Genus topology of the cosmic microwave background from WMAP

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ABSTRACT
We have independently measured the genus topology of the temperature fluctuations in the cosmic microwave background seen by the Wilkinson Microwave Anisotropy Probe (WMAP). A genus analysis of the WMAP data indicates consistency with Gaussian random-phase initial conditions, as predicted by standard inflation.

Key words: cosmic microwave background – cosmology: observations.

1 INTRODUCTION
The greatly anticipated results from the Wilkinson Microwave Anisotropy Probe (WMAP) project (Bennett et al. 2003a) have defined the state of the art in cosmic microwave background (CMB) science. The WMAP project has provided full-sky coverage of the CMB at unprecedented angular resolution. The improvement over the only previous full-sky data set (COBE, Smoot et al. 1992) is of the order of 500 in terms of resolution elements, and the sky coverage improvement over balloon and ground-based experiments (e.g. de Bernardis et al. 2000, Boomerang; Stompor et al. 2002, Maxima; Mason et al. 2003, CBI; Kovac et al. 2002, DASI) is several hundred. The experiment has resulted in dramatically improved constraints on cosmological parameters, such as the matter density, Ωm, the Hubble Constant, H0, and the cosmological constant, ΩΛ (Spergel et al. 2003).

The tool of choice for assessing the cosmological parameters is the power spectrum, where one calculates the products of the spherical harmonic coefficients aℓm and their complex conjugates (for each ℓ and m in the Yℓm expansion of the CMB sky, summed over m values to form Cℓ). This aℓm̅aℓm̅ product, however, explicitly removes phase information. And while the power spectrum is a very powerful tool for assessing cosmological parameters such as Ωm and ΩΛ, it does (explicitly) remove any phase information contained in the aℓm coefficients themselves. These phases, however, contain critical information for characterizing the primordial density fluctuations. Namely, standard inflation (e.g. Guth 1981; Albrecht & Steinhardt 1982; Linde 1982, 1983) predicts that the temperature fluctuations in the CMB, at the resolution measured by WMAP, will be characterized by spherical harmonic coefficients with Gaussian distributed amplitudes and random phases. The WMAP data provide our best opportunity to date to test that hypothesis.

The genus topology method developed by Gott, Melott & Dickinson (1986) directly tests for the Gaussian random-phase nature of a density (or temperature) distribution in three dimensions (Adler 1981; Gott, Melott & Dickinson 1986; Hamilton, Gott & Weinberg 1986; Gott, Weinberg & Melott 1987) or in two dimensions (Adler 1981; Melott et al. 1989). Coles (1988) independently developed an equivalent statistic in two dimensions. The two-dimensional case has been studied for a variety of cosmological data sets: on redshift slices (Park et al. 1992; Colley 1997), on sky maps (Gott et al. 1992; Park, Gott & Choi 2001a; Hoyle, Vogele & Gott 2002); and on the CMB, in particular (Smoot et al. 1992; Kogut 1993; Kogut et al. 1996; Colley, Gott & Park 1996; Park et al. 2001b). Watts & Coles (2003), Chiang & Coles (2000), among many others, have investigated other methods for measuring phases, such as looking at the Fourier modes directly.

The WMAP team has recently measured the genus of the WMAP sky (Komatsu et al. 2003), and demonstrated that the WMAP results are consistent with the Gaussian random-phase hypothesis. To do this, they carried out a large number of simulations of the CMB, in which the spherical harmonic coefficients were drawn from a Gaussian random-phase distribution. They then used their known beam profiles, to simulate the results in each frequency, and applied the Kp0 (Bennett et al. 2003b) mask, just as one would with the real data set. For each of these simulations, they computed the genus (as defined by Melott et al. 1989) and compared it with that of the real data set. They found that the real data set does not depart significantly from the genus of the simulated Gaussian random-phase data sets, very much in agreement with the results predicted for WMAP by Park et al. (1998). Bennett et al. (2003b) further explored Gaussianity by testing other Minkowski functionals (Minkowski 1903) and the bispectrum (related to the three-point correlation function), and found constraints on Gaussianity similar to those provided by the genus study.

We seek, first, to confirm this result using our own methods, but also to compare the genus measured from the data directly to the theoretical genus curve for structures on a sphere. Rather than simulating Gaussian random-phase realizations of the CMB, we compare directly with the theoretical prediction for the genus in two dimensions. Furthermore, we provide some details concerning the nuances of carrying out the genus calculation on the HEALPix map projection and the stereographic map projection.
The WMAP team has released the data\(^1\) in a unique map projection called the HEALPix projection (Górski et al. 2000). We have chosen to re-present the WMAP results in Fig. 1 for a few reasons. First, we will be studying the genus on stereographic projections (Sections 6 and 7), so it is useful to see the full-sky map in that projection. Secondly, we have chosen a very different colour scheme that is more relevant to genus studies than the original colour scheme used by the WMAP team. The WMAP colour scheme is pretty, informative and impressive, but is not symmetric with respect to hot and cold spots. In our colour scheme, the mean temperature contour is white. For temperatures higher than the mean the amount of red ink per pixel is linearly proportional to the hotness of a hotspot; for temperatures below the mean, the amount of blue ink per pixel on the page is linearly proportional to the coldness of a cold spot. The scale runs from \(-200\) \(\mu\)K (solid blue) to +200 \(\mu\)K (solid red), the same range as in the temperature maps produced by the WMAP team (Bennett et al. 2003a). Since the two-dimensional genus of a Gaussian random-phase field is all about the symmetry between hotspots and cold spots, we find this colour map highly instructive for our purposes.

