

APPENDIX S2: WITOMI RESULTS AND SPECIES' CODE

Table S1. Hydraulic requirement parameters of the 57 invertebrate taxa sampled in spring and autumn. Inertia = total variability; OMI = outlying mean index. *P* number of random permutations (out of 1000) that yielded a higher value than the observed marginality (OMI, WitOMIG or WitOMIG_K) (the value in **bold** characters are significant, *P* < 0.05). Tol = tolerance, Rtol = residual tolerance. I. = inertia; *G* (*G*_k) are the subset marginality WitOMIG (WitOMIG_K); – = NA. The WitOMI cannot be calculated when the OMI is not significant (See Discussion for further details). The species is the one use as example for Figure 3D, 3E, and 3F (Species code in Appendix S2; Table S3).

Season	All					Spring					Autumn					Spring					Autumn					
	Code	I.	OMI	Tol	Rtol	<i>P</i>	I.	<i>G</i>	Tol	Rtol	<i>P</i>	I.	<i>G</i>	Tol	Rtol	<i>P</i>	I.	<i>G</i> _K	Tol	Rtol	<i>P</i>	I.	<i>G</i> _K	Tol	Rtol	<i>P</i>
AFLU	5.54	1.63	1.16	2.75	0.05	5.59	2.46	0.54	2.59	0.00	5.05	4.99	0.04	0.02	0.00	5.89	2.76	0.70	2.43	0.00	5.05	4.99	0.04	0.02	0.00	
ANTO	6.26	2.86	0.48	2.92	0.01	6.26	2.86	0.48	2.92	0.00	-	-	-	-	-	7.08	3.68	0.41	3.00	0.00	-	-	-	-	-	
ATSP	5.26	0.77	1.94	2.54	0.10	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
BASP	4.95	0.15	0.92	3.88	0.00	5.22	0.37	1.28	3.57	0.01	4.02	0.95	0.86	2.21	0.00	4.89	0.04	2.11	2.74	0.01	4.02	0.95	0.86	2.21	0.00	
BIMI	6.44	2.67	1.22	2.56	0.07	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
BFAS	5.52	1.63	0.83	3.06	0.00	5.52	1.63	0.83	3.06	0.00	-	-	-	-	-	4.86	0.97	1.62	2.27	0.00	-	-	-	-	-	
CASP	4.75	2.09	0.70	1.96	0.00	3.82	2.28	0.45	1.09	0.00	5.17	2.46	0.75	1.96	0.00	3.98	2.44	0.46	1.08	0.00	5.17	2.46	0.75	1.96	0.00	
CESP	5.99	1.36	1.74	2.89	0.14	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
CERA	5.24	0.44	1.56	3.24	0.07	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
CLEP	5.03	0.44	1.38	3.21	0.00	5.04	0.48	0.96	3.60	0.01	4.99	1.70	1.40	1.89	0.00	4.68	0.12	2.26	2.30	0.00	4.99	1.70	1.40	1.89	0.00	
CMAR	5.30	2.09	1.10	2.10	0.00	5.36	2.17	0.96	2.24	0.00	4.97	3.47	0.60	0.90	0.00	4.80	1.60	1.55	1.65	0.00	4.97	3.47	0.60	0.90	0.00	
CHIR	4.44	0.49	1.52	2.42	0.12	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
CPIC	5.51	2.83	0.52	2.16	0.00	5.51	2.83	0.52	2.16	0.00	-	-	-	-	-	5.72	3.04	0.56	2.12	0.00	-	-	-	-	-	
