

Table A-1:
Physiochemical Data

Year	Study Reach	Stream Order	Elevation (ft.)	Gradient	Wet stream width (ft.)	Q (cfs)	Habitat Assessment Score	Watershed Area (ac)	OS	URB	AG	HERB	Stream temp. (°F)	pH	Dissolved Oxygen (mg/l)	Conductivity (µS)	Specific conductance (µS at 25°C)	% riparian canopy coverage
2000	C1	4	15	0.008	12.4	0.21	100	9598	80	3	17	0	65.3	7.6	11.8	1240	1416	ND
2000	C2	4	50	0.008	17.5	0.18	112	9336	83	2	15	0	68.5	6.6	11.9	1083	1190	ND
2000	C3	3	420	0.033	20.9	4.34	173	4517	100	0	0	0	60.8	8.6	7.9	539	651	ND
2000	F1	3	10	0.007	8.3	0.18	26	5905	57	15	29	0	73.0	8	9.7	1204	1257	ND
2000	SM1	2	10	0.014	8.3	0.35	26	5945	56	15	28	0	73.8	8.2	8.5	578	599	ND
2000	M1	3	40	0.008	13.8	0.14	80	6799	49	49	2	0	80.4	8	14.9	1220	1177	ND
2000	M3	3	400	0.071	13.2	0.74	157	4142	79	18	2	0	66.6	8	9.1	820	922	ND
2000	M4	2	1000	0.083	14.3	0.32	178	1376	100	0	0	0	70.2	8	8.8	810	873	ND
2000	M5	2	2600	0.167	12.1	0.14	156	440	100	0	0	0	59.9	8	8.9	587	717	ND
2000	M6	2	1550	0.125	16.5	0.00	150	1042	100	0	0	0	72.0	8	7.3	817	863	ND
2000	AB2	3	75	0.010	8.8	0.25	83	4789	60	30	9	1	72.3	8	11.8	1580	1663	ND
2000	AB3	2	300	0.025	11.6	0.39	163	2356	90	3	6	0	63.5	8	9.2	1026	1198	ND
2000	SA1	3	260	0.028	14.9	0.32	143	2867	88	3	3	6	65.5	8	9.4	992	1130	ND
2000	SA2	3	500	0.05	15.4	0.18	137	2556	94	0	2	4	68.4	8	8.7	1190	1310	ND
2000	MY1	3	130	0.017	11.0	0.39	110	3779	81	6	10	4	77.2	8	8.9	1260	1258	ND
2000	MY2	3	290	0.033	12.1	0.81	168	2213	96	0	4	0	72.3	7.8	7.4	645	679	ND
2000	MY3	3	1200	0.125	14.3	0.21	172	1316	100	0	0	0	59.7	7.2	9.3	493	604	ND
2000	SJ1	4	30	0.005	27.0	1.09	110	5169	64	14	20	1	65.1	8	8.8	1283	1468	ND
2000	SJ2	3	120	0.014	12.1	0.56	104	3836	81	1	16	1	68.9	8	10	1395	1526	ND
2000	SJ3	2	1000	0.107	18.7	0.21	182	3245	95	1	4	1	61.2	8	8.8	735	883	ND
2000	SJ4	1	1840	0.055	6.1	0.07	160	950	96	3	0	2	62.1	7	7.9	351	417	ND
2000	T1	3	30	0.01	15.8	0.49	80	3616	75	4	19	2	64.9	7.5	9.7	1705	1955	ND
2000	T2	3	110	0.017	7.7	0.35	111	3221	81	0	18	1	77.4	8	9.8	1673	1667	ND
2000	DP1	4	40	0.01	16.0	0.67	111	4574	88	0	11	1	71.2	7.5	9.2	1340	1427	ND
2000	R1	3	180	0.02	11.0	0.14	107	3391	79	12	4	4	70.5	7.8	8.3	1122	1205	ND
2000	R2	3	360	0.036	13.2	0.28	152	2840	85	15	0	0	68.4	8	9.3	785	864	ND
2000	AH1	2	150	0.040	13.8	0.49	168	2583	100	0	0	0	60.8	7.5	9.8	887	1071	ND
2000	AH2	2	240	0.16	18.7	0.67	182	2324	100	0	0	0	61.5	7.7	9.2	833	997	ND
2000	SO1	2	20	0.025	11.6	0.64	161	1316	96	0	0	4	63.3	7.8	7.6	839	981	ND
2000	SO2	2	500	0.133	16.5	0.56	189	1025	96	0	0	4	63.7	7.7	7.3	730	850	ND
2001	C1	4	15	0.008	20.4	0.35	111	9598	80	3	17	0	64.4	8.4	ND	1102	1272	ND
2001	C2	4	50	0.008	19.8	0.11	122	9336	83	2	15	0	61.0	7.5	ND	667	804	ND
2001	C3	3	420	0.033	17.6	1.76	175	4517	100	0	0	0	59.2	7.8	ND	345	425	ND
2001	AB2	3	75	0.010	14.9	0.78	70	4789	60	30	9	1	64.4	8.4	ND	838	967	ND
2001	AB3	2	300	0.025	13.2	0.78	171	2356	90	3	6	0	61.3	7.8	ND	828	993	ND
2001	AT1	4	15	0.002	31.4	1.38	60	12443	48	39	7	6	64.9	8	ND	1692	1940	ND
2001	AT2	3	50	0.007	20.9	0.81	92	2393	12	73	7	8	66.7	8.4	ND	1413	1586	ND
2001	SJ1	4	30	0.005	16.0	1.80	109	5169	64	14	20	1	63.7	7.8	ND	1055	1229	ND
2001	SJ2	3	120	0.014	11.0	1.34	97	3836	81	1	16	1	62.1	8.4	ND	992	1179	ND
2001	SJ3	2	1000	0.107	18.7	0.28	179	3245	95	1	4	1	59.5	7.8	ND	656	805	ND
2001	AH1	2	150	0.040	12.3	0.74	170	2583	100	0	0	0	65.1	7.8	ND	638	730	ND
2001	SO2	2	500	0.133	18.2	0.46	188	1025	96	0	0	4	60.3	8.1	ND	567	689	ND
2002	RIN1	3	220	0.022	14.3	0.30	128	8131	83	0	16	1	58.8	7.65	8.64	894	1108	ND
2002	C1	4	15	0.008	14.5	0.16	102	9598	80	3	17	0	56.8	7.84	6.65	1206	1521	ND

