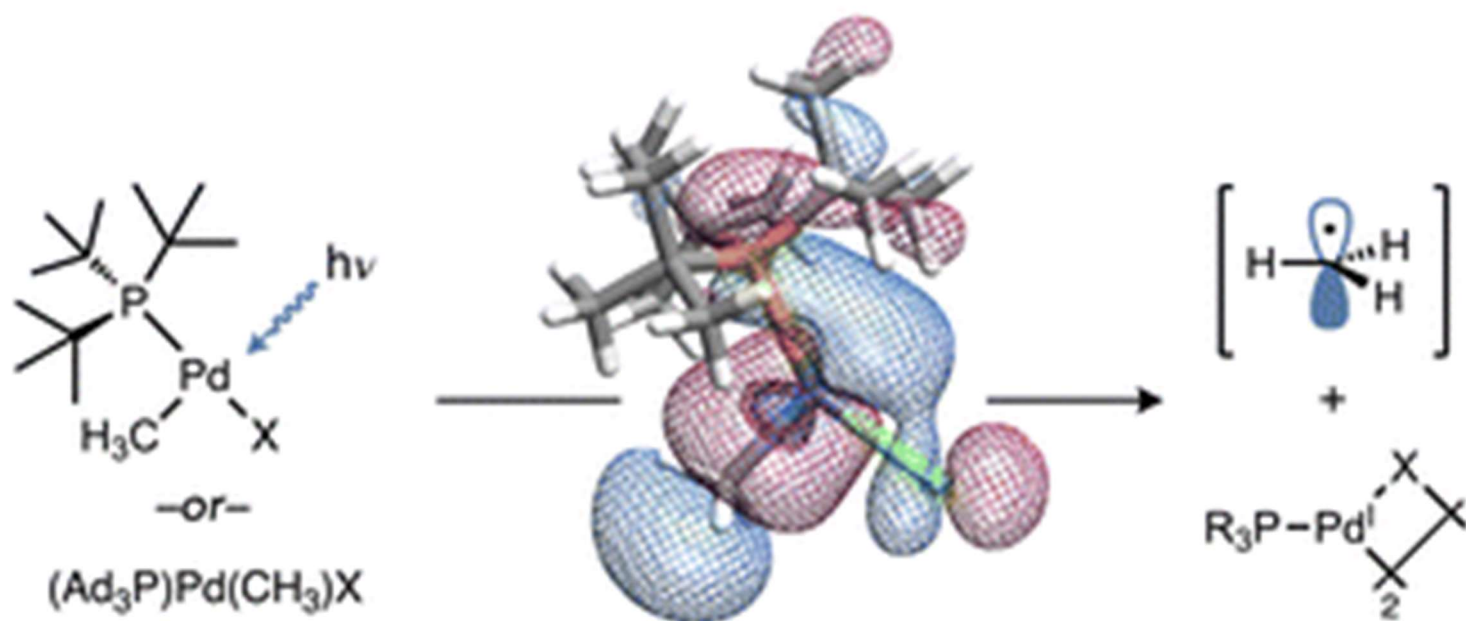


# Computational and Machine Learning Exploration of Pd-C Photocleavage in T-shaped Organopalladium Complexes

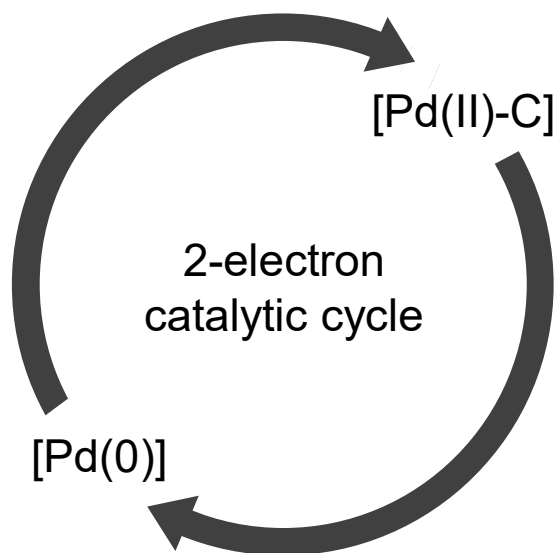


Peter M. Waddell, Ph.D.



## ACCESSING 1-ELECTRON MANIFOLDS WITH Pd?

thermal reactivity

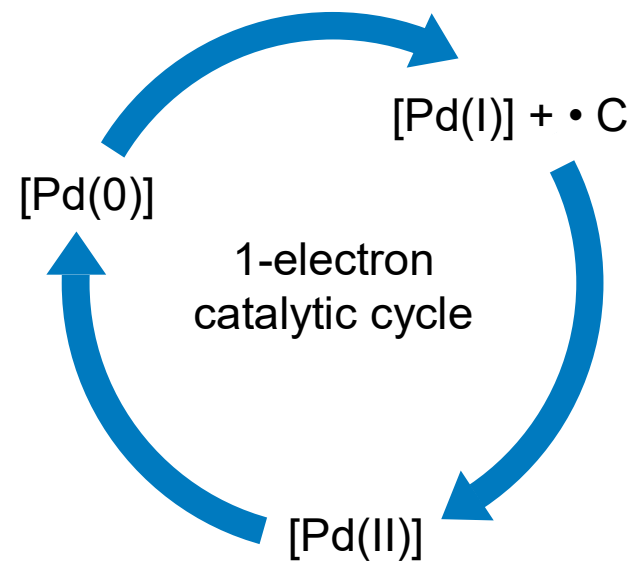


Pd is biased to this  
general manifold



Can light energy be  
used to toggle  
between manifolds?

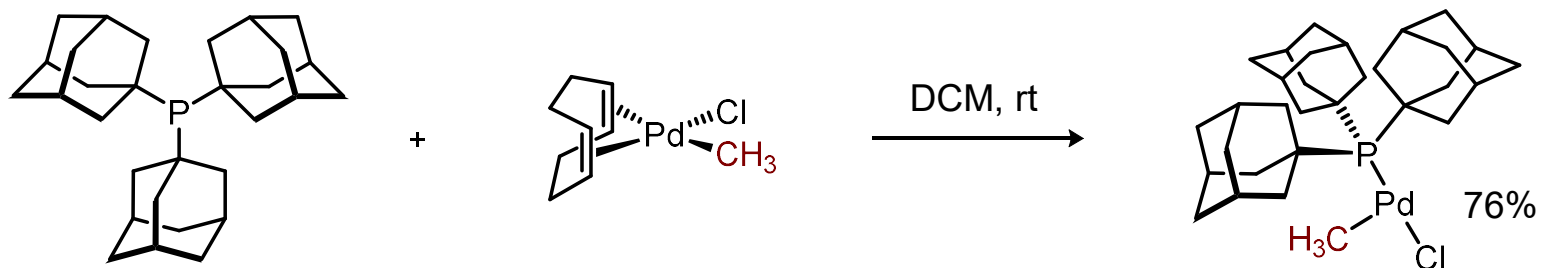
*excited state reactivity*



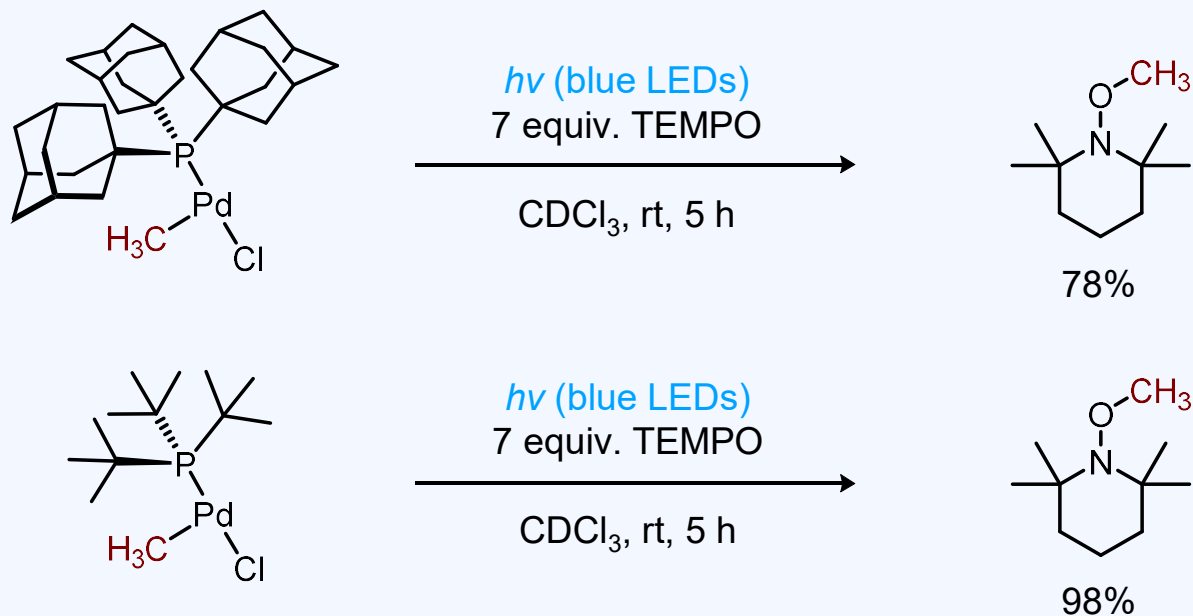
challenging to  
access



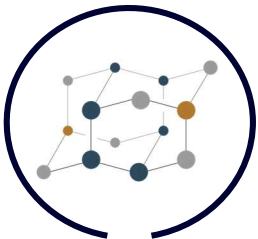
## T-SHAPED COMPLEX Pd-C PHOTOCLEAVAGE WITH VISIBLE LIGHT



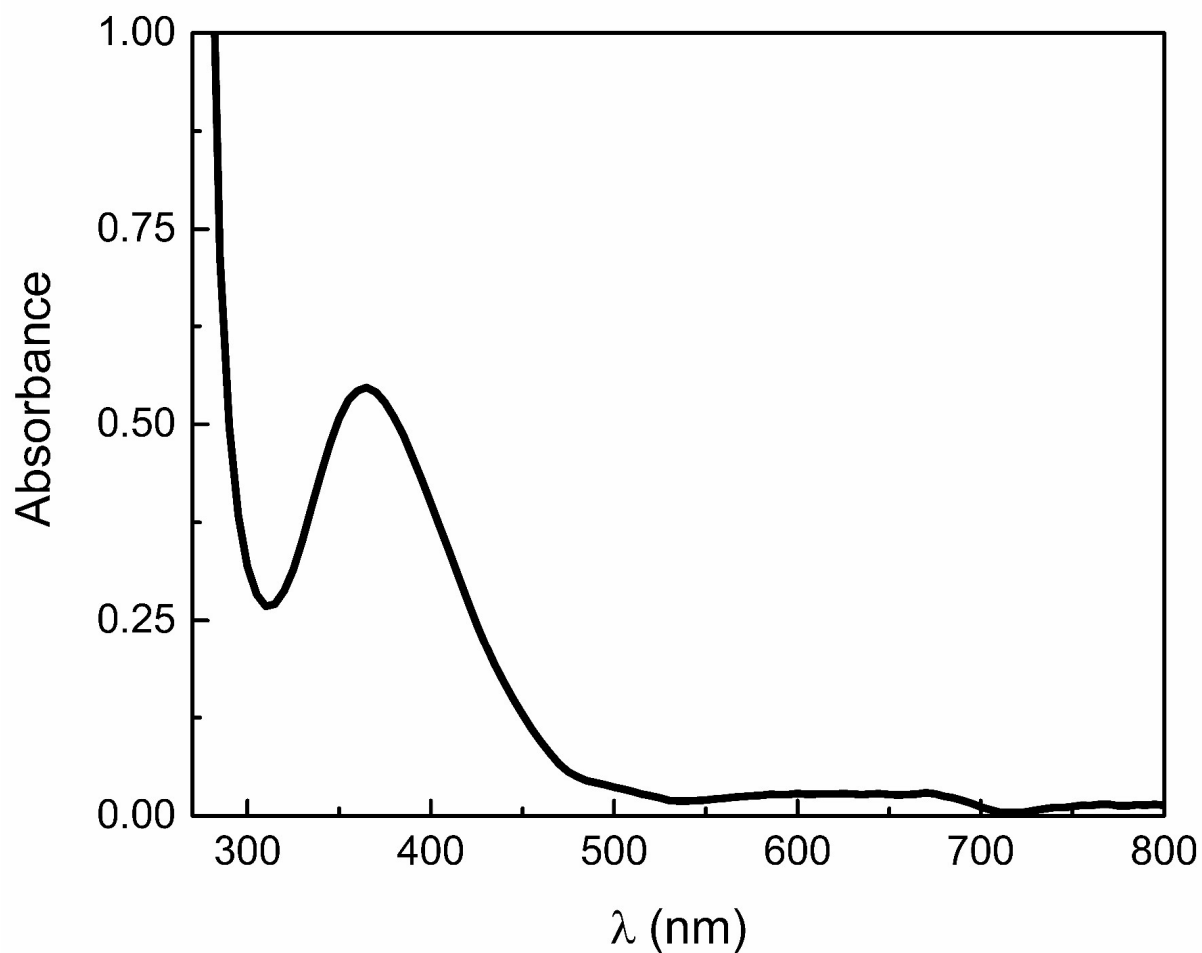
Do these T-shaped complexes exhibit any photoreactivity?



*Visible light*-induced bond weakening triggers SET chemistry in the most synthetically versatile metal and oxidation state (d<sup>8</sup>, Pd<sup>II</sup>)! Previously, this reactivity was much more limited to high-energy UV light.



## Pd-C PHOTOCLEAVAGE WITH VISIBLE LIGHT



Absorption spectrum of the T-shaped complex shows the key peak for the transition that leads to Pd-C cleavage, with a  $\lambda_{max}$  just under 400 nm.



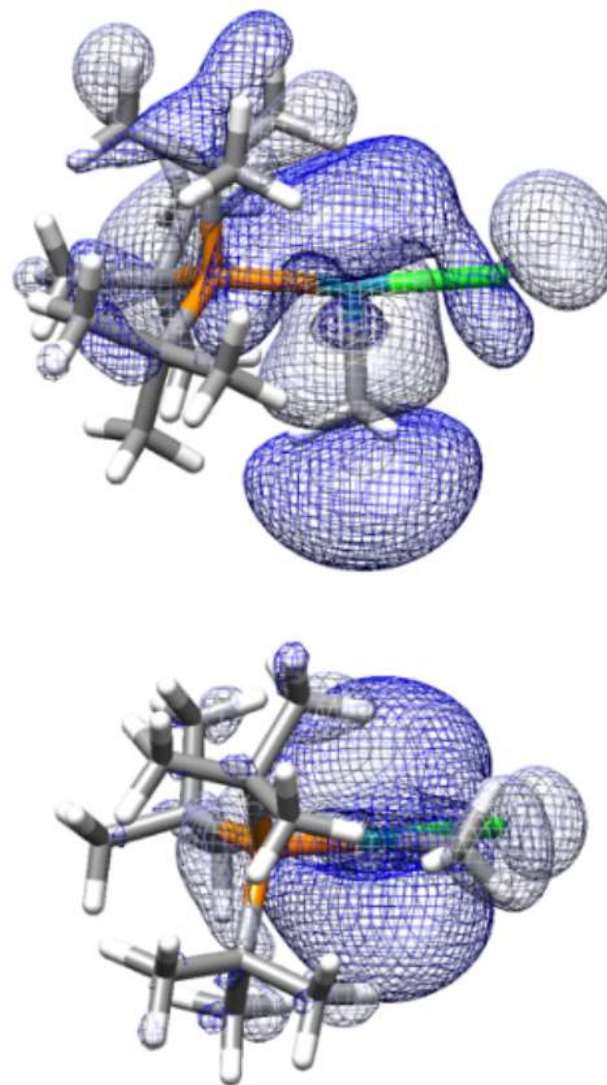
## HOMO/LUMO ANALYSIS

with Lei Tian

LUMO:  $d_{x^2-y^2}/\sigma^*$

Gap: 2.72 eV  
(456 nm)

HOMO:  $d_{z^2}$

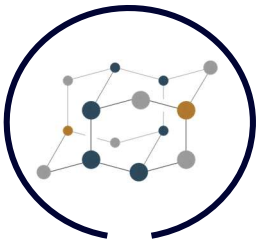


LUMO shows Pd-C  $\sigma^*$  character, ***possibly accessible with visible light due to the absence of a ligand trans to C in the fourth site of the square plane.***

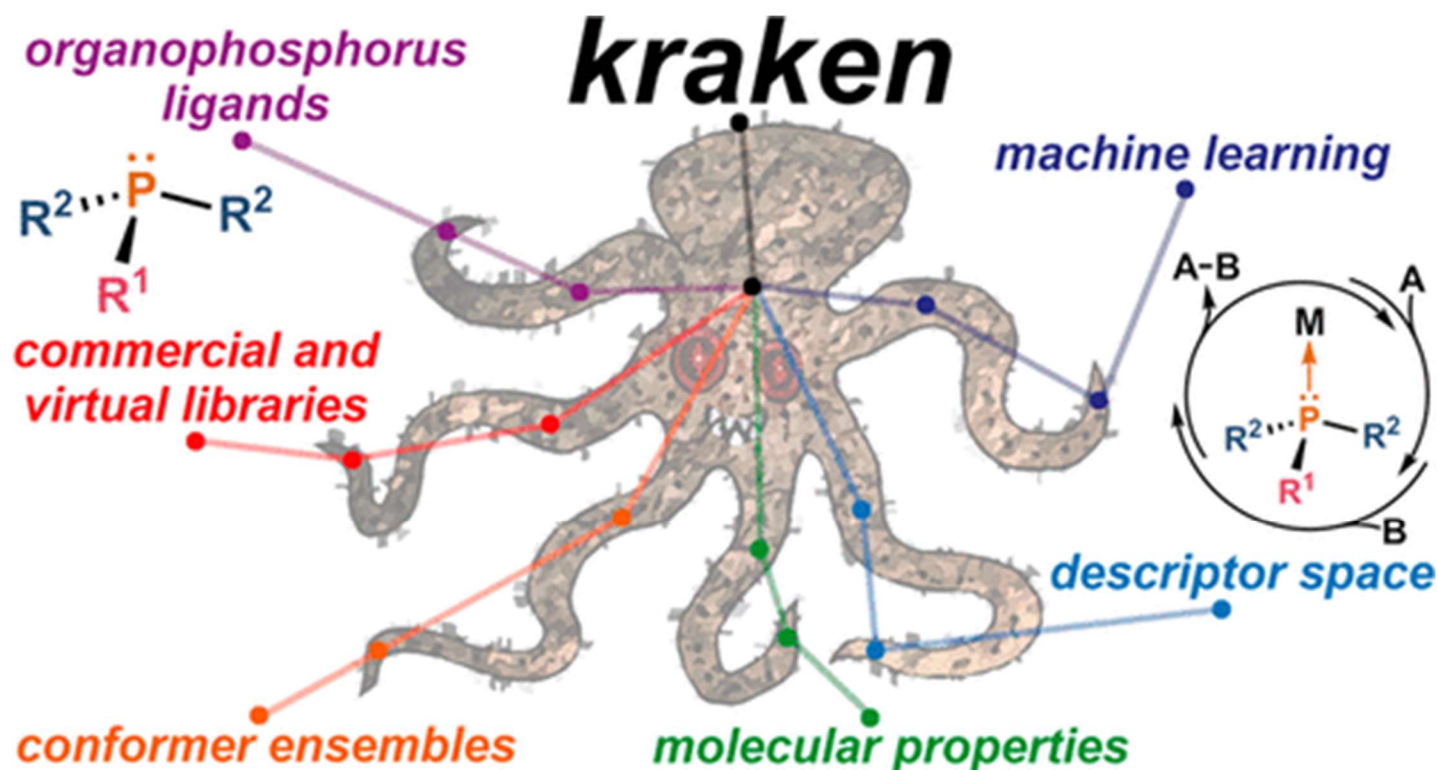


## COMPUTATIONAL STUDY OBJECTIVES

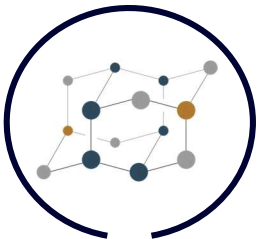
- How do the steric and electronic properties of the ancillary ligand affect the UV-vis absorption spectrum of their organopalladium complexes? What characteristics cause red or blue shifts in the  $\lambda_{max}$  of the key peak? What is the lowest energy light that might be able to access this reactivity?
- Hypothesis: stronger bonding interaction in the fourth site will cause a blue shift in  $\lambda_{max}$ .
- Can we build a model with machine learning techniques to predict the spectral profile of a complex based only on information from its ancillary ligand?
- How conserved is the Pd-C photocleavage reactivity among a broad set of organophosphorus ligands?