DRSP	4.50	1.07	1.90	1.52	0.04	8.57	3.37	2.26	2.94	0.00	3.75	2.11	0.71	0.93	0.00	7.42	2.22	2.13	3.08	0.00	3.75	2.11	0.71	0.93	0.00	
DUSP	5.33	0.14	1.74	3.45	0.05	5.85	0.59	1.64	3.61	0.00	4.78	1.01	1.72	2.05	0.00	5.38	0.12	3.08	2.18	0.00	4.78	1.01	1.72	2.05	0.00	
ECSP	5.35	0.33	1.47	3.55	0.00	5.33	0.53	1.48	3.32	0.01	5.49	0.57	1.22	3.70	0.00	4.86	0.06	2.57	2.24	0.01	5.49	0.57	1.22	3.70	0.00	
ECTE	5.46	3.09	0.12	2.25	0.12	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
ELMA	3.18	0.79	0.04	2.35	0.86	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
EPEO	5.68	1.27	0.75	3.66	0.26	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
EVIR	5.30	0.34	1.28	3.68	0.04	5.30	0.34	1.28	3.68	0.01	-	-	-	-	-	5.20	0.24	2.07	2.89	0.00	-	-	-	-	-	
EPAR	269	3.93	0.14	0.80	3.00	0.26	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
EPYG	4.77	0.14	1.02	3.61	0.02	4.77	0.32	0.50	3.95	0.01	4.77	0.47	0.54	3.75	0.00	4.54	0.09	1.10	3.36	0.00	4.77	0.47	0.54	3.75	0.00	
ESSP	4.86	0.14	0.86	3.86	0.00	5.04	0.65	1.45	2.94	0.00	4.74	0.49	0.47	3.78	0.00	5.00	0.61	1.58	2.81	0.00	4.74	0.49	0.47	3.78	0.00	
EGEN	5.05	1.91	0.86	2.28	0.00	5.05	1.91	0.86	2.28	0.00	-	-	-	-	-	5.01	1.87	0.91	2.23	0.00	-	-	-	-	-	
GASP	5.69	0.36	1.17	4.16	0.05	5.87	0.43	1.09	4.35	0.01	2.58	0.62	0.35	1.62	0.00	5.51	0.07	2.19	3.25	0.01	2.58	0.62	0.35	1.62	0.00	
HEXO	5.22	0.38	1.25	3.60	0.00	5.30	0.44	1.05	3.81	0.01	4.64	2.02	0.82	1.80	0.00	4.92	0.07	2.49	2.37	0.01	4.64	2.02	0.82	1.80	0.00	
HPEL	5.24	0.36	1.20	3.69	0.00	5.32	0.45	0.99	3.89	0.01	4.46	2.37	0.30	1.78	0.00	4.93	0.06	2.58	2.30	0.01	4.46	2.37	0.30	1.78	0.00	
HYDR	3.57	1.48	0.42	1.67	0.06	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
HYSP	5.42	0.34	1.01	4.07	0.00	5.44	0.42	0.98	4.05	0.01	5.12	0.76	1.38	2.98	0.00	5.03	0.01	1.85	3.18	0.01	5.12	0.76	1.38	2.98	0.00	
HYDS	6.18	2.75	0.83	2.59	0.03	6.18	2.75	0.83	2.59	0.00	-	-	-	-	-	7.16	3.73	0.75	2.67	0.00	-	-	-	-	-	
LEPT	5.34	2.74	0.47	2.13	0.16	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
LESP	4.42	0.31	0.90	3.21	0.04	4.61	0.45	1.09	3.07	0.01	2.63	1.24	0.26	1.13	0.00	4.40	0.24	1.31	2.85	0.00	2.63	1.24	0.26	1.13	0.00	
LOPAad	4.97	1.20	1.97	1.79	0.01	4.97	1.20	1.97	1.79	0.00	-	-	-	-	-	4.87	1.10	2.04	1.73	0.00	-	-	-	-	-	
LOPAla	4.57	0.48	1.25	2.85	0.00	5.11	0.37	0.47	4.27	0.01	3.72	1.91	0.31	1.50	0.00	4.88	0.14	1.21	3.53	0.01	3.72	1.91	0.31	1.50	0.00	
