

**Appendix D1: Quantitative  
macroinvertebrate data for streams  
undermined and sampled during the 3<sup>rd</sup>  
assessment period and re-sampled by the  
University**

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Bailey Mine

Stream Name: UNT to Wharton Run, 32507

Stream NHD#: \_\_\_\_\_

Sample Date: 4/16/2013

Pre-Mining Sampling Survey: 1 or 2 (check one)

Post-Mining Sampling Survey: 1 or X2 (check one)

Length of Sampled Reach: 100 meters

Sampler(s): G. Noble, A. Hale, K. Garmire

Comments: \_\_\_\_\_

Starting Lat/Long: 39.90799° / -80.51289°

Ending Lat/Long: 39.90848° / -80.51389°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats									
<del>XXXXXXXXXX</del>					Number of jabs				
Cobble / Gravel Substrate					2				
Snag					2				
Coarse Particulate Organic Matter					2				
Submerged Aquatic Vegetation					2				
Sand / Fine Sediment					2				
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)									
Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____			
F	G	F	G	F	G	F	G	F	G
	35								
	15								
	1								
	8								
	38								
	1								
	3								
	18								
	13								
	1								
	2								
	1								
	7								
	22								
	2								
	3								
	1								
	1								
	10								
	3								
	1								
Total Number of Individuals:		32	145						
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)									

Class or Order:	Family:	Genus:	Functional Feeding Group	Pollution Tolerance Value	Voltine Status (M) Multi, (U) uni, (S) semi			
Ephemeroptera	Ameletidae	Ameletus	CG	0				
Ephemeroptera	Siphonuridae	Siphonurus	CG	7				
Plecoptera	Nemouridae	Podmosta	SH	2				
Plecoptera	Nemouridae	Amphinemura	SH	3				
Plecoptera	Perlodidae	Isoperla	PR	2				
Plecoptera	Capniidae	Allocaenia	SH	3				
Trichoptera	Rhyacophilidae	Rhyacophila	PR	1				
Trichoptera	Uenoidae	Neophylax	SC	3				
Trichoptera	Limnephilidae	Ironoquia	SH	3				
Trichoptera	Lepidostomatidae	Lepidostoma	SH	1				
Coleoptera	Psephenidae	Ectopria	SC	5				
Diptera	Tabanidae	Tabanus	PR	5				
Diptera	Tipulidae	Pilaria	PR	7				
Diptera	Chironomidae		CG	6				
Diptera	Ceratopogonidae		PR	6				
Bivalvia	Sphaeriidae		FC	8				
Odonata	Cordulegastridae	Cordulegaster	PR	3				
Megaloptera	Corydalidae	Nigronia	PR	1				
Oligochaeta	Oligochaeta		CG	10				
Decapoda	Cambaridae		CR	6				
Gastropoda	Physidae		SC	8				

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Bailey Mine Stream Name: UNT to Wharton Run

Stream ID#: 32507 Segment ID: \_\_\_\_\_

Sampler(s): A. Hale, J. Phillips, C. Fisher (sample 1); Length of Sampled Reach: 100 meters  
G. Noble, A. Hale, K. Garmire (sample 2)

Pre-Mining

Post-Mining

Score 1 - Sample Date: 4/26/2009

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	15	49.2	49.2
Trichoptera Richness	4	38.1	38.1
Percent EPT Richness	66.7	108.3	100.0
Intolerant Taxa Richness	9	56.3	56.3
FC + PR Taxa Richness	5	37.0	37.0
Total Biological Score 1 (Mean of adjusted values):			56.1

Score 2 - Sample Date: 4/16/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	21	68.9	68.9
Trichoptera Richness	4	38.1	38.1
Percent EPT Richness	47.6	77.3	77.2
Intolerant Taxa Richness	11	68.8	68.8
FC + PR Taxa Richness	8	59.3	59.3
Total Biological Score 2 (Mean of adjusted values):			62.4

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \*10.7 %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): 59.3

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Bailey Mine  
 Stream Name: UNT to Dunkard Fork, 32532  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 4/23/13  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: 1 or X2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): T. Hann, A. Hale

Starting Lat/Long: 39.899957° / -80.492992°  
 Ending Lat/Long: 39.899157° / -80.493253°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats

	Number of jabs
<input checked="" type="checkbox"/>	
Cobble / Gravel Substrate	2
Snag	2
Coarse Particulate Organic Matter	2
Submerged Aquatic Vegetation	2
Sand / Fine Sediment	2

Comments: \_\_\_\_\_

<u>Class or Order:</u>	<u>Family:</u>	<u>Genus:</u>	<u>Functional Feeding Group</u>	<u>Pollution Tolerance Value</u>	Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)										
Voltine Status (M) Multi, (U) uni, (S) semi	Voltine Status (M) Multi, (U) uni, (S) semi				Sub. 1 - 4	Sub. _____	Sub. _____	Sub. _____	F	G	F	G	F	G	
Ephemeroptera	Ephemerellidae	Ephemerella	CG	1				2							
Ephemeroptera	Ephemerellidae	Eurylophella	SC	4				8							
Ephemeroptera	Ameletidae	Ameletus	CG	0				32							
Plecoptera	Perlodidae	Cultus	PR	2				1							
Plecoptera	Nemouridae	Amphinemura	SH	3				98							
Plecoptera	Perlodidae	Isoperla	PR	2				3							
Plecoptera	Perlodidae	Clioperla	PR	2				1							
Plecoptera	Perlidae	Unidentified	PR	3				1							
Trichoptera	Rhyacophilidae	Rhyacophila	PR	1				6							
Coleoptera	Elmidae	Optioservus	FC	4				1							
Coleoptera	Elmidae	Stenelmis	SC	5				2							
Diptera	Empididae	Clinocera	PR	6				9							
Diptera	Chironomidae		CG	6				13							
Diptera	Simuliidae	Prosimulium	FC	5				6							
Diptera	Simuliidae	Simulium	FC	6				4							
Bivalvia	Sphaeriidae		FC	8				1							
Gastropoda	Physidae		SC	8				2							
Oligochaeta	Oligochaeta		CG	10				2							
Total Number of Individuals:					18	174									

Lab sub-sample 1-4 (200 +/- 20%)  
 (Continue to sub-sample if numbers are <160 or >240.)

