



## Low Upper Limit to Methane Abundance on Mars

Christopher R. Webster *et al.*

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# Low Upper Limit to Methane Abundance on Mars

Christopher R. Webster,<sup>1\*</sup> Paul R. Mahaffy,<sup>2</sup> Sushil K. Atreya,<sup>3</sup> Gregory J. Flesch,<sup>1</sup> Kenneth A. Farley,<sup>4</sup> MSL Science Team†

By analogy with Earth, methane in the Martian atmosphere is a potential signature of ongoing or past biological activity. During the past decade, Earth-based telescopic observations reported “plumes” of methane of tens of parts per billion by volume (ppbv), and those from Mars orbit showed localized patches, prompting speculation of sources from subsurface bacteria or nonbiological sources. From in situ measurements made with the Tunable Laser Spectrometer (TLS) on Curiosity using a distinctive spectral pattern specific to methane, we report no detection of atmospheric methane with a measured value of  $0.18 \pm 0.67$  ppbv corresponding to an upper limit of only 1.3 ppbv (95% confidence level), which reduces the probability of current methanogenic microbial activity on Mars and limits the recent contribution from extraplanetary and geologic sources.

Methane is the most abundant hydrocarbon in our solar system and is found in the atmospheres of several planets and satellites (1). On Earth, 90 to 95% of atmospheric methane is biologically produced, either from extant or fossil sources, and it is easy to identify and quantify with confidence by using spectroscopic methods (2). For Mars, three possible origins have been proposed: geologic, biotic, and exogenous (3–5). Over the past decade, there have been several reports of methane detection from Earth and from Mars orbit. Observations with the Canada-France-Hawaii Telescope (CFHT) found a global average value of  $10 \pm 3$  parts per billion by volume (ppbv) (5). The Planetary Fourier Spectrometer (PFS) on the Mars Express (MEX) spacecraft found a global average abundance of  $10 \pm 5$  ppbv (4), later updated (6) to 15 ppbv, with indications of discrete localized sources (4) and a summer time maximum of 45 ppbv in the north polar region. A search for methane from the Infrared Telescope Facility (IRTF) and the Keck-2 telescope reported methane release in plumes (7) from discrete sources in Terra Sabae, Nili Fossae, and Syrtis Major, with the largest plume containing 19,000 tons of CH<sub>4</sub> in March 2003; seasonal changes with a summer time maximum of ~45 ppbv near the equator were seen. Methane abundances later retrieved (8) from a second instrument in Mars orbit, the Thermal Emission Spectrometer (TES) of the Mars Global Surveyor (MGS), reported methane abundances as intermittently present (1999 to 2003), ranging from 5 to 60 ppbv in locations where favorable geological conditions such as residual geothermal activity (Tharsis and Elysium) and strong hydration (Arabia Terra) are expected. More recent observations report methane mixing ratios that

have diminished considerably since 2004 to 2006 to upper limits of 7 to 8 ppbv (9–11), suggesting a very short lifetime for atmospheric CH<sub>4</sub> and contradicting the MEX claim that methane persisted from 2004 to 2010. Ground-based observations favor episodic injection of methane in 1999 and 2003, 10 ppbv at Valles Marineris in February 2006 (9, 11), and <8 ppbv in January 2006 (10), 2009, and 2010; whereas orbital data from PFS and TES suggest a more regular behavior with latitudinal, seasonal, and interannual variabilities. At Curiosity’s Gale Crater landing site (4.5°S, 137°E), published maps of PFS data (6) show an increase from ~15 ppbv in fall to ~30 ppbv in winter, whereas the TES trend (8) is opposite: ~30 ppbv in fall and ~5 ppbv in winter.

The Tunable Laser Spectrometer (TLS) of the Sample Analysis at Mars (SAM) (12, 13) instrument suite on the Curiosity rover has a spectral resolution— $0.0002 \text{ cm}^{-1}$ , which is far superior to those of the ground-based telescopic and orbiting spectrometers—that offers unambiguous identification of methane in a distinct fingerprint spectral pattern of three well-resolved adjacent <sup>12</sup>CH<sub>4</sub> lines in the 3.3 μm band (Fig. 1). The in situ technique of tunable laser absorption in a closed sample cell is simple, noninvasive, and sensitive. TLS is a two-channel tunable laser spectrometer that uses both direct and second harmonic detection of infrared (IR) laser light. One laser source is a near-IR tunable diode laser at 2.78 μm that can scan two spectral regions containing CO<sub>2</sub> and H<sub>2</sub>O isotopic lines that have been used to report <sup>13</sup>C/<sup>12</sup>C, <sup>18</sup>O/<sup>17</sup>O/<sup>16</sup>O, and D/H ratios in the Martian atmosphere (13). The second laser source is an interband cascade (IC) laser at 3.27 μm used for methane detection alone, scanning across seven rotational lines that includes the R(3) triplet used in this study (Fig. 1 and table S1). The IC laser beam makes 81 passes of a 20-cm-long sample cell of the Herriott design fitted with high-vacuum microvalves that allow evacuation with a turbomolecular pump for “empty cell” scans or filled to Mars ambient pressure (~8 mbar) for “full cell” runs. During data collection, the cell and other optics are kept at  $47 \pm 3^\circ\text{C}$  by using a heater that thermally stabilizes the cell but is ramped up and

down within these temperature limits in order to increase gas sensitivity by spoiling the accumulation of optical interference fringes during the 2-min period of spectrum collection. Our methane determination is made by comparing the measured methane abundances in our sample cell when filled with Mars atmosphere with those of the same cell evacuated, as detailed in (14). The laser scans every second through the methane spectral region, and each spectrum is co-added on board to downlink sequential 2-min-averaged spectra during a given run of ~1 to 2 hours in duration. Typically, we recorded 26 2-min “empty cell” spectra followed by 26 2-min “full cell” spectra, then finally five additional 2-min empty cell spectra. For each 2-min spectrum, we retrieved methane abundances from three spectral lines (14) individually and combined the results so as to produce a weighted average value. By subtracting all retrieved abundances (full and empty cell) from the empty cell mean value for that sol (one Martian solar day) run, we were left with 31 differences for the empty cell and 26 for the full cell. For our statistical analysis, we analyzed the empty cell and full cell differences for all the sols taken as one data set (14). For sols 79, 81, 106, and 292, the foreoptics chamber contained residual terrestrial air (Table 1, pressures), including CH<sub>4</sub>, that produced absorption line signals in the sample cell detector channel, as described in (14). For sols 306 and 313, the foreoptics was evacuated. Both the sample cell and foreoptics chamber have pressure and temperature sensors. This experiment has been repeated on six separate Martian sols to date (Martian sols 79, 81, 106, 292, 306, and 313 after landing in August 2012). The inlet to the TLS is a stainless steel tube (14) heated to 50°C and located on the rover side ~1 m above the Martian surface and was pointed at a variety of directions relative to the nominal wind direction. Mars atmospheric gas was ingested during the night for sols 79, 81, 106, 292, and 313 and during the day for sol 306 (Table 1). Our measurements correspond to southern spring (sols 79, 81, and 106) and mid-late summer (sols 292, 306, and 313) on Mars.

To date, we have no detection of methane. Individually (Table 1), each of our six data sets produces a mean methane value ranging from –2.2 to 1.7 ppbv. Combining the individual sol results with equal weighting yields a mean and SE of  $0.11 \pm 0.67$  ppbv. Alternatively, combining all of the individual measurements from all sols yields a grand mean and SE of  $0.18 \pm 0.67$  ppbv. At the 95% confidence level, either approach (14) yields an upper limit on Mars atmospheric methane of 1.3 ppbv. Curiosity’s low upper limit is not expected given observations only a few years ago of large methane plumes and calculations (7) that the plume dispersion should produce global values of ~6 ppbv after the 6-month period (3, 15) needed to mix uniformly across the planet, which would persist with a photochemical lifetime of several hundred years (3, 5, 16).

Before Curiosity’s landing on Mars in August 2012, observational evidence for methane on

<sup>1</sup>Jet Propulsion Laboratory, California Institute of Technology, Pasadena, CA 91109, USA. <sup>2</sup>NASA Goddard Space Flight Center, Greenbelt, MD 20771, USA. <sup>3</sup>Department of Atmospheric, Oceanic, and Space Sciences, University of Michigan, Ann Arbor, MI 48109, USA. <sup>4</sup>Department of Geological and Planetary Sciences, California Institute of Technology, Pasadena, CA 91125, USA.