MPOW	5.77	1.21	2.07	2.49	0.27	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
MYSA	4.20	3.34	0.28	0.58	0.24	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
ORHE	5.58	1.77	0.89	2.92	0.00	5.58	1.77	0.89	2.92	0.00	-	-	-	-	-	4.93	1.12	1.73	2.08	0.00	-	-	-	-	-	
ONSP	5.18	1.10	1.74	2.34	0.06	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
ORTH	5.12	0.23	1.26	3.63	0.00	5.39	0.39	0.91	4.09	0.01	4.35	0.90	1.57	1.88	0.00	5.01	0.01	2.50	2.50	0.01	4.35	0.90	1.57	1.88	0.00	
ORSP	6.83	1.32	2.33	3.18	0.07	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
OTROad	5.32	0.32	0.79	4.21	0.02	5.25	0.41	0.98	3.87	0.01	6.23	2.59	1.00	2.64	0.00	5.01	0.16	1.60	3.25	0.01	6.23	2.59	1.00	2.64	0.00	
OTROla	5.52	0.23	1.21	4.08	0.06	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
PISI	4.43	0.37	1.12	2.94	0.77	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
POLY	6.09	1.16	2.02	2.91	0.07	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
PBIF	6.95	3.46	0.59	2.90	0.01	8.24	7.16	0.33	0.76	0.00	5.87	3.73	0.79	1.36	0.00	9.48	8.39	0.35	0.74	0.00	5.87	3.73	0.79	1.36	0.00	
PPUS	5.23	0.14	0.91	4.18	0.15	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
RASP	4.76	0.13	0.81	3.82	0.97	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
RHIP	6.91	6.12	0.52	0.27	0.00	6.91	6.12	0.52	0.27	0.00	-	-	-	-	-	6.36	5.57	0.54	0.25	0.00	-	-	-	-	-	
RHYP	5.93	1.93	1.14	2.86	0.00	5.93	1.93	1.14	2.86	0.00	-	-	-	-	-	5.28	1.28	1.61	2.39	0.00	-	-	-	-	-	
SIGN	5.51	0.53	1.71	3.26	0.00	5.51	0.53	1.71	3.26	0.01	-	-	-	-	-	5.30	0.32	2.08	2.90	0.00	-	-	-	-	-	
SARG	5.69	0.24	1.27	4.19	0.02	6.12	2.03	1.66	2.42	0.00	5.24	0.83	2.00	2.41	0.00	5.44	1.36	1.34	2.74	0.00	5.24	0.83	2.00	2.41	0.00	
SIMU	5.34	0.51	1.24	3.59	0.00	5.63	0.58	0.94	4.12	0.00	4.38	2.45	0.87	1.06	0.00	5.11	0.06	1.82	3.24	0.01	4.38	2.45	0.87	1.06	0.00	
STEN	6.27	0.57	1.59	4.11	0.38	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
SCAN	6.36	1.40	1.40	3.56	0.02	6.30	3.00	0.45	2.84	0.00	6.47	0.98	0.62	4.87	0.00	6.54	3.25	0.56	2.74	0.00	6.47	0.98	0.62	4.87	0.00	
TANYP	4.88	0.73	1.69	2.46	0.02	4.88	0.73	1.69	2.46	0.01	-	-	-	-	-	4.95	0.80	1.92	2.23	0.00	-	-	-	-	-	
TANYT	5.34	0.17	1.32	3.85	0.03	5.48	0.32	0.91	4.25	0.01	4.46	1.41	0.79	2.26	0.00	5.22	0.06	1.52	3.64	0.01	4.46	1.41	0.79	2.26	0.00	
TFLU	6.94	2.03	1.56	3.36	0.11	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-

