

THE COMMUNITY SCIENCE INSTITUTE **Bio-monitoring RESULTS 2013**

	non-
	impacted
	slightly
	impacted
ſ	moderately
	impacted
ſ	severely
	impacted

Cayuga Lake Watershed

Seneca Lake Watershed

Susquehanna

Watershed

moderately impacted severely impacted	Total Family Richness	EPT Richness	Family Biotic Index	Percent Model Affinity	Density of organisms in sample	BAP Value Biological Assessment Profile
Enfield Creek - Mouth 5/4/13 42.398095, -76.546700 just upstrm of rte 13 bridge	18.5 no impact	9.5 no impact	4.57 slight impact	89% no impact	274	9.1 no impact
Cayuga Inlet 5/4/13 42.396910, -76.542953 upstream of confluence with Enfield Cr.	8.5 moderate impact	5.0 slight impact	3.39 no impact	73% no impact	1454	6.9 slight impact
W. Br. Cayuga Inlet-trib 4/24/13 42.359620, -76.598464 upstrm of Main St near Trumbulls Crnr Rd	9.5 moderate impact	8.0 no impact	3.01 no impact	52% slight impact	994	7.0 slight impact
Six Mile Creek-Slatvl600rd 9/1/13 42.396300N, 76.334333W Slaterville 600 Rd.	19.0 no impact	10.0 no impact	4.94 slight impact	66% no impact	analyzed live	8.5 no impact
Six Mile Creek -upstm GerCrs 9/1/13 42.402909 N, 76.436369 W German Cross Rd	11.0 slight impact	7.0 slight impact	5.26 slight impact	69% no impact	analyzed live	6.9 slight impact
Six Mile Creek-blw Potters Falls 8/28/13 42.419465N, 76.465312W first good riffle dwnstrm of Potters Falls	12.5 slight impact	5.5 slight impact	4.41 no impact	53% moderate impact	1022 analyzed live	6.7 slight impact
Six Mile Creek-Plain Street 8/31/13 42.434082, -76.504191 just downstream of Plain St. Bridge	9.0 moderate impact	5.5 slight impact	3.84 no impact	58% slight impact	analyzed live	6.5 slight impact
Fall Creek 10/16/13 42.457884, -76.437961 Freese Rd	11.5 slight impact	7.5	4.34 no impact	68%	1220	7.5 no/slight impct
Cascadilla Creek 6/15/13 42.447921, -76.501958 Hancock Bridge	Tier 2 Protocols Used	6.5 slight impact	jor Group Biotic Ind 5.80 moderate impact	41% moderate impact	1392	Tier 2 Protocols Used
Trumansburg Creek 6/29/13 42.541258N, 76.650769W King St (upstream of wstwtr trtmnt)	11.5 slight impact	7.5	4.60 slight impact	79%	3562	7.5 no/slight impet
Trumansburg Creek 6/28/13 42.540714N, 76.649079W just dwnstrm of Tburg wstwtr trtmnt	8.5 moderate impact	3.5 slight impact	5.60 moderate impact	44% moderate impact	6292	4.7 moderate impact
Mill Creek 5/3/13 42.635728, -76.634322 Atwater Rd, West of Lake Rd	1.5 severe impact	O severe impact	6.00 moderate impact	21% severe impact	606	1.3 severe impact
Tug Hollow (Logan) Creek 5/12/13 42.423581N, 76.852775W upstream of CR 5 bridge	14.0 no impact	9.0 no impact	3.83 no impact	78% no impact	1554	8.6
Wylie Brook 4/29/13 42.19253N, 75.61975W Off of Rte 235	14.0 no impact	9.0 no impact	3.37	67%	430	8.4
Danby Creek 8/24/13 42.32857 -76.426048 southernmost site, Halton property	14.0 no impact	6.5 slight impact	4.68 slight impact	63% slight impact	900	7.4 slight impact
****Bumps Creek 9/21/13 42.23035N, 75.53273W	12.0 slight impact	9.0 no impact	3.50 no impact	71% no impact		8.1***

Samples collected and analyzed using Hudson Basin River Watch Tier 3 Protocols (www.hudsonbasin.org) unless otherwise noted.

BAP is a composite index of aquatic habitat quality that incorporates Total Family Richness, Family Biotic Index, EPT Richness and Percent Model Affinity.

^{*} Note on Data for W. Branch Cayuga Inlet Site: Probably not all organisms were pulled out of the sub-samples for the two replicates. Sorting for each rep was done during a 45 minute period by Newfield HS students and may not be totally complete. Family Biotic Index and Percent Model Affinity (and subsequently BAP) may be off as a result.

^{**} Note: Physical habitat may not have been ideal for sampling. It is difficult to tell if this affected the results. Volunteers plan to sample a nearby, more ideal location this year to evaluate stream health.

^{***} Analysis of the two replicate samples for this site were not adequately similar

^{****} These metrics are based on data from only one of the two replicate sampes since the other replicate only had 59 organisms identified (metrics are based on 100 organism sub-sample)