Table S1. Overlap between *Pseudorasbora parva* and *Knipowitschia caucasica* isotopic niches as estimated according to Kernel utilization density (UD), bivariate ellipse and minimum convex polygon home ranges considering 40%, 75% and 95% of the data.

	OverlapID	*K. caucasica*_40	*K. caucasica*_75	*K. caucasica*_95	*P. parva*_40	*P. parva*_75	*P. parva*_95
rKin	*K. caucasica*_40	1.000	1.000	1.000	0.000	0.000	0.000
	*K. caucasica*_75	0.251	1.000	1.000	0.000	0.000	0.006
	*K. caucasica*_95	0.114	0.452	1.000	0.000	0.000	0.030
	*P. parva*_40	0.000	0.000	0.000	1.000	1.000	1.000
	*P. parva*_75	0.000	0.000	0.000	0.409	1.000	1.000
	*P. parva*_95	0.000	0.002	0.018	0.222	0.544	1.000
	OverlapID	*K. caucasica*_40	*K. caucasica*_75	*K. caucasica*_95	*P. parva*_40	*P. parva*_75	*P. parva*_95
ellipse	*K. caucasica*_40	1.000	1.000	1.000	0.000	0.000	0.000
	*K. caucasica*_75	0.368	1.000	1.000	0.000	0.000	0.000
	*K. caucasica*_95	0.171	0.463	1.000	0.000	0.000	0.000
	*P. parva*_40	0.000	0.000	0.000	1.000	1.000	1.000
	*P. parva*_75	0.000	0.000	0.000	0.368	1.000	1.000
	*P. parva*_95	0.000	0.000	0.000	0.171	0.463	1.000
	OverlapID	*K. caucasica*_40	*K. caucasica*_75	*K. caucasica*_95	*P. parva*_40	*P. parva*_75	*P. parva*_95
polygon	*K. caucasica*_40	1.000	1.000	1.000	0.000	0.000	0.000
	*K. caucasica*_75	0.227	1.000	1.000	0.000	0.000	0.000
	*K. caucasica*_95	0.010	0.364	1.000	0.000	0.000	0.000
	*P. parva*_40	0.000	0.000	0.000	1.000	1.000	1.000
	*P. parva*_75	0.000	0.000	0.000	0.280	1.000	1.000
	*P. parva*_95	0.000	0.000	0.000	0.165	0.591	1.000