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Bailey Mine Stream Name: UNT to Dunkard Fork

Stream ID#: 32532 Segment ID: \_\_\_\_\_

Sampler(s): A. Hale, J. Phillips, C. Fisher, Length of Sampled Reach: 100 meters  
A. Glassmire (sample 1); T. Hann, A. Hale (sample 2)

Pre-Mining

Post-Mining

Score 1 - Sample Date: 5/5/2009

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	24	78.7	78.7
Trichoptera Richness	5	47.6	47.6
Percent EPT Richness	66.7	108.3	100.0
Intolerant Taxa Richness	14	87.5	87.5
FC + PR Taxa Richness	7	51.9	51.9
Total Biological Score 1 (Mean of adjusted values):			73.1

Score 2 - Sample Date: 4/16/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	18	59.0	59.0
Trichoptera Richness	1	9.5	9.5
Percent EPT Richness	50	81.17	81.2
Intolerant Taxa Richness	10	62.5	62.5
FC + PR Taxa Richness	10	74.1	74.1
Total Biological Score 2 (Mean of adjusted values):			57.3

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \*24.4 %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): 65.2

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Enlow Fork Mine  
 Stream Name: UNT to Templeton Fork, 32740  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 5/9/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: 1 or X2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A. Hale, G. Noble, K. Piper, L. Kiefer  
 Comments: \_\_\_\_\_

Starting Lat/Long: 40.05283° / -80.389657°  
 Ending Lat/Long: 40.053718° / -80.389913°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats									
					Number of jabs				
Cobble / Gravel Substrate					3				
Snag					1				
Coarse Particulate Organic Matter					2				
Submerged Aquatic Vegetation					2				
Sand / Fine Sediment					2				
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)									
Sub. 1 - 4		Sub. _____		Sub. _____		Sub. _____			
F	G	F	G	F	G	F	G	F	G
	10								
	18								
	1								
	1								
	1								
	1								
	28								
	36								
	3								
	2								
	3								
	4								
	1								
	1								
	1								
	1								
	1								
	2								
	3								
	2								
	6								
	8								
	53								
	3								
	6								
Total Number of Individuals:		70	122						
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)									

Class or Order: Voltine Status (M) Multi, (U) uni, (S) semi	Family: Voltine Status (M) Multi, (U) uni, (S) semi	Genus:	Functional Feeding Group	Pollution Tolerance Value
Ephemeroptera	Ameletidae	Ameletus	CG	0
Ephemeroptera	Baetidae	Baetis	CG	6
Ephemeroptera	Baetidae	Centroptilium	CG	2
Ephemeroptera	Siphonuridae	Siphonurus	CG	7
Ephemeroptera	Ephemerellidae	Eurylophella	SC	4
Plecoptera	Chloroperlidae	Haploperla	PR	0
Plecoptera	Nemouridae	Amphinemura	SH	3
Plecoptera	Perlodidae	Isoperla	PR	2
Plecoptera	Perlidae	Perlesta	PR	4
Plecoptera	Leuctridae	Leuctra	SH	0
Trichoptera	Rhyacophilidae	Rhyacophila	PR	1
Trichoptera	Hydropsychidae	Diplectrona	FC	0
Trichoptera	Hydroptilidae	Ochrotrichia	SC	4
Trichoptera	Glossomatidae	Agapetus	SC	0
Trichoptera	Uenoidae	Neophylax	SC	3
Coleoptera	Elmidae	Optioservus	SC	4
Coleoptera	Elmidae	Stenelmis	SC	5
Coleoptera	Psephenidae	Psephenus	SC	4
Diptera	Tipulidae	Tipula	SH	4
Diptera	Tipulidae	Hexatoma	PR	2
Diptera	Tipulidae	Pseudolimnophila	PR	2
Diptera	Ephydriidae		PI	6
Diptera	Ceratopogonidae		PR	6
Diptera	Chironomidae		CG	6
Decapoda	Cambaridae		CG	6
Oligochaeta			CG	10

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Enlow Fork Mine Stream Name: UNT to Templeton Fork

Stream ID#: 32740 Segment ID: \_\_\_\_\_

Sampler(s): A. Hale, S. Tonsor, J. Phillips (sample 1); Length of Sampled Reach: 100 meters  
A. Hale, G. Noble, K. Piper, L. Kiefer (sample 2)

Pre-Mining

Post-Mining

Score 1 - Sample Date: 5/29/2009

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	16	52.5	52.5
Trichoptera Richness	4	38.1	38.1
Percent EPT Richness	68.8	111.7	100.0
Intolerant Taxa Richness	12	75.0	75.0
FC + PR Taxa Richness	4	29.6	29.6
Total Biological Score 1 (Mean of adjusted values):			59.0

Score 2 - Sample Date: 5/9/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	26	85.2	85.2
Trichoptera Richness	5	47.6	47.6
Percent EPT Richness	57.7	93.7	93.7
Intolerant Taxa Richness	18	112.5	100.0
FC + PR Taxa Richness	8	59.3	59.3
Total Biological Score 2 (Mean of adjusted values):			77.2

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \*26.6 %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): 68.1

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

### FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Cumberland Mine  
 Stream Name: Dyers Fork, 41261, DF STA 2  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 5/7/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: 1 or X2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A. Hale, G. Noble, K. Piper, L. Kiefer  
 Comments: \_\_\_\_\_

Starting Lat/Long: 39.829797, -80.122747  
 Ending Lat/Long: 39.830363, -80.123902

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats									
					Number of jabs				
<del>Cobble / Gravel Substrate</del>					4				
Snag					0				
Coarse Particulate Organic Matter					0				
Submerged Aquatic Vegetation					4				
Sand / Fine Sediment					2				
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)									
Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____			
F	G	F	G	F	G	F	G	F	G
	1								
	1								
	2								
	6								
	43								
	3								
	23								
	1								
	1								
	139								
	1								
	4								
Total Number of Individuals:		145	80						
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)									

Class or Order:		Family:		Genus:		Functional Feeding Group	Pollution Tolerance Value		
Voltine Status (M) Multi, (U) uni, (S) semi		Voltine Status (M) Multi, (U) uni, (S) semi							
Ephemeroptera		Caenidae		Caenis		CG	7		
Ephemeroptera		Baetidae		Baetis		CG	6		
Plecoptera		Perlodidae		Isoperla		PR	2		
Plecoptera		Perlidae		Perlesta		PR	4		
Trichoptera		Hydroptilidae		Ochrotrichia		SC	4		
Coleoptera		Elmidae		Dubiraphia		SC	6		
Coleoptera		Elmidae		Stenelmis		SC	5		
Coleoptera		Elmidae		Optioservus		SC	4		
Diptera		Ceratopogonidae				PR	6	1	
Diptera		Chironomidae				CG	6	139	
Bivalvia		Sphaeriidae				FC	6	1	
Oligochaeta		Oligochaeta				CG	10	4	



### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Cumberland Mine Stream Name: Dyers Fork

Stream ID#: 41261 Segment ID: DF STA 2

Sampler(s): A. Hale, S. Tonsor, J. Phillips (sample 1); Length of Sampled Reach: 100 meters  
A. Hale, G. Noble, K. Piper, L. Kiefer (sample 2)

Pre-Mining  Post-Mining

Score 1 - Sample Date: 5/12/2009

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	21	68.9	68.9
Trichoptera Richness	2	19.0	19.0
Percent EPT Richness	28.6	46.4	46.4
Intolerant Taxa Richness	9	56.3	56.3
FC + PR Taxa Richness	7	51.9	51.9
Total Biological Score 1 (Mean of adjusted values):			48.5