3 GENUS TOPOLOGY ON A SPHERE

The properties of the genus are well known in three dimensions (3D) (Gott et al. 1986; Hamilton et al. 1986; Gott et al. 1987), but require some explanation in the two-dimensional (2D) case (Melott et al. 1989), particularly in the case of the sphere (Coles & Plionis 1991; Gott et al. 1990).

For 2D topology on a plane the genus of a microwave background map is defined as (Melott et al. 1989)

\[
g_{2D} = \text{number of hot spots} - \text{number of cold spots}. \tag{1}
\]

For a Gaussian random field,

\[
g_{2D} = A\nu \exp(-\nu^2/2), \tag{2}
\]

where the amplitude, \(A\), is a function only of the power spectrum, and \(\nu\) is a parameter that measures the area fraction in the hotspots:

\[
f = (2\pi)^{-1/2} \int_0^\infty \exp(-x^2/2) \, dx. \tag{3}
\]

So for \(\nu > 0 (f < 0.5)\) there are more hotspots than cold spots, while for \(\nu < 0 (f > 0.5)\) there are more cold spots than hotspots. The genus is also equal to the integral of the curvature around the temperature contour divided by \(2\pi\). If we were to drive a truck around an isolated hotspot we would have to turn a total angle of \(2\pi\) as we complete an entire circuit around the hotspot. Driving a truck around an isolated cold spot, we would turn a total angle of \(2\pi\) with the opposite sign, with a negative turn angle defined as one that is a turn to the left when the hot region is on your right. Thus, in a temperature field that is divided into pixels we may define a pixel as hot if it is above the contour threshold and cold if it is below the contour threshold. Then the contour line is actually composed of a series of line segments with turns occurring at vertices in the pixel map. There is a certain turn occurring at each vertex (which, divided by \(2\pi\), is equal to the contribution to the genus). Around each vertex there are four pixels, the contribution to the total turn (and genus)

\(^1\) The WMAP homepage is at http://map.gsfc.nasa.gov/

\(^2\) WMAP data sets are available at http://lambda.gsfc.nasa.gov/
at that vertex depends on how many of the four surrounding pixels are hot and cold:

- 0 hot, 4 cold: genus = 0, the contour line does not intersect the vertex.
- 1 hot, 3 cold: genus = +1/4, the contour line makes a right angle turn.
- 2 hot, 2 cold: genus = 0, no turn, or turns that add to zero on average.
- 3 hot, 1 cold: genus = −1/4, right angle turn around a cold spot.
- 4 hot, 0 cold: genus = 0, contour line does not intersect the vertex.

This is carried out by an identical algorithm to that in the CONTOUR2D program as described in Melott et al. (1989).

Now we wish to define rigorously the 2D genus on a spherical surface. Here we will follow our previous derivation (Gott et al. 1990). The 2D genus is defined to be equal to minus the 3D genus of the hot regions confined within a thin spherical shell, measured in 3D. Use lead paint to paint the hot regions on the surface of a balloon – burst the balloon and you will have solid curved lead shapes that will have a certain 3D genus – take the minus of this number and that will be the 2D genus as we will define it.

Recall that we have defined the 3D genus as the number of (doughnut) holes minus the number of isolated regions. Our 3D genus is equal to the integral of the Gaussian curvature over the contour surface divided by −4π. Thus, a sphere in 3D has a genus of −1, because it is one isolated region. The Gaussian curvature is +1/r^2 while the area of the sphere is 4πr^2, so the integral of the Gaussian curvature over the sphere is 4π, independent of the size of the sphere, and the genus is −1. A cube has a genus of −1 also, because it has the same topology as a sphere. In this case, the curvature of the surface entirely consists of eight delta functions at the eight vertices where three squares meet at a point introducing a contribution to the Gaussian curvature integral of π/2 at each vertex, equal to the conical deficit angle at the vertex. (If we sandpapered off the corners and edges of the cube, we would produce at each vertex an octant conical deficit angle at the vertex. (If we sandpapered off the corners because it is one isolated region. The Gaussian curvature is −1 because this would look like a sugar bowl without any handles, which is also one isolated region. The topology in each case is identical since one can be deformed into the other, so the genus should be the same in all three cases. If the hot region ran around the equatorial regions and there were cold spots at the northern and southern polar regions, the 2D genus would be 0 because in 3D the equatorial band (confined in a thin spherical shell) is a doughnut in 3D, which has a 3D genus of 0. If the sphere had N isolated hotspots (polka dots) on a cold background, the 2D genus would be N. If the sphere had N isolated cold spots on a hot background the genus would be 2 − N. [Remember that one isolated hotspot on a cold background (a hotspot centred on the north pole; the rest cold) is the same topologically as an isolated cold spot centred on the south pole (the rest hot), because the latitude of the contour line can be simply moved downward to transform one into the other.] Suppose we have a hotspot covering the northern hemisphere with the southern hemisphere cold. This is a spherical cap of area 2πr^2. Its 2D genus is +1 because in 3D it is one isolated region. Considered as a spherical cap (part of a thin spherical shell) this isolated region must have an integral of the Gaussian curvature on its surface equal to 4π by the Gauss–Bonnet theorem. There are several contributions to this: there is a contribution of 2π from the outside (spherical) cap surface and a contribution of 2π from the inside (spherical) cap surface (a spherical shell has an inside and an outside boundary both spherical), and a net contribution of 0 from the delta function curvatures on the inside and outside edges at the equator marking the boundaries of the inner and outer spherical caps – the contribution of the outer edge is positive, while the contribution of the inner edge is saddle-shaped and therefore negative. The equator is a geodesic, so we could drive a truck around the equator without turning; hence, the integral of the turning around the contour boundary is zero. Since we are circling an isolated hotspot, why is the total turning not 2π as it is on the plane? Because we are circling a curved region, and if we parallel transport (without turning) a vector around the boundary it will suffer a deflection equal to the integral of the Gaussian curvature in the interior of the boundary, which in this case is 2π, the value we expect for a closed curve in the plane. The vector must return to its initial position after circling, so the total of the turning integral and the parallel transport deflection due to circling a curved region must add to give 2π.

