



Questions

1. **Comparing the Curves** : I don't have a great idea about how to best compare the fit between the Genetic Accumulation Curve and the two Spatial Accumulation Curve. The current r squared values give us a number and that might be enough. However there might be better options out there. Looking at the 90% accumulation as the end threshold? Normalizing the dataset rather than starting a zero? This would be a great thing to get some additional opinions on.

2. **Buffer intervals** : We want a defined set to work with. 50km is the only shoe in my book. If there are other referenced used of buffer size for gap analysis that be great to include those. Also if there are strong opinions on distance we can work those. Just being upfront and saying we don't want to to test all values between 1 and 100 or 1/2km intervals.

3. **Integration with existing workflow.** : Austin this is kinda me and you to talk though but I'm not sure attempting to building this into the current workflow you've create is my first choice.

4. **Timeline integration:** Austin, I know you've set a