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Year	Study Reach	Stream Order	Elevation (ft.)	Gradient	Wet stream width (ft.)	Q (cfs)	Habitat Assessment Score	Watershed Area (ac)	OS	URB	AG	HERB	Stream temp. (°F)	pH	Dissolved Oxygen (mg/l)	Conductivity (µS)	Specific conductance (µS at 25°C)	% riparian canopy coverage
2002	C3	3	420	0.033	19.8	1.11	170	4517	100	0	0	0	54.9	8.45	10.26	529	693	ND
2002	SY1	3	30	0.010	9.4	0.30	84	2482	45	54	2	0	59.2	8.35	8.85	2545	3140	ND
2002	M1	3	40	0.008	17.6	0.23	81	6799	49	49	2	0	72.7	8.01	15.86	1252	1313	ND
2002	M2	1	50	0.020	8.3	0.17	64	643	24	76	0	0	63.5	7.66	4.05	1172	1367	ND
2002	M3	3	400	0.071	12.7	0.08	157	4142	79	18	2	0	60.6	8.01	8.14	895	1083	ND
2002	AB1	3	25	0.008	15.4	1.58	71	5646	51	38	7	4	61.9	7.93	6.86	1350	1608	ND
2002	AB2	3	75	0.010	16.5	1.92	101	4789	60	30	9	1	56.5	7.97	8.12	976	1248	ND
2002	AB3	2	300	0.025	11.0	0.30	168	2356	90	3	6	0	56.1	8.17	8.51	1097	1409	ND
2002	AT1	4	15	0.002	18.7	0.46	75	12443	48	39	7	6	57.4	7.74	3.85	2023	2559	ND
2002	AT2	3	50	0.007	6.6	0.30	74	2393	12	73	7	8	62.4	7.85	7.65	1840	2178	ND
2002	SJ1	4	30	0.005	7.8	0.23	72	5169	64	14	20	1	51.8	8.00	7.50	1069	1459	ND
2002	SJ2	3	120	0.014	10.5	0.26	107	3836	81	1	16	1	63.3	7.87	9.99	1387	1623	ND
2002	SJ3	2	1000	0.107	22.4	0.36	177	3245	95	1	4	1	53.6	8.35	9.83	724	965	ND
2002	EC1	3	10	0.020	11.6	0.12	134	3915	93	2	0	5	56.7	7.75	7.00	938	1198	ND
2002	AH1	2	150	0.040	11.6	0.31	169	2583	100	0	0	0	52.9	8.35	9.56	773	1039	ND
2002	SO2	2	500	0.133	8.0	0.21	179	1025	96	0	0	4	56.8	8.18	9.27	793	1008	ND
2002	GAV1	3	25	0.008	13.2	1.10	140	12791	59	0	0	41	66.4	8.18	11.71	1434	1617	ND
2002	GAV2	3	260	0.020	13.2	0.95	126	10828	54	0	0	46	76.6	8.16	9.57	1542	1548	ND
2003	C1	4	15	0.008	20.8	1.29	113	9598	80	3	17	0	58.5	8.20	9.68	662	824	66
2003	C3	3	420	0.033	23.0	4.62	174	4517	100	0	0	0	57.7	8.45	9.31	497	624	81
2003	MON1	3	350	0.050	9.8	2.79	133	3465	82	17	1	0	65.1	7.90	8.35	845	968	93
2003	SY1	3	30	0.010	13.7	0.90	95	2482	45	54	2	0	60.8	8.51	11.64	1786	2158	80
2003	SY2	3	200	0.033	15.3	0.88	109	1956	52	46	2	0	57.7	8.28	9.68	1291	1624	72
2003	M1	3	40	0.008	13.1	1.69	88	6799	49	49	2	0	66.7	8.20	12.26	1210	1358	27
2003	M2	1	50	0.020	8.8	0.72	79	643	24	76	0	0	71.2	8.37	8.11	1280	1364	68
2003	M3	3	400	0.071	15.3	2.91	157	4142	79	18	2	0	60.3	8.86	9.49	802	975	93
2003	AB1	3	25	0.008	14.2	3.05	104	5646	51	38	7	4	63.3	8.57	10.61	1696	1985	40
2003	AB2	3	75	0.010	14.8	2.15	84	4789	60	30	9	1	60.8	8.43	6.84	1356	1637	87
2003	AB3	2	300	0.025	9.1	3.24	158	2356	90	3	6	0	60.3	8.74	9.73	789	960	92
2003	AT1	4	15	0.002	18.0	1.15	83	12443	48	39	7	6	69.3	8.39	9.36	2516	2744	18
2003	AT2	3	50	0.007	10.9	0.74	58	2393	12	73	7	8	70.5	8.47	11.57	1917	2060	37
2003	SA2	3	500	0.05	7.1	1.63	152	2556	94	0	2	4	65.8	8.89	8.94	898	1019	68
2003	SJ1	4	30	0.005	13.7	4.17	90	5169	64	14	20	1	54.7	8.26	10.46	724	948	39
2003	SJ2	3	120	0.014	12.6	2.48	106	3836	81	1	16	1	60.1	8.13	10.23	634	772	70
2003	SJ3	2	1000	0.107	20.3	0.44	178	3245	95	1	4	1	58.6	8.86	9.19	696	864	86
2003	AH1	2	150	0.040	13.8	2.87	174	2583	100	0	0	0	59.9	8.81	9.22	710	868	100
2003	SO2	2	500	0.133	12.0	1.15	181	1025	96	0	0	4	60.1	9.02	9.16	819	1000	93
2003	GAV1	3	25	0.008	13.1	3.84	157	12791	59	0	0	41	65.3	8.63	9.65	1397	1593	15

Note: ND = Not determined.