Now consider a hotspot centred at the north pole and for which the boundary extends to latitude 30° N. This is a spherical cap with area πr^2. It is an isolated region so the 3D genus is −1, and the sum of the integral of the Gaussian curvature over the spherical cap in 3D must be 4π. The integral on the outside of the spherical cap is πr^2, and on the inside of the spherical cap is also πr^2, and the boundary surface connecting the inside and outside cap shells is locally flat (part of a cone), so the contribution from the inside and outside edges of the boundary must be 2π, which must be equal to twice the turning integral for a truck driving on the spherical surface [since the total Gaussian integral, 4πr^2, is twice the 2D genus for an isolated boundary (turning plus deviation from parallel transport)]. If we were to parallel transport a vector around this boundary, we would find a deflection of π because that is the area in the spherical cap; to return to its original orientation, a rotation of 2πr, the truck must turn an additional angle of π. Indeed, if you would like to drive counterclockwise around the sphere at latitude 30° N, you would have to keep your steering wheel turned to the left, since this not a geodesic, and the total of that turning integral is π. Now, locally it is the turning integral that is being calculated from the formula for a Gaussian random field. So let us define the effective genus as

\[ g_{2D,\text{eff}} = g_{2D} - 2f, \]

where \( f \) is the fraction of the area of the sphere in the hotspots. In our example, where the one hotspot was the entire northern hemisphere, \( g_{2D} = 1, f = 1/2 \) and \( g_{2D,\text{eff}} = 0 \). The value of \( g_{2D,\text{eff}} \) is equal to the turning integral of a truck driving around all the individual components of the boundary contour (around all the hotspots) divided by 2π; in this case it is 0 because the truck driving around the equator drives straight ahead without turning. For the case where the hotspot...
is centred at the north pole and extends to latitude 30°N, \( f = \frac{1}{2} \), so the value of \( g_{2D, \text{eff}} = \frac{1}{2} \), because the turning integral in this case is \( \pi \). Now for a Gaussian random field on the sphere, 
\[
g_{2D, \text{eff}} \propto v \exp(-v^2/2),
\]
(5)
because the Gaussian random field locally behaves on the sphere as on the plane to produce this contribution to the turning integral. Thus, in comparing the WMAP data with the random-phase formula we will use \( g_{2D, \text{eff}} \) defined rigorously as described above.

### 4 GENUS IN THE HEALPIX PROJECTION

The WMAP data are plotted using an unusual map projection of the sphere, called the HEALPix projection (Górski et al. 2000). The sphere is effectively projected on to a rhombic dodecahedron. (This semiregular polyhedron is the cell for the face-centred cubic crystal lattice, the polyhedron representing the boundary of the set of points closer to a particular atomic nucleus than to any other.) It has 12 faces, four diamonds that meet at the top like the four sides of a pyramid, four diamonds that circle the equator, and four that meet at the bottom like the sides of an inverted pyramid. If we were to set this rhombic dodecahedron up aligned with the cardinal directions we could label the 12 faces as follows: TN (top north), TE (top east), TS (top south), TW (top west), NE (northeast, on the equator), SE (southeast, on the equator), SW (southwest, on the equator), NW (northwest, on the equator), BN (bottom north), BE (bottom east), BS (bottom south) and BW (bottom west). There are six vertices where four diamonds meet at a point (i.e. TN, TE, TS, TW all meet at the top, and similarly TN, NE, NW, BN meet at a vertex). There are eight vertices where three diamonds meet at a point (e.g. TN, TE, NE meet at a vertex). Each diamond may be subdivided into diamond-shaped pixels. This is done by factors of 4, subdividing each diamond into four diamonds, and repeating by similarly subdividing each diamond. (In a similar way we may produce a chequerboard of 64 square pixels by dividing a square first into four quadrant squares, dividing each of these into four subsquares, and each of these subsquares into four chequerboard squares). In the WMAP case these are all diamonds (rhombuses) instead of squares, but the topology is the same. The subdivisions are made so that all the pixels are of equal area, and diagonal rows of pixels touching point to point have centres that lie on circles of latitude. With \( N \) successive subdivisions of the diamonds into four subdiamonds, we have a total of \( 12 \times 4^N \) pixels of equal area covering the sphere. For example, WMAP uses \( N = 9 \), where there are 3 145 728 diamond-shaped pixels covering the sphere, each with dimensions of approximately \( 0.11 \times 0.11 \) deg\(^2\), which is adequate resolution to show the WMAP data, which has an angular resolution of approximately 0.3. To study the topology properly, one needs pixels that are at least 2.5 times smaller than the smoothing length.

Each of these diamond pixels on the sphere has angles that would be complicated to calculate, so a version of \textsc{contour2d} for the sphere would seem to be complex and require many calculations to calculate the turning angles. However, we may simplify this greatly by using the topological invariance of \( g_{2D} \) to projection (first from the sphere to the rhombic dodecahedron, then from the rhombic dodecahedron to a cube). This cube has six faces, which we will label T (top), N (north), E (east), S (south), W (west) and B (bottom). Each diamond of the rhombic dodecahedron is oriented with its short diagonal along an edge of the cube, and each of the 12 rhombuses is mapped into two triangles that appear on adjacent faces of the cube. For example, the rhombus TN is divided into two triangles, one of which is mapped on to the T map on the cube and the other is mapped on to the N face of the cube. In fact, the letter designation of each rhombus tells us which of the two cubic faces it is mapped on to. For example, the SW rhombus is divided into two triangles that are mapped on to the S and W faces of the cube. The T face of the cube is thus a square that is divided into four right triangles for which the hypotenuses form the sides of the square, and which meet in the centre; these four triangles represent the upper halves of the four rhombuses TN, TE, TS, TW, which meet at the top.