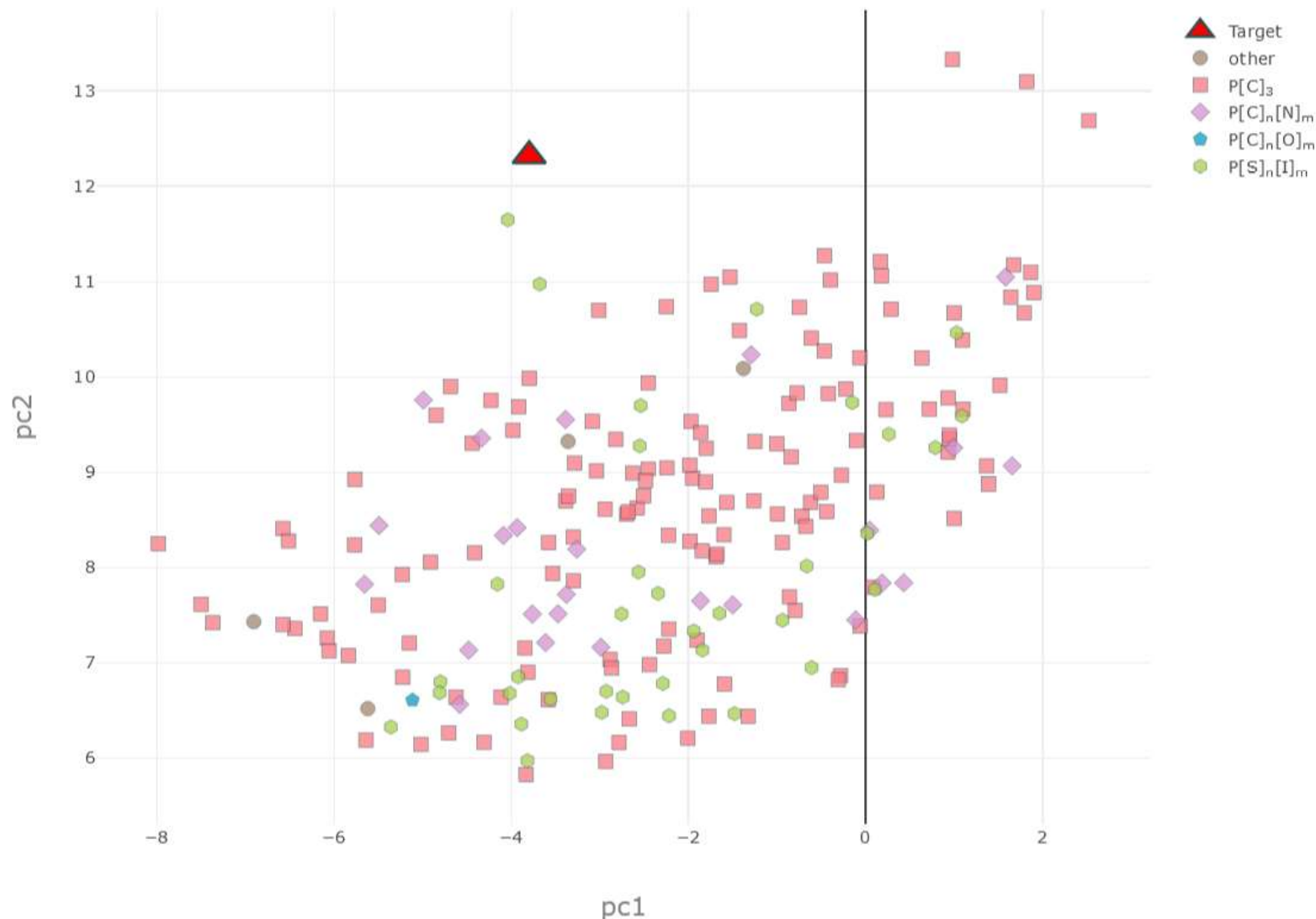
## EXPLORING PHOSPHINE LIGAND SPACE WITH KRAKEN LIBRARY



The Kraken library of phosphine ligands is a perfect starting point to explore the chemical space of bulky organophosphorus ligands.



## EXPLORING PHOSPHINE LIGAND SPACE WITH KRAKEN LIBRARY

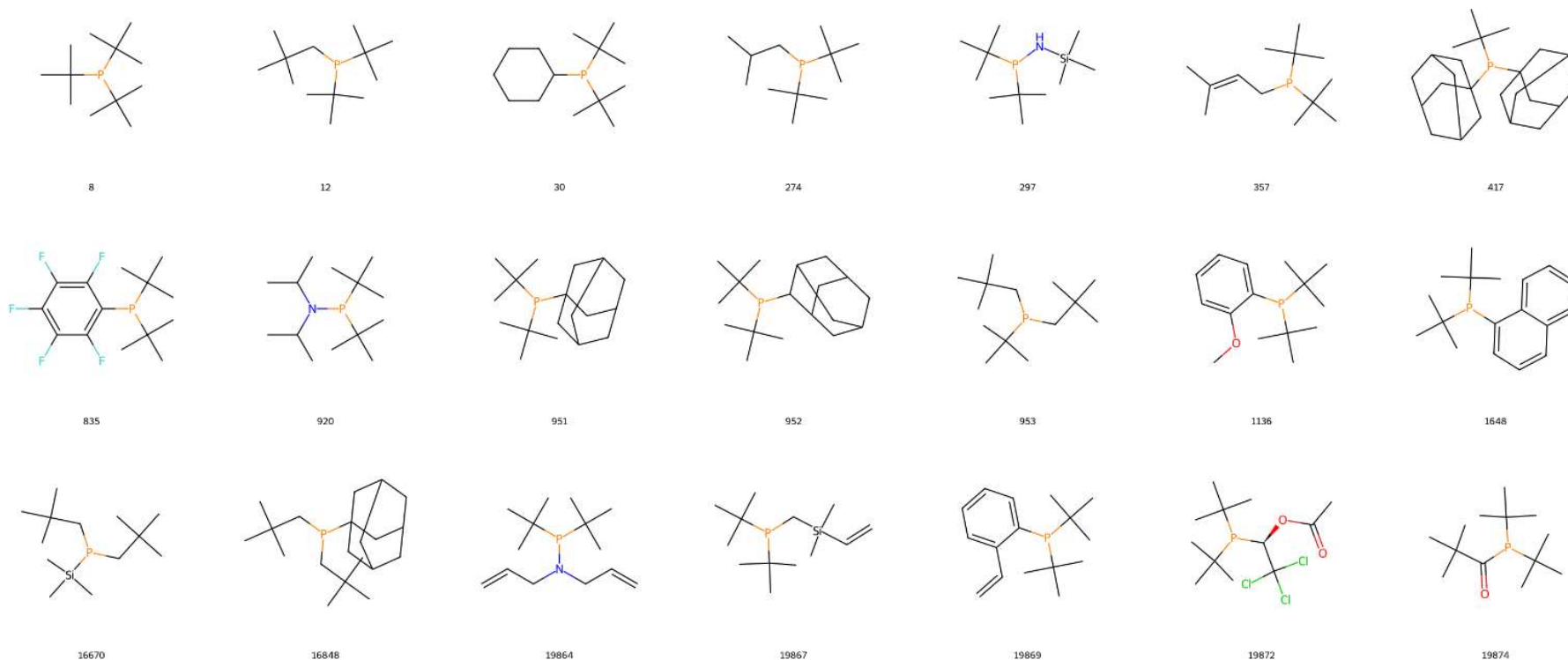


To start, I used the Kraken library to find the 210 closest (in principal component space) organophosphorus ligands to our target (tri-*t*-butylphosphine)





## EXPLORING PHOSPHINE LIGAND SPACE WITH KRAKEN LIBRARY

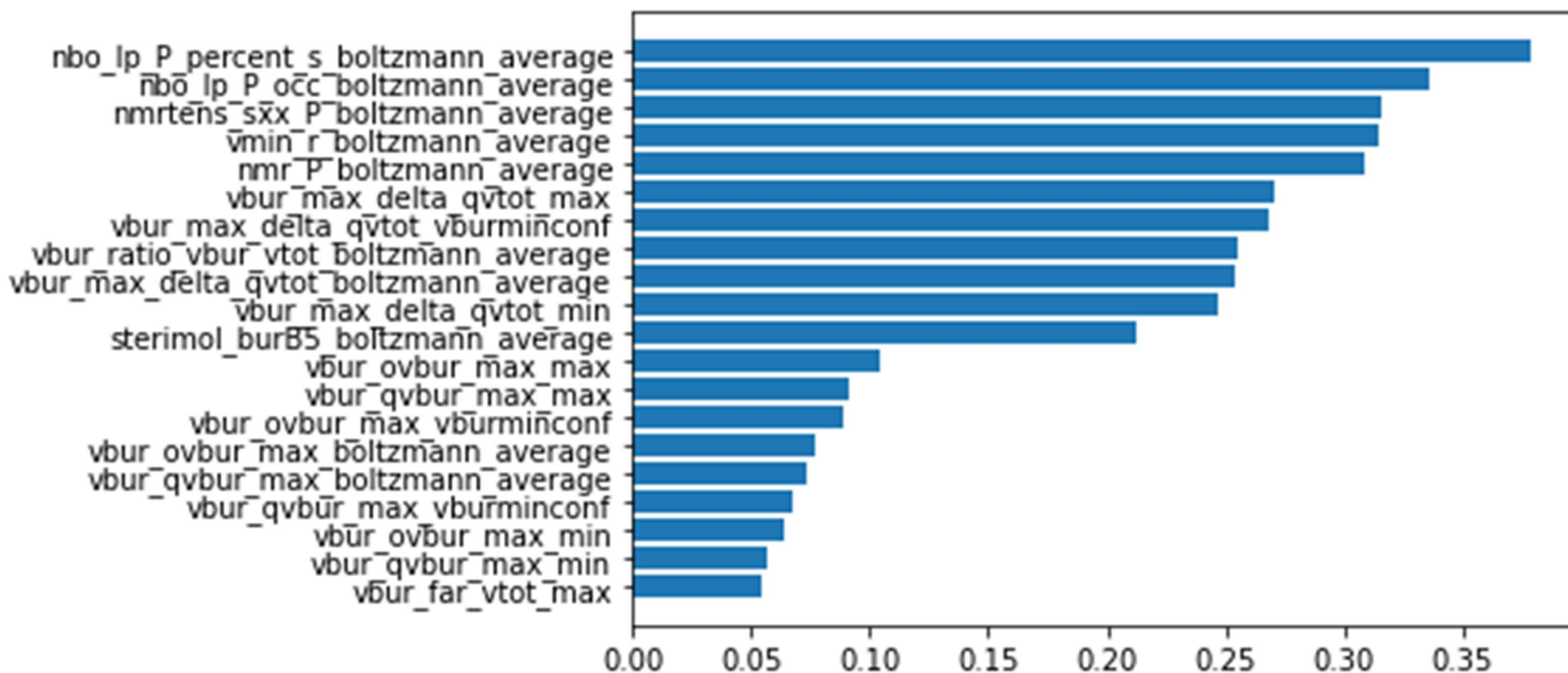


In order to ensure that the organophosphorus ligands would form T-shaped (monoligated) Pd complexes, I restricted this set to the ligands with Boltzmann-average  $\%V_{bur}$  of 68% or more, very close to the value for tri-*t*-butylphosphine.

This gave a final set of **107 ligands** (small subset pictured).



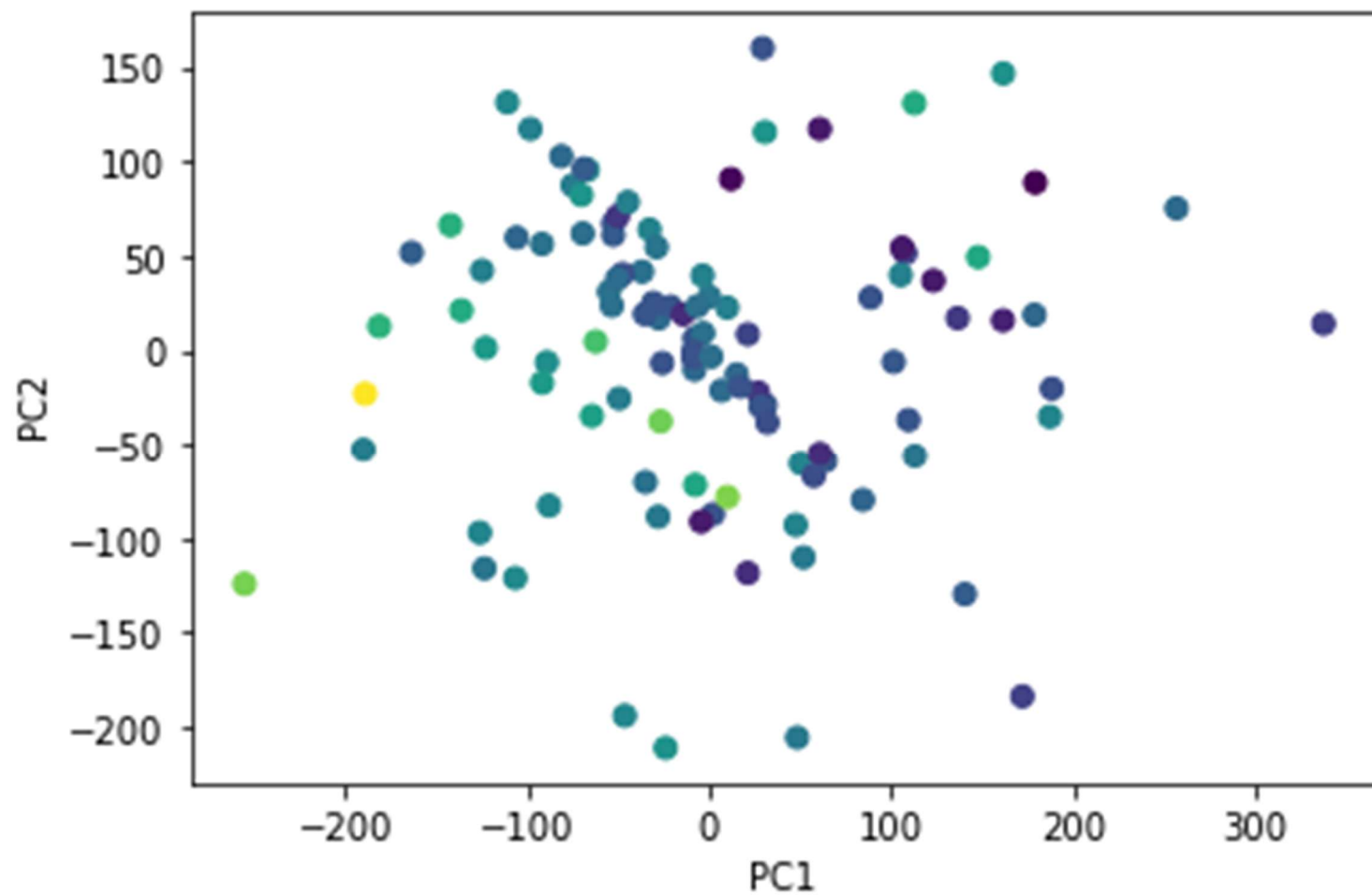
## LIGAND SET PRINCIPAL COMPONENT ANALYSIS



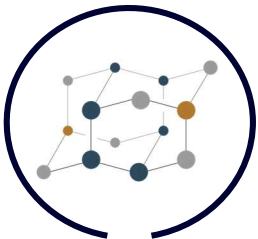
Here are the twenty most important features for the first principal component. Notably, electronic features such as Boltzmann-averaged NBO P lone pair %s and NMR values rank most highly.