Table A-2
Plant Species Observed at the Study Reaches

		Plant Species Checklist																																																
Family	Common Name	Scientific Name	RIN1	C1	C2	C3	MONT1	SY1	SY2	M1	M2	M3	M4	M5	M6	AB1	AB2	AB3	AT1	AT2	SA1	SA2	MY1	MY2	MY3	SJ1	SJ2	SJ3	SJ4	T1	T2	DP1	EC1	R1	R2	AH1	AH2	SO1	SO2	GAV1	GAV2									
Rhamnaceae	Coffee berry	<i>Rhamnus californica</i>	1	4	2	4	1	2	1	3	2	3	1	1	1	2	4	4	3	3	1	2	1	1	1	4	4	4	1	1	1	1	1	1	1	1	4	1	1	1	1	1	1							
Rhamnaceae	Redberry	<i>Rhamnus crocea</i>										1														1																								
Rhamnaceae	Hollyleaf redberry	<i>Rhamnus ilicifolia</i>																									1																							
Rosaceae	Mountain mahogany	<i>Cercocarpus betuloides</i>			1								1	1	1							1	1																											
Rosaceae	Toyon	<i>Heteromeles arbutifolia</i>			1								1	1	1							1	1																											
Rosaceae	Holly-leaf cherry	<i>Prunus ilicifolia</i>																				1	1					1	1																					
Rosaceae	California rose	<i>Rosa californica</i>										1											1																											
Rosaceae	California blackberry	<i>Rubus ursinus</i>	1	1	1	1	1			1	1	1	1			1	1	1	1	1	1	1	1		1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1			
Salicaceae	Narrow-leaved willow	<i>Salix exigua</i>																			1	1																												
Scrophulariaceae	Annual paintbrush	<i>Castilleja minor ssp. spiralis</i>																																																
Scrophulariaceae	Heart-leaved penstemon	<i>Keckeitia cordifolia</i>											1									1	1																											
Scrophulariaceae	Bush monkey flower	<i>Mimulus aurantiacus</i>				1																1			1	1																								
Scrophulariaceae	Scarlet monkey flower	<i>Mimulus cardinalis</i>				1	1						1	1								1																												
Scrophulariaceae	California figwort	<i>Scrophularia californica</i>	1	1	1	1						1	1	1							1	1	1																											
Solanaceae	White nightshade	<i>Solanum douglasii</i>	1	1	1	1	1	1	1	1	1	1	1			1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1		
Solanaceae	Purple nightshade	<i>Solanum xanti</i>				1																																												
Typhaceae	Cattails	<i>Typha spp.</i>	1	1	1	1				1	1					1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1		
Urticaceae	Stinging nettle	<i>Urtica dioica</i>		1		1				1												1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1		
Verbenaceae	Western vervain	<i>Verberna lasiostachys</i>																																																
Vitaceae	Wild grape	<i>Vitis girdiana</i>			1																																													
Non-Native Trees																																																		
Anacardiaceae	Peruvian pepper tree	<i>Schinus molle</i>																																																
Myoporaceae	Myoporum	<i>Myoporum laetum</i>			1																																													
Myrtaceae	Eucalyptus	<i>Eucalyptus spp.</i>		1	1		1	1	1	1	1	1				1	1									1	1																							
Pittosporaceae	Pittosporum	<i>Pittosporum sp.</i>		1			1	1	1	1	1																																							
Tamaricaceae	Tamarisk	<i>Tamarisk spp.</i>					1	1	1	1	1																																							
Non-Native Shrubs, Vines, and Herbs																																																		
Apiaceae	Poison hemlock	<i>Conium maculatum</i>	1	1	1	1			1		1																																							
Apiaceae	Sweet fennel	<i>Foeniculum vulgare</i>	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	
Apocynaceae	Periwinkle	<i>Vinca major</i>	1	1	1	1	1	1	1	1	1	1																																						
Araliaceae	English ivy	<i>Hedera helix</i>	1	1			1	1	1	1	1	1	1																																					
Asteraceae	Thoroughwort	<i>Ageratina ademerpha</i>	1		1	1	1	1	1	1																																								
Asteraceae	Italian thistle	<i>Cardus pycnocephalus</i>																																																
Asteraceae	Brassbuttons	<i>Cortula coronopifolia</i>																																																
Asteraceae	Cudweed	<i>Gnaphalium luteo-album</i>																																																
Asteraceae	Cape ivy (formerly German ivy)	<i>Senecio mikioides</i>	1	1	1	1	1	1	1	1	1	1				1	1																																	
Asteraceae	Common sow thistle	<i>Sonchus oleraceus</i>	1	1	1																																													
Asteraceae	Milk thistle	<i>Silybum marianum</i>																																																
Brassicaceae	Black mustard	<i>Brassica nigra</i>	1	1	1		1	1	1	1	1	1																																						
Brassicaceae	Wild radish	<i>Raphanus sativus</i>	1				1			1	1																																							
Caprifoliaceae	Japanese honeysuckle	<i>Lonicera japonica</i>					1																																											
Cyperaceae	Cyperus	<i>Cyperus spp.</i>					1			1																																								
Euphorbiaceae	Castor bean	<i>Ricinus communis</i>	1				1	1		1	1	1				1	1</																																	