In such a projection we paint the hot regions with lead paint on the surface of the cube, and the 3D genus of these lead shapes is the same as if they were painted on to the sphere, because they have only been distorted in shape, not changed in topology. The total integrated Gaussian curvature over the cube, which totals \( 4\pi \) is concentrated entirely in eight delta functions at the eight vertices, where there is an angle deficit of \( \pi/2 \) at each, as three squares instead of four meet at each of these eight vertices. Each of the 12 rhombuses of the original dodecahedron is mapped on to two right triangles meeting at their hypotenuses along an edge of the cube; these can be flattened out to make a square, with square pixels. So we have equivalently, 12 square maps of \( 4^N \) square pixels each. Thus, we can consider each rhombus as a square map with square pixels and we can calculate \( g_{2D} \) for each using \textsc{contour2d}. Vertices that are within the rhombus are included. Vertices that are along an edge with the next rhombus are shared according to a prescription where the vertices along the top and right-hand edges of the rhombus are assigned to it when laid out properly, and the bottom and left-hand edges are assigned to adjacent rhombuses, since two rhombuses meet along each edge. That accounts for all pixel vertices except for those at the corners of the rhombuses. Each rhombus has two corners where four rhombuses meet (such as TN, TE, TS, TW meet at the top) and two corners where three rhombuses meet at a vertex (such as TN, TE, NE) which occurs at a vertex of the cube where there is a \( \pi/2 \) angle deficit in our cubic map and where three pixels only meet at a point. The corners where four rhombuses meet at a point are like ordinary vertices in the plane where four square pixels meet at a point, so this is handled with \textsc{contour2d}. The value of the contribution to the genus from that vertex must then be divided and shared equally among all the rhombuses that meet there. The prescription for vertices where three pixels meet at a point (at the corner of the cube in our projection) is as follows:

| 0 hot, 3 cold | genus = 0 | the contour does not intersect the vertex |
| 1 hot, 2 cold | genus = 1/4 | added to hot rhombus, 0 added to others |
| 2 hot, 1 cold | genus = 0 | |
| 3 hot, 0 cold | genus = 1/12 | added to each of the three rhombuses. |

If we were to smooth the edges of the cube by sanding, the edges would turn into quarter cylinders (with zero Gaussian curvature) and the eight vertices (where three squares meet at a point) turn into tiny octants of a sphere each with an integrated Gaussian curvature of \( \pi/2 \). The faces of the cube are flat so all the curvature is located in the eight vertices. The edges of the three rhombuses meet at 120° angles in the spherical octant. So in the case of one hot and two cold there is a turn of 60° in the contour plus a curvature of 30° (equal to one-third of the integrated curvature over the octant of the sphere) so the genus contribution (turn plus parallel transport deviation 90°) should correspond to \(+1/4\) (or \( \pi/2 \) divided by \( 2\pi \)). This is then added to the hot rhombus as this is the hotspot we are circling. In the case of two hot, one cold, there is a \(-60° \) turn, plus two regions of 30° curvature contributing to the parallel transport.
deviation, making a total contribution of 0 to the genus. In the case of three hot, zero cold, the vertex lies entirely within the hot region but it adds $\pi/2$ to the curvature integral within that region and this must be shared equally with all three rhombuses to add $+1/12$ to the genus in each. The total genus for the sphere $g_{2D}$ is calculated by adding the genus from all 12 rhombuses. From $g_{2D}$ we can calculate $g_{2D, \text{eff}}$ by subtracting $2f$. We can calculate $f$ as simply the fraction of the diamond-shaped pixels that are hot, since these are all of equal area. Thus, from the WMAP data we can calculate the 2D topology and compare it with that expected from a Gaussian random-phase distribution.

5 THE GENUS OF THE CMB, MEASURED BY WMAP

The most direct data product from the WMAP team is the ‘internal linear combination’ (ILC) map of the CMB, which is given in the HEALPix format. This ILC map uses the optimum linear combination of the skymaps at the different frequencies to remove the Galaxy and some other foregrounds (Bennett et al. 2003b). This is the primary map distributed as the best rendering of the CMB and has a resolution (beamwidth) of $0.3 \times 0.3$ deg$^2$. This map presents a few problems for directly computing the genus, not the least of which is that the WMAP team advised against statistical studies of this map (Bennett et al. 2003b). We will address these issues in the next section, but it is worth checking the genus on this map directly.

The HEALPix rhombus projection provides a natural division of the data set (sky) into 12 independent regions. We therefore measured the genus in each of these regions and computed the total for the whole sky, which we plot in Fig. 2. In this figure $\nu$ is computed in terms of the pixelwise temperature mean and standard deviation measured over the whole sky (method 1). This method would be equivalent to the area fraction method described by equation (3) if the pixel histogram of temperature were strictly Gaussian. Fig. 3 is identical, except that $\nu$ is computed by area fraction in each individual

![Figure 2](https://example.com/figure2.png)

**Figure 2.** Total genus, $g_{2D, \text{eff}}$, for the original HEALPix projection of the WMAP data (internal linear combination) for the entire sky, with no masking. The abscissa, $\nu$, is the departure from the pixelwise temperature mean for the whole sky in terms of pixelwise temperature standard deviation for the whole sky. Inner error bars are computed from the standard deviation of the 12 HEALPix diamonds; the outer error bars reflect the sigma value scaled for 95.4 per cent confidence in the student’s $t$ distribution.