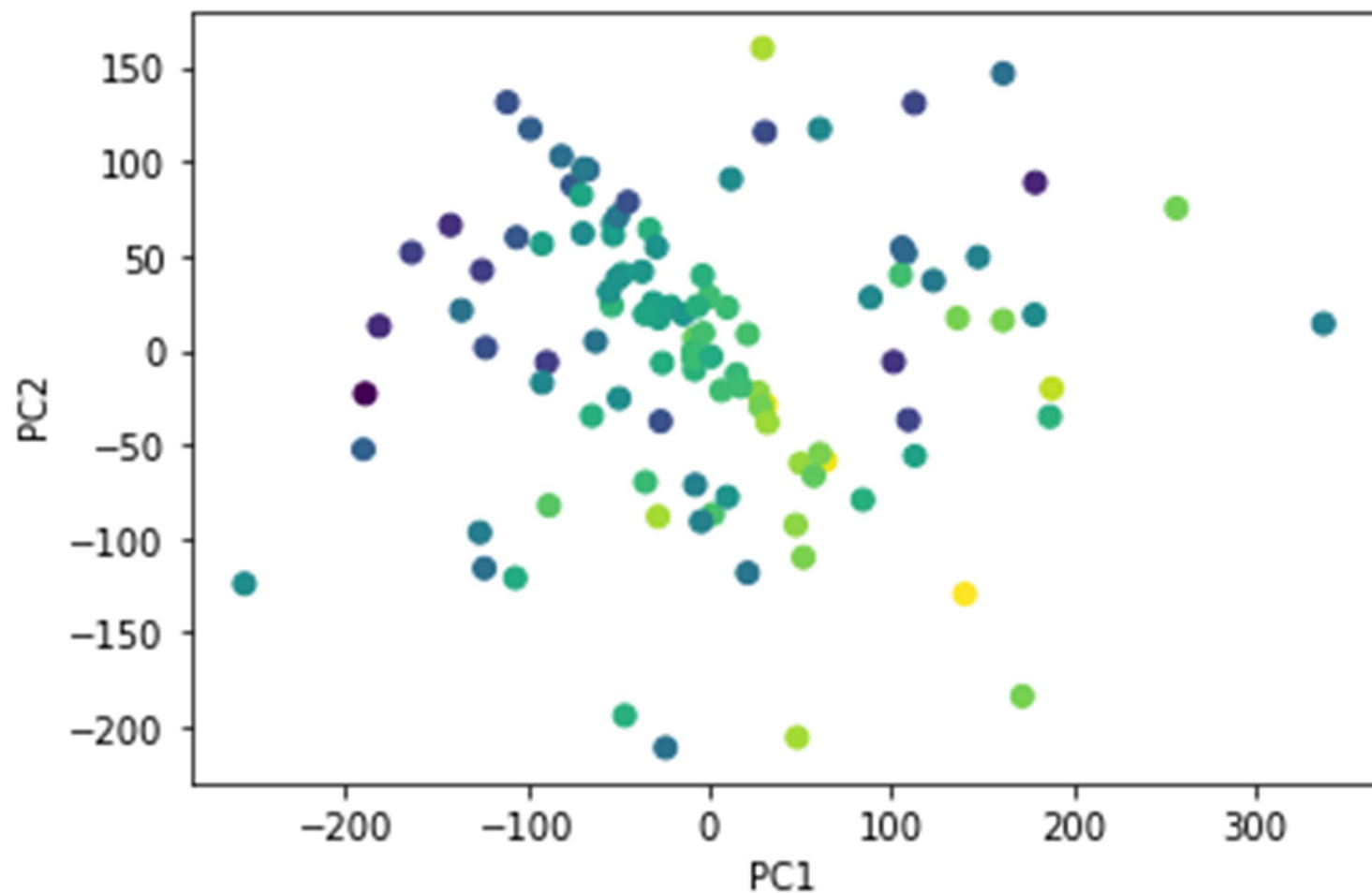
## LIGAND SET PRINCIPAL COMPONENT ANALYSIS



PCA on our ligand set, colored according to Boltzmann-average NBO P lone pair %s.



## LIGAND SET PRINCIPAL COMPONENT ANALYSIS



PCA on our ligand set, colored according to highest total volume difference in  $\%V_{bur}$  between two neighboring quadrants.



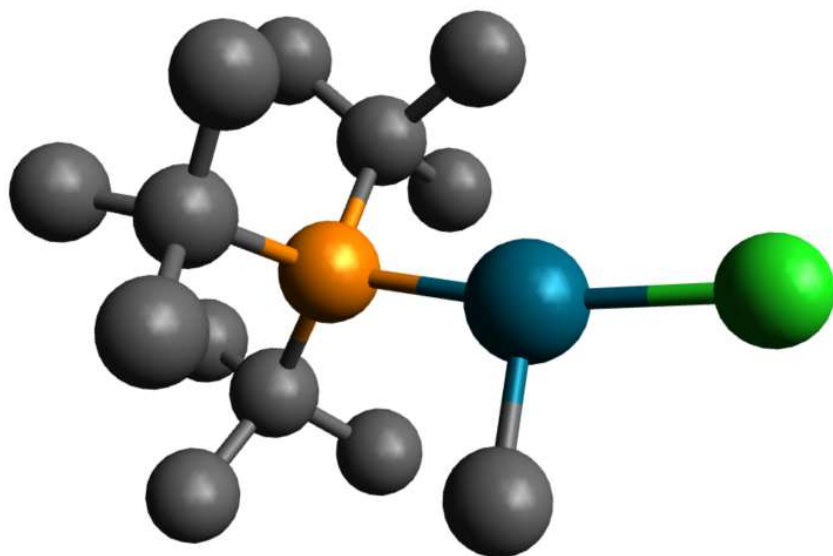
## COMPUTATIONAL WORKFLOW

- Gather and prune ligand set from Kraken database
- Generate SMILES strings for methylpalladium chloride T-shaped complexes for each ligand
- Use RDKit to run conformer searches for each complex
- Perform geometry optimizations based on the **lowest energy conformer only** for each complex (note that currently I am neglecting all other conformers at this time, which I intend to address later with more compute time)
- Perform TDDFT calculations and simulate the UV-vis spectra for each complex, determine the  $\lambda_{max}$  for the relevant transition
- Find trends in  $\lambda_{max}$  and ligand/complex properties
- Use machine learning to build and validate models to predict  $\lambda_{max}$  for other organophosphorus ligands, extract insights
- See the repo at: <https://github.com/pmwaddell/pd-c-photochem-ML> for the source code, all steps are automated with scripts



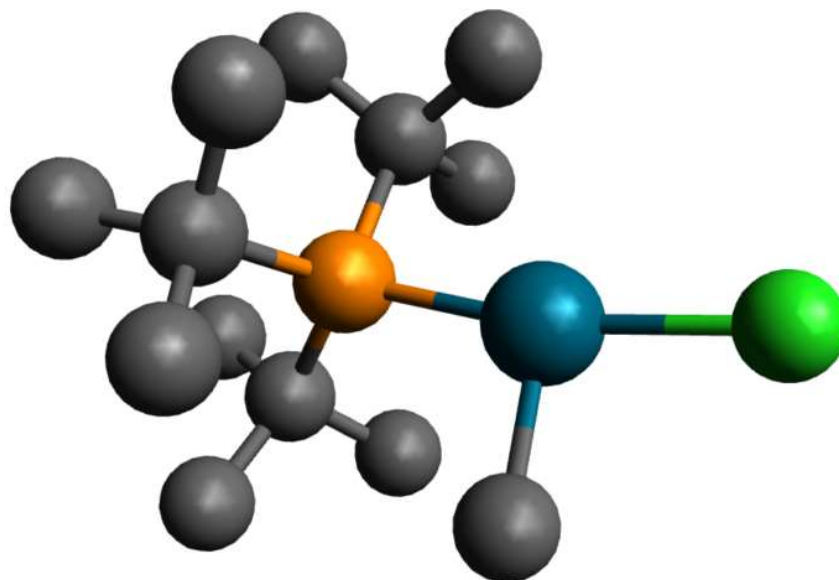
## DFT: LEVEL OF THEORY, GEOMETRY OPTIMIZATION

Literature X-ray structure



From Nozaki *Organometallics* **2006**, 25, 4588.

DFT geometry optimized structure



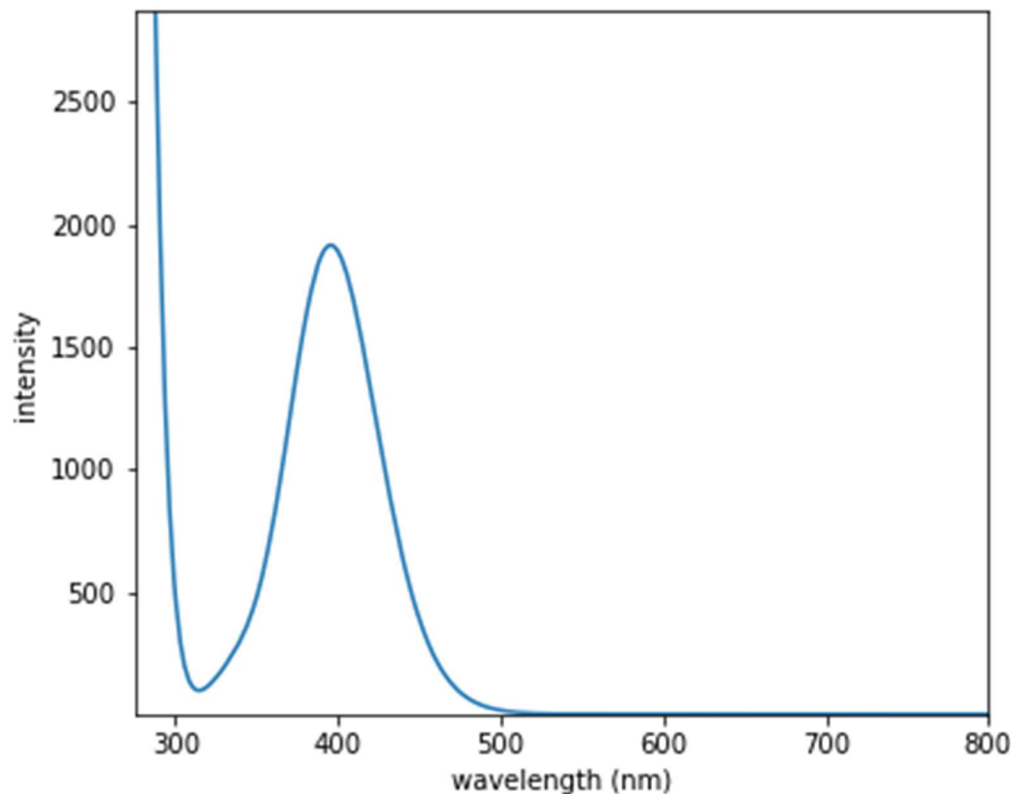
(hydrogens omitted)

After screening various basis sets and functionals for geometry optimization, **B3LYP-D3/def2-TZVP with CPCM( $\text{CHCl}_3$ )** was found to give good agreement (key bond lengths within 0.05 Å, angles within 1°) with the reported X-ray structure while being relatively time-efficient.

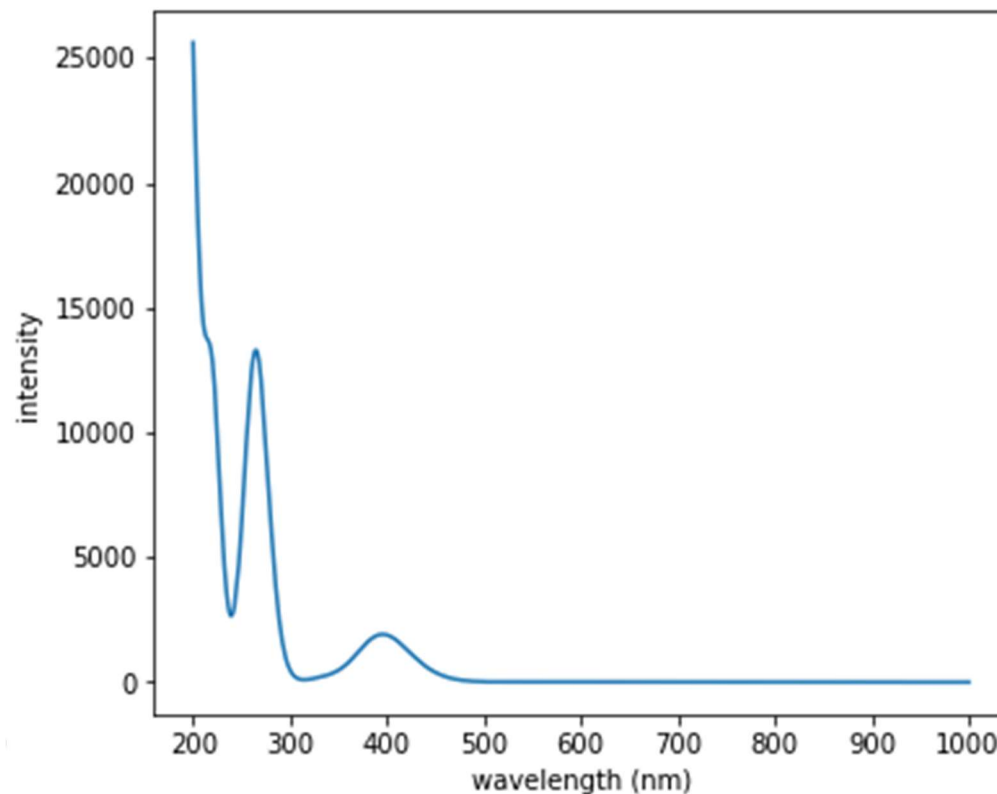


## DFT: LEVEL OF THEORY, TDDFT

TDDFT UV-vis spectrum



TDDFT UV-vis spectrum, full

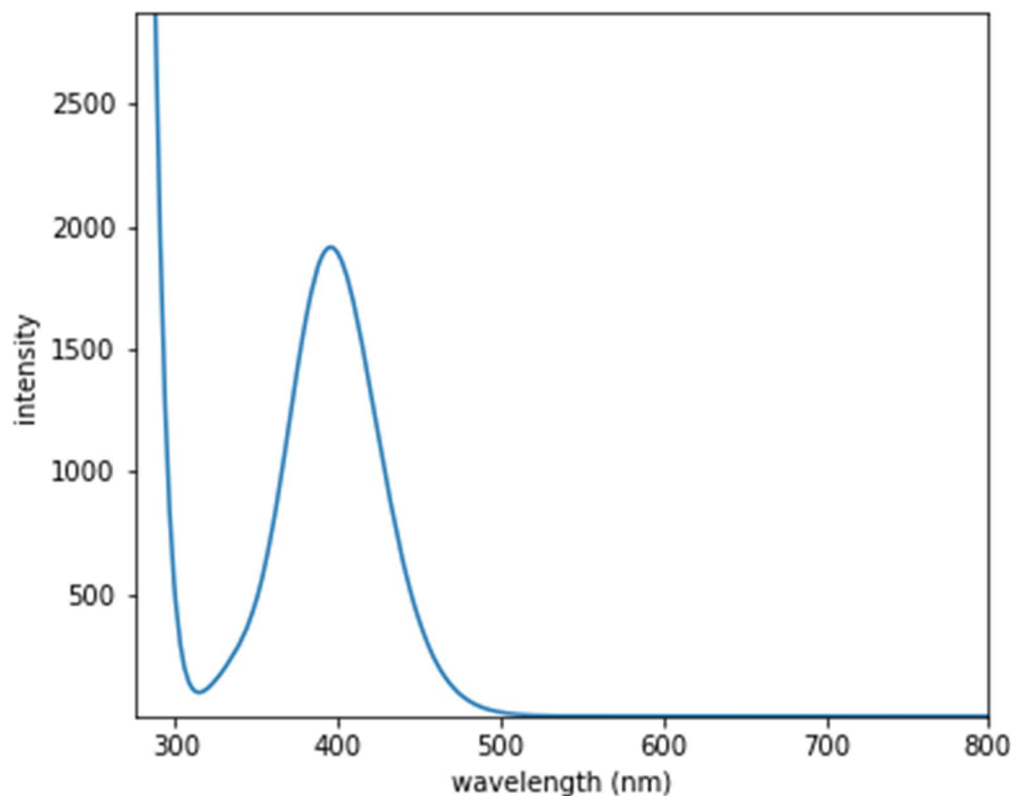


For TDDFT calculations, agreement with the experimental UV-vis absorption spectrum was used to benchmark performance. Similarly, **B3LYP/def2-TZVPP with CPCM(CHCl<sub>3</sub>)** was found to give the closest agreement.