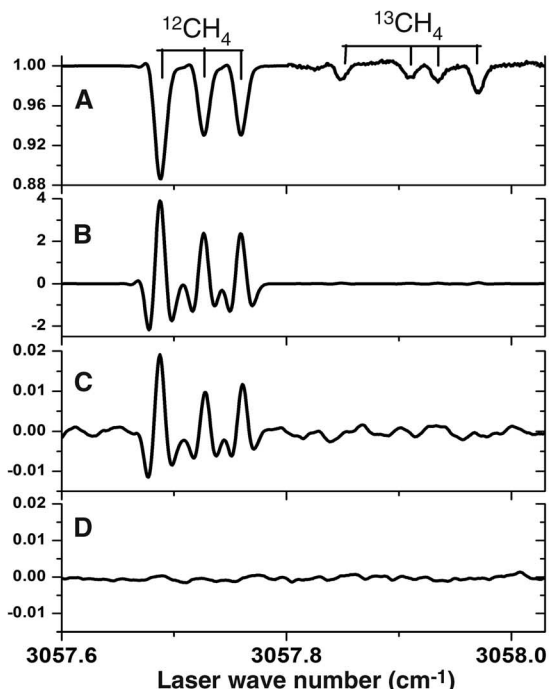
\*Corresponding author. E-mail: chris.r.webster@jpl.nasa.gov  
†MSL Science Team authors and affiliations are listed in the supplementary materials.

Mars was questioned in the published literature (15, 17, 18). Contradictions were noted between the locations of maxima reported from ground-based observations and maps inferred by PFS and TES from Mars orbit. The plume results (7) were questioned (17) on the basis of a possible misinterpretation from methane lines whose positions coincided with those of terrestrial isotopic  $^{13}\text{CH}_4$  lines. Krasnopolsky (19) argued that cometary and volcanic contributions were not sufficient to explain high methane abundances, calculating a cometary contribution of only  $\sim 0.1$  ppbv and noting the lack of current volcanism, lack of hot spots in thermal imaging (20), and the extremely

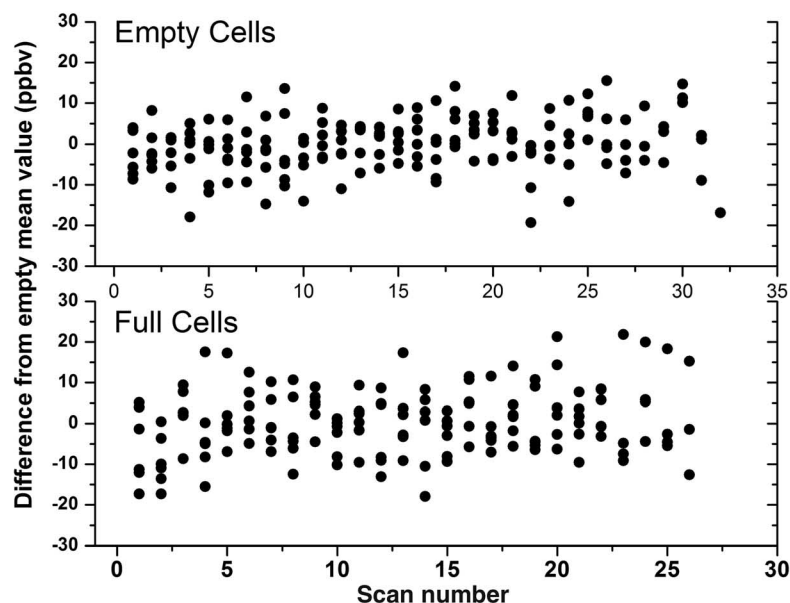
low upper limit for Mars  $\text{SO}_2$  (9, 21) that in Earth's volcanic emissions is orders of magnitude more abundant than  $\text{CH}_4$  (5).

The very short methane lifetime of 0.4 to 4 years derived from the 2003–2006 observations (7) requires powerful destruction mechanisms that have not been identified to date. Although models have been proposed for rapid removal of methane by oxidants—such as hydrogen peroxide and perchlorates, or by superoxides derived from their mineral reactions (22–24) and directly by electric fields generated in dust devils (25)—there remains no evidence for their existence at Mars. Moreover, it has not been demonstrated that any of these

processes can reduce the lifetime of methane by the required factor of 100 or more compared with its photochemical lifetime. Our reported upper limit of 1.3 ppbv is substantially lower than the methane abundances reported from Mars remote sensing spacecraft observations and those from Earth telescopic observations, including both the earlier high values of typically tens of ppbv and the more recently reported upper limits of 7 to 8 ppbv (9, 10). Although TLS samples only the very lowest part ( $\sim 1$  m) of the Mars atmosphere as compared with those of the other observations that are vertical column-integrated results, the atmospheric scale height ( $\sim 10$  km) and mixing



**Fig. 1. The TLS-SAM methane measurements.** (Left) Examples of flight spectra downloaded from Curiosity. (A) Spectrum recorded during an unrelated Evolved Gas Analysis (EGA) run (14) showing location of  $^{12}\text{CH}_4$  and  $^{13}\text{CH}_4$  lines, in which the second half has been vertically expanded by  $\times 20$  to show the weaker  $^{13}\text{CH}_4$  lines. (B) Same as (A) but second harmonic (2f) spectrum (14), without vertical expansion. (C) Averaged full cell 2f spectrum for sol 106 (nighttime ingest), with foreoptics contribution (14). (D) Averaged



full cell 2f spectrum for sol 306 (daytime ingest), with foreoptics evacuated. [Spectra (A) and (B) are shown here in part because they were taken after the atmospheric runs and show that our  $\text{CH}_4$  lines have not moved and that the instrument continued to work well with consistent capability to detect methane.] (Right) Individual 2-min data points from 6 sols. (Top) Empty cell data with mean value of 0.0 ppbv. (Bottom) Full cell data with mean value of 0.18 ppbv.

**Table 1. Curiosity SAM-TLS methane measurements at Gale Crater (4.5°S, 137.4°E) over an 8-month period.** SEM, standard error from the mean; Ls, solar longitude.

Martian sol after landing on 6 August 2012	Earth date	Ls (degrees)	Gas ingest time/cell pressure (mbar)/foreoptics pressure (mbar)	Mean value $\pm$ 1 SEM (ppbv)
79	25 October 2012	195.0	Night/8.0/11.5	1.62 $\pm$ 2.03
81	27 October 2012	196.2	Night/8.0/11.5	1.71 $\pm$ 2.06
106	27 November 2012	214.9	Night/8.5/10.9	-0.55 $\pm$ 1.45
292	1 June 2013	328.6	Night/8.7/9.2	0.60 $\pm$ 1.74
306	16 June 2013	336.5	Day/8.1/0.0	-2.21 $\pm$ 0.94
313	23 June 2013	340.5	Night/8.7/0.0	-0.50 $\pm$ 0.94
Mean of individual sol results				0.11 $\pm$ 0.56
Mean for entire aggregated data set				0.18 $\pm$ 0.67

time (approximately a few months) suggests that our measured upper limit is representative of the global mean background level. With an expected photochemical lifetime of methane in the Martian atmosphere of hundreds of years (3, 5, 16), there currently remains no accepted explanation (15, 17) for the existence and distribution of the reported plumes nor of the apparent disappearance of methane over the past few years. Our result sets an upper limit that is ~6 times lower than other recent measurements and greatly reduces the probability of substantial methanogenic microbial activity on Mars and recent methane production through serpentinization or from exogenous sources, including meteoritic, interplanetary dust and cometary infall.

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#### Supplementary Materials

www.sciencemag.org/content/342/6156/355/suppl/DC1  
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## Genomically Recoded Organisms Expand Biological Functions

Marc J. Lajoie,<sup>1,2</sup> Alexis J. Rovner,<sup>3,4</sup> Daniel B. Goodman,<sup>1,5</sup> Hans-Rudolf Aerni,<sup>4,6</sup> Adrian D. Haimovich,<sup>3,4</sup> Gleb Kuznetsov,<sup>1</sup> Jaron A. Mercer,<sup>7</sup> Harris H. Wang,<sup>8</sup> Peter A. Carr,<sup>9</sup> Joshua A. Mosberg,<sup>1,2</sup> Nadin Rohland,<sup>1</sup> Peter G. Schultz,<sup>10</sup> Joseph M. Jacobson,<sup>11,12</sup> Jesse Rinehart,<sup>4,6</sup> George M. Church,<sup>1,13\*</sup> Farren J. Isaacs<sup>3,4\*</sup>

We describe the construction and characterization of a genomically recoded organism (GRO). We replaced all known UAG stop codons in *Escherichia coli* MG1655 with synonymous UAA codons, which permitted the deletion of release factor 1 and reassignment of UAG translation function. This GRO exhibited improved properties for incorporation of nonstandard amino acids that expand the chemical diversity of proteins in vivo. The GRO also exhibited increased resistance to T7 bacteriophage, demonstrating that new genetic codes could enable increased viral resistance.