![Figure 3](https://example.com/figure3.png)

**Figure 3.** As in Fig. 2, except the $\nu$ values are computed by area fraction (see the text).

HEALPix rhombus (method 2); see equation (3). Using the area fraction method measures the random-phase nature of the distribution more directly, since it separates that from any departures from a Gaussian histogram of the temperature (which could be measured directly). In method 1, we use $\nu$ defined in terms of the standard deviation in temperature over the whole sky; in method 2, we define $\nu$ by area fraction in each rhombus, thus treating each of the 12 rhombuses completely independently. The two methods provide a contrast of data treatment, and yet, as we shall see, give essentially identical results.

The inner error bars in Figs 2 and 3 are the standard deviation of the mean in the genus for the whole sky, estimated from those of the 12 independent regions of the sky (adding the values of $g_{2D, \text{eff}}$ obtained from each). The standard 1$\sigma$ error bars are shown as the inner error bars. The outer error bars show the 95.4 per cent (Gaussian 2$\sigma$) confidence interval for a student’s $t$ variable with 12 degrees of freedom, which we will explain shortly.

The solid curves in Figs 2 and 3 give the Gaussian random-phase genus curve, according to equation (2), with the best-fitting amplitude applied. The amplitudes, $A$, [$g_{2D, \text{eff}} = Av\exp(-\nu^2/2)$], in Figs 2 and 3 are $A = 3432$ and 3657, respectively. These are by far the highest in any genus measurements of cosmological structure to date. As expected, the first value of $A$ is lower than the second one, because the power at large scales (quadrupole and octopole) lifts some regions (rhombuses) to higher temperature, consolidating some cold spots therein, but lowers other regions (rhombuses), consolidating some cold spots therein. Method 2 sets a median in each rhombus separately and thus discovers more structures.

The student’s $t$ formulation is necessary since we are estimating the standard error with the distribution itself. Namely, if the true genus for the whole sky at each point on the curve were $\mu$ and the true standard deviation in the mean expected for that number of structures were $\sigma$ (we expect the distribution to be Gaussian for Gaussian random-phase fields, Gott et al. 1990), then our estimator for $\mu$ is the arithmetic average estimated from the 12 rhombuses (the values from each rhombus are multiplied by 12, then they are averaged), $\bar{x}$, and our estimator for $\sigma$ is $s/\sqrt{n-1}$, where $s$ is the root-mean-square difference from the mean (as usual). We expect that $\mu$ is given exactly by equation (2) and that our best fit for the amplitude has negligible error compared with any one point on the curve.
expression \((\bar{x} - \mu)/(s/\sqrt{n-1})\) is defined to be a student’s \(t\) variate (Lupton 1993).

On average, we would expect only one of the 21 points in each of the genus curves in Figs 2 and 3 to miss the curve outside of the 95.4 per cent confidence error bar. In fact, not even one misses. From the properties of the student’s \(t\) distribution with 12 d.o.f., one would expect to find that 33.7 per cent (\(\sim 7\)) of the 21 points miss the curve by more than the inner error bars. In both cases, the agreement is again better than expected, with only three misses in each figure. This better-than-expected agreement can easily arise by chance from correlations between the genus measurements at neighbouring values of \(v\), as shown by Colley (1997) (these correlations mean that the 21 data points are not completely independent—neighbouring genus values measure many of the same structures).

None the less, the genus of temperature fluctuations in the CMB, as observed by WMAP, is certainly consistent with the Gaussian random-phase hypothesis.

6 GENUS ON A STEREOGRAPHIC MAP PROJECTION

In the ILC HEALPix data release, there are a few issues that must be addressed. The first issue is foreground emission. Bennett et al. (2003b) discuss in quite a bit of detail the foregrounds in the WMAP results. They provide a very useful sky mask that should be applied to avoid foreground contaminants, the principal one of which is the Galaxy. In a second study, we have therefore used their \(K\) region (most conservative) sky mask, which also removes some 200 point sources. Furthermore, we have restricted ourselves to Galactic latitudes \(|b| > 18^\circ\). Since the mask must remove much of the area near the Galactic equator, the HEALPix projection, with its four equatorial diamonds, ceases to be ideal. Instead, we re-project the map, using the stereographic projection for both the Northern and Southern Galactic Hemispheres (as in Fig. 1, in which the \(|b| > 18^\circ\) cut occurs at 72.7 per cent of the total radius in each hemisphere).

We can calculate \(g_{2D}\) from the north and south galactic caps by using stereographic map projections of the caps. Stereographic map projections of a hemisphere are conformal and preserve angles. A stereographic projection of the north galactic cap is a circle, in which azimuthal angle in the map is the galactic longitude in the cap. The radius in the map, \(r = 2 \tan((90^\circ - b)/2)\) (see Fig. 1). The outside boundary is a circle of radius \(r_s = 2 \tan((90^\circ - 18^\circ)/2)\). The contours on the sphere can be approximated as spherical polygons—geodesic segments joined at vertices where turning on the sphere occurs. The stereographic projection preserves angles, so the turning that occurs at vertices is mapped properly. However, geodesics (great circles) are mapped into arcs of circles rather than straight lines, which turn on the map but not on the sphere so the total genus \(g_{2D}\) on the sphere is not automatically calculable from the total turning that comes from circling the density contour on the map.

So instead of calculating \(g_{2D}\) for the sphere from the stereographic map directly, which would be complicated, let us instead calculate the relation between \(g_{2D,\text{eff}}\), the spherical cap, and \(g_{2D,\text{map}}\) measured by the \textsc{contour2D} program on the flat stereographic map of the cap. [Of course once we have \(g_{2D,\text{eff}}\) we can calculate the contribution to \(g_{2D}\) due to the structures in the north galactic cap by adding \(2f \text{area}_{\text{cap}}/(4\pi)\).] However, it is really \(g_{2D,\text{eff}}\) we are interested in anyway because it is what will be compared with the theoretical formula. Thus, we can skip a step by going directly after \(g_{2D,\text{eff}}\).