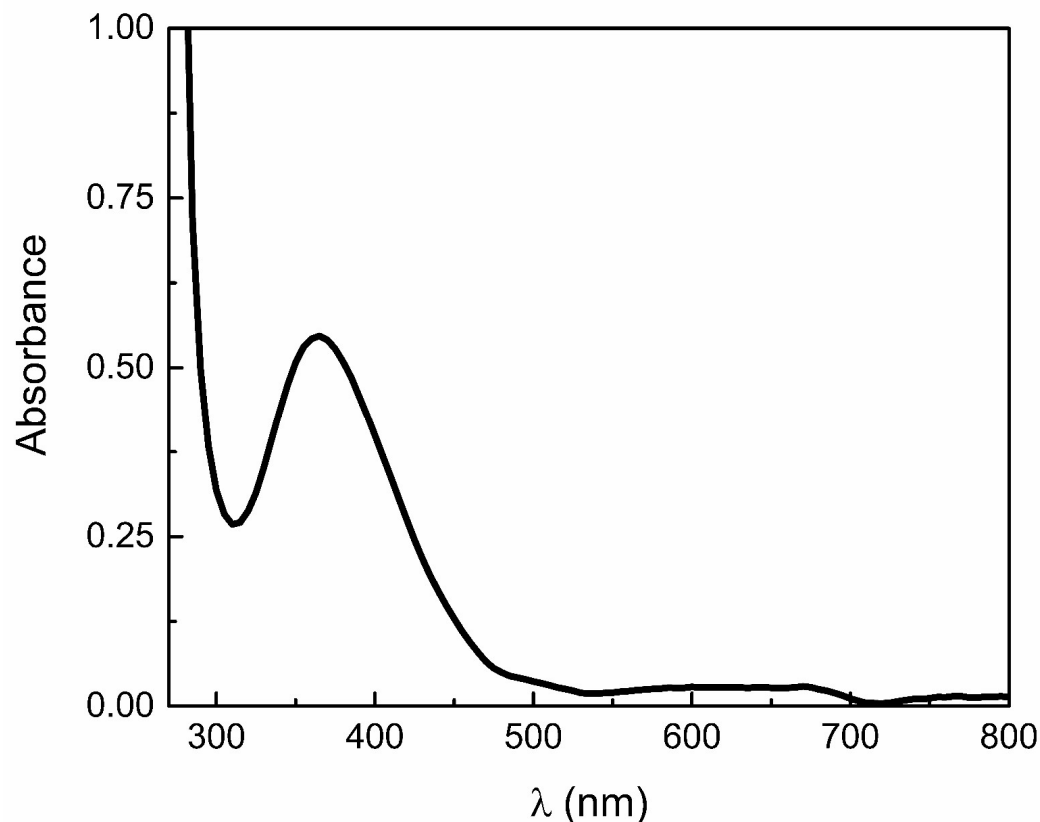


## DFT: LEVEL OF THEORY, TDDFT

TDDFT UV-vis spectrum

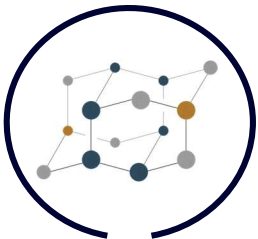


experimental UV-vis spectrum

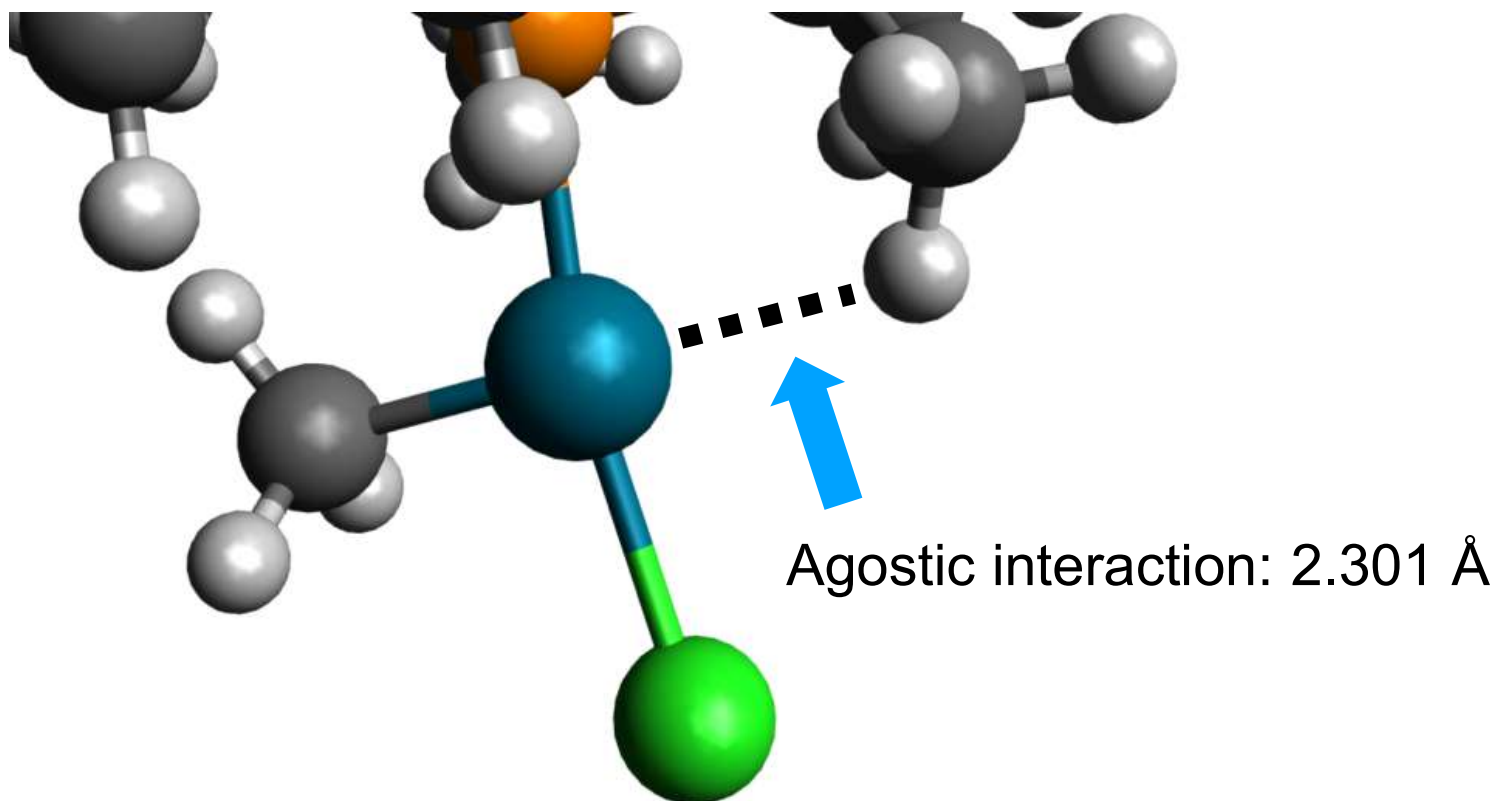


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## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{\text{MAX}}$ , AGOSTIC INTERACTIONS

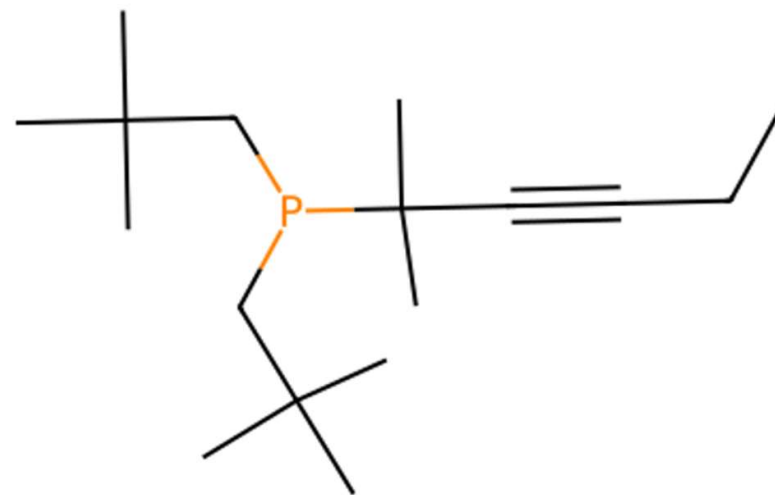
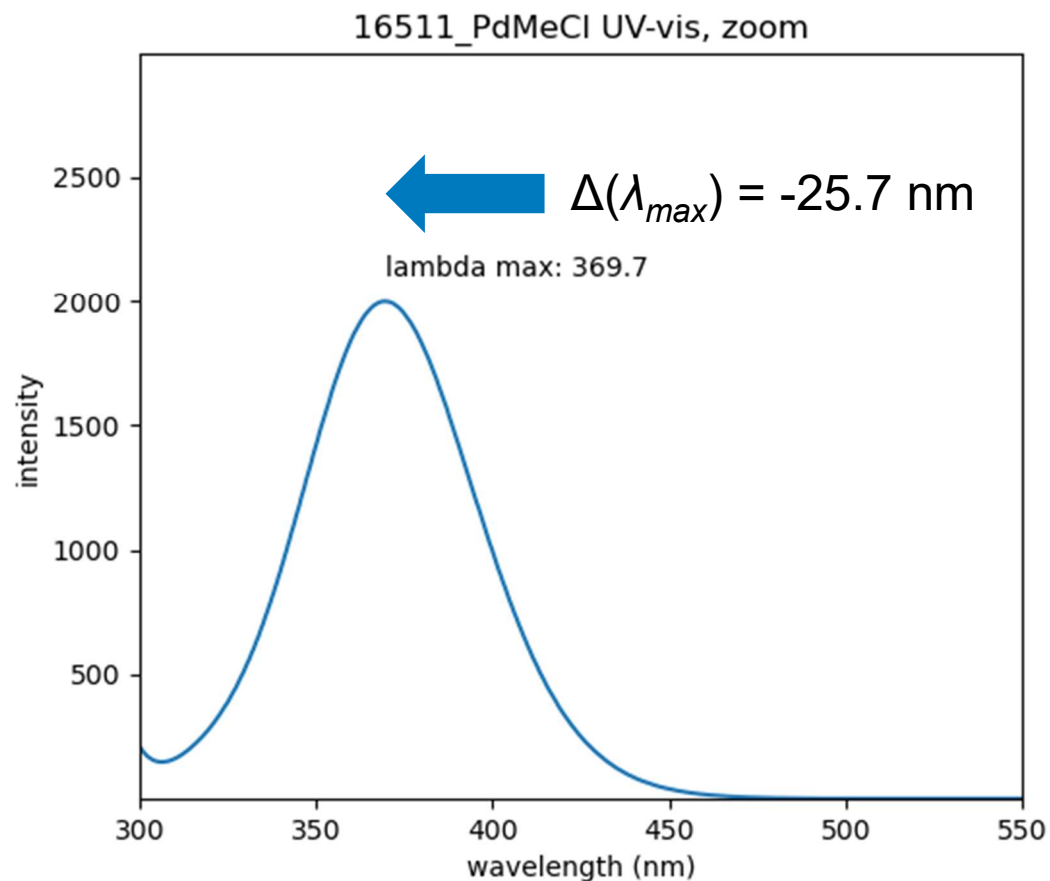


While we refer to these complexes as T-shaped, they often feature an agostic interaction in the fourth site. The distance between Pd and H correlates with the strength of the interaction.

My hypothesis is that the lack of a strong bonding interaction in this site lowers the LUMO ( $\sigma^*$ ) energy. Thus,  $\lambda_{\text{max}}$  should correlate with agostic interaction strength.



## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{max}$ , AGOSTIC INTERACTIONS

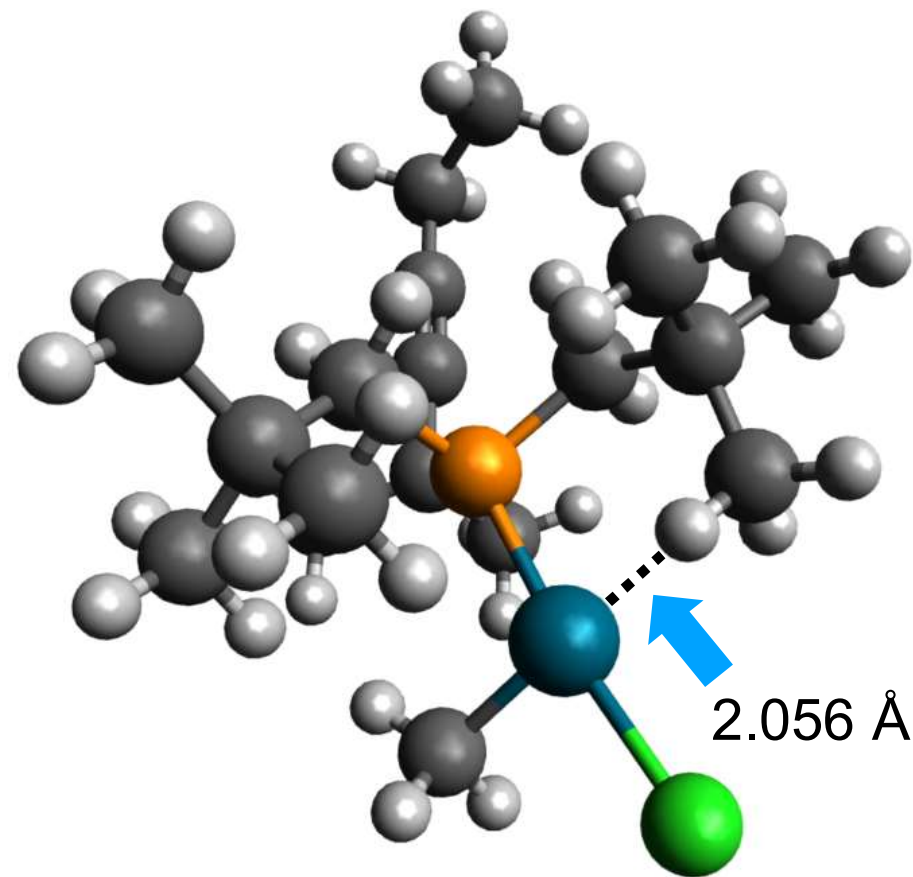
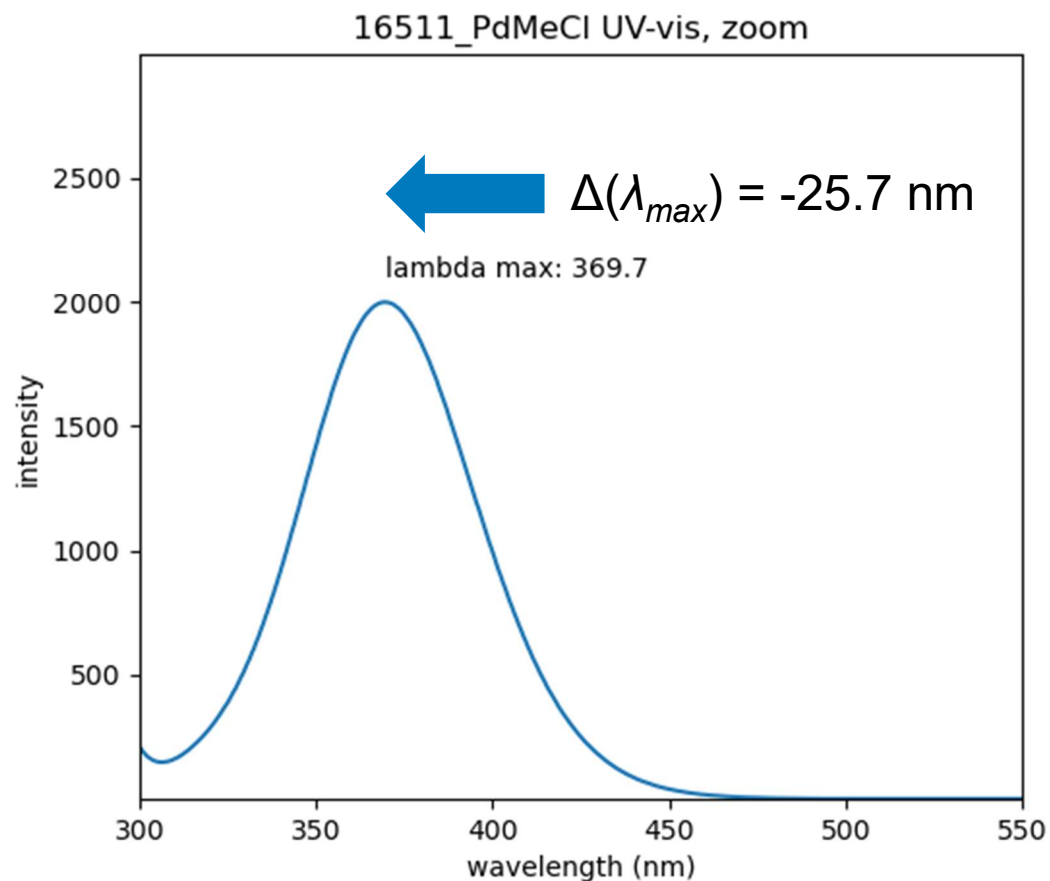


Ligand 16511 (Kraken designation)

As an example, this ligand features neopentyl groups which allow the H to come in closer proximity to Pd. As a result, the stronger agostic interaction leads to a  $\lambda_{max}$  blueshift of over 25 nm!



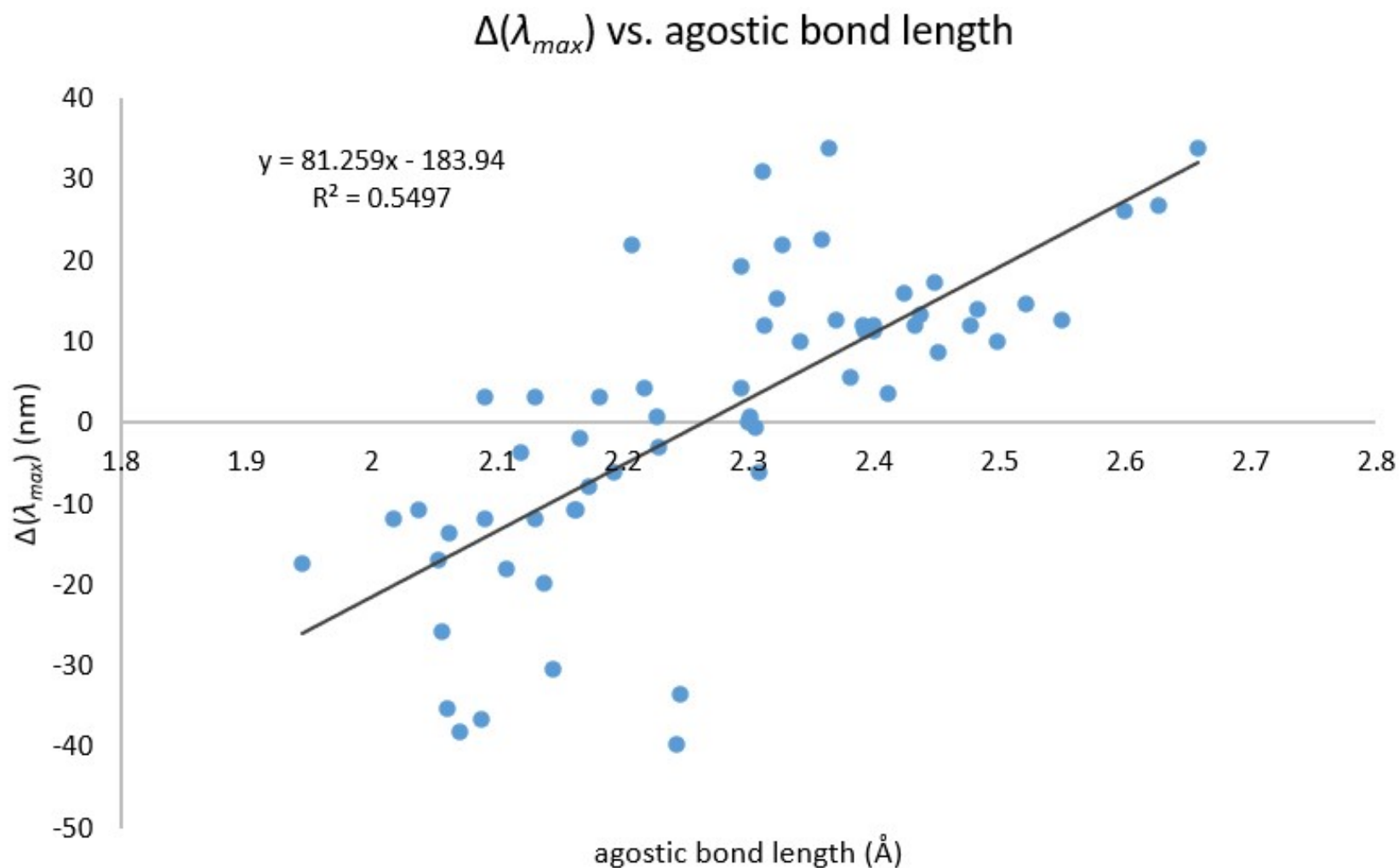
## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{max}$ , AGOSTIC INTERACTIONS



