The statistical reliability of 267 GHz JCMT observations of Venus: No significant evidence for phosphine absorption

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ABSTRACT

In the light of the recent announcement of the discovery of the potential biosignature phosphine in the atmosphere of Venus I present an independent reanalysis of the original JCMT data to assess the statistical reliability of the detection. Two line detection methods are explored, low order polynomial fits and higher order multiple polynomial fits. A non-parametric bootstrap analysis reveals that neither line detection method is able to recover a statistically significant detection. Similar to the results of other reanalyses of ALMA Venus spectra, the polynomial fitting process results in false positive detections in the JCMT spectrum. There is thus no significant evidence for phosphine absorption in the JCMT Venus spectra.

Key words: planets and satellites: atmospheres – methods:statistical- astrobiology

1 INTRODUCTION

Phosphine (PH₃) is a well-known trace constituent of gas giant atmospheres, first identified within Jupiter in the mid-infrared by Ridgway et al. (1976) and within Saturn in the millimetre and far-infrared by Weisstein & Serabyn (1994) and Davis et al. (1996). The main terrestrial sources of PH₃ are either biogenic (via anaerobic processes) or anthropogenic (via agricultural production and semiconductor doping), see further references in Sousa-Silva et al. (2020) for more details. The biogenic production route and the short chemical lifetime of PH₃ in an oxidising planetary atmosphere has led to the suggestion that PH₃ may be a potential biosignature in exoplanet atmospheres (Sousa-Silva et al. 2020).

Recently, Greaves et al. (2020) presented the potential detection of PH₃ in the atmosphere of Venus via a candidate spectral feature at 267 GHz with the JCMT and ALMA. PH₃ is not expected to be prevalent in the highly oxidised Venusian atmosphere. If this feature is truly identified with PH₃, it implies a concentration several orders of magnitude larger than abiogenic Venusian PH₃ production could sustain Bains et al. (2020). Excluding a wide range of asteroidal delivery and abiogenic production processes Greaves et al. (2020) and Bains et al. (2020) conclude that an unknown, potentially biogenic, atmospheric process could be at work.

The existence of a Venusian atmospheric biosphere was first proposed by Morowitz & Sagan (1967). It has been noted by many authors that at an altitude of 50 km, the Venusian atmosphere has a remarkably similar temperature and pressure to that of Earth. In the words of Landis (2003), "... viewed in a different way, the problem with Venus is merely that the ground level is too far below the one atmosphere level". If the candidate PH₃ feature identified by Greaves et al. (2020) is confirmed to be real and all abiogenic production routes can be ruled out this would lend important support to the Mo-

rowitz & Sagan (1967) hypothesis and be a potential step towards the discovery of a novel non-terrestrial biosphere.

Since the Greaves et al. (2020) claim, there has been much speculation as to the implication and significance of the results (e.g. Lingam & Loeb 2020; Siraj & Loeb 2020). Hein et al. (2020) have proposed a precursor mission for a balloon-based in-situ search for Venusian life, which could launch as early as 2022. However, the Greaves et al. (2020) results are not without controversy, due to the complex and difficult nature of the observations and assumptions within the chemical network of Bains et al. (2020).

An independent reanalysis of the ALMA Venus data was performed by Snellen et al. (2020) who showed that the analysis technique of Greaves et al. (2020) as applied to a spectrum dominated by non-Gaussian noise can result in false positive detections. A more conservative lower order baseline removal resulted in a candidate feature in the ALMA Venus spectrum of only 2σ . A further independent reanalysis of the ALMA Venus data has been conducted by Villanueva et al. (2020), who again concluded that the ALMA line detection is a false positive. Atmospheric modelling with a more realistic SO₂ vertical profile additionally shows that the depth of the JCMT line feature is entirely consistent with the nearby SO₂ transition (Villanueva et al. 2020).

However, the JCMT Venus spectra from Greaves et al. (2020) are also analysed by fitting multiple high order polynomials and the reported JCMT line feature may also be a false positive detection. Motivated by the investigation of Snellen et al. (2020) into the ALMA Venus spectrum this letter seeks to determine the statistical reliability of the original JCMT single-dish detection and reveal whether this line feature is a significant absorption line or not.

