FDEP Macroinvertebrate Taxonomic Internal QC Resolution

Sample #			
Rep #		Type	
Original Analysis		**QC Analysis**	
Analyst		QC Analyst	
ID Date		QC Date	

(1) None = If difference is no more than one taxon level.
(2) Level = If difference is >1 taxon level. (ie. Family to species)
(3) Mis-ID = If taxon is identified incorrect.
(4) Count = If count difference is > 2 individuals or 5% of original count
(5) New = If taxon is unaccounted for by original analyst (missed bugs).
(6) Unresolved = If no consensus (error will count against original analyst)

ID # Taxa		QC # Taxa		Resolution # Taxa		Taxa Error	0	**Precision Taxa**	**#DIV/0!**
ID # Ind	0	QC # Ind	0	Resolution # Ind	0	Ind Error	0	**Precision Ind**	**#DIV/0!**

OrigTaxonID	**Orig Count**	**QCTaxonID**	**QC Count**	**Resolution ID**	**Resol. Count**	**Resol. Type**	**Comments**	**SBIO chg'd**
Total # individuals	0	Total # individuals	0	Total # individuals	0			

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Precision	
Individuals	Taxa
#DIV/0!	#DIV/0!

Check count difference			Individuals Effected		0	0
Original - Resolution	Count difference	>2 or 5% original, but not error 5	Count for # errors 2, 3, 4	Count for ID errors 2, 3, 5	**Total Ind**	**Taxa Error**
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
					0	0