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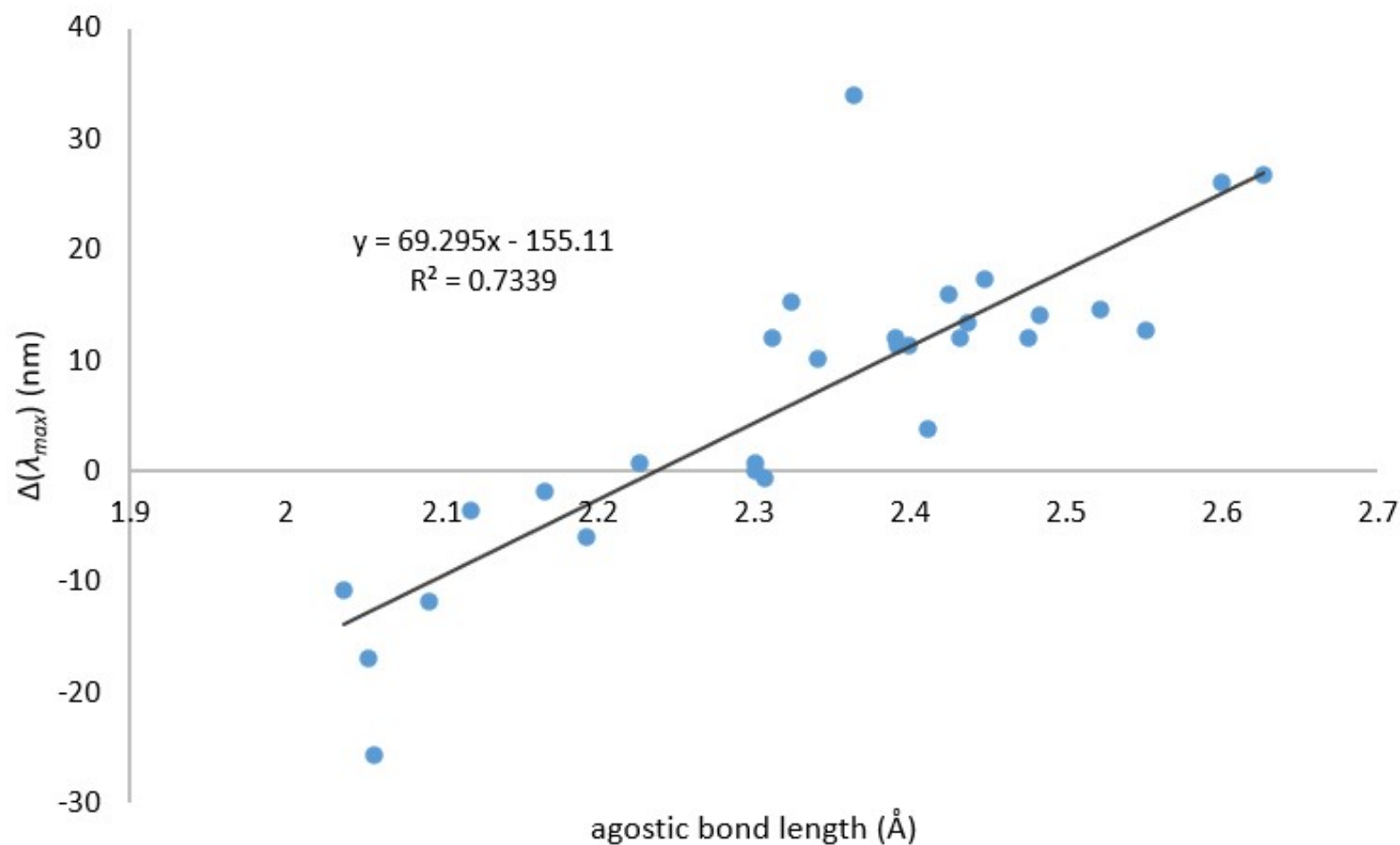


Across all ligands that apparently have an agostic interaction, the correlation between agnostic Pd-H bond length and the change in  $\lambda_{max}$  relative to tri-*t*-butylphosphine (which we will generally refer to as  $\Delta(\lambda_{max})$ ) is pretty good.



## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{MAX}$ , AGOSTIC INTERACTIONS

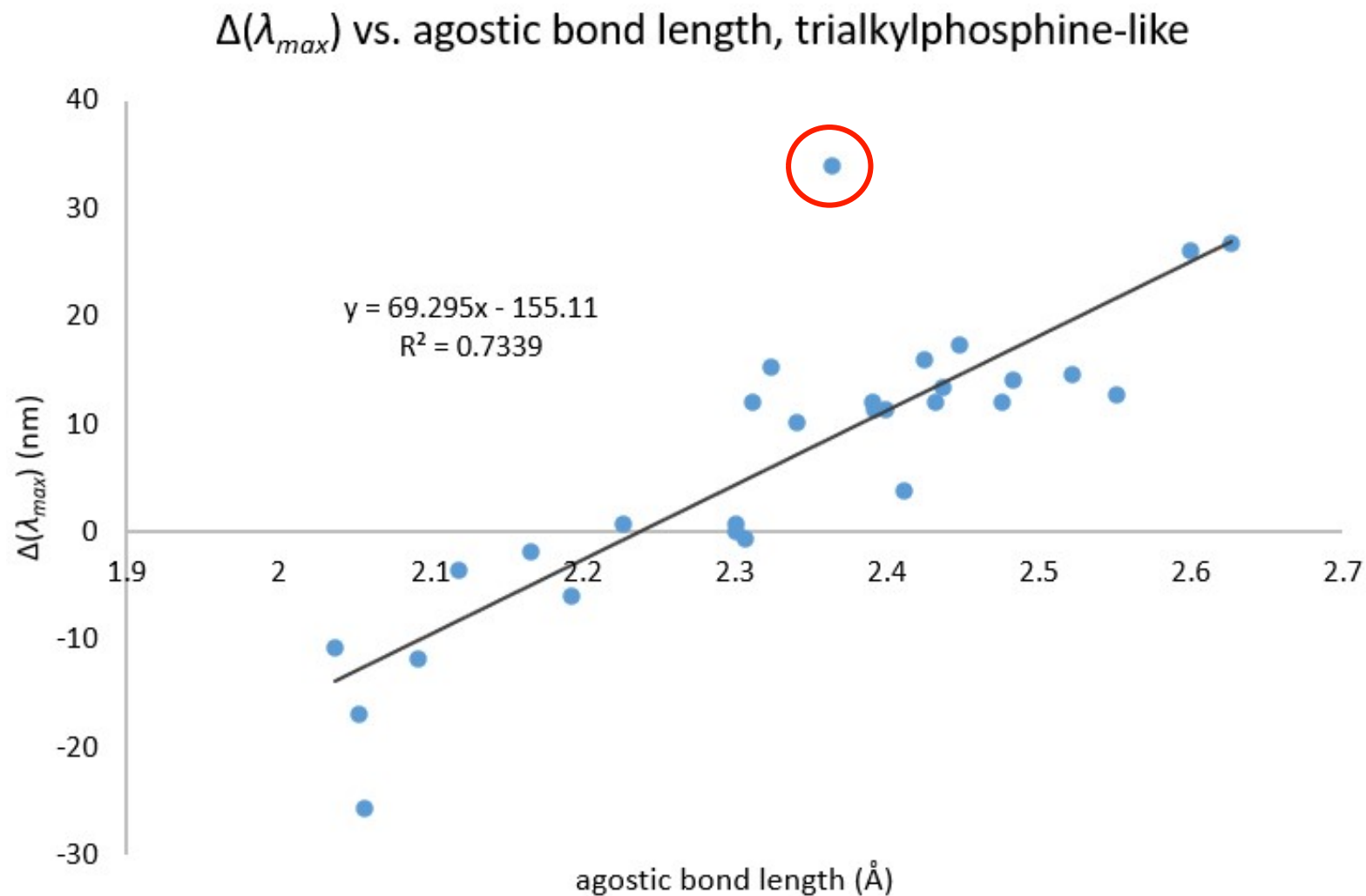
$\Delta(\lambda_{max})$  vs. agostic bond length, trialkylphosphine-like



If we limit ourselves to trialkylphosphine-like ligands (also including Pd-Si and non-aryl P-Csp<sup>2</sup>), the correlation improves significantly.



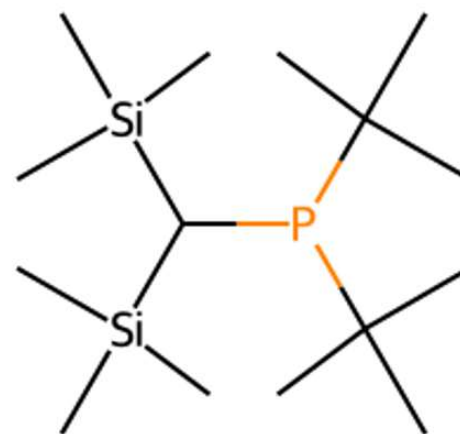
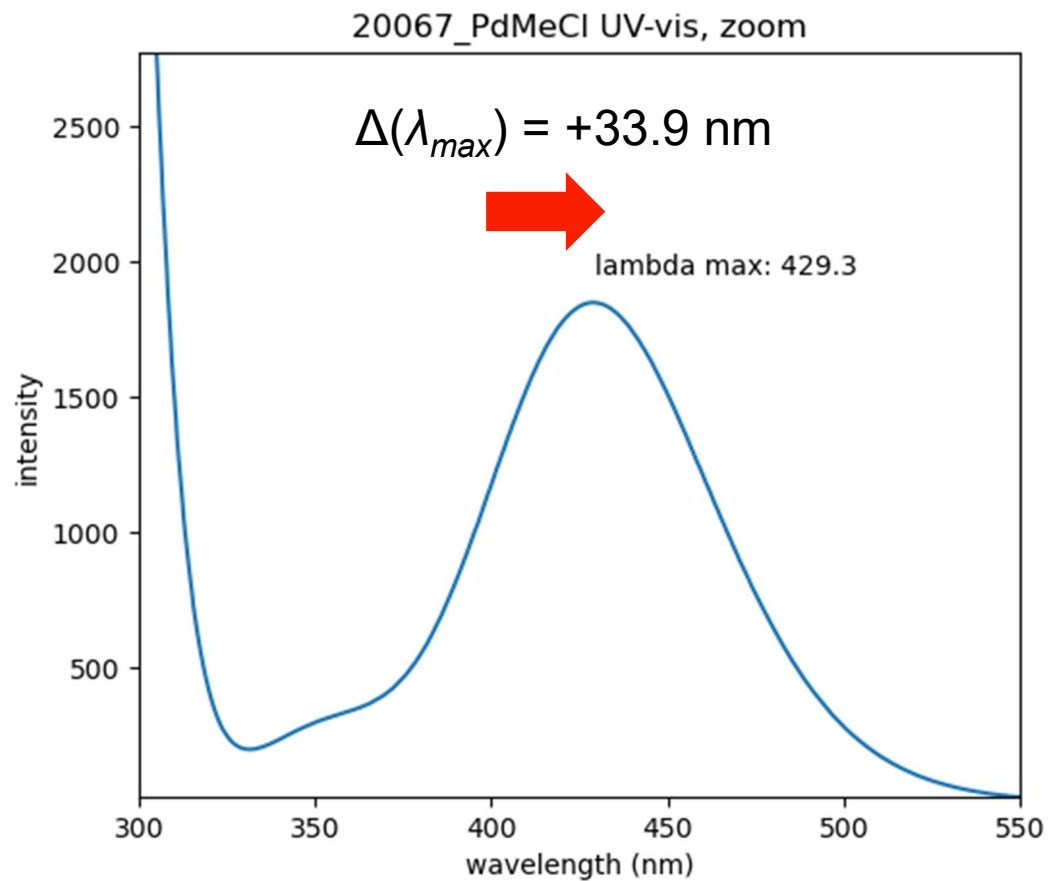
## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{max}$ , AGOSTIC INTERACTIONS



One particular case deviates strongly from this relationship, showing a much higher  $\lambda_{max}$  than expected. Can we figure out why?

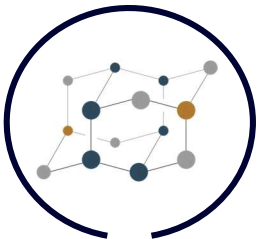


## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{\text{MAX}}$ , AGOSTIC INTERACTIONS

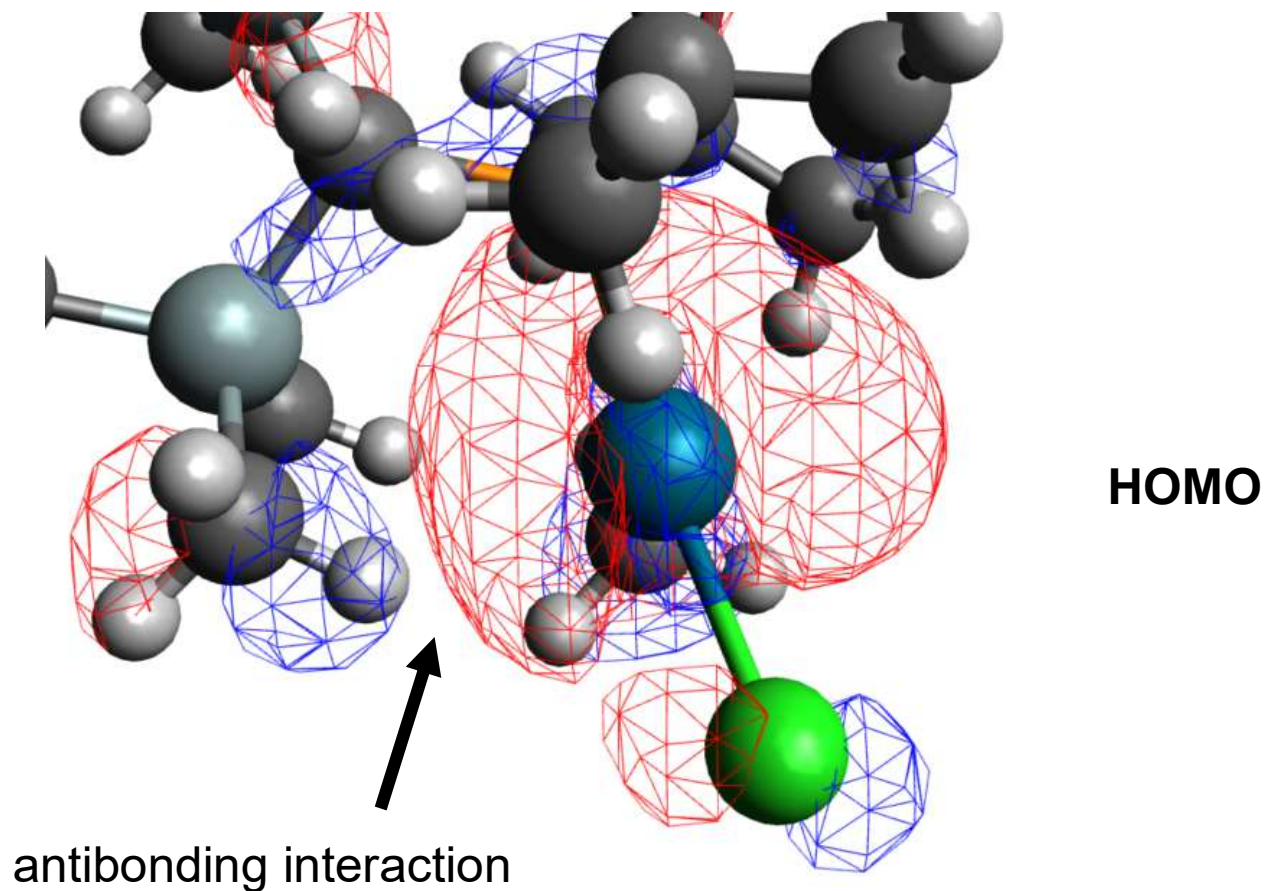


Ligand 20067

This is the ligand in question, which features large TMS groups on one substituent.