Table A-5
Biological Metric Screening ANOVAs Summary

Parameter	r-square	p	Group	n	Mean	Std Error	Lower 95%	Upper 95%
# I Orders-field	0.58	<.0001	HIGH DIST	21	3.90476	0.2533	3.3915	4.418
			MOD DIST	11	6.36364	0.34999	5.6545	7.0728
			REF	8	6.75	0.4104	5.9184	7.5816
# I Fams-field	0.74	<.0001	HIGH DIST	21	5.8571	0.58892	4.664	7.05
			MOD DIST	11	13.9091	0.8137	12.26	15.558
			REF	8	15.5	0.95415	13.567	17.433
BMI density	0.01	0.7371	HIGH DIST	40	1773.92	226.73	1322.6	2225.2
			MOD DIST	18	2038.57	337.99	1365.8	2711.3
			REF	24	1704.54	292.71	1121.9	2287.2
# BMI orders	0.16	0.001	HIGH DIST	40	9.075	0.28556	8.507	9.643
			MOD DIST	18	11.0556	0.42569	10.208	11.903
			REF	24	9.5417	0.36865	8.808	10.275
# BMI families	0.57	<.0001	HIGH DIST	40	17.025	0.8737	15.286	18.764
			MOD DIST	18	30.3333	1.3024	27.741	32.926
			REF	24	28.9583	1.1279	26.713	31.203
# non-insect taxa	0.3	<.0001	HIGH DIST	40	4.075	0.24631	3.5847	4.5653
			MOD DIST	18	4.22222	0.36718	3.4914	4.9531
			REF	24	1.91667	0.31799	1.2837	2.5496
# insect orders	0.47	<.0001	HIGH DIST	40	5	0.19978	4.6023	5.3977
			MOD DIST	18	6.83333	0.29782	6.2405	7.4261
			REF	24	7.625	0.25792	7.1116	8.1384
# insect families	0.62	<.0001	HIGH DIST	40	12.95	0.8727	11.213	14.687
			MOD DIST	18	26.1111	1.3009	23.522	28.701
			REF	24	27.0417	1.1266	24.799	29.284
# EPT families	0.64	<.0001	HIGH DIST	40	3.75	0.51974	2.715	4.785
			MOD DIST	18	11.6111	0.77478	10.069	13.153
			REF	24	12.9167	0.67098	11.581	14.252
% EPT	0.36	<.0001	HIGH DIST	40	22.7	2.9969	16.735	28.665
			MOD DIST	18	45.8333	4.4675	36.941	54.726
			REF	24	53.5	3.869	45.799	61.201
% PT	0.44	<.0001	HIGH DIST	40	2.975	1.1284	0.729	5.221
			MOD DIST	18	12.7778	1.6822	9.429	16.126
			REF	24	16.9167	1.4568	14.017	19.816
% sens EPT	0.66	<.0001	HIGH DIST	40	1.1	1.0281	-0.946	3.146
			MOD DIST	18	8.3889	1.5325	5.338	11.439
			REF	24	21.7917	1.3272	19.15	24.433
BI score	0.53	<.0001	HIGH DIST	40	5.89975	0.10881	5.6832	6.1163
			MOD DIST	18	5.12222	0.16221	4.7994	5.4451
			REF	24	4.23458	0.14047	3.955	4.5142
% sens BMIs	0.67	<.0001	HIGH DIST	40	1.15	1.0572	-0.954	3.254
			MOD DIST	18	8.7778	1.576	5.641	11.915
			REF	24	23.0833	1.3649	20.367	25.8
% tol BMIs	0.13	0.0049	HIGH DIST	40	20.5	2.7133	15.099	25.901
			MOD DIST	18	14.7778	4.0447	6.727	22.829
			REF	24	5.5417	3.5028	-1.43	12.514
% dominant taxa	0.37	<.0001	HIGH DIST	40	50.8	2.5205	45.783	55.817
			MOD DIST	18	26.2778	3.7574	18.799	33.757
			REF	24	27.0833	3.254	20.606	33.56
% two dominant taxa	0.46	<.0001	HIGH DIST	40	69.475	2.349	64.799	74.151
			MOD DIST	18	42.7222	3.5017	35.752	49.692
			REF	24	42.7083	3.0326	36.672	48.745
% Diptera as Chironomidae	0.17	0.0008	HIGH DIST	40	78.925	3.0069	72.94	84.91
			MOD DIST	18	62.4444	4.4824	53.522	71.366
			REF	24	62.1667	3.8819	54.44	69.893
% non-insect BMIs	0.2	0.0001	HIGH DIST	40	21.425	2.6253	16.199	26.651
			MOD DIST	18	10.9444	3.9136	3.155	18.734