The conservation of the genetic code permits organisms to share beneficial traits through horizontal gene transfer (1) and enables the accurate expression of heterologous genes in nonnative organisms (2). However, the

common genetic code also allows viruses to hijack host translation machinery (3) and compromise cell viability. Additionally, genetically modified organisms (GMOs) can release functional DNA into the environment (4). Virus resistance (5) and biosafety (6) are among today's major unsolved problems in biotechnology, and no general strategy exists to create genetically isolated or virus-resistant organisms. Furthermore, biotechnology has been limited by the 20 amino acids of the canonical genetic code, which use all 64 possible triplet codons, limiting efforts to expand the chemical properties of proteins by means of nonstandard amino acids (NSAAs) (7, 8).

Changing the genetic code could solve these challenges and reveal new principles that explain how genetic information is conserved, encoded, and exchanged (fig. S1). We propose that genomically recoded organisms (GROs), whose codons have been reassigned to create an alternate genetic code) would be genetically isolated from natural organisms and viruses, as horizontally transferred genes would be mistranslated, pro-

ducing nonfunctional proteins. Furthermore, GROs could provide dedicated codons to improve the purity and yield of NSAA-containing proteins, enabling robust and sustained incorporation of more than 20 amino acids as part of the genetic code.

We constructed a GRO in which all instances of the UAG codon have been removed, permitting the deletion of release factor 1 (RF1; terminates translation at UAG and UAA) and, hence, eliminating translational termination at UAG codons. This GRO allows us to reintroduce UAG codons, along with orthogonal translation machinery [i.e., aminoacyl-tRNA synthetases (aaRSs) and tRNAs] (7, 9), to permit efficient and site-specific incorporation of NSAAs into proteins (Fig. 1). That is, UAG has been transformed from a nonsense codon (terminates translation) to a sense codon (incorporates amino acid of choice), provided the appropriate translation machinery is present. We selected UAG as our first target for genome-wide codon reassignment because UAG is the rarest codon in *Escherichia coli* MG1655 (321 known instances), prior studies (7, 10) demonstrated the feasibility of amino acid incorporation at UAG, and a rich collection of translation machinery capable of incorporating NSAAs has been developed for UAG (7).

We used an in vivo genome-editing approach (11), which is more efficient than de novo genome synthesis at exploring new genotypic landscapes and overcoming genome design flaws. Although a single lethal mutation can prevent transplantation of a synthetic genome (12), our approach allowed us to harness genetic diversity and evolution to overcome any potential deleterious mutations at a cost considerably less than de novo genome synthesis (supplementary text section B, "Time and cost"). In prior work, we used multiplex automated genome engineering [MAGE (13)] to remove all known UAG codons in groups of 10

<sup>1</sup>Department of Genetics, Harvard Medical School, Boston, MA 02115, USA. <sup>2</sup>Program in Chemical Biology, Harvard University, Cambridge, MA 02138, USA. <sup>3</sup>Department of Molecular, Cellular and Developmental Biology, Yale University, New Haven, CT 06520, USA. <sup>4</sup>Systems Biology Institute, Yale University, West Haven, CT 06516, USA. <sup>5</sup>Program in Medical Engineering and Medical Physics, Harvard–Massachusetts Institute of Technology (MIT) Division of Health Sciences and Technology, Cambridge, MA 02139, USA. <sup>6</sup>Department of Cellular and Molecular Physiology, Yale University, New Haven, CT 06520, USA. <sup>7</sup>Harvard College, Cambridge, MA 02138, USA. <sup>8</sup>Department of Systems Biology, Columbia University, College of Physicians and Surgeons, New York, NY 10032, USA. <sup>9</sup>MIT Lincoln Laboratory, Lexington, MA 02420, USA. <sup>10</sup>Department of Chemistry, The Scripps Research Institute, La Jolla, CA 92037, USA. <sup>11</sup>Center for Bits and Atoms, MIT, Cambridge, MA 02139, USA. <sup>12</sup>MIT Media Lab, MIT, Cambridge, MA 02139, USA. <sup>13</sup>Wyss Institute for Biologically Inspired Engineering, Harvard University, Boston, MA 02115, USA.

\*Corresponding author. E-mail: farren.isaacs@yale.edu (F.J.I.); gchurch@genetics.med.harvard.edu (G.M.C.)



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## Supplementary Material for

### **Low Upper Limit to Methane Abundance on Mars**

Christopher R. Webster,\* Paul R. Mahaffy, Sushil K. Atreya, Gregory J. Flesch,  
Kenneth A. Farley, the MSL Science Team

\*To whom correspondence should be addressed. E-mail: [chris.r.webster@jpl.nasa.gov](mailto:chris.r.webster@jpl.nasa.gov)

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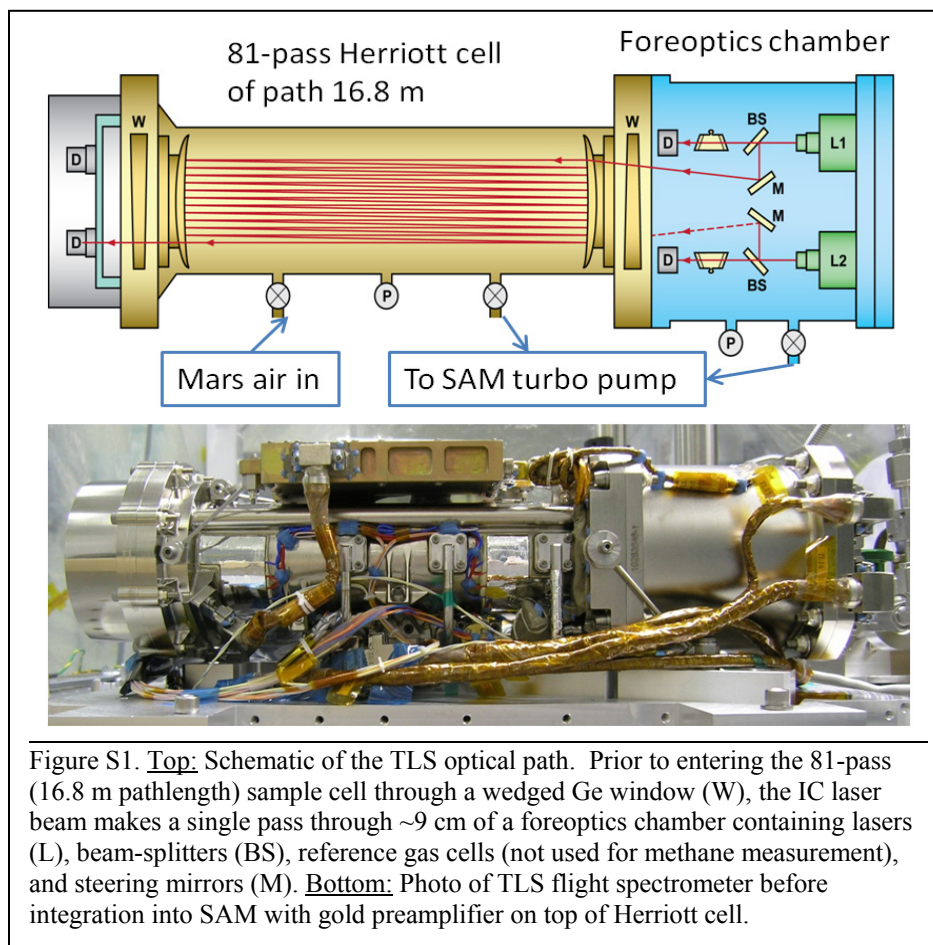
**SUPPLEMENTAL INFORMATION:**

**Low Upper Limit to Methane Abundance on Mars**

C. R. Webster, P. R. Mahaffy, S. K. Atreya, G. J. Flesch and K. A. Farley

**The Tunable Laser Spectrometer (TLS) in the Sample Analysis at Mars (SAM) instrument suite on the Curiosity Rover**

This instrument has been previously described in detail (12, 13). Fig. S1 below emphasizes the optical layout for the methane measurement.



**Methane spectroscopy and laser parameters:**

The TLS interband cascade (IC) laser scans through a unique fingerprint of seven spectral lines in the  $\nu_3$  band: three  $^{12}\text{CH}_4$  lines associated with R(3) and four subsequent  $^{13}\text{CH}_4$  lines associated with R(3) transitions. Table S1 below lists the three  $^{12}\text{CH}_4$  lines used for this study, as identified by both the HITRAN data base (26) and laboratory measurements. We create the labels e, f, g for these three lines, where the g line is strongest, and both e and f are about half the intensity of the g line.