2 REANALYSIS OF JCMT DATA

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The reduction and analysis procedure for the JCMT Venus data is comprehensively described in Greaves et al. (2020). Here the basic steps are briefly given to illustrate our reanalysis. The aim is to follow as closely as possible the steps carried out by Greaves et al. (2020) by modifying the data reduction scripts supplied in that publication. The raw data were obtained from the JCMT Science Archive as described in Greaves et al. (2020) and are stored as time series spectral datacubes calibrated on the T_A^* scale. 512 channels were blanked from the edges of each spectrum to remove noise present at the edge of the passband. The top panel of Figure 1 shows a composite native velocity-resolution (0.0347 km s⁻¹ channels) spectrum obtained by integrating the Venus spectra along the time axis and dividing by the mean continuum value to obtain a line-to-continuum spectrum. The bottom panel of Figure 1 shows the same spectrum with a 3rd-order polynomial baseline fitted to and subtracted from the entire spectrum, additionally rebinned to 3.5 km⁻¹ channels to maintain a common velocity resolution with Greaves et al. (2020). It must be noted that these baseline-subtracted spectra should strictly be referred to as the "line-to-continuum ratio - 1", as they are zero-centred. However to maintain consistency with Greaves et al. (2020) we will refer to these spectra as line-to-continuum in the following text.

As Figure 1 shows, the JCMT spectra are affected by baseline ripples, which are likely to be caused by reflections between the main and cold-load dewar, an unidentified surface in the receiver cabin and the well-known "JCMT 16-MHz ripple" caused by a standing wave between the secondary mirror and the receiver cabin¹. This underlines the extreme difficulty of identifying faint spectral features in the presence of a strong background continuum source.

2.1 Low order polynomial fits

The simplest way to identify a spectral line in a spectrum with a complex baseline is to fit a low order polynomial to spectral channels bracketing the suspected line position. Greaves et al. (2020) carried out such a simple approach as an independent test of their high order polynomial fits by integrating (or collapsing in JCMT parlance) the time-series spectral cubes along the time axis and fitting a low order polynomial. It is reported that the phosphine line was recovered with a lower signal-to-noise-ratio than the higher order fitting approach (Greaves et al. 2020).

The data reduction scripts presented by Greaves et al. (2020) were modified to repeat this simple approach and investigate the significance of the line recovery with a single low order polynomial. Polynomials of 3rd and 4th-order were fitted to the spectrum shown in the top panel of Figure 1. To investigate the effects of different fitting ranges these fits were carried out over varying width spectral regions centred on the PH₃ Venus rest-frame velocity. A line region of $|\Delta v| = 5 \text{ km s}^{-1}$ centred on the PH₃ J=1–0 velocity was excluded from the fit, following Greaves et al. (2020). The fitted spectra were then rebinned to a common channel width of 3.5 km s⁻¹.

Figure 2 shows the results of these polynomial fits. The line search window of $|\Delta v| = 5 \text{ km s}^{-1}$ that was excluded from the polynomial fit is indicated by the central unshaded region in each plot. As can be seen, the result of the fit is sensitive to both the velocity range of channels used in the fitting process and the order of the polynomial used. A low-significance single-channel line feature at the PH₃ rest frame Venus velocity (with SNR of 2.6) is only recovered with a 4th-order fit over our smallest velocity range ($|\Delta v| \le 30 \text{ km s}^{-1}$).