Table A-5
Biological Metric Screening ANOVAs Summary

Parameter	r-square	p	Group	n	Mean	Std Error	Lower 95%	Upper 95%
			REF	24	2.6667	3.3893	-4.08	9.413
% non-insects + diptera	0.47	<.0001	HIGH DIST	40	72.375	3.1607	66.084	78.666
			MOD DIST	18	40.5556	4.7117	31.177	49.934
			REF	24	31.8333	4.0805	23.711	39.955
% non-insects+chironimidae	0.46	<.0001	HIGH DIST	40	63.375	3.3476	56.712	70.038
			MOD DIST	18	29.8889	4.9904	19.956	39.822
			REF	24	21.5417	4.3218	12.939	30.144
EPT/Chironomidae	0.14	0.0027	HIGH DIST	40	1.69725	0.52426	0.6537	2.7408
			MOD DIST	18	3.375	0.78151	1.8194	4.9306
			REF	24	4.71208	0.67681	3.3649	6.0592
% cg	0.03	0.2615	HIGH DIST	40	40.05	3.0218	34.035	46.065
			MOD DIST	18	46.9444	4.5046	37.978	55.911
			REF	24	47.0833	3.9011	39.318	54.848
%sc	0.1	0.0153	HIGH DIST	40	2.4	0.66276	1.0808	3.7192
			MOD DIST	18	5.77778	0.98799	3.8112	7.7443
			REF	24	4.33333	0.85562	2.6303	6.0364
%sh	0.41	<.0001	HIGH DIST	40	0.4	0.63362	-0.8612	1.6612
			MOD DIST	18	4.55556	0.94455	2.6755	6.4356
			REF	24	7.875	0.818	6.2468	9.5032
%cf	0.03	0.3188	HIGH DIST	40	8.025	1.09	5.8555	10.195
			MOD DIST	18	8.61111	1.6248	5.377	11.845
			REF	24	5.70833	1.4071	2.9075	8.509
%pred	0.45	<.0001	HIGH DIST	40	4.9	0.60081	3.704	6.096
			MOD DIST	18	11.4444	0.89563	9.662	13.227
			REF	24	11.875	0.77564	10.331	13.419
%sc+sh	0.31	<.0001	HIGH DIST	40	2.8	1.0429	0.7242	4.876
			MOD DIST	18	10.3333	1.5547	7.2389	13.428
			REF	24	12.25	1.3464	9.5701	14.93
%sc+sh+pred	0.52	<.0001	HIGH DIST	40	7.775	1.2028	5.381	10.169
			MOD DIST	18	21.8889	1.793	18.32	25.458
			REF	24	24.125	1.5528	21.034	27.216
%cg+sc+sh	0.14	0.0027	HIGH DIST	40	42.8	3.1502	36.53	49.07
			MOD DIST	18	57.4444	4.6961	48.097	66.792
			REF	24	59.25	4.0669	51.155	67.345
%cg+cf	0.03	0.3524	HIGH DIST	40	48	3.0764	41.876	54.124
			MOD DIST	18	55.5556	4.5861	46.427	64.684
			REF	24	52.6667	3.9717	44.761	60.572
%cg+pred	0.12	0.0064	HIGH DIST	40	44.875	3.0481	38.808	50.942
			MOD DIST	18	58.6111	4.5439	49.567	67.655
			REF	24	59.0417	3.9351	51.209	66.874
%pred+sh	0.61	<.0001	HIGH DIST	40	5.325	0.8412	3.651	6.999
			MOD DIST	18	16.2222	1.2539	13.726	18.718
			REF	24	19.7917	1.0859	17.63	21.953
EPHEMEROPTERA	0.19	0.0003	HIGH DIST	40	59.05	7.919	43.288	74.81
			MOD DIST	18	99.333	11.805	75.837	122.83
			REF	24	110.083	10.223	89.734	130.43
Heptagenidae	0.41	<.0001	HIGH DIST	40	0.175	0.9138	-1.6439	1.994
			MOD DIST	18	3.1667	1.3623	0.4551	5.878
			REF	24	11.125	1.1798	8.7768	13.473
Leptophlebiidae	0.34	<.0001	HIGH DIST	40	0.825	2.383	-3.918	5.568
			MOD DIST	18	8.1667	3.5524	1.096	15.237
			REF	24	25.75	3.0764	19.626	31.874
PLECOPTERA	0.43	<.0001	HIGH DIST	40	0.025	1.2281	-2.419	2.469
			MOD DIST	18	1.8889	1.8308	-1.755	5.533
			REF	24	15.1667	1.5855	12.011	18.322
TRICOPTERA	0.31	<.0001	HIGH DIST	40	8.95	3.1743	2.632	15.268

Table A-5
Biological Metric Screening ANOVAs Summary

Parameter	r-square	p	Group	n	Mean	Std Error	Lower 95%	Upper 95%
			MOD DIST	18	36.2222	4.732	26.803	45.641
			REF	24	35.2083	4.098	27.051	43.365
COLEOPTERA	0.27	<.0001	HIGH DIST	40	9.875	2.5245	4.85	14.9
			MOD DIST	18	25.7222	3.7633	18.232	33.213
			REF	24	30.9583	3.2591	24.471	37.445
Elmidae	0.25	<.0001	HIGH DIST	40	1.9	1.6475	-1.379	5.179
			MOD DIST	18	6.5	2.4559	1.612	11.388
			REF	24	15.7083	2.1269	11.475	19.942
DIPTERA	0.22	<.0001	HIGH DIST	40	152.925	9.798	133.42	172.43
			MOD DIST	18	89.222	14.606	60.15	118.29
			REF	24	87.167	12.649	61.99	112.34
Chironomidae	0.25	<.0001	HIGH DIST	40	125.85	9.494	106.95	144.75
			MOD DIST	18	57.222	14.153	29.05	85.39
			REF	24	57	12.257	32.6	81.4
ODONATA	0.18	0.0005	HIGH DIST	40	2.075	0.6767	0.7281	3.4219
			MOD DIST	18	5.61111	1.0088	3.6032	7.619
			REF	24	6.20833	0.8736	4.4695	7.9472
EPHEMEROPTERA LOGY+1	0.28	<.0001	HIGH DIST	40	1.364	0.08338	1.198	1.53
			MOD DIST	18	1.97111	0.12429	1.7237	2.2185
			REF	24	2.02417	0.10764	1.8099	2.2384
Heptagenidae LOGY+1	0.56	<.0001	HIGH DIST	40	0.039	0.05235	-0.0652	0.1432
			MOD DIST	18	0.447778	0.07803	0.29245	0.6031
			REF	24	0.883333	0.06758	0.74882	1.0178
Leptophlebiidae LOGY+1	0.63	<.0001	HIGH DIST	40	0.1415	0.0587	0.0247	0.2583
			MOD DIST	18	0.79556	0.0875	0.6214	0.9697
			REF	24	1.22667	0.07578	1.0758	1.3775
PLECOPTERA LOGY+1	0.65	<.0001	HIGH DIST	40	0.0075	0.05026	-0.0925	0.1075
			MOD DIST	18	0.316667	0.07492	0.16755	0.4658
			REF	24	0.99375	0.06488	0.86461	1.1229
TRICOPTERA LOGY+1	0.45	<.0001	HIGH DIST	40	0.55275	0.07966	0.3942	0.7113
			MOD DIST	18	1.44	0.11875	1.2036	1.6764
			REF	24	1.46292	0.10284	1.2582	1.6676
COLEOPTERA LOGY+1	0.32	<.0001	HIGH DIST	40	0.84	0.06317	0.7143	0.9657
			MOD DIST	18	1.30167	0.09417	1.1142	1.4891
			REF	24	1.42042	0.08156	1.2581	1.5827
Elmidae LOGY+1	0.44	<.0001	HIGH DIST	40	0.25125	0.06131	0.12922	0.3733
			MOD DIST	18	0.67167	0.09139	0.48976	0.8536
			REF	24	1.03375	0.07915	0.87621	1.1913
DIPTERA LOGY+1	0.16	0.0011	HIGH DIST	40	2.11275	0.03683	2.0394	2.1861
			MOD DIST	18	1.915	0.0549	1.8057	2.0243
			REF	24	1.915	0.04754	1.8204	2.0096
Chironomidae LOGY+1	0.2	0.0001	HIGH DIST	40	1.99525	0.04928	1.8972	2.0933
			MOD DIST	18	1.69722	0.07346	1.551	1.8435
			REF	24	1.67958	0.06362	1.5529	1.8062
ODONATA LOGY+1	0.25	<.0001	HIGH DIST	40	0.33775	0.0547	0.22886	0.44664
			MOD DIST	18	0.683333	0.08155	0.52101	0.84565
			REF	24	0.751667	0.07062	0.61109	0.89224
E+P+T LOGY+1	0.33	<.0001	HIGH DIST	40	1.42475	0.08424	1.2571	1.5924
			MOD DIST	18	2.11556	0.12558	1.8656	2.3655
			REF	24	2.19417	0.10875	1.9777	2.4106
P+T LOGY+1	0.53	<.0001	HIGH DIST	40	0.55375	0.07802	0.3985	0.709
			MOD DIST	18	1.46167	0.1163	1.2302	1.6932
			REF	24	1.65333	0.10072	1.4529	1.8538
Hept+Leptophl LOGY+1	0.7	<.0001	HIGH DIST	40	0.161	0.05938	0.0428	0.2792
			MOD DIST	18	0.89278	0.08852	0.7166	1.069
			REF	24	1.44458	0.07666	1.292	1.5972