Spectral line center (cm <sup>-1</sup> )	Line-strength at 296 K (cm <sup>-1</sup> /molecule·cm <sup>-2</sup> )	Ground-state energy (cm <sup>-1</sup> )	Assignment	Label
3057.687285	2.085E-19	62.8781	R(3)	g
3057.726529	1.245E-19	62.8768	R(3)	f
3057.760524	1.245E-19	62.8757	R(3)	e

The IC laser was developed at JPL, and operated near 245 K stabilized by a two-stage TEC cooler producing single-mode (>99%) continuous-wave output power with a linewidth retrieved from low-pressure (Doppler limited) spectra of ~10 MHz. This light was collimated using an efficient triple-lens collimator to produce ~1 mW laser power that passes through the foreoptics chamber then into the sample (Herriott) cell. Prior to entering the Herriott cell, the beam was attenuated by a factor of ~20 by a thin mylar sheet (not shown in Fig. S1) to reduce optical fringing and detector non-linearity. We note that the pre-launch settings for the TEC and laser current scans (used for calibration also) have not been changed and the target spectral line positions remain in our scan window. Very small (~linewidth) variations in the spectral line position are seen depending on the Curiosity heat ramp behavior, but we observe and track the methane lines continually for each spectrum through the simultaneously-recorded reference cell detector; the tracked methane spectrum arises from residual methane gas in the foreoptics chamber.

#### Evolved Gas Analysis Spectrum of Figure 1:

In Figure 1 we presented actual flight spectra downloaded from Curiosity rather than showing calculated HITRAN spectra. Spectra C and D are from the methane analysis presented here, but spectra A and B are from a different unrelated experiment on Mars in which a rock sample is heated (Evolved Gas Analysis (EGA) run) in a pyrolysis oven to produce evolved gas fed to TLS that was observed to contain methane. These spectra are shown here in part because they were taken AFTER the atmospheric methane runs and show that our CH<sub>4</sub> lines have not moved, and the instrument continues to work well with consistent capability to detect methane. The EGA spectra of Fig 1 (A, B) show the location of <sup>12</sup>CH<sub>4</sub> and <sup>13</sup>CH<sub>4</sub> lines in both the direct absorption and second harmonic (2f) spectrum.

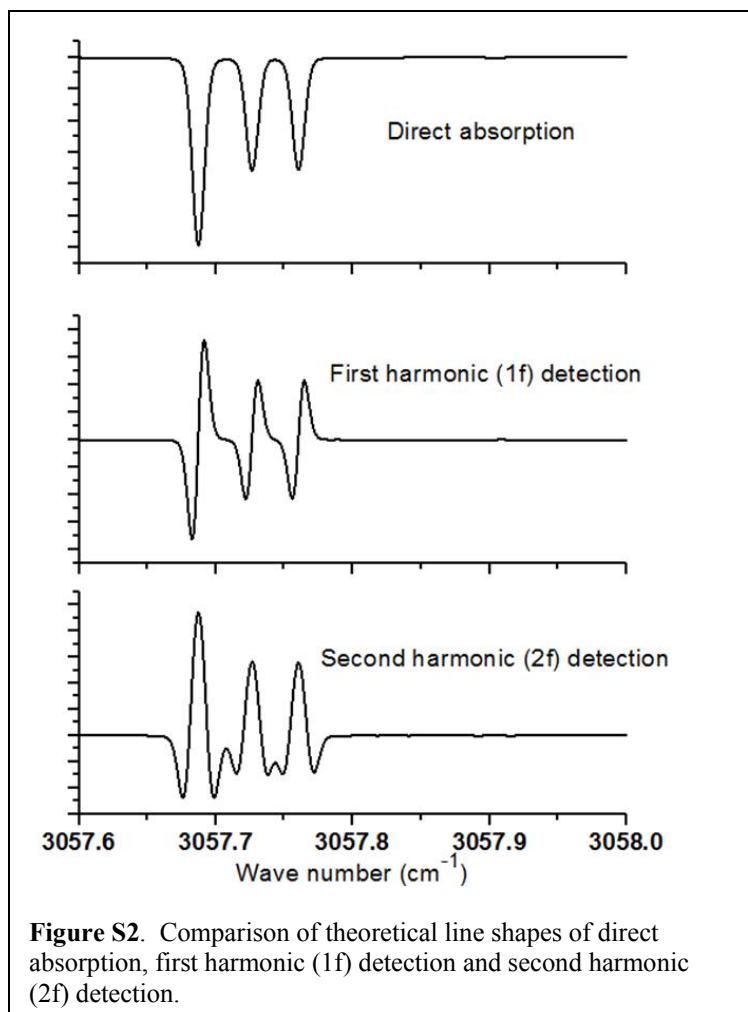
#### Description of the Difference Method:

We determine Mars methane abundances by differencing full cell and empty cell results (not spectra), as described below. In a typical run on one sol, we collect (downlink) 26 empty cell spectra (2 minutes on board averaged each) followed by 26 full cell, then a few more empty for return-to-zero check. Cell temperatures and pressures are extremely stable during the complete sol run and contribute negligibly to our results (see later). We chose to record relatively long periods of continuous empty or full spectra to make sure that no drift (growth or loss) in retrieved methane abundance was observed during the run. We record sequential 2-minute empty cell spectra for ~1 hour followed by ~1 hour of sequential full cell spectra. We do not difference full-empty spectra before processing. Rather, with powerful computing resources now available, we process each of our 3 methane lines separately in each and all of our 2-minute spectra (by comparison with HITRAN calculations described below), then produce a combined efg-line average abundance for each spectrum that becomes a single raw 2-minute data point. Then, after applying common calibration factors and error contributions, we compare statistically the empty and full cell results for all the sols after normalizing to the empty cell mean values.

#### Direct and Second-harmonic (2f) Spectra

TLS is designed to simultaneously produce both direct absorption and second harmonic (2f) spectra, as is standard for commercial and laboratory tunable laser spectrometers (27). Tunable laser spectrometers “scan” through spectral lines by applying a current ramp (usually saw-tooth) to the laser that through junction heating changes the wavelength by a small amount, the ramp repeated typically every one second (as done in TLS).

In direct absorption, absorption line depths that indicate gas abundance are measured as dips in the large light level on the detector as the laser is scanned. For very weak absorptions of ~1% or less (due to low gas amounts, too small path lengths or gas pressures, etc., and as expected for low methane (<20 ppbv) amounts) it is challenging for electronics and dynamic range to measure small changes in a large signal, and a “harmonic” detection is preferred. In harmonic detection, the very narrow laser linewidth (much narrower than the gas absorption line) is modulated (“dithered”) at high frequency (say 10 kHz) by applying a sinusoidal component to the laser current ramp (increasing laser current is the normal method of tuning the laser across the spectral scan) with an amplitude that is small compared to the gas linewidth. So, if we modulate at 10 kHz and look at only the component of the detector signal at 10 kHz (using phase-sensitive detection), we would record a first-harmonic or first-derivative 1f spectrum as shown in Fig. S2. Outside the spectral line and at the line center, the laser is jiggling left and right where no difference exists, so it records zero in these places, but has its maximum signals (negative and positive) at the side of the line where the slope is maximum.



**Figure S2.** Comparison of theoretical line shapes of direct absorption, first harmonic (1f) detection and second harmonic (2f) detection.

If we now modulate at 10 kHz, but look at the component of the laser light on the detector that is at 20 kHz, we would record (as we do on TLS) the second-harmonic or second-derivative (2f) spectrum seen in Fig. S2. Both 1f and 2f spectral signals are zero-based in amplitude (electronics gain likes that) and move the detection frequency to higher frequency (kHz) compared to the direct (DC) spectrum, where 1/f noise is lower. Thus the harmonic method produces higher signal-to-noise spectra. The 1f spectrum is not usually used since it can have small vertical offsets and the line center position is a zero-crossing rather than a peak. The 2f spectrum is preferred since it has its peak in the same place as the direct absorption spectrum, and moves the detection regime to the higher (20 kHz) frequency.

### Spectral Data Processing

The Beer-Lambert law models the optical transmission of light through an absorbing medium:

$$I_{\nu} = I_0 e^{-k(\nu)\rho l}$$

where  $I_{\nu}$  is the transmitted light intensity at frequency  $\nu$ ,  $I_0$  is the incident light intensity,  $k(\nu)$  is a line shaping function that may be Doppler, Lorentzian, or Voigt, although the Doppler lineshape is a close approximation at Mars atmospheric pressures.  $\rho$  is the number density and  $l$  is the path length in cm. We use this model to determine the abundances of individual absorption lines present in our sampled measurements. The model needs many input spectral parameters for temperature dependence, air broadening, ground state energy, etc., and we use the HITRAN database for this information (26). Direct absorption spectra produce good results for gases that have line center absorption depths of ~1% or greater. For higher sensitivity, we add a modulation to the laser current and then demodulate the returning detector signal at twice that frequency. This effectively gives us a second harmonic or 2f



spectrum in which sensitivities of up to 2 parts in  $10^5$  are possible. See the section above and also Webster et al. (27) for a complete discussion.