However, the ability of a polynomial fitting routine to incorrectly identify a negative or positive baseline deviation as a spectral line is



Figure 1. *Top:* Line-to-continuum spectrum of Venus obtained by integrating the individual time-series spectra and dividing by the mean continuum level of each spectrum. *Bottom:* the above line-to-continuum spectrum with a 3rd-order polynomial subtracted and denoted as line-to-continuum ratio – 1. The inset lower velocity resolution spectrum is the result of rebinning to 3.5 km s^{-1} channels. The dotted orange lines represent the $\pm 1\sigma$ standard deviation of each 3.5 km s^{-1} channel, calculated from the spread of the values from the individual spectra, revealing substantial dispersion within each channel over time.

magnified by giving the routine fewer constraints and a higher degree of freedom (e.g. a smaller velocity range and/or a higher order fit; Snellen et al. 2020). As also noted by Snellen et al. (2020), the non-Gaussian nature of the spectral noise makes line identifications more difficult. The usual approach of using the statistical *z*-score (or signal-to-noise ratio) of the peak line temperature becomes meaningless when the noise spectrum does not follow a normal or Gaussian distribution.

Non-parametric methods are an ideal way to investigate the significance of measurements in a non-Gaussian sample, of which one example is the bootstrap resampling procedure. This process was used to analyse the significance of the recovered line feature in the following way. A Python wrapper was written around the modified **Greaves et al.** (2020) scripts to permit polynomial baseline fits to be repeatedly carried out for different $|\Delta v| = 30 \text{ km s}^{-1}$ velocity windows centred on each channel of the native-resolution Venus spectrum. Within each window we identified the minimum value within the central $|\Delta v| = 5 \text{ km s}^{-1}$ line region excluded from the fit.

This approach is analogous to the non-parametric bootstrap method, resampling velocity windows from the original raw spectrum and carrying out the same baseline-fitting and line-search ap-

¹ https://www.eaobservatory.org/jcmt/instrumentation/heterodyne/observingmodes/



Figure 2. The results of 3rd-order (*left column*) and 4th-order (*right column*) polynomial fits to velocity windows centred on the PH₃ Venus rest-frame velocity. Spectra are offset vertically for clarity. The width of each velocity window is indicated in the top right of each spectrum and the signal-to-noise ratio of any identified absorption feature in the central line window is indicated in the top left. Standard deviations were determined from the same line-free channels in each spectrum, as shown by shaded regions. Horizontal and vertical dashed lines indicate the baseline and velocity origins

proach to each window. As such, this method is more powerful than the arbitrary sample of velocities presented in Figure 2 of Snellen et al. (2020), as the fraction of realisations that exceed a set peak line-to-continuum ratio can thus be used to estimate the probability that a particular line feature is significant (as a pseudo probability density function or PDF).

It is important to note that the statistical power of this technique is limited due to the fact that there are only ~4500 possible contiguous velocity windows that can be resampled from the native-resolution 0.0347 km s^{-1} spectrum. Greater statistical power could be obtained by relaxing the constraint that each window must be contiguously drawn from the original spectrum, instead filling each window with randomly chosen channels from the entire original spectrum. However this would randomize any underlying ripples in the baseline structure and it was felt that greater statistical power was not warranted in this case.

Figure3 shows a histogram created from our bootstrap realisations, plotting the frequency of the minimum values of line-to-continuum obtained in the line-search region. The distribution of these values is distinctly non-Gaussian, however it can be seen that the observed depth of the candidate PH₃ absorption line lies towards the centre of the distribution. In fact 65% of our bootstrap realisations produce a deeper feature than the candidate PH₃ absorption line. The null hypothesis that the candidate PH₃ feature is drawn from a random sample from the spectrum thus cannot be excluded.



Figure 3. Histogram of the peak line-to-continuum features extracted from the bootstrap process. The vertical dotted line represents the peak line-to-continuum ratio of the line feature detected at the PH₃ rest velocity in the 4th-order polynomial fit shown in Figure 2 (-1.66×10^{-4}) .

2.2 Higher order polynomial fits

The main analysis procedure used in Greaves et al. (2020) to identify the candidate PH₃ absorption line involves repeated multiple fitting of polynomials to remove a complex spectral baseline. The chosen order of each polynomial are motivated by particular instrumental and observational effects. An initial 4th order baseline is fitted to each observed spectrum to remove a ripple caused by reflections between main and cold load dewars. A further 9th order polynomial is subtracted from a boxcar-smoothed spectrum to remove a reflection from within the receiver cabin. Finally, an 8th order polynomial is subtracted from a 100 km s⁻¹ wide window between -36-64 km s⁻¹ to remove baseline ripples of similar width to the expected line.