Table A-5
Biological Metric Screening ANOVAs Summary

Parameter	r-square	p	Group	n	Mean	Std Error	Lower 95%	Upper 95%
Hept.+Leptophl+Pleco LOGY+1	0.74	<.0001	HIGH DIST	40	0.1635	0.05995	0.0442	0.2828
			MOD DIST	18	0.94333	0.08937	0.7654	1.1212
			REF	24	1.61375	0.0774	1.4597	1.7678
Dipt+NIs LOGY+1	0.46	<.0001	HIGH DIST	40	2.3035	0.02736	2.249	2.358
			MOD DIST	18	2.06833	0.04079	1.9871	2.1495
			REF	24	1.95167	0.03533	1.8813	2.022
Dom. Taxon LOGY+1	0.41	<.0001	HIGH DIST	40	2.15375	0.02697	2.1001	2.2074
			MOD DIST	18	1.865	0.04021	1.785	1.945
			REF	24	1.88667	0.03482	1.8174	1.956
Two Dom. Taxa LOGY+1	0.45	<.0001	HIGH DIST	40	2.3085	0.0192	2.2703	2.3467
			MOD DIST	18	2.08944	0.02863	2.0325	2.1464
			REF	24	2.09208	0.02479	2.0427	2.1414

Note: standard error uses a pooled estimate of error variance
P-values 0.05 or less in **bold**.