Table S2. Physical habitat preference parameters of the 12 fish taxa sampled in ten Mediterranean tributaries of the Rhône River. Inertia = total variability; OMI = outlying mean index. *P* number of random permutations (out of 1000) that yielded a higher value than the observed marginality (OMI, WitOMIG or WitOMIG_{*K*}) (the value in **bold** characters are significant, *P* < 0.05). Tol = tolerance, Rtol = residual tolerance. I. = inertia; *G* (*G_k*) are the subset marginality WitOMIG (WitOMIG_{*K*}). – = NA. The WitOMI cannot be calculated when the OMI is not significant (See Discussion for further details). The species is the one use as example for Figure 4D, 4E, and 4F (Species code in Table S4 in appendix S2).

Altitude	All					Upstream					Downstream					Upstream					Downstream						
	Code	I.	OMI	Tol	Rtol	<i>P</i>	I.	<i>G</i>	Tol	Rtol	<i>P</i>	I.	<i>G</i>	Tol	Rtol	<i>P</i>	I.	<i>G_K</i>	Tol	Rtol	<i>P</i>	S.I.	<i>G_K</i>	Tol	Rtol	<i>P</i>	
SCU	5.50	2.15	1.50	1.85	0.08	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
OTR	5.71	0.08	2.43	3.21	0.00	4.74	1.12	1.62	2.01	0.00	8.44	4.14	1.09	3.21	0.00	3.62	0.00	0.96	2.66	0.00	4.47	0.18	1.69	2.60	0.00		
YTR	5.77	0.26	1.69	3.81	0.00	4.92	1.15	1.67	2.10	0.00	9.17	5.58	1.59	1.99	0.00	3.77	0.00	0.75	3.02	0.00	4.98	1.40	1.22	2.37	0.00		
<u>MIN</u>	5.32	0.45	2.52	2.35	0.00	2.82	0.33	0.96	1.54	0.00	8.51	4.61	0.81	3.08	0.00	2.79	0.30	1.01	1.48	0.00	3.93	0.04	0.88	3.01	0.00		
STO	5.91	0.77	2.41	2.73	0.00	2.86	0.20	0.67	1.98	0.00	9.28	4.93	0.88	3.48	0.00	3.17	0.51	0.78	1.87	0.00	4.38	0.03	1.24	3.11	0.00		
BLA	4.99	0.46	2.27	2.26	0.00	2.31	0.20	0.71	1.40	0.00	8.61	4.53	0.88	3.19	0.00	2.51	0.40	0.73	1.38	0.00	4.09	0.02	0.68	3.40	0.00		
SBA	3.71	0.07	0.51	3.13	0.77	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	
SON	6.38	4.89	0.75	0.75	0.03	-	-	-	-	-	6.38	4.89	0.75	0.75	0.00	-	-	-	-	-	2.80	1.31	0.13	1.37	0.00		
NAS	12.60	10.20	1.62	0.78	0.00	-	-	-	-	-	12.60	10.20	1.62	0.78	0.00	-	-	-	-	-	4.40	2.00	1.33	1.07	0.00		
GUD	8.43	4.31	0.95	3.17	0.00	1.16	0.25	0.47	0.44	0.00	9.47	5.52	0.51	3.44	0.00	2.53	1.62	0.02	0.89	0.00	4.04	0.09	0.98	2.97	0.00		
CHU	5.58	1.58	1.69	2.31	0.00	1.27	0.03	0.13	1.11	0.00	8.61	4.53	0.88	3.19	0.00	2.34	1.10	0.28	0.96	0.00	4.09	0.02	0.68	3.40	0.00		
STR	9.92	5.28	1.19	3.46	0.00	-	-	-	-	-	11.07	6.86	0.58	3.62	0.00	-	-	-	-	-	4.63	0.43	0.84	3.37	0.00		
BAR	8.34	3.83	1.30	3.21	0.00	1.66	0.52	0.07	1.07	0.00	9.46	5.26	0.58	3.62	0.00	2.60	1.45	0.88	0.27	0.00	4.28	0.07	0.60	3.60	0.00		

Table S3. Invertebrate code from Mériçoux and Dolédec (2004).