Score 2 - Sample Date: 5/7/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	12	39.3	39.3
Trichoptera Richness	1	9.5	9.5
Percent EPT Richness	41.7	67.6	67.6
Intolerant Taxa Richness	4	25.0	25.0
FC + PR Taxa Richness	3	22.2	22.2
Total Biological Score 2 (Mean of adjusted values):			32.7

- 1.) Quality Assurance Check (% difference between Score 1 and Score 2): \*38.7 %
- 2.) Mean Total Biological Score (Average of Score 1 and Score 2): 40.6

\*If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Cumberland Mine  
 Stream Name: Dutch Run, 41246, DR STA 5  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 5/1/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: 1 or X2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A.Hale, K.Piper, G.Noble, S. Iannacchione  
 Comments: \_\_\_\_\_

Starting Lat/Long: 39.817813° / -80.089375°  
 Ending Lat/Long: 39.818703° / -80.089533°

Composite of 10 jobs from 10 sampling locations that effectively represents the observed habitats											
Number of jobs											
Cobble / Gravel Substrate						3					
Snag						3					
Coarse Particulate Organic Matter						0					
Submerged Aquatic Vegetation						2					
Sand / Fine Sediment						2					
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)											
Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____					
F	G	F	G	F	G	F	G	F	G	G	G
1											
	4										
	16										
	17										
	22										
	3										
	2										
	1										
3											
104											
4											
<b>Total Number of Individuals:</b>						112	65				
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)											

Class or Order: Voltine Status (M) Multi, (U) uni, (S) semi	Family: Voltine Status (M) Multi, (U) uni, (S) semi	Genus:	Functional Feeding Group	Pollution Tolerance Value
Ephemeroptera	Leptophlebiidae		CG	4
Plecoptera	Nemouridae	Amphinemura	SH	3
Plecoptera	Perlidae	Perlesta	PR	4
Plecoptera	Perlodidae	Isoperla	PR	2
Trichoptera	Hydroptilidae	Ochrotrichia	SC	4
Coleoptera	Elmidae	Stenelmis	SC	5
Diptera	Ephydriidae		PI	6
Diptera	Tipulidae	Molophilus	SH	4
Diptera	Ceratopogonidae		PR	6
Diptera	Chironomidae		CG	6
Oligochaeta	Oligochaeta		CG	10

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Cumberland Mine Stream Name: Dutch Run

Stream ID#: 41246 Segment ID: DR STA 5

Sampler(s): S. Tonsor, J. Phillips (sample 1); Length of Sampled Reach: 100 meters  
A. Hale, G. Noble, K. Piper, S. Iannacchione (sample 2)

Pre-Mining

Post-Mining

Score 1 - Sample Date: 5/15/2009

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	20	65.6	65.6
Trichoptera Richness	1	9.5	9.5
Percent EPT Richness	30	48.7	48.7
Intolerant Taxa Richness	7	43.8	43.8
FC + PR Taxa Richness	7	51.9	51.9
Total Biological Score 1 (Mean of adjusted values):			43.9

Score 2 - Sample Date: 5/1/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	11	36.1	36.1
Trichoptera Richness	1	9.5	9.5
Percent EPT Richness	45.5	73.9	73.9
Intolerant Taxa Richness	6	37.5	37.5
FC + PR Taxa Richness	3	22.2	22.2
Total Biological Score 2 (Mean of adjusted values):			35.8

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \*20.2 %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): 39.9

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.



**Appendix D2: Quantitative  
macroinvertebrate data for streams  
undermined during the 4<sup>th</sup> assessment  
period and sampled by the University**

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Bailey Mine  
 Stream Name: Strawn Hollow, 32547- BSW39  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 4/18/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A. Hale, K. Garmire  
 Comments: \_\_\_\_\_

Starting Lat/Long: 39.8653° / -80.48559°  
 Ending Lat/Long: 39.86501° / -80.48651°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats									
					Number of jabs				
Cobble / Gravel Substrate					2				
Snag					2				
Coarse Particulate Organic Matter					2				
Submerged Aquatic Vegetation					2				
Sand / Fine Sediment					2				
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)									
Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____			
F	G	F	G	F	G	F	G	F	G
	51								
	1								
	25								
	23								
	3								
	6								
	1								
	2								
	6								
	1								
	1								
	1								
	1								
	2								
	1								
	1								
	4								
	6								
	1								
	4								
	2								
	35								
	1								
Total Number of Individuals:		36	131						
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)									

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Bailey Mine Stream Name: Strawn Hollow

Stream ID#: 32547 Segment ID: BSW39

Sampler(s): A. Hale, K. Garmire Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 4/18/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	19	62.3	62.3
Trichoptera Richness	3	28.6	28.6
Percent EPT Richness	52.6	85.4	85.4
Intolerant Taxa Richness	11	68.8	68.8
FC + PR Taxa Richness	7	51.9	51.9
Total Biological Score 1 (Mean of adjusted values):			59.4

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

**FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY**

Mine Name: Blacksville #2  
 Stream Name: Blockhouse Run, 41812, BSW23  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 4/25/13  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A. Hale, K. Garmire  
 Comments: \_\_\_\_\_

Starting Lat/Long: 39.77347° / -80.34307°  
 Ending Lat/Long: 39.77438° / -80.34301°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats									
								Number of jabs	
Cobble / Gravel Substrate								2	
Snag								2	
Coarse Particulate Organic Matter								2	
Submerged Aquatic Vegetation								2	
Sand / Fine Sediment								2	
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)									
Sub. 1 - 6		Sub. ____		Sub. ____		Sub. ____			
F	G	F	G	F	G	F	G		
		Total Number of Individuals:		189		28			
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)									

<u>Class or Order:</u> Voltine Status (M) Multi, (U) uni, (S) semi	<u>Family:</u> Voltine Status (M) Multi, (U) uni, (S) semi	<u>Genus:</u>	Functional Feeding Group	Pollution Tolerance Value
Ephemeroptera	Caenidae	Caenis	CG	7
Ephemeroptera	Baetidae	Acentrella	SC	4
Plecoptera	Perlidae		PR	3
Plecoptera	Perlodidae	Isoperla	PR	2
Plecoptera	Nemouridae	Amphinemura	SH	3
Trichoptera	Uenoidae	Neophylax	SC	3
Trichoptera	Limnephilidae	Anabolia	SH	5
Trichoptera	Hydroptilidae	Ochrotrichia	SC	4
Amphipoda	Hyalellidae	Hyallega	CG	8
Coleoptera	Elmidae	Optioservus	SC	4
Coleoptera	Elmidae	Dubiraphia	SC	6
Coleoptera	Elmidae	Stenelmis	SC	5
Odonata	Calopterygidae	Calopteryx	PR	6
Diptera	Chironomidae		CG	6
Bivalvia	Sphaeriidae		FC	8
Oligochaeta	Oligochaeta		CG	10



### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Blacksville #2 Stream Name: Blockhouse Run

Stream ID#: 41812 Segment ID: BSW23

Sampler(s): A. Hale, K. Garmire Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	16	52.5	52.5
Trichoptera Richness	3	28.6	28.6
Percent EPT Richness	50	81.2	81.2
Intolerant Taxa Richness	7	43.8	43.8
FC + PR Taxa Richness	4	29.6	29.6
Total Biological Score 1 (Mean of adjusted values):			47.1

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Blacksville #2  
 Stream Name: Roberts Run, 41813, BSW22  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 4/25/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A. Hale, K. Garmire  
 Comments: \_\_\_\_\_

Starting Lat/Long: 39.77194° / -80.36198°  
 Ending Lat/Long: 39.77248° / -80.36292°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats										
					Number of jabs					
Cobble / Gravel Substrate					3					
Snag					2					
Coarse Particulate Organic Matter					2					
Submerged Aquatic Vegetation					0					
Sand / Fine Sediment					3					
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)										
Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____				
F	G	F	G	F	G	F	G	F	G	
	2									
	3									
	2									
	2									
	5									
	3									
	1									
	3									
	49									
	3									
	31									
	2									
	1									
	3									
	4									
	1									
	2									
	2									
	1									
	1									
	8									
	7									
	2									
	7									
	4									
	6	57								
	6	1								
Total Number of Individuals:					58	127				
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)										

Class or Order:	Family:	Genus:	Functional Feeding Group	Pollution Tolerance Value
Voltine Status (M) Multi, (U) uni, (S) semi	Voltine Status (M) Multi, (U) uni, (S) semi			
Ephemeroptera	Ameletidae	Ameletus	CG	0
Ephemeroptera	Baetidae	Baetis	CG	6
Ephemeroptera	Baetidae	Dipheter	CG	6
Ephemeroptera	Baetidae	Acentrella	SC	4
Ephemeroptera	Ephemerellidae	Ephemerella	CG	1
Ephemeroptera	Ephemerellidae	Eurylophella	SC	4
Plecoptera	Chloroperlidae	Sweltsa	PR	0
Plecoptera	Perlodidae	Diploperla	PR	2
Plecoptera	Perlodidae	Isoperla	PR	2
Plecoptera	Leuctridae	Leuctra	SH	0
Plecoptera	Nemouridae	Amphinemura	SH	3
Trichoptera	Hydropsychidae	Diplectrona	FC	0
Trichoptera	Lepidostomatidae	Lepidostoma	SH	1
Trichoptera	Limnephilidae	Pychnopsyche	SH	4
Trichoptera	Rhyacophilidae	Rhyacophila	PR	1
Trichoptera	Uenoidae	Neophylax	SC	3
Megaloptera	Sialidae	Sialis	PR	6
Coleoptera	Elmidae	Optioservus	SC	4
Coleoptera	Psephenidae	Ectopria	SC	5
Diptera	Stratiomyidae	Caloparyphus	CG	8
Diptera	Tabanidae	Chrysops	PI	7
Diptera	Tipulidae	Pseudolimnophila	PR	2
Diptera	Tipulidae	Pilaria	PR	7
Diptera	Tipulidae	Tipula	SH	4
Diptera	Chironomidae		CG	6
Decapoda	Cambaridae		CG	6

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Blacksville #2 Stream Name: Roberts Run

Stream ID#: 41813 Segment ID: BSW22

Sampler(s): A.Hale, K. Garmire Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 4/25/13

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	26	85.2	85.2
Trichoptera Richness	5	47.6	47.6
Percent EPT Richness	61.5	99.8	99.8
Intolerant Taxa Richness	17	106.3	100.0
FC + PR Taxa Richness	8	59.3	59.3
Total Biological Score 1 (Mean of adjusted values):			78.4

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

**FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY**

Mine Name: Cumberland Mine  
 Stream Name: Dyers Fork, 41261, DF STA 1  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 5/3/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A. Hale, S. Iannacchione, G. Noble, K. Piper  
 Comments: Pasture surrounds stream

Starting Lat/Long: 39.82233° / -80.11618°  
 Ending Lat/Long: 39.82252° / -80.11699°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats	
	Number of jabs
Cobble / Gravel Substrate	3
Snag	3
Coarse Particulate Organic Matter	0
Submerged Aquatic Vegetation	2
Sand / Fine Sediment	2
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)	
Sub. 1 - 4      Sub. ____      Sub. ____      Sub. ____	
F    G      F    G      F    G      F    G	

  

Class or Order:	Family:	Genus:	Functional Feeding Group	Pollution Tolerance Value	Sub. 1 - 4	Sub. ____	Sub. ____	Sub. ____
Voltine Status (M) Multi, (U) uni, (S) semi	Voltine Status (M) Multi, (U) uni, (S) semi				F	G	F	G
Ephemeroptera	Caenidae	Caenis	CG	7		1		
Ephemeroptera	Siphonuridae	Siphonurus	CG	7		1		
Ephemeroptera	Heptageniidae	Stenonema	SC	3		1		
Plecoptera	Perlidae	Perlesta	PR	4		5		
Plecoptera	Perlodidae	Isoperla	PR	2		17		
Coleoptera	Elmidae	Stenelmis	SC	5		2		
Coleoptera	Elmidae	Dubiraphia	SC	6		3		
Diptera	Culcidae	Anopheles	FC	8		1		
Diptera	Chironimidae		CG	6	118			
Diptera	Ceratopogoniidae		PR	6	1			
Bivalvia	Sphaeriidae		FC	8	1			
Oligochaeta	Oligochaeta		CG	10	26			
Total Number of Individuals:					146	31		
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)								

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Cumberland Mine Stream Name: Dyers Fork

Stream ID#: 41261 Segment ID: DF STA 1

Sampler(s): A. Hale, G. Noble, K. Piper, S. Iannacchione Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 5/3/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	12	39.3	39.3
Trichoptera Richness	0	0	0
Percent EPT Richness	41.7	67.7	67.7
Intolerant Taxa Richness	3	18.8	18.8
FC + PR Taxa Richness	5	37.0	37.0
Total Biological Score 1 (Mean of adjusted values):			32.6

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

**FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY**

Mine Name: Cumberland Mine  
 Stream Name: Dyers Fork, 41261, DF STA 21  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 5/3/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A. Hale, S. Iannacchione, G. Noble, K. Piper  
 Comments: \_\_\_\_\_

Starting Lat/Long: 39.81441° / -80.0991°  
 Ending Lat/Long: 39.814903° / -80.100013°

Composite of 10 jobs from 10 sampling locations that effectively represents the observed habitats	
Substrate	Number of jobs
Cobble / Gravel Substrate	3
Snag	1
Coarse Particulate Organic Matter	2