Biological Parameters	r-square	p (whole model)	Regressor p-values, positive (+) or negative (-) relationship					
	Whole Model	Whole Model	order	elevation	wet width	temperature	pH	sp cond.
# BMI orders	0.28	0.4472	0.1858 (-)	0.9668 (+)	0.5554 (+)	0.2312 (+)	0.4815 (-)	0.7536 (-)
# BMI families	0.19	0.7121	0.5215 (+)	0.9104 (-)	0.3947 (+)	0.4092 (-)	0.5618 (-)	0.5250 (+)
# non-insect taxa	0.26	0.4903	0.5944 (-)	0.2517 (+)	0.2794 (+)	0.8999 (-)	0.8982 (-)	0.5403 (-)
# insect orders	<i>0.51</i>	<i>0.0512</i>	0.4354 (-)	<i>0.0884</i> (-)	0.4644 (-)	<i>0.0732</i> (+)	0.3221 (-)	0.6598 (+)
# insect families	0.28	0.4367	0.3087 (+)	0.643 (-)	0.4857 (+)	0.3158 (-)	0.5176 (-)	0.3592 (+)
# EPT families	0.56	0.0223	0.0842 (+)	0.4664 (-)	0.4133 (+)	0.0104 (-)	0.337 (-)	0.5749 (+)
% EPT	0.21	0.6404	0.7263 (+)	0.9306 (+)	0.9648 (-)	0.3518 (-)	0.6343 (+)	0.3346 (+)
% PT	0.43	0.1244	0.3647 (+)	0.4864 (-)	0.4967 (-)	0.1711 (-)	0.8474 (-)	<i>0.0546</i> (+)
% sens EPT	0.5	<i>0.0555</i>	0.8301 (-)	0.1166 (+)	<i>0.0519</i> (-)	0.2701 (-)	0.5675 (+)	0.1704 (+)
BI score	0.24	0.5658	0.9684 (-)	0.7619 (+)	0.342 (+)	0.417 (+)	0.737 (-)	0.3147 (-)
% sens BMIs	<i>0.47</i>	<i>0.0761</i>	0.7528 (-)	0.1815 (+)	<i>.0657</i> (-)	0.2879 (-)	0.5924 (+)	0.1942 (+)
% tol BMIs	0.44	0.1062	<i>0.0843</i> (+)	0.4407 (+)	0.9928 (-)	0.0303 (+)	0.4756 (+)	0.8634 (-)
% dominant taxa	<i>0.48</i>	<i>0.0727</i>	<i>0.0815</i> (-)	0.6817 (-)	0.7767 (+)	0.7779 (-)	0.1634 (-)	0.0408 (-)
% two dominant taxa	0.57	0.0207	0.0374 (-)	0.7388 (+)	0.5462 (+)	0.9193 (+)	0.2245 (-)	0.0235 (-)
% Diptera as Chironomidae	0.57	0.019	0.8838 (-)	<i>0.0654</i> (+)	0.4633 (-)	0.0104 (-)	<i>0.07</i> (-)	0.9024 (+)
% non-insect BMIs	0.39	0.1787	0.5103 (-)	0.0472 (+)	<i>0.0572</i> (+)	0.4707 (+)	0.8243 (-)	0.6896 (+)
% non-insects + diptera	0.33	0.3112	0.1214 (-)	0.7781 (-)	0.7848 (-)	0.5894 (-)	0.6387 (-)	0.1022 (-)
% non-insects+chironimidae	0.53	0.0386	0.1536 (-)	0.5136 (+)	0.4882 (-)	0.138 (-)	0.2016 (-)	0.1292 (-)
EPT/Chironomidae	0.27	0.456	0.5885 (+)	0.4382 (-)	0.6039 (+)	0.2634 (+)	0.3442 (+)	0.9713 (-)
%sh	0.37	0.2189	0.9132 (+)	0.8936 (+)	0.3833 (-)	0.3741 (-)	0.7316 (+)	<i>0.0741</i> (+)
%pred	0.3	0.3885	0.3058 (+)	0.1714 (-)	0.5802 (-)	0.6577 (+)	0.7207 (+)	0.8375 (-)
%sc+sh	0.44	0.1123	0.7727 (+)	0.9824 (-)	0.3826 (-)	0.1746 (-)	0.8234 (-)	0.034 (+)
%sc+sh+pred	0.4	0.1637	0.4285 (+)	0.5135 (-)	0.2802 (-)	0.3206 (-)	0.994 (-)	0.0639 (+)
%cg+sc+sh	0.25	0.5277	0.3242 (+)	0.6055 (+)	0.8241 (+)	0.4385 (+)	0.2457 (+)	0.3348 (+)
%cg+pred	0.38	0.1952	0.2601 (+)	0.8761 (+)	0.5523 (+)	0.1042 (+)	0.1607 (+)	0.76 (-)
%pred+sh	0.31	0.2176	0.5289 (+)	0.4835 (-)	0.2772 (-)	0.5163 (-)	0.539 (+)	0.1318 (+)
EPHEMEROPTERA	0.06	0.9802	0.8474 (-)	0.6458 (+)	0.7311 (+)	0.9365 (-)	0.5673 (+)	0.8699 (-)
Heptagenidae	0.59	0.0145	0.2663 (+)	<i>0.0604</i> (-)	0.1896 (-)	0.0100 (-)	0.4015 (+)	0.6462 (-)
Leptophlebiidae	0.63	0.0067	0.7873 (-)	0.0066 (+)	0.0207 (-)	0.8763 (-)	0.7631 (+)	0.6563 (+)
PLECOPTERA	0.34	0.2784	0.1974 (-)	<i>0.0667</i> (-)	0.7993 (-)	0.4771 (-)	0.854 (-)	0.8189 (+)
TRICOPTERA	0.32	0.3199	<i>0.0896</i> (+)	0.6369 (+)	0.5919 (-)	0.3063 (-)	0.9172 (-)	<i>0.0693</i> (+)
COLEOPTERA	0.49	0.0634	<i>0.0514</i> (+)	0.8700 (+)	0.4769 (+)	0.0258 (+)	0.7824 (+)	0.4665 (+)
Elmidae	0.54	0.0339	0.1339 (+)	0.9901 (-)	0.2024 (+)	0.0100 (+)	0.4375 (+)	0.8996 (+)
DIPTERA	0.38	0.2854	0.1407 (-)	0.5953 (-)	0.5708 (-)	0.4859 (-)	0.6265 (-)	<i>0.0828</i> (-)
Chironomidae	0.55	0.0287	0.1624 (-)	0.7166 (+)	0.3119 (-)	<i>0.0834</i> (-)	0.1760 (-)	<i>0.0915</i> (-)
ODONATA	0.11	0.9032	0.2365 (+)	0.4252 (+)	0.567 (-)	0.6312 (+)	0.8859 (+)	0.3576 (+)
EPHEMEROPTERA LOGY+1	0.1	0.9286	0.7886 (-)	0.6065 (+)	0.739 (+)	0.7796 (-)	0.4446 (+)	0.7733 (-)
Heptagenidae LOGY+1	0.67	0.0035	0.306 (+)	0.0124 (-)	0.5957 (-)	0.026 (-)	0.7979 (+)	0.1555 (+)
Leptophlebiidae LOGY+1	<i>0.45</i>	<i>0.0599</i>	0.3821 (-)	0.1465 (+)	<i>0.09</i> (-)	0.5949 (-)	0.9989 (-)	0.5094 (+)
PLECOPTERA LOGY+1	<i>0.48</i>	<i>0.0707</i>	0.1091 (-)	0.0047 (-)	0.5406 (-)	0.2404 (-)	0.5692 (-)	0.5452 (-)
TRICOPTERA LOGY+1	0.4	0.16	0.0344 (+)	0.2477 (+)	0.6955 (-)	0.4057 (-)	0.7855 (+)	0.035 (+)
COLEOPTERA LOGY+1	0.55	0.0273	<i>0.0745</i> (+)	0.4016 (-)	0.1766 (+)	<i>0.0599</i> (+)	0.9024 (-)	0.2053 (+)
Elmidae LOGY+1	0.62	0.0094	0.1681 (+)	0.9436 (-)	0.0177 (+)	0.0111 (+)	0.4679 (+)	0.6167 (+)
DIPTERA LOGY+1	0.29	0.4235	0.2653 (-)	0.9461 (-)	0.5505 (-)	0.4376 (-)	0.6392 (-)	0.183 (-)
Chironomidae LOGY+1	<i>0.48</i>	<i>0.0718</i>	0.3018 (-)	0.4288 (+)	0.4241 (-)	<i>0.0503</i> (-)	0.1928 (-)	0.4738 (-)
ODONATA LOGY+1	0.06	0.9823	0.4181 (+)	0.9502 (+)	0.8069 (-)	0.9637 (-)	0.9846 (+)	0.5892 (+)
E+P+T LOGY+1	0.24	0.5619	0.668 (+)	0.7895 (+)	0.9977 (-)	0.3368 (-)	0.6561 (+)	0.2469 (+)
P+T LOGY+1	<i>0.48</i>	<i>0.068</i>	0.2487 (+)	0.5533 (-)	0.655 (-)	0.2258 (-)	0.8171 (-)	0.0213 (+)
Hept+Leptopl LOGY+1	<i>0.45</i>	<i>0.0994</i>	0.6971 (-)	0.263 (+)	0.103 (-)	<i>0.0537</i> (-)	0.9971 (-)	0.5094 (+)
Hept.+Leptopl+Pleco LOGY+1	<i>0.46</i>	<i>0.0881</i>	0.2235 (-)	0.8984 (+)	0.1664 (-)	<i>0.0526</i> (-)	0.7982 (-)	0.7354 (+)
Dipt+Nis LOGY+1	0.27	0.4573	0.2627 (-)	0.8244 (+)	0.7763 (-)	0.5056 (-)	0.6331 (-)	0.2272 (-)
Dom. Taxon LOGY+1	0.42	0.1434	0.1646 (-)	0.7816 (-)	0.7313 (+)	0.6283 (-)	0.1497 (-)	<i>0.0801</i> (-)
Two Dom. Taxa LOGY+1	<i>0.49</i>	<i>0.0631</i>	<i>0.0707</i> (-)	0.825 (+)	0.3875 (+)	0.9153 (+)	0.2387 (-)	0.0468 (-)