Snellen et al. (2020) have shown that 12th-order fits to the ALMA Venus spectrum result in false positive detections at arbitrarily chosen line-free velocities. This effect is caused by the exclusion of the central line region in the polynomial fit. A polynomial of progressively higher order has more freedom to interpolate across the excluded region and thus magnify the significance of spectral features.

A similar bootstrapping process to that described in Section 2.1 was carried out in order to assess the reliability of the candidate PH₃ spectral feature presented in Greaves et al. (2020) and to test whether the same false positive effects are present in the JCMT data analysis. A Python wrapper script was used to repeatedly apply the Greaves et al. (2020) reduction script to 100 km s⁻¹ wide windows centred on different central channels drawn from the original spectra. The minimum of each feature within the $|\Delta v| = 5$ km s⁻¹ line region excluded from the fit was then stored.

The results of this bootstrap analysis are displayed in Figures 4 and 5. Figure 4 presents a composite heat-map spectrum of a sample of realisations drawn from the bootstrap analysis. Only 10% of realisations are shown for clarity, drawn uniformly across the distribution of central velocities. The PH₃ Venus rest-frame velocity



Figure 4. Composite heat-map spectrum of different realisations from the bootstrap resampling process. Only 10% of realisations are shown for clarity. All spectra have been shifted to a common velocity frame centred on the middle of each 100 km s^{-1} window used for the baseline fitting. The spectrum at the PH₃ Venus rest-frame velocity is indicated by a thick solid black line.

realisation is shown as a solid thick black line, and can be seen to be a close match to that in Greaves et al. (2020) with minor differences likely to be caused by slightly different channelisations. However, it can also be seen that the depth of the candidate absorption line in this realisation does not appear significant compared to the other realisations. There are many realisations with deeper candidate absorption line features and, indeed, candidate emission line features.

Figure 5 shows the results of all the bootstrap realisations, again as a histogram of the frequency of the minimum values of line-tocontinuum obtained in the line-search region. It can be seen that the depth of the candidate PH₃ feature is towards the centre of the distribution. The peak line-to-continuum ratio of this feature is more significant than that obtained by low order polynomial fitting. However 25% of our bootstrap realisations produce deeper spectral features than the candidate and the null hypothesis cannot be excluded with any significance. It must be concluded that the candidate line detection presented by Greaves et al. (2020) is not statistically significant.

3 CONCLUSIONS

An independent reanalysis of the 267 GHz JCMT observations of Venus (Greaves et al. 2020) is presented. Two different line recovery methods were tested: a simple approach fitting low-order polynomials to line-free channels around the position of the PH₃ candidate; and the more complex multiple high-order polynomial approach used by Greaves et al. (2020). Similarly to the independent ALMA reanalysis of Snellen et al. (2020) and Villanueva et al. (2020) it is found that the high-order polynomial fitting applied to the JCMT spectra also results in false positive detections.

The statistical significance of identified line features was explored by a bootstrap analysis with the result that neither low-order fitting



Figure 5. Histogram of the peak line-to-continuum features extracted from the bootstrap resampled multiple polynomial fits described in Section 2.2. The vertical dotted line represents the peak line-to-continuum ratio of the line feature detected at the PH₃ rest velocity in the 4th-order polynomial fit shown in Figure 4 (2.38×10^{-4}).

or the high-order fitting method result in statistically significant detections. The null hypothesis that the line candidates are drawn from a random line-to-continuum sample within the spectrum cannot be excluded. This means that there is no significant evidence for either PH_3 or SO_2 absorption within this data.

The investigation carried out in this paper (and also Snellen et al. 2020; Villanueva et al. 2020) underlines the difficulty of identifying line features against bright continuum sources and the need for careful and rigorous analysis to exclude false positive detections.

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DATA AVAILABILITY

The raw data analysed here was observed under JCMT Service Program S16BP007 and can be obtained from the JCMT Science archive hosted at the Canadian Astronomical Data Centre.

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