Submerged Aquatic Vegetation	2
Sand / Fine Sediment	2

Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)

Sub. 1 - 8		Sub. ____		Sub. ____		Sub. ____	
F	G	F	G	F	G	F	G

Class or Order: Voltine Status (M) Multi, (U) uni, (S) semi	Family: Voltine Status (M) Multi, (U) uni, (S) semi	Genus:	Functional Feeding Group	Pollution Tolerance Value
Ephemeroptera	Baetidae	Acentrella	SC	4
Plecoptera	Capniidae	Allocaenia	SH	3
Plecoptera	Nemouridae	Amphinemura	SH	3
Plecoptera	Perlidae	Perlesta	PR	4
Plecoptera	Perlodidae	Isoperla	PR	2
Trichoptera	Hydroptilidae	Ochrotrichia	SC	4
Coleoptera	Elmidae	Dubiraphia	SC	6
Coleoptera	Elmidae	Optioservus	SC	4
Coleoptera	Elmidae	Stenelmis	SC	5
Diptera	Ceratopogonidae		PR	6
Diptera	Chironomidae		CG	6
Gastropoda	Physidae		SC	8
Oligochaeta	Oligochaeta		CG	10

Total Number of Individuals:	75	80						
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Lab sub-sample 1-4 (200 +/- 20%)  
 (Continue to sub-sample if numbers are <160 or >240.)

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Cumberland Mine Stream Name: Dyers Fork

Stream ID#: 41261 Segment ID: DF STA 21

Sampler(s): A. Hale, S. Iannacchione, G. Noble, K. Piper Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	13	42.6	42.6
Trichoptera Richness	1	9.5	9.5
Percent EPT Richness	46.1	74.8	74.8
Intolerant Taxa Richness	7	43.8	43.8
FC + PR Taxa Richness	3	22.2	22.2
Total Biological Score 1 (Mean of adjusted values):			38.6

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Cumberland Mine  
 Stream Name: Maple Run, 40607, MR4  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 4/30/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A. Hale, G. Noble, K. Piper  
 Comments: \_\_\_\_\_

Starting Lat/Long: 39.82333° / -80.23994°  
 Ending Lat/Long: 39.82257° / -80.2405°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats										
					Number of jabs					
Cobble / Gravel Substrate					3					
Snag					2					
Coarse Particulate Organic Matter					1					
Submerged Aquatic Vegetation					2					
Sand / Fine Sediment					2					
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)										
Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____				
F	G	F	G	F	G	F	G	F	G	
	22									
	1									
	3									
	1									
	137									
	1									
	6									
	1									
	1									
	3									
	1									
	1									
	1									
	1									
	37									
	11									
Total Number of Individuals:					49	178				
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)										

Class or Order: Voltine Status (M) Multi, (U) uni, (S) semi	Family: Voltine Status (M) Multi, (U) uni, (S) semi	Genus:	Functional Feeding Group	Pollution Tolerance Value
Ephemeroptera	Ameletidae	Ameletus	CG	0
Ephemeroptera	Baetidae	Acentrella	SC	4
Ephemeroptera	Siphonuridae	Siphonurus	CG	7
Ephemeroptera	Leptophlebiidae	Paraleptophlebia	CG	1
Plecoptera	Perlodidae	Isoperla	PR	2
Plecoptera	Leuctridae	Leuctra	SH	0
Plecoptera	Nemouridae	Amphinemura	SH	3
Plecoptera	Perlidae	Perlesta	PR	4
Trichoptera	Limnephilidae	Anabolia	SH	5
Trichoptera	Rhyacophilidae	Rhyacophila	PR	1
Coleoptera	Elmidae	Stenelmis	SC	5
Diptera	Tabanidae	Chrysops	PI	7
Diptera	Ceratopogonidae		PR	6
Diptera	Chironomidae		CG	6
Oligochaeta			CG	10



### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Cumberland Mine Stream Name: Maple Run

Stream ID#: 40607 Segment ID: MR 4

Sampler(s): A. Hale, G. Noble, K. Piper Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 4/30/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	15	49.2	49.2
Trichoptera Richness	2	19.0	19.0
Percent EPT Richness	66.7	108.3	100.0
Intolerant Taxa Richness	8	50.0	50.0
FC + PR Taxa Richness	4	29.6	29.6
Total Biological Score 1 (Mean of adjusted values):			49.6

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Cumberland Mine  
 Stream Name: Maple Run, 40607, MR5  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 4/30/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): G. Noble, K. Piper, A. Hale  
 Comments: \_\_\_\_\_

Starting Lat/Long: 39.81926° / -80.24236°  
 Ending Lat/Long: 39.81831° / -80.2427°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats

	Number of jabs
Cobble / Gravel Substrate	4
Snag	1
Coarse Particulate Organic Matter	0
Submerged Aquatic Vegetation	3
Sand / Fine Sediment	2

Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)

Class or Order: Voltine Status (M) Multi, (U) uni, (S) semi	Family: Voltine Status (M) Multi, (U) uni, (S) semi	Genus:	Functional Feeding Group	Pollution Tolerance Value										
					Sub. 1 - 4				Sub. _____		Sub. _____			
					F	G	F	G	F	G	F	G		
Ephemeroptera	Ameletidae	Ameletus	CG	0		131								
Ephemeroptera	Ephemerellidae	Ephemerella	CG	1		3								
Ephemeroptera	Siphonuridae	Siphonurus	CG	7		15								
Plecoptera	Nemouridae	Amphinemura	SH	3		2								
Plecoptera	Perlodidae	Isoperla	PR	2		45								
Trichoptera	Limnephilidae	Anabolia	SH	5		1								
Trichoptera	Uenoidae	Neophylax	SC	3		1								
Diptera	Tipulidae	Tipula	SH	4		2								
Diptera	Chironomidae		CG	6	2									
Oligochaeta	Oligochaeta		CG	10	1									
<b>Total Number of Individuals:</b>					3	200								

Lab sub-sample 1-4 (200 +/- 20%)  
 (Continue to sub-sample if numbers are <160 or >240.)