To begin, let us excise the spherical cap from the sphere by cutting it off and separating it from the rest of the sphere. This will add boundary terms. Each hotspot (think of a curved lead sheet) that intersects the boundary of the cap will be closed off by a curve along the boundary that will make it an isolated hotspot. Think about these hotspots in 3D when the cap is excised from the rest of the sphere: any hotspot that hits the boundary will become isolated, floating in space, because beyond the boundary there is nothing. This spherical isolated cap will have a value of \(g_{2D,\text{excised cap}} = g_{2D,\text{excised map}}\); the total genus of the isolated flat stereographic map sitting isolated in the plane—think of flat lead sheets lying in a plane, because one can be continuously deformed into the other. We can calculate the terms contributing to \(g_{2D,\text{excised cap}}\):

\[
g_{2D,\text{excised cap}} = (2\tau)^{-1} \sum (\text{turns on spherical cap})
+ f \text{area}_{\text{cap}}/(2\pi)
+ (2\tau)^{-1} \sum (\text{turns at cap boundary})
+ f_b[1 - \text{area}_{\text{cap}}/(2\pi)].
\]  

The first term, \((2\tau)^{-1} \sum (\text{turns on spherical cap})\), is equal to \(g_{2D,\text{eff, cap}}\); the interior of the spherical cap. It is the total number of turns made on the sphere around all the contours within the spherical cap, and excludes any boundary terms produced by excising the spherical cap. Now \(g_{2D,\text{eff, cap}} \propto \nu \exp(-\nu^2/2)\), so it is what we want to calculate to compare with the theoretical genus formula. The term \(f \text{area}_{\text{cap}}/(2\pi)\), where \text{area}_{\text{cap}}\) is measured in steradians, is the integrated Gaussian curvature within the hotspots in the cap divided by \(2\pi\), which contributes to the genus by the 3D argument presented above. The term \((2\tau)^{-1} \sum (\text{turns at cap boundary})\) is the sum of the turns that are made when you are driving along a contour and you encounter the cap boundary and then must turn a sharp corner there to continue driving along the cap boundary to complete circling that hotspot as an isolated region. In general, if along the boundary there are \(N\) hot segments, there will be \(2N\) such turns, as we enter and then exit each of these \(N\) hot segments along the boundary. Each of these \(2N\) turns will be some angle between 0 and \(2\pi\). The final term, \(f_b[1 - \text{area}_{\text{cap}}/(2\pi)]\), represents the total turning on the sphere that occurs during driving along the \(N\) hot segments of the boundary of the cap. If the entire boundary were hot then driving around it would have a total turn of \(2\pi - \text{area}_{\text{cap}}\) (For example, if this were a hemispherical cap the area of the cap would be \(2\pi\) and the total turn circling the boundary would be zero, because the boundary is the equator and it is a geodesic so no turning is required to circle the boundary.) As the area of the cap goes to zero, the curvature within the cap becomes unimportant to parallel transport and the total turn required to circle the boundary approaches \(2\pi\). In general, the angle deflection suffered by parallel transport around the closed boundary curve is equal to the integral of the Gaussian curvature inside, which in this case is equal to the area of the cap. When we have completed the circuit, we have returned to where we have started so we have suffered a total rotation of \(2\pi\), so the turning we do on the sphere circling the boundary plus the area of the cap is \(2\pi\). Thus, turning \(= 2\pi - \text{area}_{\text{cap}}\). The number \(f_b\) is the fraction of the boundary that is hot, and so it is this fraction of the boundary that we drive around. Dividing this by \(2\pi\), gives the contribution to the genus derived by driving along the hot segments of the boundary: \(f_b[1 - \text{area}_{\text{cap}}/(2\pi)]\). Using the Copernican principle, one would not expect the boundary to be special, so on average the fraction of the cap boundary that was hot should be equal to the fraction of the sphere that is in hotspots. On average then, we expect \(f_b = f\).
Substituting we find:
\[
\begin{align*}
\mathcal{G}^{2D,\text{excised\,cap}} &= \mathcal{G}^{2D,\text{eff,\,cap}} + \frac{f \, \text{area}_{\text{cap}}}{2\pi} \\
&\quad + \frac{1}{2\pi} \sum (\text{turns at cap boundary}) + f \left(1 - \frac{\text{area}_{\text{cap}}}{2\pi}\right) \\
&\quad + \frac{1}{2\pi} \sum (\text{turns at cap boundary}).
\end{align*}
\]

(7)

\[
\begin{align*}
\mathcal{G}^{2D,\text{excised\,map}} &= \mathcal{G}^{2D,\text{eff,\,map}} + f \\
&\quad + \frac{1}{2\pi} \sum (\text{turns at map boundary}).
\end{align*}
\]

(8)

Now let us measure the genus on the excised stereographic map \(\mathcal{G}^{2D,\text{excised\,map}}\), recalling that the excised stereographic map is topologically equivalent to the excised cap:
\[
\begin{align*}
\mathcal{G}^{2D,\text{excised\,map}} &= \frac{1}{2\pi} \sum (\text{turns on stereographic map}) \\
&\quad + \frac{1}{2\pi} \sum (\text{turns at map boundary}) + f_b.
\end{align*}
\]

(9)