### **Laser Power Normalization and Wave Number Scale**

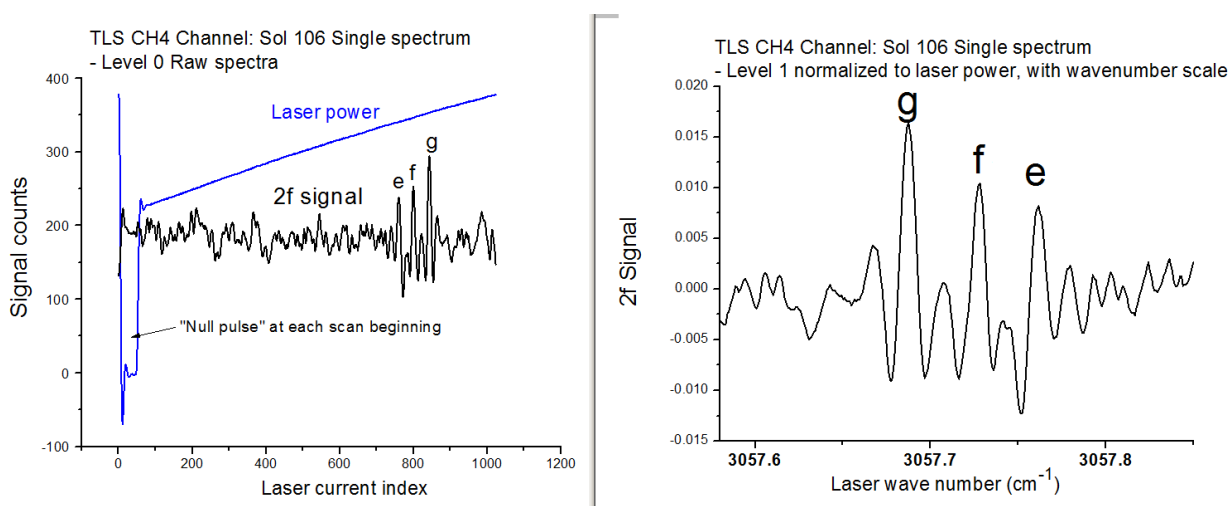
For a given channel (either CH<sub>4</sub> or CO<sub>2</sub>/H<sub>2</sub>O), TLS returns 3 spectra from the Herriott cell “science” detector, and 3 spectra from the reference channel detector. For both the Herriott cell and reference channel spectra, these 3 spectra are the direct absorption spectrum, the 2f spectrum, and a high-gain 2f spectrum. Our methane analysis is done using the 2f spectrum that is normalized to laser power from the direct absorption spectrum and mapped to a wave number scale using the reference detector signals. The high-gain 2f spectrum is not used since with only moderate gain increase (x16) it duplicates the 2f spectrum in signal-to-noise ratio but suffers from dynamic range restriction.

TLS also returns reference detector spectra recorded simultaneously with those from the science detector, and these are used to track the methane lines to provide the wave number scale for later processing. The methane signal (spectra) detected by the reference detector (located inside the foreoptics, as shown in Fig. S1) is due to residual methane in the foreoptics. The foreoptics contribution to the science spectrum is equivalent to about 90 ppbv for sols 79-292. The 2-stage thermoelectric cooler on the IC laser keeps the lines in the same position during the scans, with drifts in line positions over all sols of only about 1-2 linewidths that are tracked successfully.

For an amount of gas at a given pressure and temperature, the model will predict the depth and width (distribution in wave number) of the absorption by the gas sample for all sampled frequencies, allowing us to then compare our recorded spectra to the spectra produced by the model. But, in order to make this comparison, we must first normalize the recorded data. This process that takes level 0 data (spectra) and produces level 1 data (spectra) entails:

1. Removing a “null pulse” which is a measurement of the background light taken with the laser off, and recorded during every one second spectrum that is averaged on board for our 2-minute downlinked spectrum. This allows us to determine the direct absorption with respect to a percentage of transmitted light (i.e. 1% absorption: 99% transmission).
2. Removing any DC offsets in the harmonic spectra (described below).
3. Fit the baseline of the spectra. This sloping baseline results from the fact that the laser output power increases as it tunes through different wave numbers.
4. Assign a wave number (cm<sup>-1</sup>) scale to the real spectra. We do this by using easily identifiable peaks of known wave number.

Once the raw spectra (level 0 data) are normalized (Fig. S3) as level 1 data, we can then use the HITRAN model to scale our real world data.



**Fig. S3.** Example of normalization of a real single spectrum (2 min.) downloaded for sol 106. The methane triplet lines e, f, g can be identified from Table S1 above. The left panel is the complete level 0 spectra, whereas the right panel that shows level 1 data (same 2-min. spectrum normalized to power and given wave number scale) has been

expanded in wave number to show the methane lines used and the occurrence of optical interference fringes that limit the detection method for a single 2-min. spectrum.

### ***Producing Abundances***

Using temperatures and pressures from our instrument for input, we iteratively run the model, varying the abundance in a converging algorithm until the synthetic spectra for the single line is the same size as our real spectrum (within some determined threshold). The convergence criteria are set to optimize for the 2f spectra.

For the methane analysis, we generate two results, one named “peak-to-peak” that returns the peak-to-peak signal amplitude (actually central peak to lobe-average) values, and a second named “integral” that returns the area of the 2f line between and above the bottom lobe minima positions (wave number). The peak-to-peak method finds the signal amplitude of the 2f maximum and lobe minima average, and is our preferred method since it produces somewhat lower scatter in our data, although results for either method are very close. The integral method, which is used for retrieving H, C, O isotope ratios (19) uses the following algorithm:

1. Find the global max of the 2f absorption spectra (peak)
2. Find the two local minima (2f lobes)
3. Fit a line between the two lobes
4. Using the lobes as integration boundaries, find the area between the fitted line and the spectra for both the direct and 2f spectra. Ratio this area between real and synthetic spectra and if ratio is outside the convergence threshold, iterate with new abundance.

Once the measurements converge, we ratio the resulting areas of the real spectra to the synthetic spectra which has a known abundance. For both methods, using the same laser modulation and gain throughout (pre-launch calibration and all Mars measurements), we relate the 2f signal size to the direct absorption size through calibration as described below, and like any flight project, we rigorously run our experiment as tested and calibrated pre-launch.

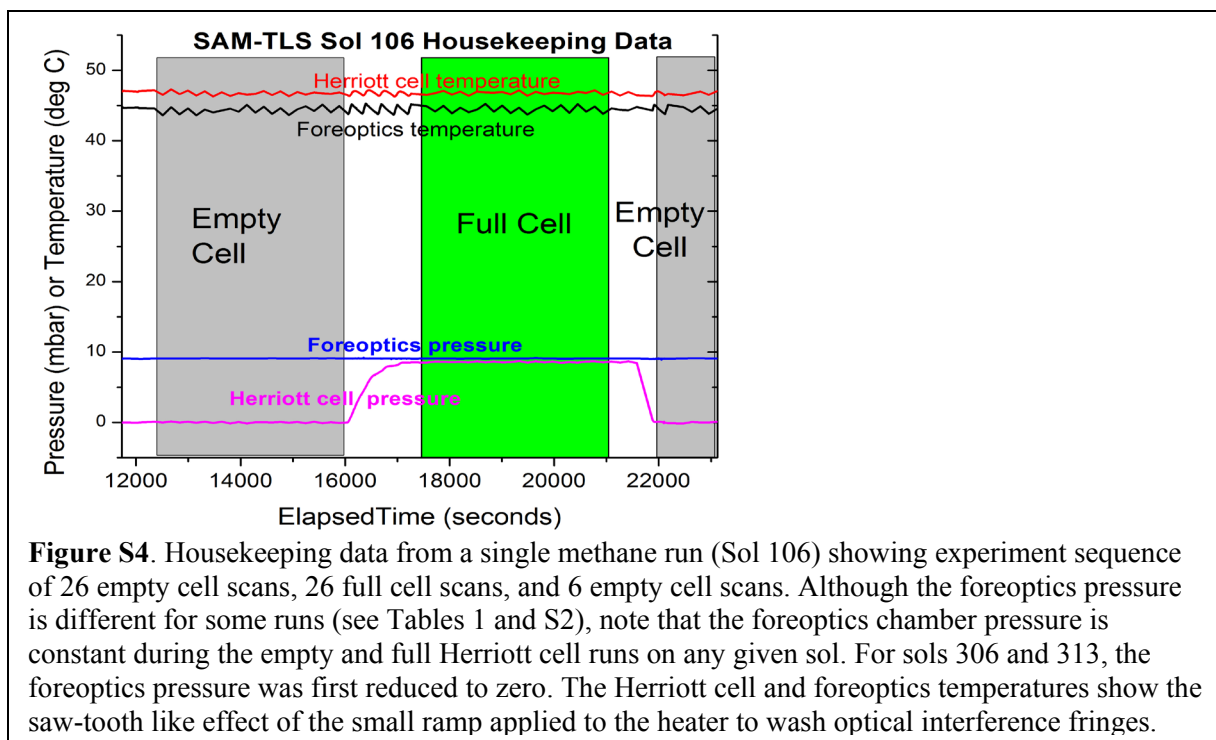
### **Calibration:**

When analyzing direct absorption spectra with known pressure, temperature and pathlength, a Beer’s law calculation using spectral line parameters from HITRAN can in theory provide the gas abundance without the need for calibration gases (i.e. someone else did the work when they created the data base). However, calibration gases serve the dual purpose of verifying the spectrometer response (a check of pathlength or number of passes in a cell, laser linewidth, pressure, mode purity, temperature, saturation, etc.) and also giving a direct calibration (relationship) between the direct absorption and the 2f channel with its various different gain stages.