Species	Code
<i>Ancyclus fluviatilis</i> Müller	AFLU
<i>Antocha</i> sp.	ANTO
<i>Athripsodes</i> sp.	ATSP
<i>Baetis</i> sp.	BASP
<i>Bidessus minutissimus</i>	BIMI
<i>Blepharicera fasciata</i> (Westwood)	BFAS
<i>Caenis</i> sp.	CASP
<i>Ceraclea</i> sp.	CESP
<i>Ceratopogoninae</i>	CERA
<i>Cheumatopsyche lepida</i> (Pictet)	CLEP
<i>Chimarra marginata</i> (Linnaeus)	CMAR
Chironomini	CHIR
<i>Choroterpes picteti</i> Eaton	CPIC
<i>Dryops</i> sp.	DRSP
<i>Dugesia</i> sp.	DUSP
<i>Ecdyonurus</i> sp.	ECSP
<i>Ecnomus tenellus</i>	ECTE
<i>Elmis maugetii</i> (l)	ELMA
<i>Epeorus</i> sp.	EPEO
<i>Ephoron virgo</i> (Olivier)	EVIR
<i>E. parallelepipedus</i> (a)	EPAR
<i>Esolus</i> spp. (l)	ESSP
<i>E. pygmaeus</i> (a) (Ph. Müller)	EPYG
<i>Euleuctra geniculata</i> Stephens	EGEN
<i>Gammarus</i> spp.	GASP
<i>Hydropsyche</i> spp.	HYSP
<i>H. excellata</i> Dufour	HEXO
<i>H. pellucidula</i> (Curtis)	HPEL
<i>Hydra</i> sp.	HYDR
<i>Hydroptila</i> sp.	HYDS
<i>Leptocerus tineiformis</i>	LEPT
<i>Leuctra</i> sp.	LESP
<i>Limnius opacus</i> (a) Ph. Müller	LOPAad
<i>L. opacus</i> (l) Ph. Müller	LOPAa
<i>Micronecta poweri</i>	MPOW
<i>Mystacides azurea</i>	MYSA
<i>Oligoneuriella rhenana</i> (Imhoff)	ORHE
<i>Onychogomphus</i> sp.	ONSP
Orthoclaadiinae	ORTH
<i>Orthotrichia</i> sp.	ORSP
<i>Oulimnius troglodytes</i> (a) (Gyllenhal)	OTROad
<i>O. troglodytes</i> (l)	OTROa
<i>Pisidium</i> sp.	PISI
<i>Polycentrus flavomaculatus</i>	POLY
<i>Proclaeon bifidum</i> (Bengtsson)	PBIF
<i>Psychomyia pusilla</i>	PPUS
<i>Radix</i> sp.	RASP
<i>Rhyacophila</i> sp.	RHYP
<i>Rhithrogena</i> sp.	RHIP
<i>Serratella ignita</i> (Poda)	SIGN
<i>Setodes argentipunctellus</i> McLachlan	SARG
Simuliidae	SIMU
<i>Stenelmis canaliculata</i> (a)	STEN
<i>S. canaliculata</i> (l)	SCAN
Tanypodinae	TANYP
Tanytarsini	TANYT
<i>Theodoxus fluviatilis</i>	TFLU

Table S4. Fish code from Dolédec et al. (2000); † Young of the year.

Species	Code
Sculpin (<i>Cottus gobio</i>)	SCU
Older trout (<i>Salmo trutta</i>)	OTR
Y-O-Y† trout (<i>Salmo trutta</i>)	YTR
Minnnow (<i>Phoxinus phoxinus</i>)	MIN
Stone loach (<i>Nemacheilus barbatulus</i>)	STO
Blageon (<i>Telestes soufia</i>)	BLA
Southwestern barbel (<i>Barbus meridionalis</i>)	SBA
Southwestern nase (<i>Chondrostoma toxostoma</i>)	SON
Nase (<i>Chondrostoma nasus</i>)	NAS
Gudgjeon (<i>Gobio gobio</i>)	GUD
Chub (<i>Leuciscus cephalus</i>)	CHU
Streambleak (<i>Alburnoides bipunctatus</i>)	STR
Barbel (<i>Barbus barbus</i>)	BAR

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