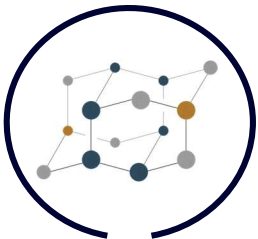


## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{\text{MAX}}$ , AGOSTIC INTERACTIONS

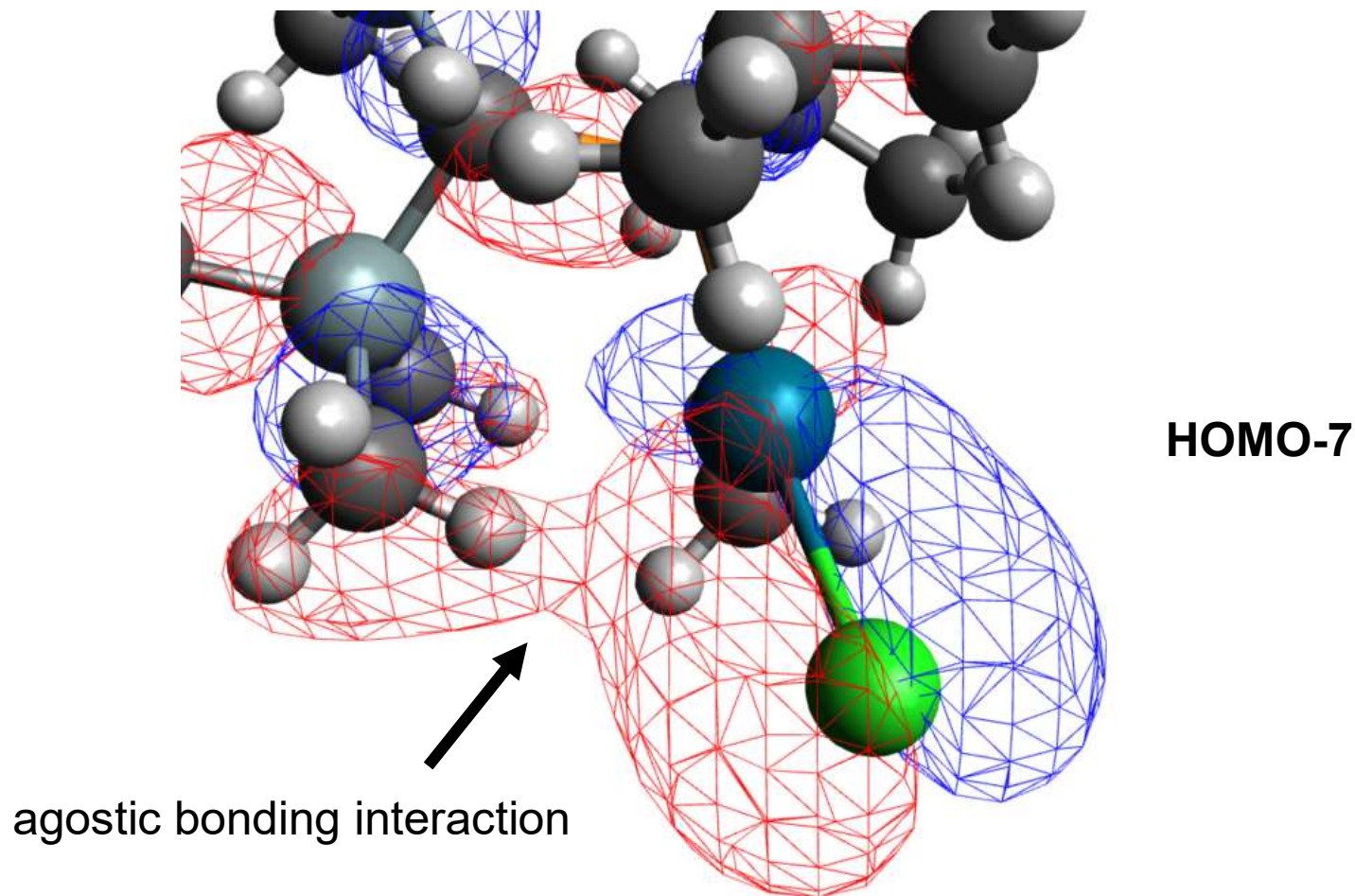


Atoms-in-molecules (AIM) analysis reveals an interaction in the apical position of the coordination sphere, suggesting an agostic interaction. Since the HOMO appears to antibonding character with respect to this interaction, a stronger apical agostic interaction should raise the HOMO energy and thus lower the HOMO-LUMO gap, redshifting  $\lambda_{\text{max}}$ .





## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{\text{MAX}}$ , AGOSTIC INTERACTIONS

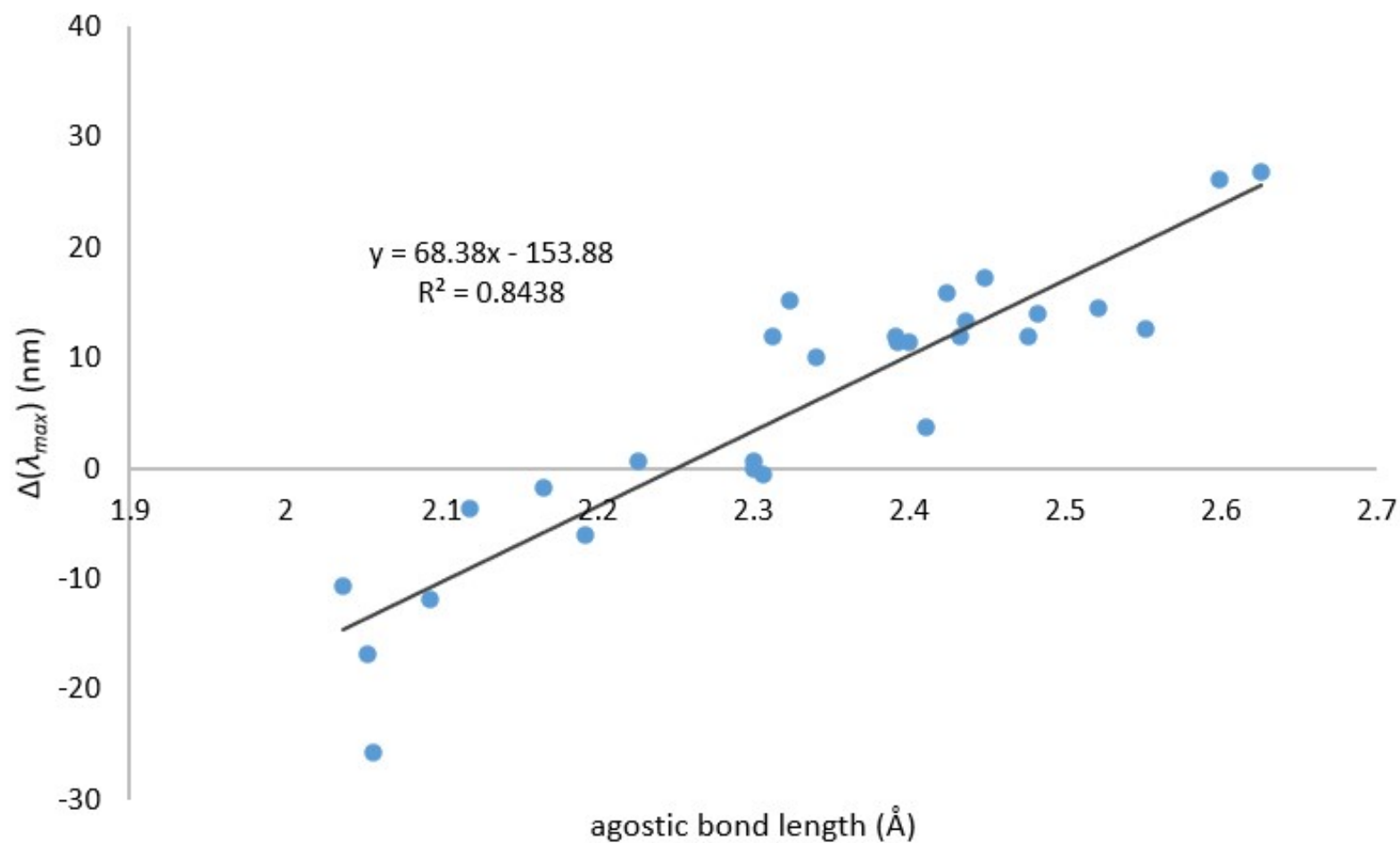


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## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{\text{MAX}}$ , AGOSTIC INTERACTIONS

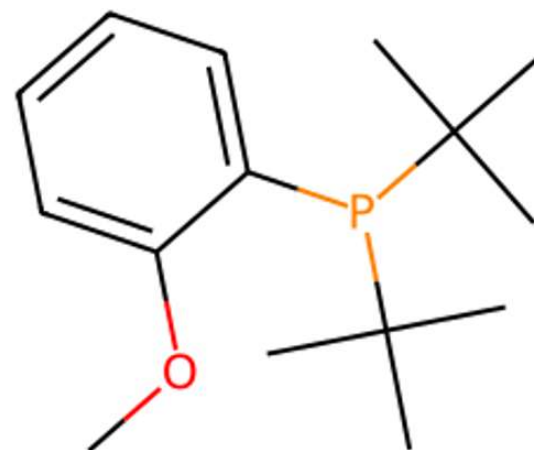
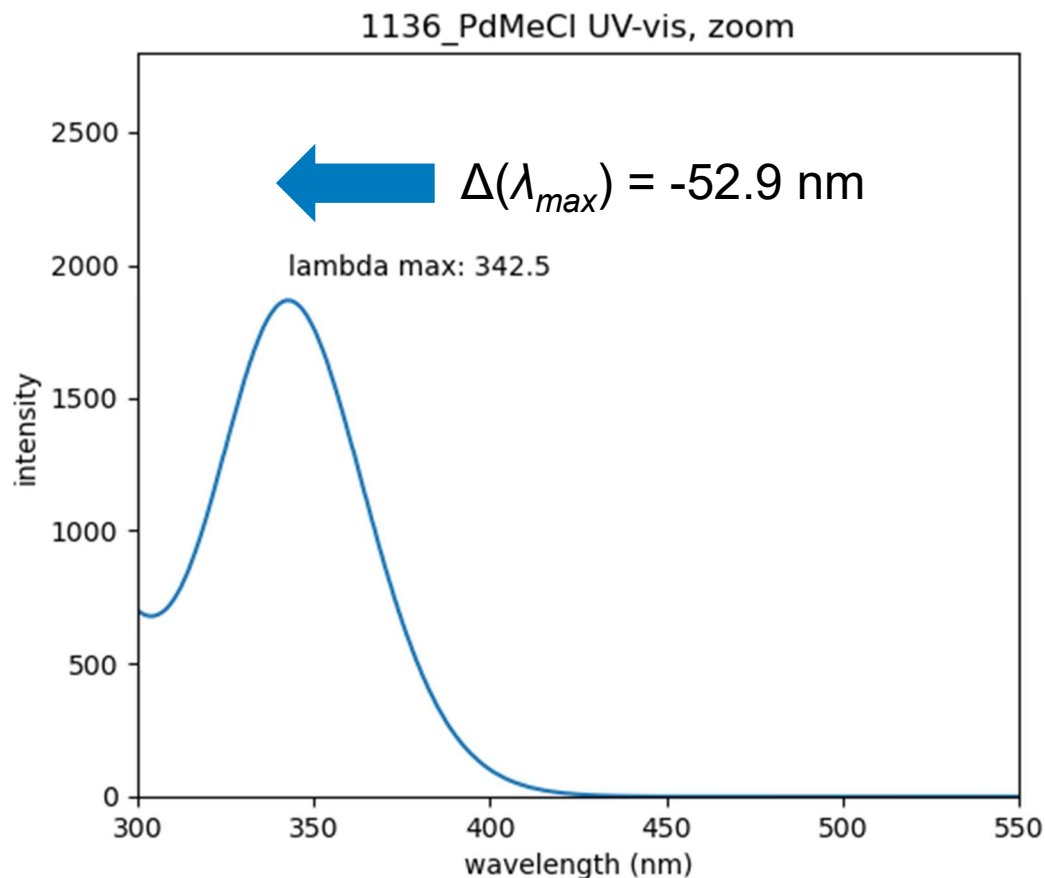
$\Delta(\lambda_{\text{max}})$  vs. agostic bond length, trialkylphosphine-like



Removing this outlier results in a much stronger correlation, of course.

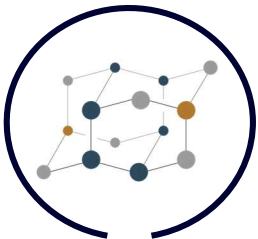


## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{MAX}$ , CHELATION

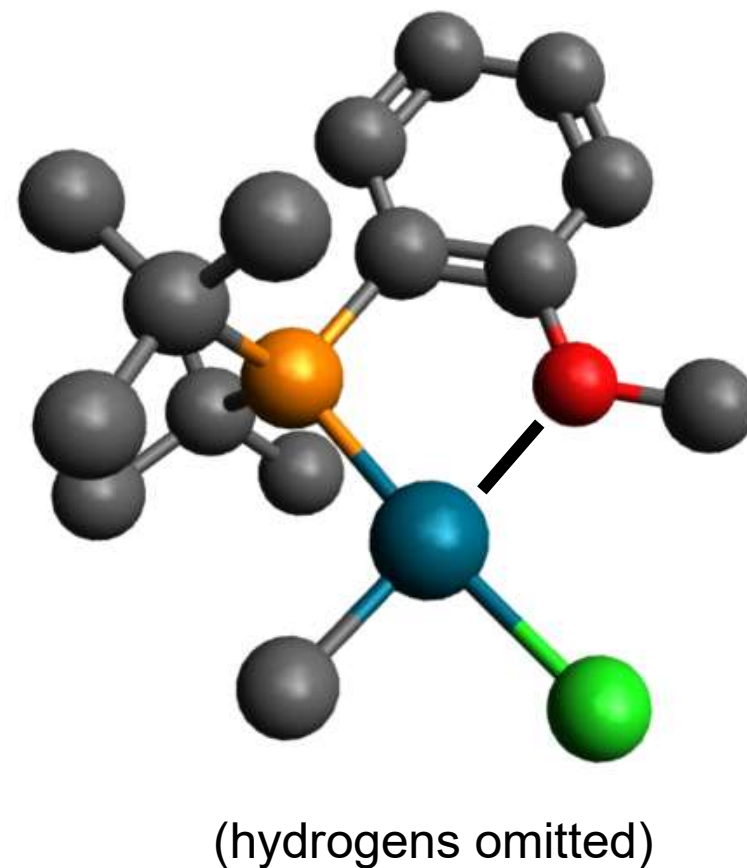
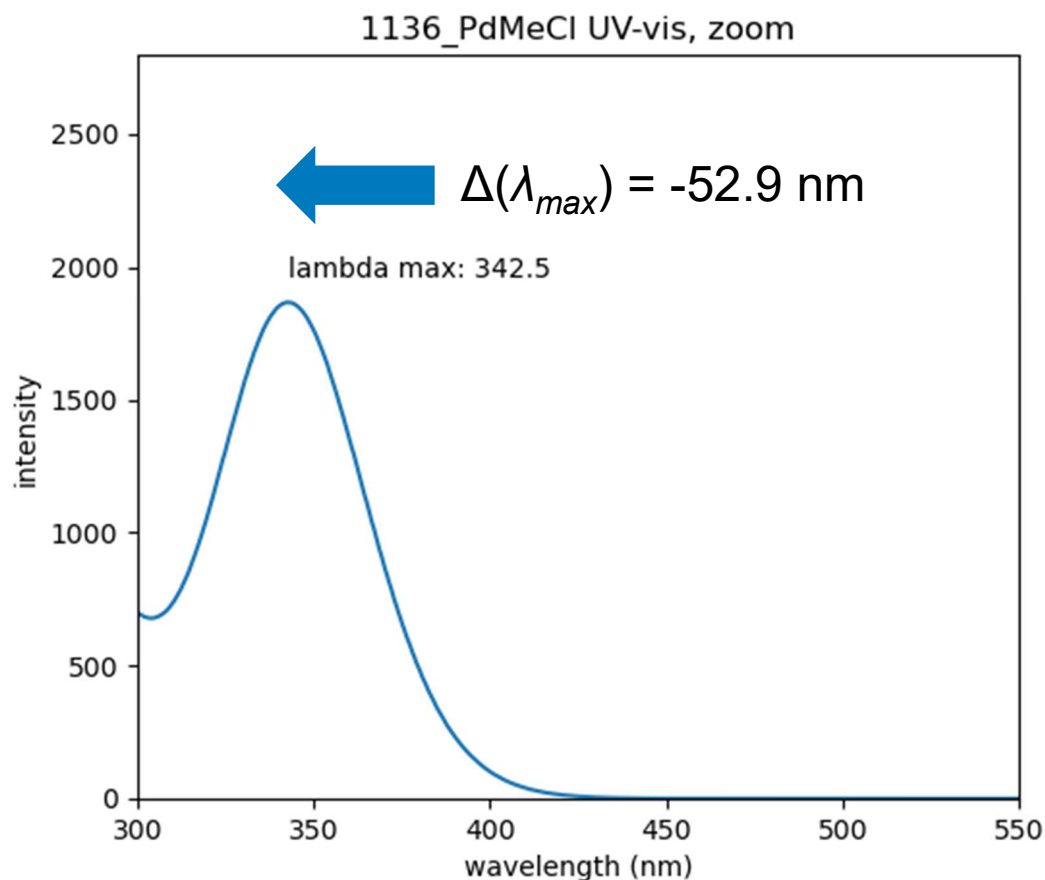


Ligand 1136

Furthermore, for ligands which have pendant heteroatoms that allow chelation, very large  $\lambda_{max}$  blueshifts are observed, again consistent with my hypothesis. Indeed, these complexes are in fact square planar rather than T-shaped.