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Cumberland Mine Stream Name: Maple Run

Stream ID#: 40607 Segment ID: MR5

Sampler(s): G. Noble, K. Piper, A. Hale Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 4-30-2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	10	32.8	32.8
Trichoptera Richness	2	19.0	19.0
Percent EPT Richness	70	113.6	100.0
Intolerant Taxa Richness	6	37.5	37.5
FC + PR Taxa Richness	1	7.4	7.4
Total Biological Score 1 (Mean of adjusted values):			39.3

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

- 1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %
- 2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Cumberland Mine

Stream Name: UNT to Maple Run, 40608, MR T12

Stream NHD#: \_\_\_\_\_

Sample Date: 3/8/2013

Pre-Mining Sampling Survey: 1 or 2 (check one)

Post-Mining Sampling Survey: X1 or 2 (check one)

Length of Sampled Reach: 100 meters

Sampler(s): G. Noble, L. Powell, T. Hann, K. Garmire, A. Hale

Comments: \_\_\_\_\_

Starting Lat/Long: 39.8199° / -80.24544°

Ending Lat/Long: 39.81978° / -80.2466°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats

	Number of jabs
Cobble / Gravel Substrate	3
Snag	3
Coarse Particulate Organic Matter	2
Submerged Aquatic Vegetation	0
Sand / Fine Sediment	2

Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)

Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____	
F	G	F	G	F	G	F	G

Class or Order:	Family:	Genus:	Functional Feeding Group	Pollution Tolerance Value
Voltine Status (M) Multi, (U) uni, (S) semi	Voltine Status (M) Multi, (U) uni, (S) semi			
Ephemeroptera	Ameletidae	Ameletus	CG	0
Ephemeroptera	Heptageniidae	Stenonema	SC	3
Ephemeroptera	Ephemerellidae	Ephemerella	CG	1
Ephemeroptera	Ephemerellidae	Eurylophella	SC	4
Plecoptera	Perlodidae	Isoperla	PR	2
Plecoptera	Nemouridae	Amphinemura	SH	3
Plecoptera	Nemouridae	Podmosta	SH	2
Plecoptera	Capniidae	Allocaenia	SH	3
Trichoptera	Rhyacophilidae	Rhyacophila	PR	1
Trichoptera	Uenoidae	Neophylax	SC	3
Diptera	Dixidae	Dixa	CG	1
Diptera	Simuliidae	Prosimulium	FC	5
Diptera	Tipulidae	Tipula	SH	4
Diptera	Tipulidae	Hexatoma	PR	2
Diptera	Tipulidae	Molophilus	SH	4
Diptera	Simulidae	Stegopterna	FC	6
Diptera	Chironomidae		CG	6
Diptera	Ceratopogonidae		PR	6
Oligochaeta	Oligochaeta		CG	10

Total Number of Individuals:	39	136					
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Lab sub-sample 1-4 (200 +/- 20%)  
(Continue to sub-sample if numbers are <160 or >240.)

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Cumberland Mine Stream Name: UNT to Maple Run

Stream ID#: 40608 Segment ID: MR T12

Sampler(s): G. Noble, L. Powell, T. Hann, K. Garmire, A. Hale Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 3/8/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	19	62.3	62.3
Trichoptera Richness	2	19.0	19.0
Percent EPT Richness	52.6	85.4	85.4
Intolerant Taxa Richness	14	87.5	87.5
FC + PR Taxa Richness	6	44.4	44.4
Total Biological Score 1 (Mean of adjusted values):			59.7

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Enlow Fork Mine

Stream Name: Templeton Fork, 32708, BSW19

Stream NHD#: \_\_\_\_\_

Sample Date: 5/9/2013

Pre-Mining Sampling Survey: 1 or 2 (check one)

Post-Mining Sampling Survey: X1 or 2 (check one)

Length of Sampled Reach: 100 meters

Sampler(s): A. Hale, G. Noble, K. Piper, L. Kiefer

Comments: \_\_\_\_\_

Starting Lat/Long: 40.06039° / -80.37785°

Ending Lat/Long: 40.06106° / -80.37704°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats									
					Number of jabs				
Cobble / Gravel Substrate					4				
Snag					0				
Coarse Particulate Organic Matter					0				
Submerged Aquatic Vegetation					4				
Sand / Fine Sediment					2				
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)									
Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____			
F	G	F	G	F	G	F	G	F	G
	2								
	6								
	2								
	16								
	2								
	3								
	1								
	1								
	4								
	1								
	1								
	2								
	183								
	1								
Total Number of Individuals:		186	40						
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)									

Class or Order: Voltine Status (M) Multi, (U) uni, (S) semi	Family: Voltine Status (M) Multi, (U) uni, (S) semi	Genus:	Functional Feeding Group	Pollution Tolerance Value
Ephemeroptera	Baetidae	Baetis	CG	6
Ephemeroptera	Caenidae	Caenis	CG	7
Plecoptera	Nemouridae	Amphinemura	SH	3
Plecoptera	Perlidae	Perlesta	PR	4
Plecoptera	Perlodidae	Isoperla	PR	2
Trichoptera	Hydropsychidae	Hydropsyche	FC	5
Odonata	Calopterygidae	Calopteryx	PR	6
Coleoptera	Elmidae	Dubiraphia	SC	6
Coleoptera	Elmidae	Stenelmis	SC	5
Coleoptera	Elmidae	Optioservus	SC	4
Diptera	Empididae	Hemerodromia	PR	6
Diptera	Tipulidae	Hexatoma	PR	2
Diptera	Ceratopogonidae		PR	6
Diptera	Chironomidae		CG	6
Gastropoda	Physidae		SC	8

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Enlow Fork Mine Stream Name: Templeton Fork

Stream ID#: 32708 Segment ID: BSW19

Sampler(s): A. Hale, G. Noble, K. Piper, L. Kiefer Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 5/9/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	15	49.2	49.2
Trichoptera Richness	1	9.5	9.5
Percent EPT Richness	40.0	64.9	64.9
Intolerant Taxa Richness	5	31.3	31.3
FC + PR Taxa Richness	7	51.9	51.9
Total Biological Score 1 (Mean of adjusted values):			41.3

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Enlow Fork Mine  
 Stream Name: Crafts Creek, 40938, BSW20  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 4/5/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): G. Noble, A. Hale, T. Hann, L. Powell  
 Comments: \_\_\_\_\_

Starting Lat/Long: 40.05588° / -80.33527°  
 Ending Lat/Long: 40.05523° / -80.33606°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats	
	Number of jabs
Cobble / Gravel Substrate	2
Snag	2
Coarse Particulate Organic Matter	2
Submerged Aquatic Vegetation	2
Sand / Fine Sediment	2

Class or Order:	Family:	Genus:	Functional Feeding Group	Pollution Tolerance Value	Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)													
					Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____							
					F	G	F	G	F	G	F	G						
Plecoptera	Nemouridae	Amphinemura	SH	3		1												
Plecoptera	Perlodidae	Isoperla	PR	2		7												
Trichoptera	Uenoidae	Neophylax	SC	3		2												
Trichoptera	Rhyacophilidae	Rhyacophila	PR	1		1												
Isopoda	Asellidae	Caecidotea	CG	6		2												
Coleoptera	Elmidae	Stenelmis	SC	5		3												
Diptera	Tabanidae	Chrysops	PI	7		2												
Diptera	Tipulidae	Pilaria	PR	7		1												
Diptera	Ceratopogonidae		PR	6	15													
Diptera	Chronomidae		CG	6	102													
Bivalvia	Sphaeriidae		FC	8	20													
Gastropoda	Physidae		SC	8	2													
Gastropoda	Planorbidae		SC	6	1													
Oligochaeta	Oligochaeta		CG	10	1													
<b>Total Number of Individuals:</b>					141	19												

Lab sub-sample 1-4 (200 +/- 20%)  
 (Continue to sub-sample if numbers are <160 or >240.)



### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Enlow Fork Mine Stream Name: Crafts Creek

Stream ID#: 40938 Segment ID: BSW20

Sampler(s): G. Noble, A. Hale, T. Hann, L. Powell Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 4/5/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	14	45.9	45.9
Trichoptera Richness	2	19.0	19.0
Percent EPT Richness	28.6	46.4	46.4
Intolerant Taxa Richness	4	25.0	25.0
FC + PR Taxa Richness	5	37.0	37.0
Total Biological Score 1 (Mean of adjusted values):			34.7

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

- 1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %
- 2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Enlow Fork Mine

Stream Name: UNT to Crafts Creek, 40941, BSW24

Stream NHD#: \_\_\_\_\_

Sample Date: 4/5/2013

Pre-Mining Sampling Survey: 1 or 2 (check one)

Post-Mining Sampling Survey: X1 or 2 (check one)

Length of Sampled Reach: 100 meters

Sampler(s): A. Hale, G. Noble, T. Hann, L. Powell

Comments: \_\_\_\_\_

Starting Lat/Long: 40.058445° / -80.334175°

Ending Lat/Long: 40.059292° / -80.334442°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats

	Number of jabs
Cobble / Gravel Substrate	2
Snag	3
Coarse Particulate Organic Matter	0
Submerged Aquatic Vegetation	2
Sand / Fine Sediment	3

Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)

Sub. 1 - 4		Sub. ____		Sub. ____		Sub. ____	
F	G	F	G	F	G	F	G

Class or Order:	Family:	Genus:	Functional Feeding Group	Pollution Tolerance Value
Voltine Status (M) Multi, (U) uni, (S) semi	Voltine Status (M) Multi, (U) uni, (S) semi			
Plecoptera	Nemouridae	Amphinemura	SH	3
Trichoptera	Uenoidae	Neophylax	SC	3
Trichoptera	Limnephilidae	Limnephilus	SH	3
Trichoptera	Lepidostomatidae	Lepidostoma	SH	1
Trichoptera	Rhyacophilidae	Rhyacophilia	PR	1
Amphipoda	Crangonyctidae	Crangonyx	CG	4
Diptera	Tipulidae	Molophilus	SH	4
Diptera	Tipulidae	Pseudolimnophila	PR	2
Diptera	Simuliidae	Stegopterna	FC	6
Diptera	Tabanidae	Tabanus	PR	5
Diptera	Tabanidae	Chrysops	PI	7
Diptera	Ceratopogonidae		PR	6
Diptera	Chironomidae		CG	6
Bivalvia	Sphaeriidae		FC	8
Insecta	Collembola		CG	9
Decapoda	Cambaridae		CG	6
Oligochaeta	Oligochaeta		CG	10

Total Number of Individuals:	34	136					
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Lab sub-sample 1-4 (200 +/- 20%)  
(Continue to sub-sample if numbers are <160 or >240.)

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Enlow Fork Mine Stream Name: UNT to Crafts Creek

Stream ID#: 40941 Segment ID: BSW24

Sampler(s): A. Hale, G. Noble, T. Hann, L. Powell Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 4/5/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	17	55.7	55.7
Trichoptera Richness	4	38.1	38.1
Percent EPT Richness	29.4	47.7	47.7
Intolerant Taxa Richness	8	50.0	50.0
FC + PR Taxa Richness	6	44.4	44.0
Total Biological Score 1 (Mean of adjusted values):			47.2

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

- 1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %
- 2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

**FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY**

Mine Name: Enlow Fork Mine  
 Stream Name: UNT to Crafts Creek, 40944, BSW13  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 3/29/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): T. Hann, G. Noble, A. Hale, K. Garmire  
 Comments: \_\_\_\_\_

Starting Lat/Long: 40.056469° / -80.355087°  
 Ending Lat/Long: 40.05727° / -80.355423°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats									
						Number of jabs			
Cobble / Gravel Substrate						2			
Snag						2			
Coarse Particulate Organic Matter						2			
Submerged Aquatic Vegetation						2			
Sand / Fine Sediment						2			
Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)									
Sub. 1 - 7		Sub. ____		Sub. ____		Sub. ____			
F	G	F	G	F	G	F	G	F	G
Total Number of Individuals:		61	128						
Lab sub-sample 1-4 (200 +/- 20%) (Continue to sub-sample if numbers are <160 or >240.)									

Class or Order: Voltine Status (M) Multi, (U) uni, (S) semi	Family: Voltine Status (M) Multi, (U) uni, (S) semi	Genus:	Functional Feeding Group	Pollution Tolerance Value	Sub. 1 - 7		Sub. ____		Sub. ____		Sub. ____	
					F	G	F	G	F	G	F	G
Ephemeroptera	Ameletidae	Ameletus	CG	0		14						
Plecoptera	Capniidae	Allocaenia	SH	3		2						
Plecoptera	Nemouridae	Amphinemura	SH	3		30						
Plecoptera	Nemouridae	Nemoura	SH	1		1						
Plecoptera	Perlodidae	Isoperla	PR	2		31						
Trichoptera	Limnephilidae	Ironoquia	SH	3		3						
Trichoptera	Rhyacophilidae	Rhyacophila	PR	1		1						
Trichoptera	Uenoidae	Neophylax	SC	3		2						
Amphipoda	Crangonyctidae	Crangonyx	CG	4		12						
Diptera	Simuliidae	Prosimulium	FC	5		15						
Diptera	Simuliidae	Stegopterna	FC	6		5						
Diptera	Tabanidae	Tabanus	PR	5		1						
Diptera	Tabanidae	Chrysops	PI	7		7						
Diptera	Tipulidae	Molophilus	SH	4		2						
Diptera	Tipulidae	Tipula	SH	4		2						
Diptera	Ceratopogonidae		PR	6		2						
Diptera	Chironomidae		CG	6		46						
Bivalvia	Sphaeriidae		FC	8		2						
Decapoda	Cambaridae		CG	6		1						
Oligochaeta	Oligochaeta		CG	10		10						

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Enlow Fork Mine Stream Name: UNT to Crafts Creek

Stream ID#: 40944 Segment ID: BSW13

Sampler(s): T. Hann, G. Noble, A. Hale, K. Garmire Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 3/29/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	20	65.6	65.6
Trichoptera Richness	3	28.6	28.6
Percent EPT Richness	40	64.9	64.9
Intolerant Taxa Richness	11	68.8	68.8
FC + PR Taxa Richness	7	51.9	51.9
Total Biological Score 1 (Mean of adjusted values):			55.9

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness			
Trichoptera Richness			
Percent EPT Richness			
Intolerant Taxa Richness			
FC + PR Taxa Richness			
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

## FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Emerald Mine  
 Stream Name: Mount Phoebe Run, 41268, MP STA 1  
 Stream NHD#: \_\_\_\_\_  
 Sample Date: 11/8/2013  
 Pre-Mining Sampling Survey: 1 or 2 (check one)  
 Post-Mining Sampling Survey: X1 or 2 (check one)  
 Length of Sampled Reach: 100 meters  
 Sampler(s): A. Hale, L. Kiefer  
 Comments: \_\_\_\_\_

Starting Lat/Long: 39.84277° / -80.12548°  
 Ending Lat/Long: 39.84368° / -80.12538°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats

Cobbles / Gravel Substrate		Number of jabs
	Cobble / Gravel Substrate	1
	Snag	3
	Coarse Particulate Organic Matter	1
	Submerged Aquatic Vegetation	2
	Sand / Fine Sediment	3

Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)

Sub. 1 - 4		Sub. _____		Sub. _____		Sub. _____	
F	G	F	G	F	G	F	G
<b>189</b>	<b>36</b>						

Lab sub-sample 1-4 (200 +/- 20%)  
(Continue to sub-sample if numbers are <160 or >240.)

Class or Order:	Family:	Genus:	Functional Feeding Group	Pollution Tolerance Value									
Voltine Status (M) Multi, (U) uni, (S) semi	Voltine Status (M) Multi, (U) uni, (S) semi	Genus:											
Ephemeroptera	Baetidae	Baetis	CG	6			3						
Plecoptera	Capniidae	Allocaenia	SH	3			1						
Trichoptera	Hydropsychidae	Cheumatopsyche	FC	6			12						
Trichoptera	Hydropsychidae	Hydropsyche	FC	5			4						
Coleoptera	Elmidae	Dubiraphia	SC	6			2						
Coleoptera	Elmidae	Optioservus	SC	4			1						
Coleoptera	Elmidae	Stenelmis	SC	5			8						
Odonata	Calopterygidae	Calopteryx	PR	6			1						
Diptera	Tipulidae	Tipula	SH	4			3						
Diptera	Tabanidae	Chrysops	PI	7			1						
Diptera	Chironomidae		CG	6	179								
Diptera	Ceratopogonidae		PR	6	6								
Gastropoda	Physidae		SC	8	1								
Oligochaeta	Oligochaeta		CG	10	3								

### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Emerald Mine Stream Name: Mount Phoebe Run

Stream ID#: 41268 Segment ID: MP STA 1

Sampler(s): A. Hale, L. Kiefer Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 11/8/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	14	45.9	45.9
Trichoptera Richness	2	19.0	19.0
Percent EPT Richness	28.6	46.4	46.4
Intolerant Taxa Richness	3	18.8	18.8
FC + PR Taxa Richness	4	29.6	29.6
Total Biological Score 1 (Mean of adjusted values):			32.0

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.

# FORM 8.8C: QUANTITATIVE MULTI-HABITAT BIOASSESSMENT OF DIVERSE COMMUNITY

Mine Name: Emerald Mine

Stream Name: Muddy Creek, 41014, MC B2

Stream NHD#: \_\_\_\_\_

Sample Date: 11/15/2013

Pre-Mining Sampling Survey: 1 or 2 (check one)

Post-Mining Sampling Survey: X1 or 2 (check one)

Length of Sampled Reach: 100 meters

Sampler(s): L. Kiefer, S. Tonsor, A. Hale

Comments: \_\_\_\_\_

Starting Lat/Long: 39.87832°/ -80.09304°

Ending Lat/Long: 39.877965°/ -80.09382°

Composite of 10 jabs from 10 sampling locations that effectively represents the observed habitats	
Substrate	Number of jabs
Cobble / Gravel Substrate	0
Snag	2
Coarse Particulate Organic Matter	2
Submerged Aquatic Vegetation	3
Sand / Fine Sediment	3

Class or Order: Voltine Status (M) Multi, (U) uni, (S) semi	Family: Voltine Status (M) Multi, (U) uni, (S) semi	Genus:	Functional Feeding Group	Pollution Tolerance Value	Enter the number of individuals for each Genus identified in lab. (F = Family / G = Genus)														
					Sub. 1 - 4		Sub. ___		Sub. ___		Sub. ___								
F	G	F	G	F	G	F	G	F	G										
Ephemeroptera	Baetidae	Callibaetis	CG	9		5													
Trichoptera	Phryganeidae	Ptilostomis	SH	5		4													
Trichoptera	Limnephilidae	Limnephilus	SH	3		1													
Coleoptera	Elmidae	Dubiraphia	SC	6		1													
Coleoptera	Elmidae	Optioservus	SC	4		1													
Odonata	Coenagrionidae	Argia	PR	6		1													
Megaloptera	Sialidae	Sialis	PR	6		1													
Diptera	Tipulidae	Hexatoma	PR	2		1													
Diptera	Tabanidae	Chrysops	PI	7		1													
Diptera	Ceratopogonidae		PR	6	1														
Diptera	Chironomidae		CG	6	124														
Bivalvia	Sphaeriidae		FC	8	6														
Gastropoda	Lymnaidae		SC	7	5														
Oligochaeta	Oligochaeta		CG	10	59														
Total Number of Individuals:					195	16													

Lab sub-sample 1-4 (200 +/- 20%)  
 (Continue to sub-sample if numbers are  
 <160 or >240.)



### FORM 8.8D: BIOMETRIC AND TOTAL BIOLOGICAL SCORE SUMMARY

Mine Name: Emerald Mine Stream Name: Muddy Creek

Stream ID#: 41014 Segment ID: MC B2

Sampler(s): L. Kiefer, S. Tonsor, A. Hale Length of Sampled Reach: 100 meters

Pre-Mining

Post-Mining

Score 1 - Sample Date: 11/15/2013

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness	14	45.9	45.9
Trichoptera Richness	2	19.0	19.0
Percent EPT Richness	21.4	34.7	34.7
Intolerant Taxa Richness	3	18.8	18.8
FC + PR Taxa Richness	5	37.0	37.0
Total Biological Score 1 (Mean of adjusted values):			31.1

Score 2 - Sample Date:

Biological Metric	Observed Value	Normalized Score (observed value / 95 <sup>th</sup> percentile value) * 100	Adjusted Value
Taxa Richness		<b>30.5</b>	
Trichoptera Richness		<b>10.5</b>	
Percent EPT Richness		<b>61.6</b>	
Intolerant Taxa Richness		<b>16.0</b>	
FC + PR Taxa Richness		<b>13.5</b>	
Total Biological Score 2 (Mean of adjusted values):			

1.) Quality Assurance Check (% difference between Score 1 and Score 2): \* \_\_\_\_\_ %

2.) Mean Total Biological Score (Average of Score 1 and Score 2): \_\_\_\_\_

\* If percentage difference is greater than 16%, reach should be re-sampled to obtain additional set of metrics.