The relative methane abundances reported here are calibrated using NIST-traceable methane in air provided by the NOAA-CMDL laboratory (provided by Jim Elkins group) specified to contain  $88 \pm 0.5$  ppbv. By injecting this gas into the TLS Herriott cell during pre-launch calibration runs of TLS and SAM in the NASA GSFC environmental chamber, we record both direct absorption and 2f signal sizes using the same conditions (e.g. laser scan, modulation, flight electronics and software, Herriott cell temperature and pressure, ramp heater) used on Mars. During the calibration run, the foreoptics is pumped out so that there is no contribution from foreoptics gas. The path length of the Herriott cell was verified to be 81 passes based on direct absorption measurements of these same methane lines using a second calibration cylinder (same provider) at 1800 ppbv. In addition, by adding pure methane gas at low pressures so that the lines are bleached to zero light transmission at line centers, the mode purity during the scan is verified. No change in alignment or detector signal sizes has been detected since pre-launch. Normalizing the mean value retrieved to 88 ppbv gives us a calibration result and uncertainty of  $88.0 \pm 1.13$  ppbv. We note that this absolute uncertainty of  $\pm 1.13$  ppbv does not carry forward in our difference method described below, since it would only serve to change the mean value and upper limit slightly (by  $\sim 1$  part in 88).

### **The foreoptics contribution to the difference method:**

The difference method is described in the body of the main paper, and the sequence shown in Fig. S4 below. During the empty or full cell periods, the foreoptics and Herriott cell pressures are very stable; during a typical run (Sol 106) the temperatures and foreoptics pressure are stable to 0.02%, and the Herriott cell pressure during the full cell section is stable to 0.1%.



**Table S2. Ingest and foreoptics pressures.**

Martian Sol after landing	Earth date	Ls (deg)	Gas ingest time and cell pressure (mbar)	Foreoptics pressure (mbar)
79	Oct 25 <sup>th</sup> 2012	195.0	Night/8.0	11.5
81	Oct 27 <sup>th</sup> 2012	196.2	Night/8.0	11.5
106	Nov 27 <sup>th</sup> 2012	214.9	Night/8.5	10.9
292	June 1 <sup>st</sup> 2013	328.6	Night/8.7	9.2
306	June 16 <sup>th</sup> 2013	336.5	Day/8.1	~0
313	June 23 <sup>rd</sup> 2013	340.5	Night/8.7	~0

During the long pre-launch and cruise phase to Mars, the foreoptics chamber leaked up to a significant pressure (~76 mbar) by the time we arrived at Mars. This pressure included terrestrial “Florida air” from the launch site that contained significant terrestrial methane gas (~10 ppmv) that showed up as a large methane signal (spectrum) on the Herriott cell science detector for both “empty” and “full” Herriott cell data, since the beam made one pass through the 9-cm length of the foreoptics. Moreover, our first attempts to measure methane on Mars showed methane spectra that increased in size with time during the empty and full cell scans that we attributed to diffusion (leakage) of the foreoptics methane gas into the Herriott cell during the run. Results from these runs made before sol 79 were discarded and not included in the analysis. To reduce the foreoptics contribution, we pumped down the foreoptics chamber in a series of steps for subsequent sol runs (80, 33, 11.5 mbar) until at 11.5 mbar we observed no detectable increase (or reduction) in the empty or full cell spectra with time over the run, so that we were confident that the leakage was negligible during the runs to follow. As a matter of good practice, for the last two data sets of sols 306 and 313, we further reduced the foreoptics pressure to close to zero by pumping on the chamber. As the data in Table S3 shows, the results for all 6 sols are consistent within measurement uncertainty. Note that the low foreoptics contribution in sols 306 and 313 reduces the scatter in the data somewhat, as shown in Table S3.

Because of the foreoptics contribution, all of our spectra (empty and full Herriott cell) look somewhat like those in Fig. S3 since (in the absence of significant Martian methane) they are dominated by the foreoptics contribution. We then process them as described above, and then look for differences in the empty and full cell results. Specifically, the “full” cell methane spectra are first processed as if the observed methane spectrum came only from the Herriott

cell, that is, we use the measured Herriott cell pressure and temperature to retrieve a “full cell” methane mixing ratio by comparison with HITRAN. Then for the “empty” cell spectra, we use the same mean temperatures and pressures of the full cell and process the empty cell spectra to reveal the “empty cell” methane mixing ratio. This method makes the difference method most sensitive to Herriott cell methane from Mars, should it be there. If there was no methane on Mars, the empty and full cell results would be identical. If there was 20 ppbv methane on Mars, the full cell result would be 20 ppbv larger than the empty cell result. For sols 79-292, for example, both the empty and full cell results are close to 90 ppbv, and for sols 306 and 313 it is <20 ppbv. For the difference data given in Table 1 and S1, the mean empty cell values for that specific run have been subtracted from the full cell values to provide the resulting Martian methane mixing ratio.

The inlet to the SAM-TLS instrument is a 3/16” internal diameter stainless steel tube heated to 50°C containing a dust filter of sintered Inconel 0.5 micron particles that is located on the rover side ~1 m above the Martian surface, and was pointed at a variety of directions relative to the nominal wind direction.

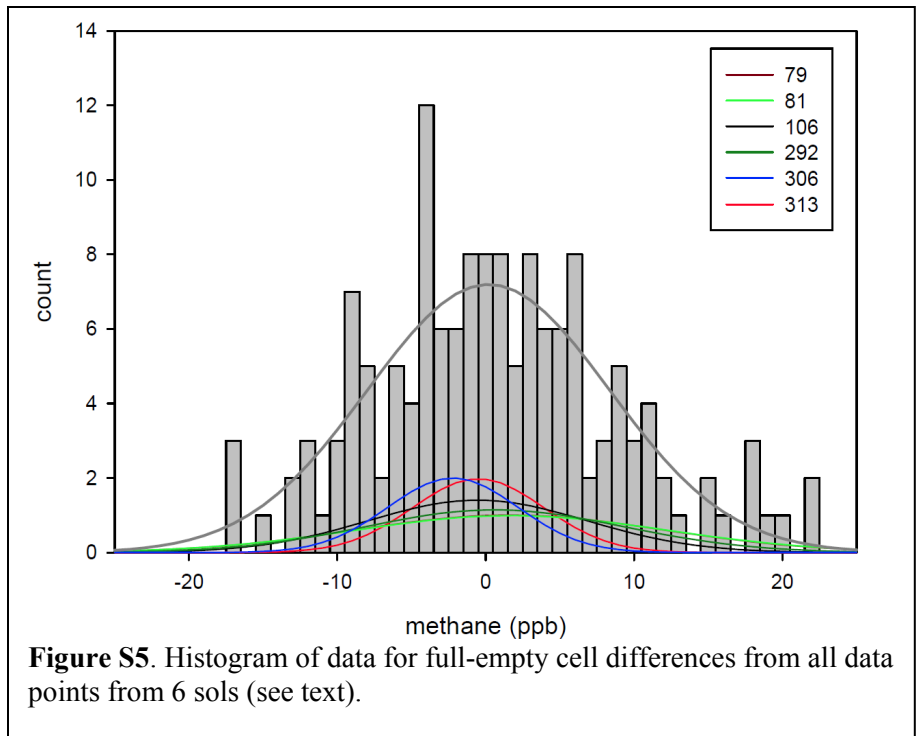
**Statistical analysis of data:**

The spacecraft returns two-minute averaged signals for each of the three spectral absorption lines given in Table S1. The TLS measurements include methane absorption occurring both in the Herriott cell and along the optical path of the foreoptics prior to entry into the cell, as described in the main text. In our first four sol runs, the foreoptics region had terrestrial air with methane in it, allowing confident identification of the methane absorption lines and continuous monitoring of scan-to-scan line shifts. The foreoptics methane signal also introduces a substantial “blank” signal which must be removed to compute the amount of methane in the Mars atmosphere in the Herriott cell. For the later two sols 306 and 313, the foreoptics were pumped out before hand, and the measured background signals although much lower were treated in exactly the same way for all six sol data comparisons.