The first term, \((2\pi)^{-1} \sum (\text{turns on stereographic\,map}) = g_{2D,\text{map}}\), is the quantity measured by CONTOUR2D on the flat map. CONTOUR2D examines only the Cartesian vertices within the stereographic map region and does not include any boundary effects. There is no ‘f area’ term because the map is flat and there is zero curvature inside the hotspots. The next term, \((2\pi)^{-1} \sum (\text{turns at map boundary})\), adds the turns that are taken when the contour hits the outer circular boundary of the map. If we use a stereographic projection, which is conformal and preserves angles, each and every turn at the boundary in the map will be equal to that encountered on the sphere, so
\[
\frac{1}{2\pi} \sum (\text{turns at map boundary}) = \frac{1}{2\pi} \sum (\text{turns at cap boundary}).
\]

(10)

The final term, \(f_b\), is the total turn taken on the \(N\) hot segments of the boundary of the map divided by \(2\pi\). If we circled the entire boundary we would have to turn a total of \(2\pi\), because the map is flat and Euclidean geometry applies. The \(N\) hot segments cover a fraction \(f_b\) of the boundary (galactic longitude is mapped on to azimuth in the stereographic map, so the fraction of the map boundary occupied by the \(N\) hot segments is exactly the same as on the sphere). Thus, the total turn is \(2\pi f_b\), and dividing this by \(2\pi\) gives \(f_b\). As we have remarked above, if the boundary is not special we expect \(f_b = f\). Since the map is excised and sits alone on the plane, these outer boundary segments must be included to create isolated hotspots in the plane out of the hot regions that hit the boundary. Equating \(g_{2D,\text{excised\,cap}}\) and \(g_{2D,\text{excised\,map}}\) (because the cap can be deformed into the map), and substituting from the above equations we find
\[
\begin{align*}
\mathcal{G}^{2D,\text{excised\,cap}} &= \mathcal{G}^{2D,\text{excised\,map}} \\
\mathcal{G}^{2D,\text{eff,\,cap}} + f + \frac{1}{2\pi} \sum (\text{turns at cap boundary}) \\
&= \mathcal{G}^{2D,\text{map}} + \frac{1}{2\pi} \sum (\text{turns at map boundary}) + f \\
\mathcal{G}^{2D,\text{eff,\,cap}} &= \mathcal{G}^{2D,\text{map}}.
\end{align*}
\]

(11)

(12)

(13)

So ‘what you see is what you get’. If we measure the 2D topology in the interior of the stereographic map using CONTOUR2D (which ignores the boundary) our result will equal, on average, \(g_{2D,\text{eff,\,cap}}\), which is what we were looking for. This works for any radius spherical cap, since terms proportional to the area of the cap cancel out.

And since the map is conformal, the turns at the boundary are equal in both cases and cancel out as well. Finally, since we expect the boundary to be typical, its coverage with hotspots should be equal to the fraction \(f\) of the sphere covered by hotspots, and this causes other terms to cancel as well, leaving the above simple result.

7 GENUS TOPOLOGY ON STEREOGRAPHIC WMAP RESULTS

We may now compute the 2-D genus on a stereographic re-projection of the WMAP data, in which we address the issues of foreground contamination and noise variability arising from the scanning strategy of the WMAP spacecraft.

First, to alleviate contamination from the Galaxy, we completely exclude the equator region up to \(|b| < 18^\circ\). The Kp0 mask does include removal of local regions for point source avoidance, and some additional Galaxy masking above the \(|b| > 18^\circ\) cut, particularly near the Galactic Centre, the Gum Nebula and Orion Complex. Any vertex in the stereographic map projection that falls within the mask is simply ignored by CONTOUR2D.

We proceed by dividing the stereographic projection into octants that correspond to octants on the sphere. These octants form eight independent regions in which to measure the genus. A slight adjustment for the actual solid angle in each octant is made, because the Kp0 mask changes the solid angle of each octant very slightly (a few per cent variation). We normalize the genus to the mean solid angle of the masked octants, and proceed treating each masked octant as one of eight independent measurements of the genus per average masked octant.

In carrying out the re-projection to the stereographic map, we have applied a slight smoothing to a total smoothing FWHM of 0.35. Figs 4 and 5 show the genus for the masked stereographic projection, computed by methods 1 and 2 (the same methods as in the HEALPix genus computation). The error bars are now computed for eight octants instead of 12 diamonds, which changes the relative size of the 95.4 per cent error bars and the direct 1σ error bars.

![Figure 4](https://academic.oup.com/mnras/article-abstract/344/3/686/1011161/25/1011161)
somewhat. The best-fitting theoretical genus curve (equation 2) is plotted in each figure, with respective amplitudes of $A = 2057$ and 2186 (as with the HEALPix genus curves, the first value is slightly less than the second, as expected). The amplitudes have come down somewhat from the HEALPix maps due to the exclusion of the equatorial band (so the total genus covers only a fraction of the whole sphere), the additional masked pixels in the $Kp0$ prescription, and the additional smoothing applied in the stereographic projection.

In each of the figures, the agreement with the theoretical genus curve is striking. Quantitatively, we would expect on average that one of the 21 points sampling the curve would miss the theoretical fit outside the 95.4 per cent error bar. With method 1 (Fig. 4), no points miss the theoretical curve outside their 95.4 per cent error bar, while with method 2 (Fig. 5), two points miss by a small margin. The 1σ (inner error bars) for the eight degrees of freedom of the octants reside at the 65.3 per cent confidence interval, so on average, one would expect approximately seven of the 21 points on each of Figs 4 and 5 to miss the theoretical curve outside their 1σ error bars. Using method 1, we find that three points miss at this margin and nine miss using method 2. Though the method 2 results show slightly more than the average number of misses expected at both the 1σ and 95.4 per cent level, the excess is certainly within the uncertainty arising from small-number statistics in each case.

