Additional File 2. Amino acid polymorphism in 40 unique partial Pokey transposase sequences.

The sequences were derived from 53 *Pokey* elements obtained from members of the *Daphnia pulex* complex. Numbers at each position represent the number of sequences that carry this amino acid. Not all values sum to 40 as indels occurred in some sequences. The pink bar represents the amino sequence coded by the partial exon 1. The blue bar represents the amino sequence coded by exon 2. The dark blue regions with greek numbers represent estimated regions of recombination identified using maximum chi-square analysis (Additional File 3).

	5	10	15	20	25	30	35	40
A40 F 39R40	V40A40S40N	140Y40 <mark>H40G40</mark> S40D26I35	V40K40V40R4	40R40K39Q40R40D	40 G 40 S 40 F 40 R 35 S 3	35 K 40 S 40 C 40 P 40 K	40A40I40D40D40Y	38 V 40 N 40 N 40
V_1		N_4 L_4		R_1	K_4 T_2	4		
	45	50	55	60	65	70	75	80
$M_{40}G_{40}G_{39}$	$V_{40}D_{40}T_{40}P_{40}$	A40N40Q39L40R40S40Y40	$Y_{40}E_{40}R_{40}D_{4}$	40R40K40A40K40K	40W40W40H39R40L	40L40Y40S40L40L	40E40T40C40L40V	40N40S40W40
	_	R_1			Q_1			
	85	90	95	100	105	110	115	120
$I_{40}C_{40}F_{40}$	$N_{40}D_{40}M_{37}V$	7 ₄₀ E ₄₀ E ₄₀ N ₄₀ Y ₃₉ L ₄₀ E ₂₃			40E40F40K40R40N			$_{40}{ m N}_{40}{ m E}_{40}{ m N}_{23}$
	Ιз			$_{4}$ Q_{18} $P_{19}L_{17}$		A_1	Y ₄ S ₁₇	K_{17}
		V_3					Q_2	
	105		N ₇ D ₂		1 4 5	1.50	155	1.00
	125	130	135	140	145	150	155	160
~		A40G40R40M40M25P38T34	1 1 40 H 40 P 40 S 4		40K40K40K39K40S: R			
$E_{10}K_{19}$		K ₁₅ R ₂ S ₆		G_1	K:	2	$egin{array}{cccc} Q_8 \ F_{17} & L \ E_7 \end{array}$	15 0 16 0 19
N_4			I				7 نـــا	
IN 4	165	170	175	180	185	190	195	200
G40N19H40		740G40E40A40R40G40R40						
I ₁₅	F ₁	40 40 11 40 140 140 140	S ₂			402 402 402 400	55 C 401 140 Q 40 C 4011	V ₃
L_6			Y		5 T 5			
				Н		${f \Pi}$		
	205	210	215	220	225	230	235	240
C40L40 G 40	K40K40R40 N	I ₄₀ C ₄₀ F ₄₀ V ₄₀ E ₄₀ Y ₂₃ H ₄₀	D40E26 <mark>N40</mark> Y3	38 F 16Q16S19E40E	32E5 A5 P40L40E	20E 19V40E 40A40V	39Y40S40 A 40S39E	40E40S40D40
		\mathbf{F}_{17}	D_{14} C_{1}	V ₁₅ D	7 D:	13 D 17		
		_		\mathbf{F}_{5} G	1 A	5 G 3		
			${f II}$ and ${f IV}$	A_1				
	245	250	255	260	265	270	275	280
		C40P40T40N40K23E40C33					$_4$ M_4 I_4 D_4 D_4 Y	4 D4 D4 F4
V_{17}		A_{15} F_3	T_1	$P_1 Y_9 G$	9 P4 D1 Y11S12N	19		
3.7		E_2			$H_1 P_{10}$			
V	005	VI	005		I 1			
	285	290	295 E 0 0					
~		V ₂₅ I ₃₉ D ₄₀ E ₃₁ Y ₄₀ D ₃₀ E ₂₅	~ ~					
-	E ₈ Y ₁₉ N	$I_8 F_1 D_9 N_{10} D_{15}$	L6 E17					