We treat each of these lines as a separate estimate of the absorption attributable to methane somewhere along the optic path. These absorptions were converted into an apparent methane mixing ratio in the Herriott cell by assuming that this is the only region in which methane occurs. As shown in Table S3, all of our sol runs except sol 306 were executed to produce 26 full cell points and 32 empty cell points. For sols 292 and 313 the heat ramp monitor showed that we had not quite reached temperature, and the first one and four points, respectively were removed from the analysis. For sol 306, the daytime run and rover power demand meant that the number of full, empty data points was limited to 22 and 18 points. No data points were removed from this analysis. Results for each of the six sol measurements (see Tables 1 and S3) show mean values ranging from -2.2 to 1.7 ppbv. In the absence of any notable difference in the atmospheric methane abundances retrieved on the six sols, we chose to merge the six individual data sets to obtain best estimates of the atmospheric methane amount, its uncertainty, and our upper limit methane value. We thus obtained 147 full cell points and 167 empty cell points that were then statistically analyzed as a single data set. For this calculation we subtracted the mean blank signal on each sol from the measured signals on that sol. The results are shown in Figure S5 as a histogram for the aggregated 6-sol data set. Each individual sol defines a broadly Gaussian distribution and all sols have statistically equivalent variance. These Gaussian distributions are also indicated in Figure S5. The mean empty cell value is by definition zero, while the blank-corrected full cell mean is  $0.18 \pm 0.67$  ppbv. These data imply an upper limit with 95% confidence of 1.3 ppbv for the methane volume mixing ratio of the Martian atmosphere.

**Table S3. Statistical data for each sol and 6-sol data treated as single data set.**

Martian Sol after landing on Aug 6 <sup>th</sup> 2012	Number of Full Cell points	Number of Empty Cell points	Mean CH4 value $\pm$ 1SEM (ppbv)
79	26	26	$1.62 \pm 2.03$
81	26	32	$1.71 \pm 2.06$
106	26	31	$-0.55 \pm 1.45$
292	25	31	$0.60 \pm 1.74$
306	22	18	$-2.21 \pm 0.94$
313	22	27	$-0.50 \pm 0.94$
All six sols treated as one data set	147	165	$0.18 \pm 0.67$



## **Mars Science Laboratory (MSL) Science Team**

### **Aalto University**

Osku Kempainen

### **Applied Physics Laboratory (APL) at Johns Hopkins University**

Nathan Bridges, Jeffrey R. Johnson, Michelle Minitti

### **Applied Research Associates, Inc. (ARA)**

David Cremers

### **Arizona State University (ASU)**

James F. Bell III, Lauren Edgar, Jack Farmer, Austin Godber, Meenakshi Wadhwa, Danika Wellington

### **Ashima Research**

Ian McEwan, Claire Newman, Mark Richardson

### **ATOS Origin**

Antoine Charpentier, Laurent Peret

### **Australian National University (ANU)**

Penelope King

### **Bay Area Environmental Research Institute (BAER)**

Jennifer Blank

### **Big Head Endian LLC**

Gerald Weigle

### **Brock University**

Marie Schmidt

### **Brown University**

Shuai Li, Ralph Milliken, Kevin Robertson, Vivian Sun

### **California Institute of Technology (Caltech)**

Michael Baker, Christopher Edwards, Bethany Ehlmann, Kenneth Farley, Jennifer Griffes, John Grotzinger, Hayden Miller, Megan Newcombe, Cedric Pilorget, Melissa Rice, Kirsten Siebach, Katie Stack, Edward Stolper

### **Canadian Space Agency (CSA)**

Claude Brunet, Victoria Hipkin, Richard Léveillé, Geneviève Marchand, Pablo Sobrón Sánchez

### **Capgemini France**

Laurent Favot

### **Carnegie Institution of Washington**

George Cody, Andrew Steele

### **Carnegie Mellon University**

Lorenzo Flückiger, David Lees, Ara Nefian

### **Catholic University of America**

Mildred Martin

### **Centre National de la Recherche Scientifique (CNRS)**

Marc Gailhanou, Frances Westall, Guy Israël

**Centre National d'Etudes Spatiales (CNES)**

Christophe Agard, Julien Baroukh, Christophe Donny, Alain Gaboriaud, Philippe Guillemot, Vivian Lafaille, Eric Lorigny, Alexis Paillet, René Pérez, Muriel Saccoccio, Charles Yana

**Centro de Astrobiología (CAB)**

Carlos Armians-Aparicio, Javier Caride Rodríguez, Isaias Carrasco Blázquez, Felipe Gómez Gómez, Javier Gómez-Elvira, Sebastian Hettrich, Alain Lepinette Malvitte, Mercedes Marín Jiménez, Jesús Martínez-Frías, Javier Martín-Soler, F. Javier Martín-Torres, Antonio Molina Jurado, Luis Mora-Sotomayor, Guillermo Muñoz Caro, Sara Navarro López, Verónica Peinado-González, Jorge Pla-García, José Antonio Rodríguez Manfredi, Julio José Romeral-Planelló, Sara Alejandra Sans Fuentes, Eduardo Sebastian Martinez, Josefina Torres Redondo, Roser Urqui-O'Callaghan, María-Paz Zorzano Mier

**Chesapeake Energy**

Steve Chipera

**Commissariat à l'Énergie Atomique et aux Énergies Alternatives (CEA)**

Jean-Luc Lacour, Patrick Mauchien, Jean-Baptiste Sirven

**Concordia College**

Heidi Manning

**Cornell University**

Alberto Fairén, Alexander Hayes, Jonathan Joseph, Steven Squyres, Robert Sullivan, Peter Thomas

**CS Systemes d'Information**

Audrey Dupont

**Delaware State University**

Angela Lundberg, Noureddine Melikechi, Alissa Mezzacappa

**Denver Museum of Nature & Science**

Julia DeMarines, David Grinspoon

**Deutsches Zentrum für Luft- und Raumfahrt (DLR)**

Günther Reitz

**eINFORMe Inc. (at NASA GSFC)**

Benito Prats

**Finnish Meteorological Institute**

Evgeny Atlaskin, Maria Genzer, Ari-Matti Harri, Harri Haukka, Henrik Kahanpää, Janne Kauhanen, Osku Kempainen, Mark Paton, Jouni Polkko, Walter Schmidt, Tero Siili

**GeoResources**

Cécile Fabre

**Georgia Institute of Technology**

James Wray, Mary Beth Wilhelm

**Géosciences Environnement Toulouse (GET)**

Franck Poitrasson

**Global Science & Technology, Inc.**

Kiran Patel

**Honeybee Robotics**

Stephen Gorevan, Stephen Indyk, Gale Paulsen

**Imperial College**

Sanjeev Gupta

**Indiana University Bloomington**

David Bish, Juergen Schieber

**Institut d'Astrophysique Spatiale (IAS)**

Brigitte Gondet, Yves Langevin

**Institut de Chimie des Milieux et Matériaux de Poitiers (IC2MP)**

Claude Geffroy

**Institut de Recherche en Astrophysique et Planétologie (IRAP), Université de Toulouse**

David Baratoux, Gilles Berger, Alain Cros, Claude d'Uston, Olivier Forni, Olivier Gasnault, Jérémie Lasue, Qiu-Mei Lee, Sylvestre Maurice, Pierre-Yves Meslin, Etienne Pallier, Yann Parot, Patrick Pinet, Susanne Schröder, Mike Toplis

**Institut des Sciences de la Terre (ISTerre)**

Éric Lewin

**inXitu**

Will Brunner

**Jackson State University**

Ezat Heydari

**Jacobs Technology**

Cherie Achilles, Dorothy Oehler, Brad Sutter

**Laboratoire Atmosphères, Milieux, Observations Spatiales (LATMOS)**

Michel Cabane, David Coscia, Guy Israël, Cyril Szopa

**Laboratoire de Géologie de Lyon : Terre, Planète, Environnement (LGL-TPE)**

Gilles Dromart

**Laboratoire de Minéralogie et Cosmochimie du Muséum (LMCM)**

François Robert, Violaine Sautter

**Laboratoire de Planétologie et Géodynamique de Nantes (LPGN)**

Stéphane Le Mouélic, Nicolas Mangold, Marion Nachon

**Laboratoire Génie des Procédés et Matériaux (LGPM)**

Arnaud Buch

**Laboratoire Interuniversitaire des Systèmes Atmosphériques (LISA)**

Fabien Stalport, Patrice Coll, Pascaline François, François Raulin, Samuel Teinturier