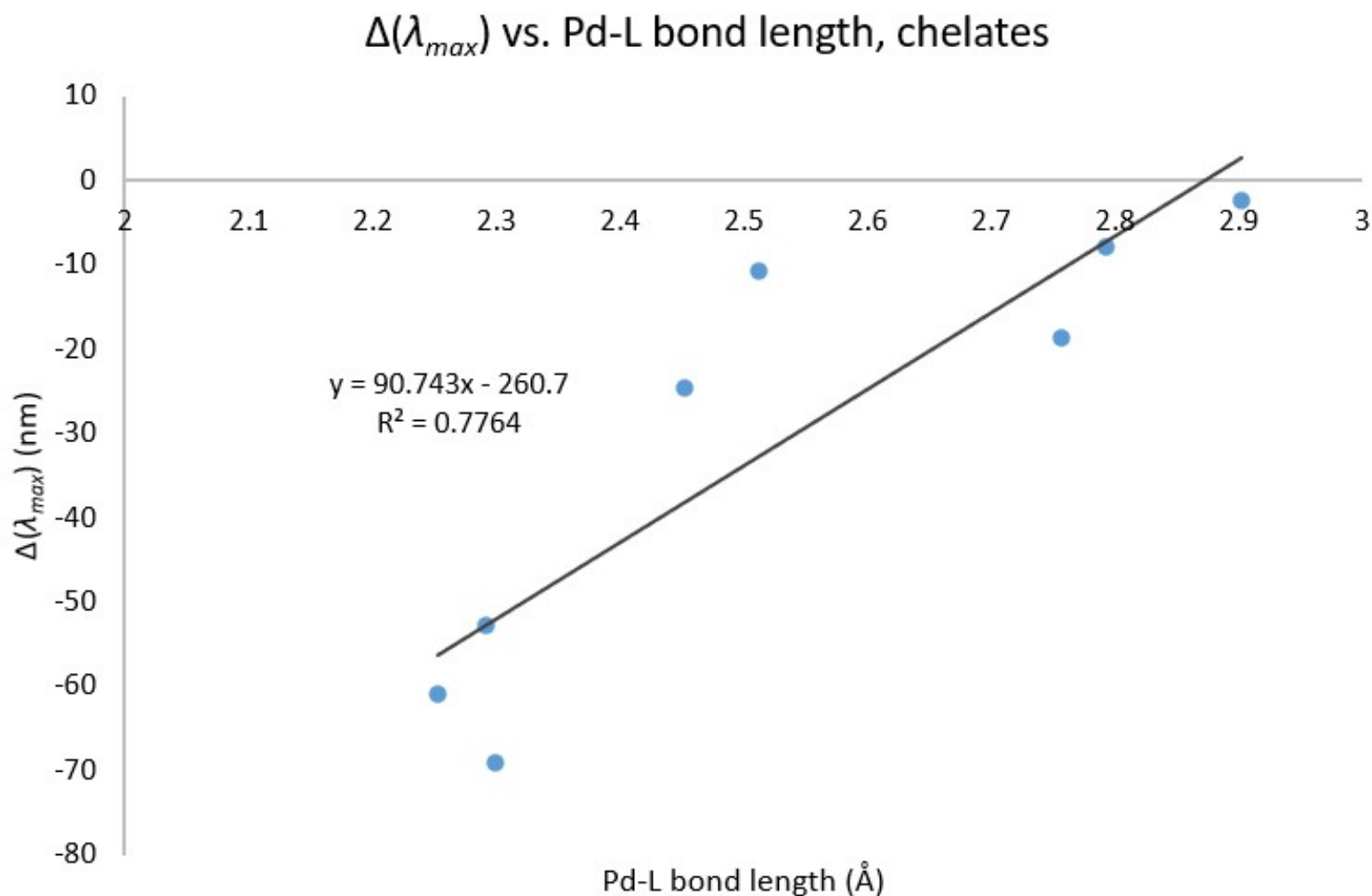
## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{max}$ , CHELATION



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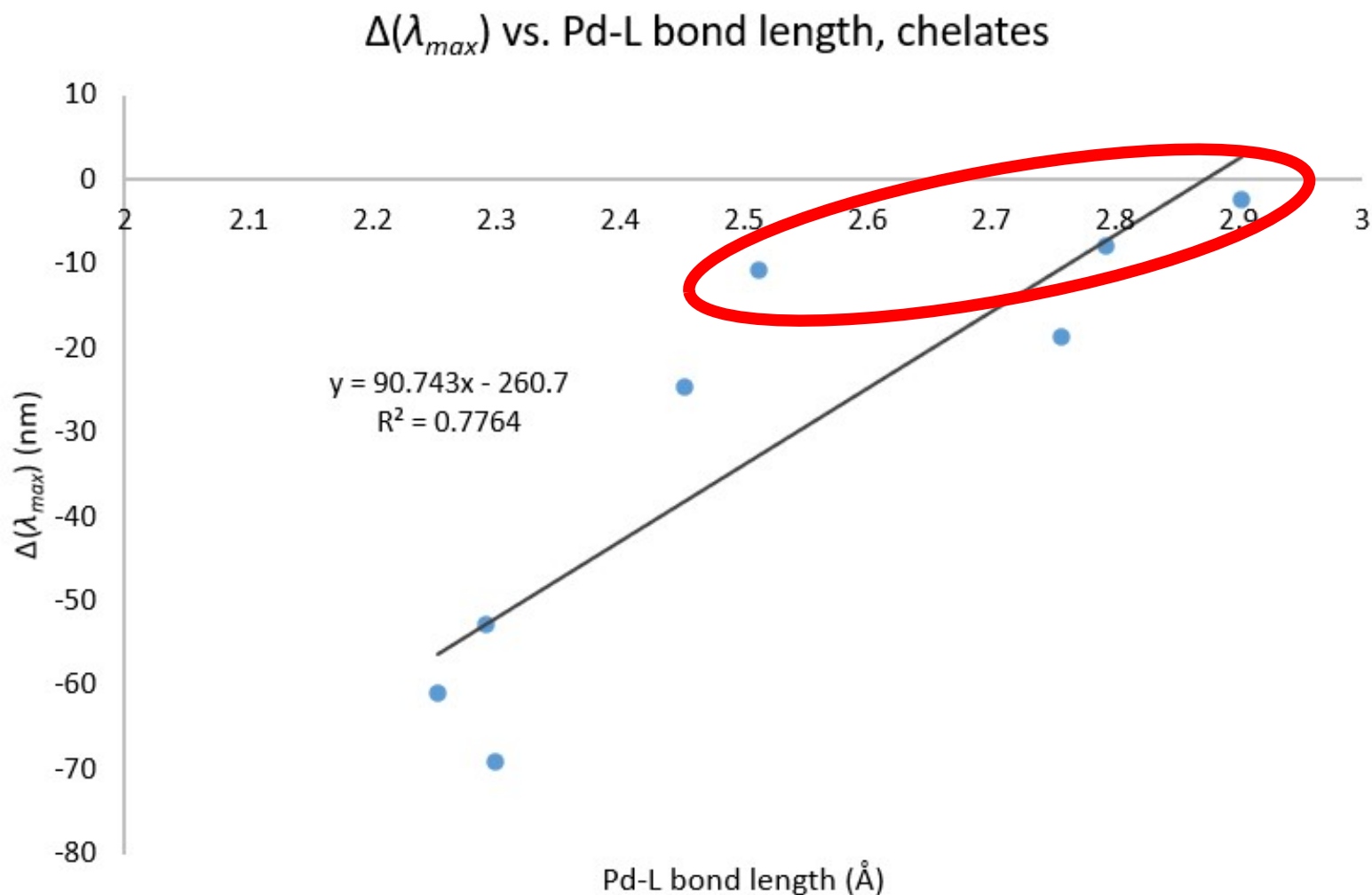
## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{max}$ , CHELATION



For ligands that form chelates with pendant L-type donors, the distance between Pd and the pendant donor atom correlates pretty well with  $\lambda_{max}$ , even across different types of donors (N, O, S, Cl).



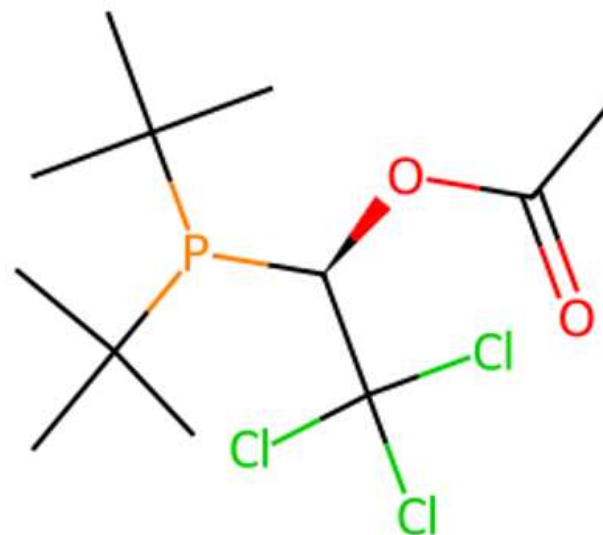
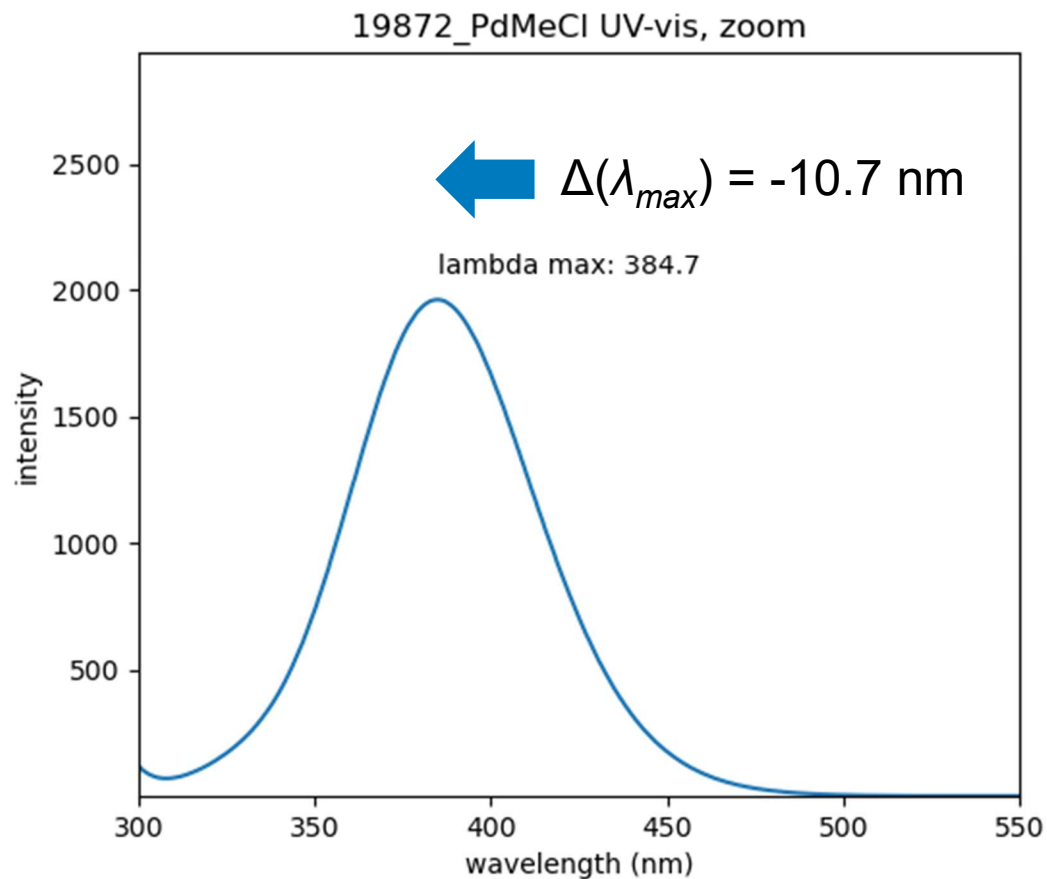
## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{max}$ , CHELATION



Interestingly, several of the apparently weaker chelates show a minimal blueshift in  $\lambda_{max}$ ! This suggests that Pd-C photocleavage could be accessible with visible light from *square planar complexes*!



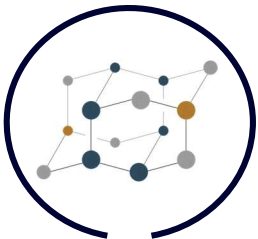
## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{\text{MAX}}$ , CHELATION



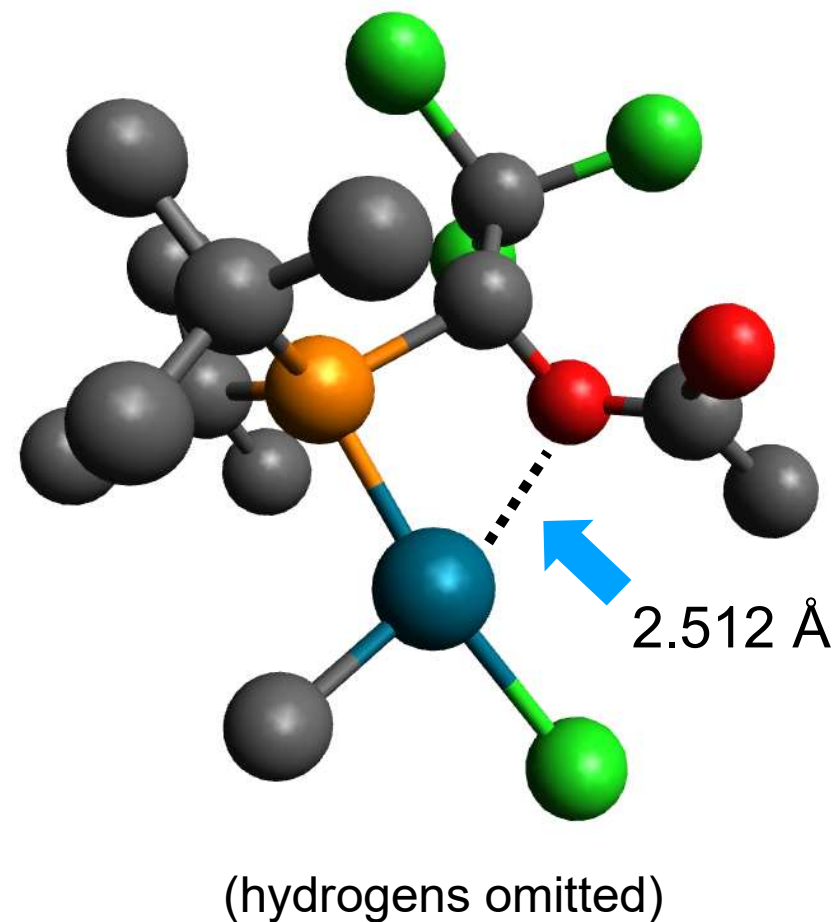
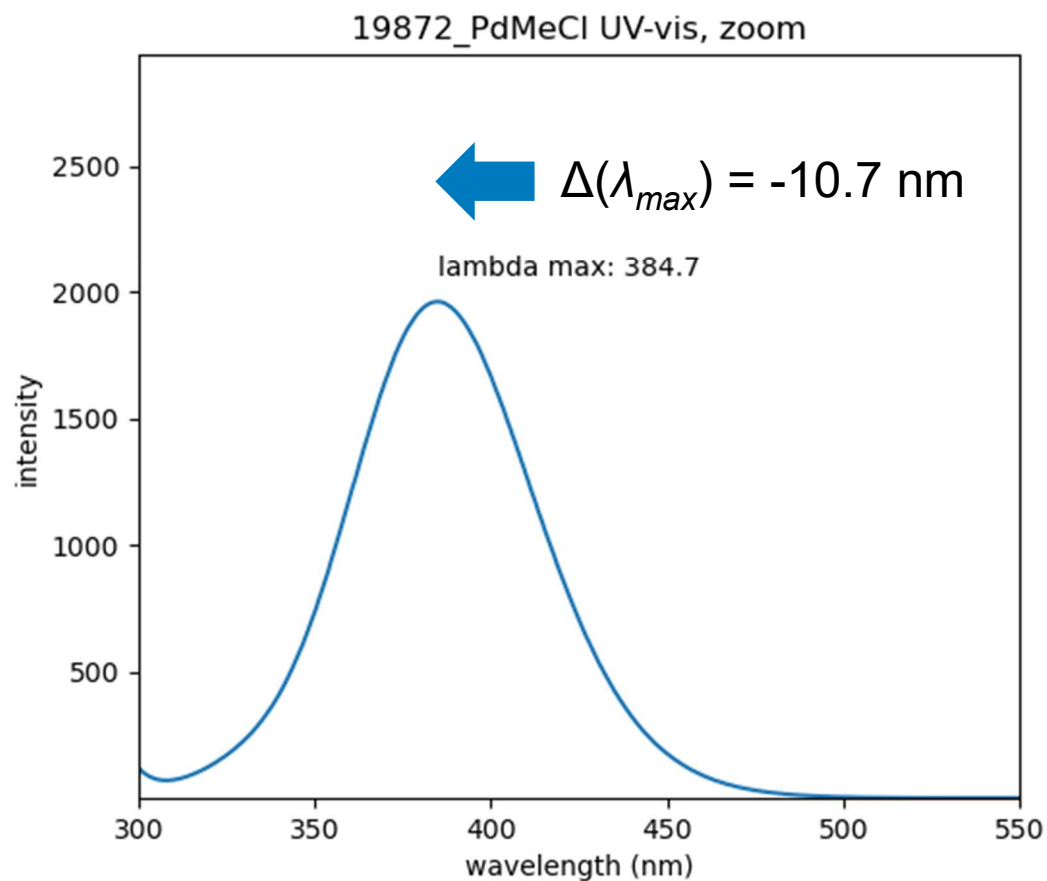
Ligand 19872

For example, consider this complex with a pendant ester has a  $\lambda_{\text{max}}$  which is blueshifted by only about 11 nm compared to tri-*t*-butylphosphine.



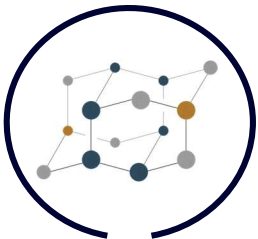


## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{max}$ , CHELATION

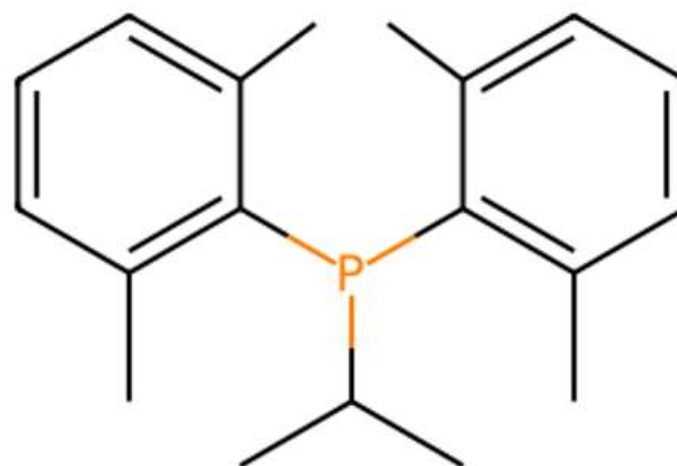
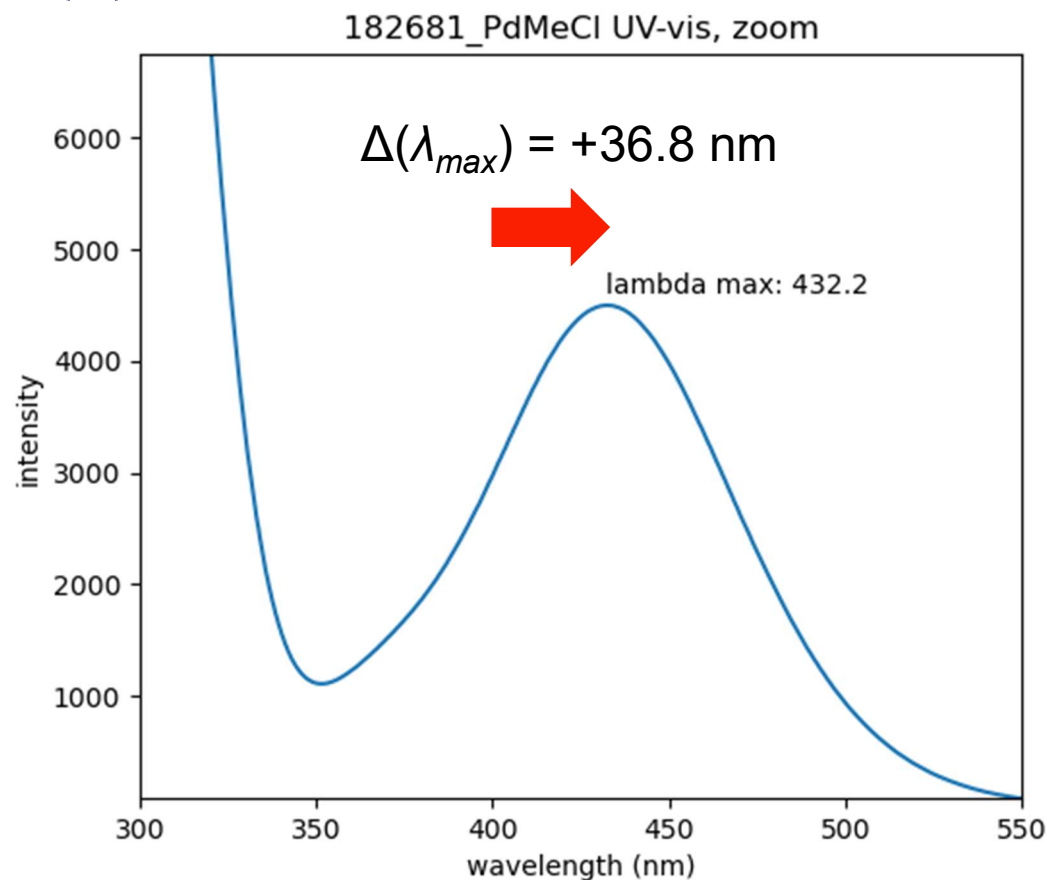


For example, consider this complex with a pendant ester has a  $\lambda_{max}$  which is blueshifted by only about 11 nm compared to tri-*t*-butylphosphine.





## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{\text{MAX}}$ , OPEN SITE

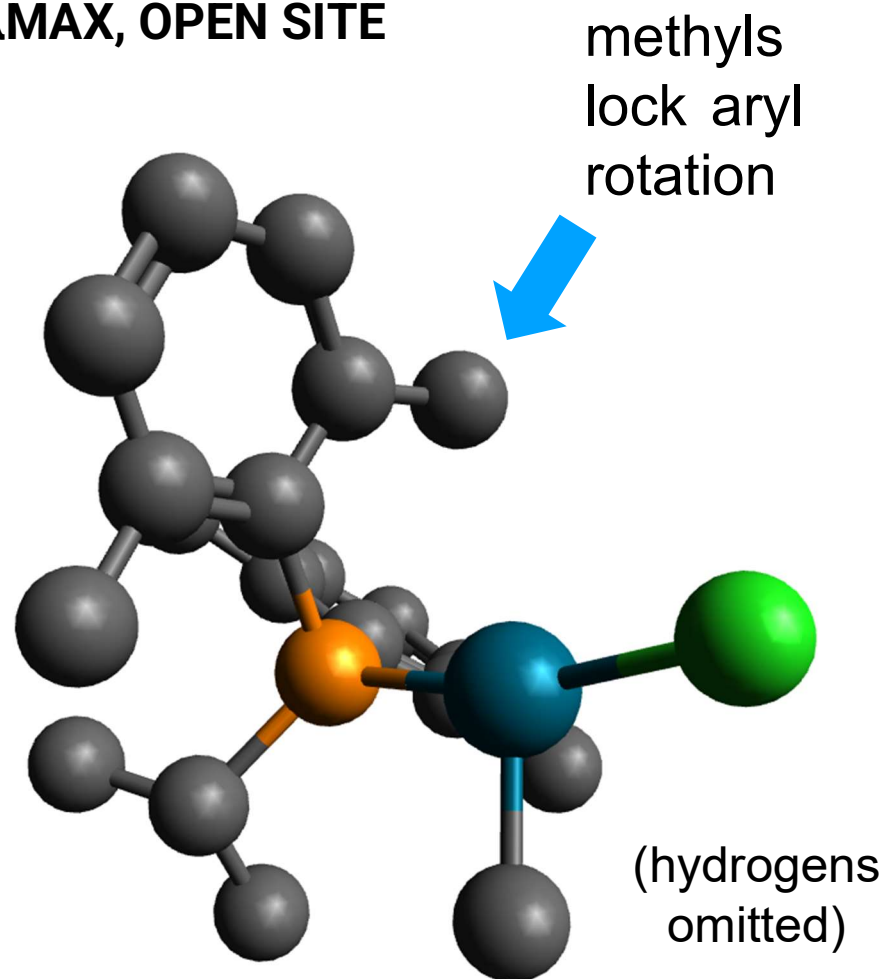
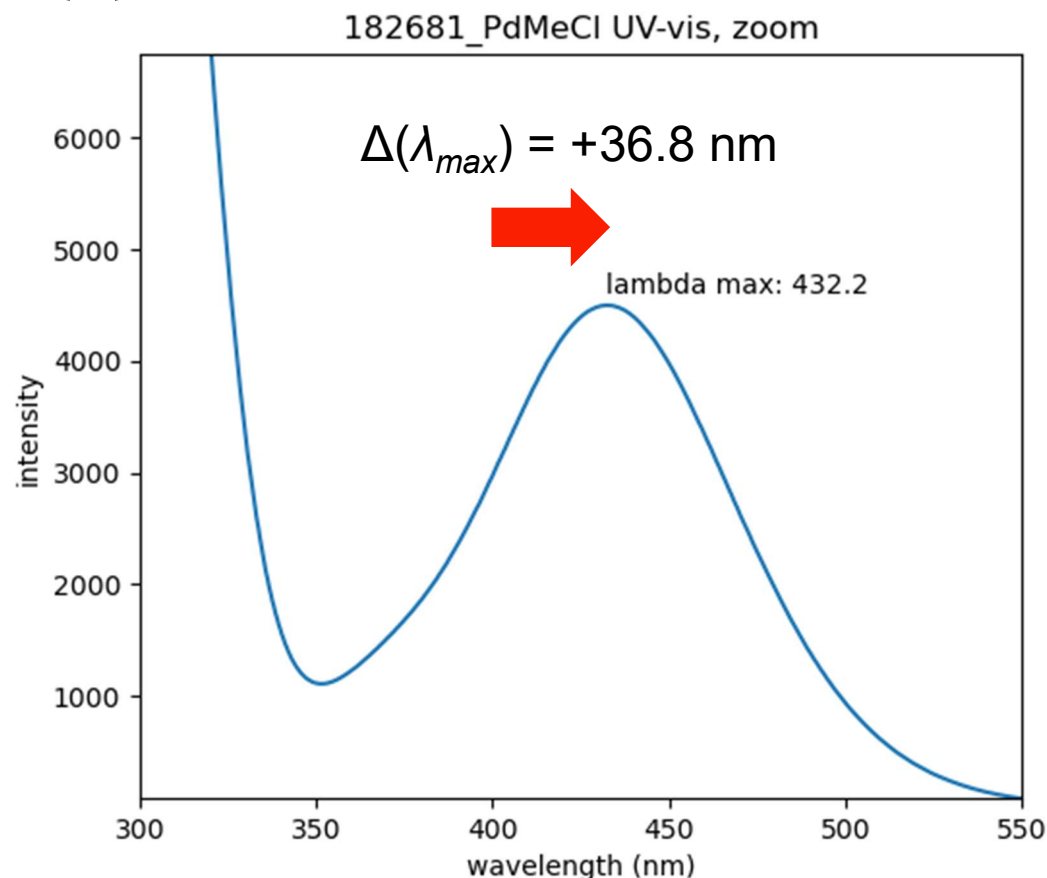


Ligand 182681

In some cases, the favored ligand conformation appears to leave the fourth site open entirely. In this limiting case,  $\lambda_{\text{max}}$  tends to be around 420-430 nm. However, in the condensed phase, this site would be open to occupation by solvent or e.g. bridging Cl ligand, so they may not be a good choice in practice for triggering this photochemistry with low energy light.



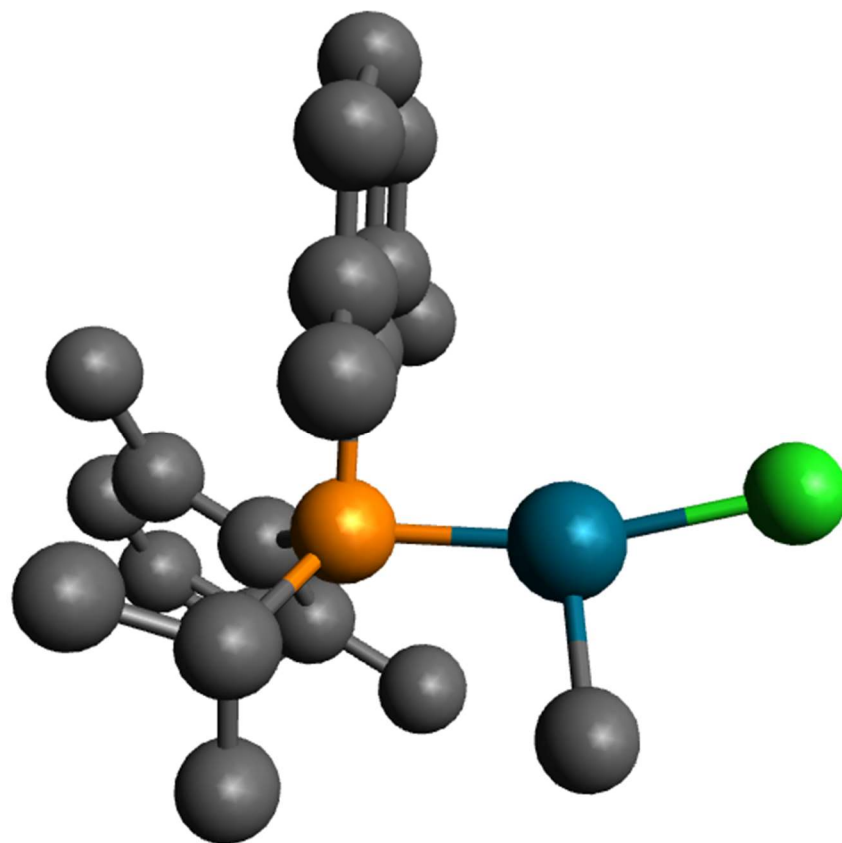
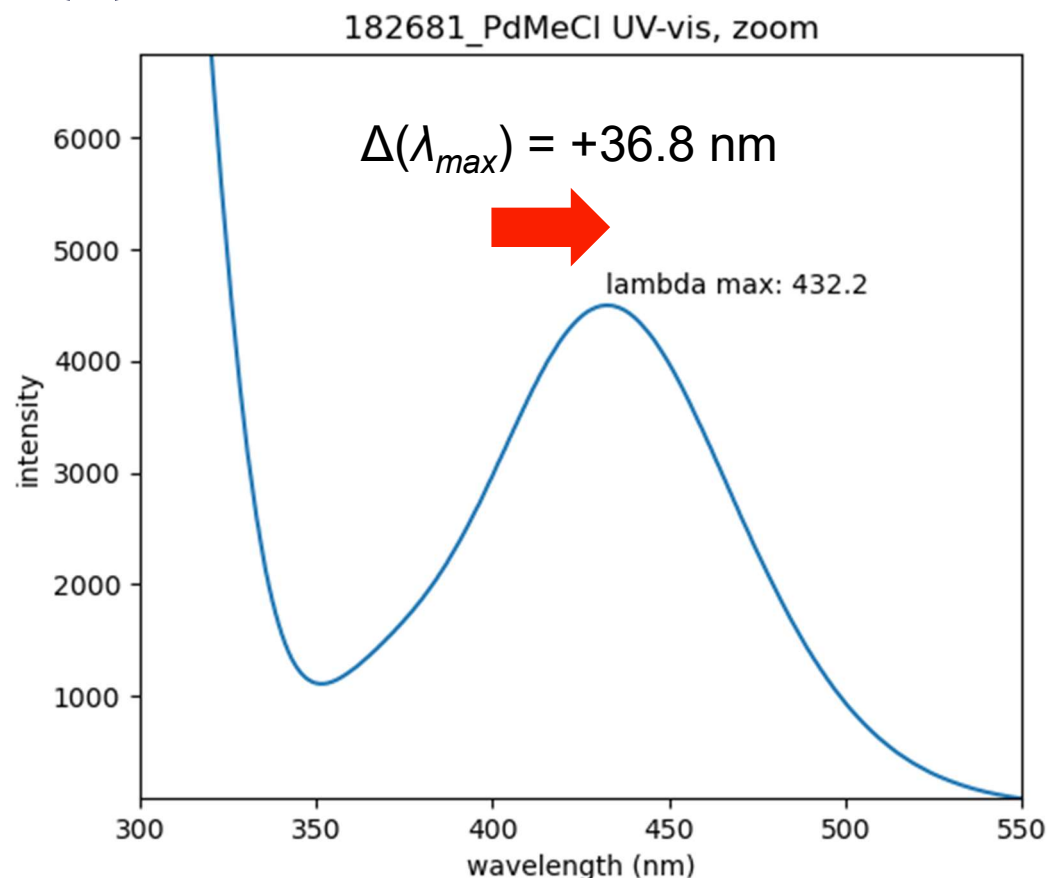
## INITIAL OBSERVATIONS: TRENDS IN $\lambda_{\text{MAX}}$ , OPEN SITE



In some cases, the favored ligand conformation appears to leave the fourth site open entirely. In this limiting case,  $\lambda_{\text{max}}$  tends to be around 420-450 nm. However, in the condensed phase, this site would be open to occupation by solvent or e.g. bridging Cl ligand, so they may not be a good choice in practice for triggering this photochemistry with low energy light.



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So, my hypothesis that the strength of the bonding interaction in the fourth site strongly influences the position of the peak in the absorption spectrum corresponding to the HOMO-LUMO electronic transition (and thus, the wavelength of light that can be used to cause Pd-C homolytic cleavage) seems to be very well validated.

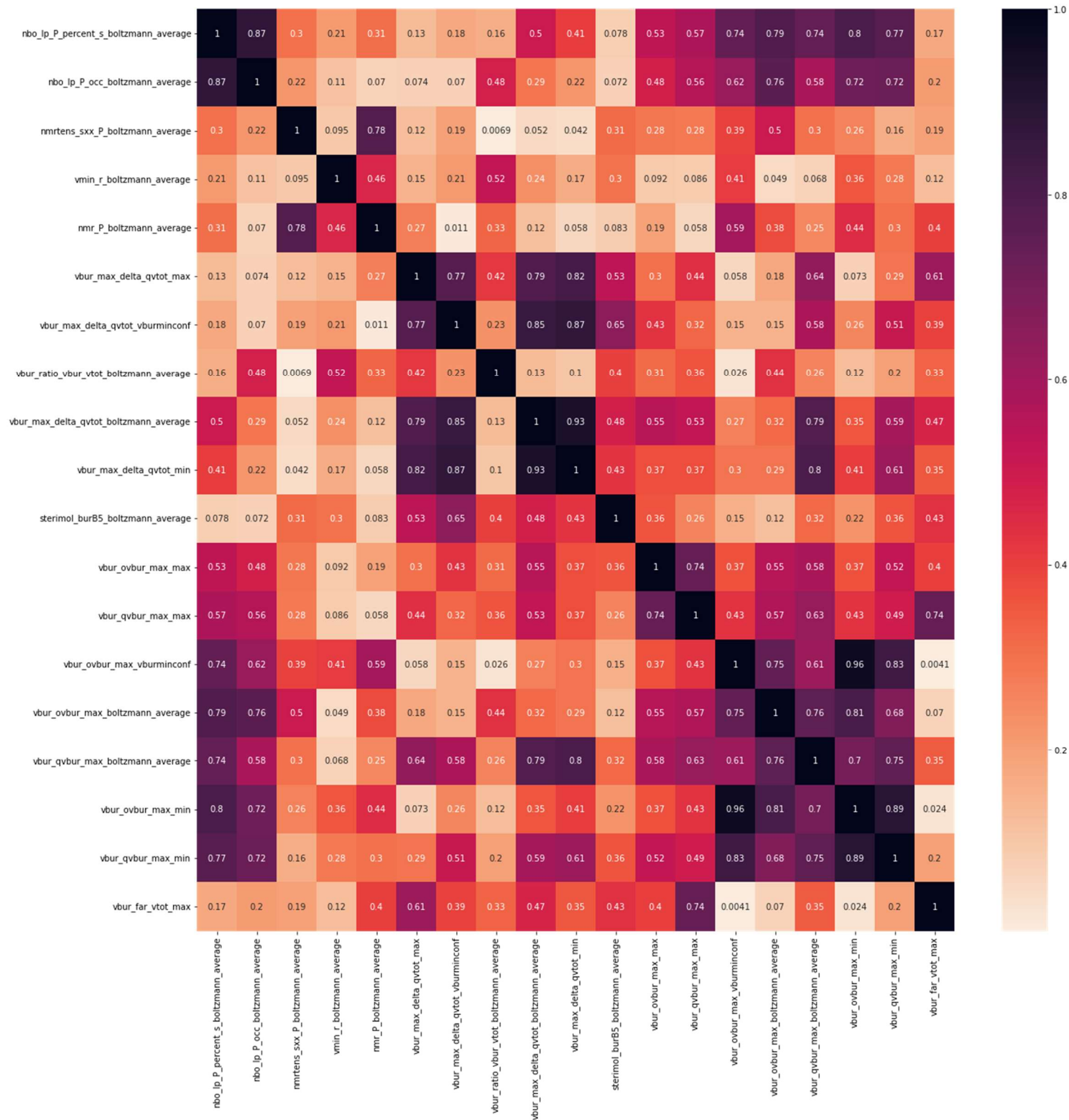
Now, what other factors can we find that influence  $\lambda_{max}$ ?



## MACHINE LEARNING MODELS TO PREDICT $\lambda_{\text{MAX}}$ : ELASTIC NET

- Our case is highly dimensional: each ligand has 192 features, while we have at most around 100 data points.
- To begin with, let us restrict ourselves to the top 20 features from our principal component analysis from earlier.
- Furthermore, there is significant multicollinearity in our feature set (see next slide)
- **Results to come!!**

For simplicity, the absolute correlation is shown here, with darker meaning more highly correlated.





**To be continued...**