Note: standard error uses a pooled estimate of error variance
P-values 0.05 or less in **bold**, p-values between 0.05 and 0.10 in *italics*.

Table A-7
IBI Scores and Classifications of Biological Integrity by Study Reach

Study Reach	2000		2001		2002		2003		No. years sampled	IBI Score Range	Classification Range
	IBI Score	Classification	IBI Score	Classification	IBI Score	Classification	IBI Score	Classification			
RIN1	NS	NS	NS	NS	40	Fair	NS	NS	1	NA	NA
C1	20	Very Poor	16	Very Poor	14	Very Poor	18	Very Poor	4	14-20	Very Poor
C2	42	Fair	18	Very Poor	NS	NS	NS	NS	2	18-42	Very Poor to Fair
C3	48	Good	44	Fair	56	Excellent	50	Good	4	44-56	Fair to Excellent
F1	12	Very Poor	NS	NS	NS	NS	NS	NS	1	NA	NA
SM1	14	Very Poor	NS	NS	NS	NS	NS	NS	1	NA	NA
MONT1	NS	NS	NS	NS	NS	NS	50	Good	1	NA	NA
SY1	NS	NS	NS	NS	18	Very Poor	32	Poor	2	18-32	Very Poor to Poor
SY2	NS	NS	NS	NS	NS	NS	32	Poor	1	NA	NA
M1	14	Very Poor	NS	NS	14	Very Poor	14	Very Poor	3	14	Very Poor
M2	NS	NS	NS	NS	14	Very Poor	14	Very Poor	2	14	Very Poor
M3	50	Good	NS	NS	48	Good	42	Fair	3	42-50	Fair to Good
M4	52	Good	NS	NS	NS	NS	NS	NS	1	NA	NA
M5	48	Good	NS	NS	NS	NS	NS	NS	1	NA	NA
AB1	NS	NS	NS	NS	22	Very Poor	26	Poor	2	22-26	Very Poor to Poor
AB2	26	Poor	18	Very Poor	18	Very Poor	24	Poor	4	18-26	Very Poor to Poor
AB3	44	Fair	44	Fair	44	Fair	36	Fair	4	36-44	Fair
AT1	NS	NS	16	Very Poor	12	Very Poor	12	Very Poor	3	12 to 16	Very Poor
AT2	NS	NS	18	Very Poor	16	Very Poor	24	Poor	3	18-24	Very Poor to Poor
SA1	36	Fair	NS	NS	NS	NS	NS	NS	1	NA	NA
SA2	40	Fair	NS	NS	NS	NS	52	Good	2	40-52	Fair to Good
MY1	40	Fair	NS	NS	NS	NS	NS	NS	1	NA	NA
MY2	42	Fair	NS	NS	NS	NS	NS	NS	1	NA	NA
MY3	50	Good	NS	NS	NS	NS	NS	NS	1	NA	NA
SJ1	22	Very Poor	24	Poor	22	Very Poor	28	Poor	4	22-28	Very Poor to Poor
SJ2	36	Fair	26	Poor	32	Poor	34	Poor	4	26-36	Poor to Fair
SJ3	48	Good	52	Good	48	Good	48	Good	4	48-52	Good
SJ4	42	Fair	NS	NS	NS	NS	NS	NS	1	NA	NA
T1	34	Poor	NS	NS	NS	NS	NS	NS	1	NA	NA
T2	42	Fair	NS	NS	NS	NS	NS	NS	1	NA	NA
DP1	44	Fair	NS	NS	NS	NS	NS	NS	1	NA	NA
EC1	NS	NS	NS	NS	37	Fair	NS	NS	1	NA	NA
R1	38	Fair	NS	NS	NS	NS	NS	NS	1	NA	NA
R2	50	Good	NS	NS	NS	NS	NS	NS	1	NA	NA
AH1	60	Excellent	36	Fair	54	Excellent	60	Excellent	4	36-60	Fair-Excellent
AH2	46	Fair	NS	NS	NS	NS	NS	NS	1	NA	NA
SO1	56	Excellent	NS	NS	NS	NS	NS	NS	1	NA	NA
SO2	52	Good	56	Excellent	58	Excellent	58	Excellent	4	52-58	Good to Excellent
GAV1	NS	NS	NS	NS	42	Fair	38	Fair	2	38-42	Fair
GAV2	NS	NS	NS	NS	38	Fair	NS	NS	1	NA	NA

Note: NS indicates the study reach was not sampled in that year. NA indicates not applicable, the study reach was sampled in one year only, there is no range.