**Lightstorm Entertainment Inc.**

James Cameron

**Los Alamos National Lab (LANL)**

Sam Clegg, Agnès Cousin, Dorothea DeLapp, Robert Dingler, Ryan Steele Jackson, Stephen Johnstone, Nina Lanza, Cynthia Little, Tony Nelson, Roger C. Wiens, Richard B. Williams



**Lunar and Planetary Institute (LPI)**

Andrea Jones, Laurel Kirkland, Allan Treiman

**Malin Space Science Systems (MSSS)**

Burt Baker, Bruce Cantor, Michael Caplinger, Scott Davis, Brian Duston, Kenneth Edgett, Donald Fay, Craig Hardgrove, David Harker, Paul Herrera, Elsa Jensen, Megan R. Kennedy, Gillian Krezoski, Daniel Krysak, Leslie Lipkaman, Michael Malin, Elaina McCartney, Sean McNair, Brian Nixon, Liliya Posiolova, Michael Ravine, Andrew Salamon, Lee Saper, Kevin Stoiber, Kimberley Supulver, Jason Van Beek, Tessa Van Beek, Robert Zimdar

**Massachusetts Institute of Technology (MIT)**

Katherine Louise French, Karl Iagnemma, Kristen Miller, Roger Summons

**Max Planck Institute for Solar System Research**

Fred Goesmann, Walter Goetz, Stubbe Hviid

**Microtel**

Micah Johnson, Matthew Lefavor, Eric Lyness

**Mount Holyoke College**

Elly Breves, M. Darby Dyar, Caleb Fassett

**NASA Ames**

David F. Blake, Thomas Bristow, David DesMarais, Laurence Edwards, Robert Haberle, Tori Hoehler, Jeff Hollingsworth, Melinda Kahre, Leslie Keely, Christopher McKay, Mary Beth Wilhelm

**NASA Goddard Space Flight Center (GSFC)**

Lora Bleacher, William Brinckerhoff, David Choi, Pamela Conrad, Jason P. Dworkin, Jennifer Eigenbrode, Melissa Floyd, Caroline Freissinet, James Garvin, Daniel Glavin, Daniel Harpold, Andrea Jones, Paul Mahaffy, David K. Martin, Amy McAdam, Alexander Pavlov, Eric Raaen, Michael D. Smith, Jennifer Stern, Florence Tan, Melissa Trainer

**NASA Headquarters**

Michael Meyer, Arik Posner, Mary Voytek

**NASA Jet Propulsion Laboratory (JPL)**

Robert C. Anderson, Andrew Aubrey, Luther W. Beegle, Alberto Behar, Diana Blaney, David Brinza, Fred Calef, Lance Christensen, Joy A. Crisp, Lauren DeFlores, Bethany Ehlmann, Jason Feldman, Sabrina Feldman, Gregory Flesch, Joel Hurowitz, Insoo Jun, Didier Keymeulen, Justin Maki, Michael Mischna, John Michael Morookian, Timothy Parker, Betina Pavri, Marcel Schoppers, Aaron Sengstacken, John J. Simmonds, Nicole Spanovich, Manuel de la Torre Juarez, Ashwin R. Vasavada, Christopher R. Webster, Albert Yen

**NASA Johnson Space Center (JSC)**

Paul Douglas Archer, Francis Cucinotta, John H. Jones, Douglas Ming, Richard V. Morris, Paul Niles, Elizabeth Rampe

**Nolan Engineering**

Thomas Nolan

**Oregon State University**

Martin Fisk

**Piezo Energy Technologies**

Leon Radziemski

**Planetary Science Institute**

Bruce Barraclough, Steve Bender, Daniel Berman, Eldar Noe Dobrea, Robert Tokar, David Vaniman, Rebecca M. E. Williams, Aileen Yingst

**Princeton University**

Kevin Lewis

**Rensselaer Polytechnic Institute (RPI)**

Laurie Leshin

**Retired**

Timothy Cleghorn, Wesley Huntress, Gérard Manhès

**Salish Kootenai College**

Judy Hudgins, Timothy Olson, Noel Stewart

**Search for Extraterrestrial Intelligence Institute (SETI I)**

Philippe Sarrazin

**Smithsonian Institution**

John Grant, Edward Vicenzi, Sharon A. Wilson

**Southwest Research Institute (SwRI)**

Mark Bullock, Bent Ehresmann, Victoria Hamilton, Donald Hassler, Joseph Peterson, Scot Rafkin, Cary Zeitlin

**Space Research Institute**

Fedor Fedosov, Dmitry Golovin, Natalya Karpushkina, Alexander Kozyrev, Maxim Litvak, Alexey Malakhov, Igor Mitrofanov, Maxim Mokrousov, Sergey Nikiforov, Vasily Prokhorov, Anton Sanin, Vladislav Tretyakov, Alexey Varenikov, Andrey Vostrukhin, Ruslan Kuzmin

**Space Science Institute (SSI)**

Benton Clark, Michael Wolff

**State University of New York (SUNY) Stony Brook**

Scott McLennan

**Swiss Space Office**

Oliver Botta

**TechSource**

Darrell Drake

**Texas A&M**

Keri Bean, Mark Lemmon

**The Open University**

Susanne P. Schwenger

**United States Geological Survey (USGS) Flagstaff**

Ryan B. Anderson, Kenneth Herkenhoff, Ella Mae Lee, Robert Sucharski

**Universidad de Alcalá**

Miguel Ángel de Pablo Hernández, Juan José Blanco Ávalos, Miguel Ramos

**Universities Space Research Association (USRA)**

Myung-Hee Kim, Charles Malespin, Ianik Plante

**University College London (UCL)**

Jan-Peter Muller

**University Nacional Autónoma de México (UNAM)**

Rafael Navarro-González

**University of Alabama**

Ryan Ewing

**University of Arizona**

William Boynton, Robert Downs, Mike Fitzgibbon, Karl Harshman, Shauna Morrison

**University of California Berkeley**

William Dietrich, Onno Kortmann, Marisa Palucis

**University of California Davis**

Dawn Y. Sumner, Amy Williams

**University of California San Diego**

Günter Lugmair

**University of California San Francisco**

Michael A. Wilson

**University of California Santa Cruz**

David Rubin

**University of Colorado Boulder**

Bruce Jakosky

**University of Copenhagen**

Tonci Balic-Zunic, Jens Frydenvang, Jaqueline Kløvgaard Jensen, Kjartan Kinch, Asmus Koefoed, Morten Bo Madsen, Susan Louise Svane Stipp

**University of Guelph**

Nick Boyd, John L. Campbell, Ralf Gellert, Glynis Perrett, Irina Pradler, Scott VanBommel

**University of Hawai'i at Manoa**

Samantha Jacob, Tobias Owen, Scott Rowland

**University of Helsinki**

Evgeny Atlaskin, Hannu Savijärvi

**University of Kiel**

Eckart Boehm, Stephan Böttcher, Sönke Burmeister, Jingnan Guo, Jan Köhler, César Martín García, Reinhold Mueller-Mellin, Robert Wimmer-Schweingruber

**University of Leicester**

John C. Bridges

**University of Maryland**

Timothy McConnochie

**University of Maryland Baltimore County**

Mehdi Benna, Heather Franz

**University of Maryland College Park**

Hannah Bower, Anna Brunner

**University of Massachusetts**

Hannah Blau, Thomas Boucher, Marco Carmosino

**University of Michigan Ann Arbor**

Sushil Atreya, Harvey Elliott, Douglas Halleaux, Nilton Rennó, Michael Wong

**University of Minnesota**

Robert Pepin

**University of New Brunswick**

Beverley Elliott, John Spray, Lucy Thompson

**University of New Mexico**

Suzanne Gordon, Horton Newsom, Ann Ollila, Joshua Williams

**University of Queensland**

Paulo Vasconcelos

**University of Saskatchewan**

Jennifer Bentz

**University of Southern California (USC)**

Kenneth Neelson, Radu Popa

**University of Tennessee Knoxville**

Linda C. Kah, Jeffrey Moersch, Christopher Tate

**University of Texas at Austin**

Mackenzie Day, Gary Kocurek

**University of Washington Seattle**

Bernard Hallet, Ronald Sletten

**University of Western Ontario**

Raymond Francis, Emily McCullough

**University of Winnipeg**

Ed Cloutis

**Utrecht University**

Inge Loes ten Kate

**Vernadsky Institute**

Ruslan Kuzmin

**Washington University in St. Louis (WUSTL)**

Raymond Arvidson, Abigail Fraeman, Daniel Scholes, Susan Slavney, Thomas Stein, Jennifer Ward

**Western University**

Jeffrey Berger

**York University**

John E. Moores

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