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SDI FINAL EVALUATION FORM 1.1

PART 1:

Journal Name:	International Journal of Plant & Soil Science
Manuscript Number:	2014_IJPSS_13067
Title of the Manuscript:	An understory comparison of the exotic PhellodendronamurenseRupr. (RUTACEAE) and adjacent native canopy species in an urban and suburban woodland

PART 2:

FINAL		
	EVALUATOR'S comments on revised paper (if any)	Authors' response to final evaluator's comments
1.	Need to place p-values for your significant & non-significant results in the ABSRACT	
	section;	
2.	L 97 How long were the transects? What was the minimum distance between plots?;	
3.	L112 documents <i>B. lenta</i> as a major component of both forests in terms of importance	
	value. Thus, please resolve with L95 that states Forest Park is dominated by <i>Quercus</i>	
	spp.;	
4.	L123 'sited' should be 'sites';	
5.	L132 'assure' should be 'ensure'; L142 It is probably more important to note that 1-tailed t-tests were run, given your	
0.	hypotheses;	
7.	L153-154 Provide Standard Error (SE) or 95% confidence interval (CI) estimates for	
, .	your derived means;	
8.	L154-156 The authors reverted back to the original results; which I had issue with	
	in my original review. It is rather simple, however from my original comments the	
	first time through:	
L140	Please be explicit with the statistical analyses used. I'll assume that t-tests were	
	run. However, looking at the degrees of freedom it looks like you used 'plot' as a	
	sampling point. Given that 4 quadrats were drawn from under the same tree, I	
	question the independence of each plot (nb., a stipulation of any parametric	
	analysis). It would perhaps be better to obtain the mean from the four plots	
	under each tree & run your t-tests using these averaged values. Thus, 72 plots at	
	Bartlett becomes n=18; & the 96 at Forest Park becomes 24, etc. As before, you	
	could nest cardinal direction (aspect) into your design to see if any differences in	
	regeneration patterns existed. You could then perhaps respond to some of your	
	Qs concerning mechanisms.	
	Re-analysing your dataset may not change the results any, but at least you will	
	have run analyses correctly & have no worry that your results are marred by	
	pseudoreplication (Hurlbert 1984) I have provided you the link to an important	
	article here:	
	http://www.masterenbiodiversidad.org/docs/asig3/Hurlbert_1984_Pseudoreplic	
	ation.pdf	
0	I 167 170 As above with analyzed also add CE as Classica to means	
	L167-170 As above with analyses; also, add SE or CI values to means; L187-190 As above with analyses; add SE or CI values to means;	
	L202-203 Add Se or CI values to means;	
	L207 F &df values missing;	
	Results & Discussion – I would still like to know what the species actually were under	
	the canopies of each (i.e., add a species list for each understory group). Because, what	
	is potentially more important is if there were differences in species composition	

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rather than total number of species. This may help provide insight into developing hypotheses for the *mechanisms* involved. For example, what if understory plant composition was dominated by different species under *P. amurense* canopy or excluded certain species normally found under native canopy? This could potentially alter successional patterns & thus, species diversity across the landscape over time. Providing us (the reader) with only info on abundance simply scrapes the surface of what you can do/provide the reader with the information you collected. The reader is told that diversity is lower under invasive spp canopy. But is it always the same 8.95 (± xxSE) understory species found (at Bartlett) or are different mixes of species also found under native canopy established there? i.e., *are there any understory species only found under invasive spp canopy or only found under native species canopy*?

14. Please actually do the revisions this time. Again, if running analyses is an issue, seek guidance from one of your peers. Also, addressing my last comment (providing species lists) would help immensely, not so much so that we know what species are present (as most people not in NY or CT wouldn't care), but more so to determine if species composition differed under invasive vs native canopies

Reviewer Details:

Name:	Anonymous
Department, University & Country	Canada

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