATTAGATGGTGATGTTAATGGGCACAAATT						
20b	GATGTTAATGGGCACAAATT						
	III Secondary structure and temperature studies, trityl-on / off						
S11	ACCAGATGGCTGTTCTAGAACCGCTGACGCTGGTTCTCCAGCTGGTTCTGCTTGA						
S12	GTAAGGCTATCCAATTGCACCCATTGGTTTGTGCTGCTTACAACGCTGACTTCGATGGTG						
S13	GTCACTACTTTCTCTTATGGTGTTCAATGCTTTTCAAGATACCCAGATCATATGAAACGG						
S14	TTGGAATACAACTATAACTCACACAATGTATACATCATGGCAGACAAACAA						
	IV PAGE purified samples (IDT), trityl-off						
40b	TTCTTGTTGAATTAGATGGTGATGTTAATGGGCACAAATT						
Biotin	Biotin-TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT						
Amine	Amine-TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT						
Phosphate	Phosphate-TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT						

Table 1 Sequences for oligonucleotides used for this research (5'-3')



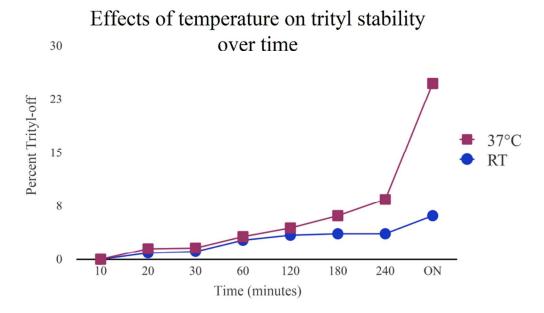
**Fig. 1** Control 5' truncated 'failure' strands processed with ion-pairing RP HPLC. This chromatogram shows oligonucleotide strand resolution between 40b - 39b, 39b - 38b, 38b - 35b, 35b - 31b and 31b - 20b, where gradient and method have been optimized for enhanced peak-peak resolution relative to the full-length 40b strand.

		30m	30m	30m	30m	30m	30m	30m	30m
		30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ
			0.5 ug SSBP		0.5 ug SSBP		0.5 ug SSBP		0.5 ug SSBP
		Total digested (pm)	Total digested (pm)	Failures digested (pm)	Failures digested (pm)	Percent failure decrease	Percent failure decrease	Increase purity	Increase purity
Sa	ample								
	S1	20.4	59.2	4	12	15	43	0.8	2.6
	S2	11	34.3	10	31	15	47	2.1	7.9
	S3	17.4	35.7	13	20	33	52	3.2	5.1
	S4	14	30	6	22	20	81	1.3	5.8
	S5	18.6	26	17	24	29	40	4.6	6.3
	S6	14.1	40	9	13	34	45	2.4	3.2
	S7	7.9	30.9	8	30	8	28	0.8	6
	S8	8.1	19.6	10	20	27	56	2	4.2
	S9	8.3	34	9	29	10	32	1.4	5.7
	S10	17.9	54.5	22	44	31	61	4.8	9.7
Av	erage	14	36	11	25	22	49	2	6
	andard viation	5	12	5	9	10	15	1	2

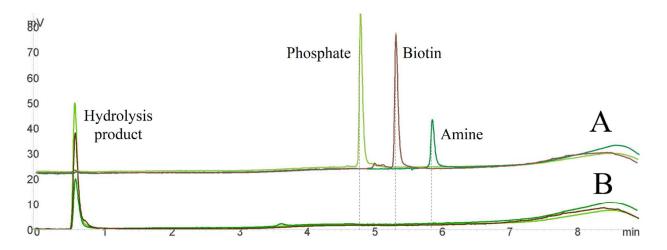
**Table 2** Digestive performance of 30 U RecJ (+ / - 0.5 ug SSBP) on 10 unique 40b samples (crude trityl-on 500 pm) over 30 minutes incubation at 37°C.

	60m	60m	60m	60m	60m	60m	60m	60m
	30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ	30 U RecJ
		0.5 ug SSBP		0.5 ug SSBP		0.5 ug SSBP		0.5 ug SSBP
	Total digested (pm)	Total digested (pm)	Failures digested (pm)	Failures digested (pm)	Percent failure decrease	Percent failure decrease	Increase purity	Increase purity
Sample								
S1	35	68	15	10	55	36	3.8	1.9
S2	27.4	53.9	25	49	38	73	6	13.3
S3	23.1	52.4	20.8	45.6	30.2	55.3	5.3	8
S4	11.4	39.9	18.4	35.3	29	62.1	3.9	8.5
S5	22.1	59.7	15.6	34.5	28.3	52.1	5	7.5
S6	28.2	49.3	14.1	27.3	29.9	47	4.7	7.3
<b>S7</b>	4.2	48.5	6	47	6	44	0.1	10
S8	22.7	60	13	24	37	66	2.7	5
S9	12.7	38.9	16	38	17	41	3.1	7.8
S10	28.2	57.2	25	40	35	56	5.5	8.7
Average	22	53	17	35	31	53	4	8
Standard deviation	9	9	6	12	13	12	2	3

**Table 3** Digestive performance of 30 U RecJ (+ / - 0.5 ug SSBP) on 10 unique 40b samples (crude trityl-on 500 pm) over 60 minutes incubation at 37°C.



**Fig. 2** Effects of time and temperature on the rate of detritylation. Each plot point (time of sample collection) represents an average of three samples (Supplementary Table 1 III) where the percent detritylation was measured (percent peak area from RP HPLC analysis report). (RT, room temperature; ON, overnight; 5 uM ssDNA in water and 1x Buffer 2 at pH 7).



**Fig. 3** RecJ-SSBP digestion of oligonucleotides modified at the 5' end with phosphate, biotin and amine (each 20 pm). This figure shows overlapping RP HPLC chromatograms before (A) and after (B) treatment with 30 U RecJ and 0.5 ug SSBP for 60 min at 37°C.