The second contaminant in the ILC HEALPix map is that the noise behaviour is very complicated, and differs substantially from pixel to pixel. First, the beams at each frequency in the combination have different widths. Secondly, the scan method of the WMAP satellite yields many more samples at the ecliptic poles than at the ecliptic equator (see Bennett et al. 2003a, Fig. 3). To alleviate some of these problems, we have downloaded from the WMAP website maps in the same five frequencies as used in the ILC map, but ones that have been pre-smoothed to the same effective beamwidth (0.82). These are then added with the same weights as used in producing the ILC map. At the cost of the number of structures, these maps significantly enhance the signal-to-noise ratio of the remaining structures, thus reducing the significance of the noise differences arising from the scanning strategy. This is, therefore our most conservative map.

In projecting these maps into stereographic maps, we have smoothed to 0.9, slightly more than the 0.82 smoothing of the pre-smoothed maps. Figs 6 and 7 show the genus measured in these stereographic projections. The best-fitting amplitudes have dropped to $A = 772$ for method 1 and $A = 831$ for method 2, due to the smoothing.

From these final stereographic projections, the genus measured at the 21 points should also be expected to miss the theoretical curve outside of the 95.4 per cent error bars once on each figure. What we find is that with method 1 (Fig. 6), no points miss the curve outside the 95.4 per cent error bar, while with method 2 (Fig. 7), one point misses by a small amount. The points miss the curve outside their 1σ error bars twice for method 1 and four times for method 2, again a bit better than expected. The final figure, Fig. 7, represents our most conservative estimate of the true genus available in the WMAP distribution, and shows excellent agreement with the Gaussian random-phase hypothesis.
8 DISCUSSION

As we have shown, topologically, within the errors, the CMB measured by WMAP appears to be consistent with Gaussian random phase. Indeed, the overall visual impression is strikingly random phase. The stereographic projection is conformal, so shapes are preserved locally, though the scale grows as one approaches the outer boundary (by a factor of 2 from the pole to the equator, as seen in Fig. 1). We may ask whether there are any features visible in the map at all that look to be non-random phase in any way.

Our choice of map projection has a miscellaneous advantage over the Mollweide projection used by the WMAP team, which is that it shows the Galactic polar regions very well. In the south polar region, there is one possibly non-random feature, a fairly narrow red (hot) feature stretching from approximately the 8:30 position toward the 2:30 position curving like a bow below the South Galactic Pole. This feature is more visually apparent when viewed in landscape orientation. It has the appearance of a scanning error approximately along a great circle, but is not coincident with any particular locus in the WMAP scanning geometry.

We have considered several possibilities to explain such a feature. First, there is the simple phenomenon of the ‘canals’ on Mars, the linear features reported in naked-eye observations of the red planet in the late 19th century. We now know that there are no such canals, and that the highly evolved ability of our eyes to detect linear features or edges was simply creating an illusion. Most likely the same thing is going on here, particularly since the feature is more apparent when the figure is viewed in landscape rather than in portrait orientation. In fact, if one had a true Gaussian random-phase distribution and there were no features that looked unusual to the eye, that would be unusual. So this feature may be merely coincidental.

What are the other possibilities? It is not coincident with the Magellanic Stream measured at 21 cm (Brooks 2000). Furthermore, we would expect the stream to have similar microwave ‘colour’ to the gas in the Galaxy, and hence be subtracted about as well as the Galaxy is in the ILC map. To test this, we constructed a three-colour map, using the K band as red, Q as green and W as blue, and found that while the foregrounds were distinctly pink to orange to yellow, the CMB was quite grey, and so was the feature.

Another possibility that comes to mind is cosmic strings. Cosmic strings, however, do not amplify temperature along a line. Instead, a moving cosmic string produces a cliff, which is hotter on one side (the trailing side) than the other side (the leading side). Thus a hot stripe would require two parallel cosmic strings, having just passed each other (an interesting theoretical possibility (Gott 1991), but not a likely one, since we are unlikely to find the two most prominent cosmic strings within the horizon in such an alignment by chance). Also, there is no obvious evidence for cosmic string cliffs anywhere else in the WMAP data, which we have tested for by making gradient maps as suggested by Gott et al. (1990).

The feature could also be due to the integrated Sachs–Wolfe effect from a great wall seen edge-on, but in looking at several galaxy catalogues and X-ray maps, we could detect no obvious feature with appropriate geometry.

Finally, Chiang et al. (2003) have shown that the ‘cleaned’ map of Tegmark, de Oliveira-Costa & Hamilton (2003) exhibits correlated phases that could produce locally linear structures, though the correlations are presumably an artefact of the cleaning technique, since the Weiner filtered map of Tegmark et al. (2003) (where only smoothing is involved, changing amplitudes but not phases), appears to be approximately random phase.

This list of possibilities is by no means exhaustive, and, in the end, it may be most likely that we are simply falling prey to ‘Martian canals’ in an overall Gaussian random-phase map. Since there are countless possibilities for what one may see and deem non-random (circles, squares, triangles, even letters or words), the a posteriori statistical significance of any such a weak feature is difficult to estimate.

The overall consistency of the genus topology in the WMAP data with that predicted a priori from a Gaussian random-phase distribution is quite spectacular. The unprecedented combination of angular resolution and all-sky coverage of the cosmic microwave background of WMAP has presented by far the best confirmation to date of the standard inflationary prediction of Gaussian random-phase initial conditions.

In standard inflation, the structures seen in the CMB are fossil remains of random quantum fluctuations occurring just $10^{-33}$ s after the big bang. In galaxy clustering surveys, these fossils have become somewhat distorted due to non-linear effects and biased galaxy formation. In the CMB, however, we are seeing these fluctuations while they are still in the linear regime, making the WMAP measurements of the CMB a very attractive data set for studying these primordial perturbations.

With the hundreds of structures observed even at the 0.9 scale, the WMAP results have provided the largest and best data set for studying the primordial structures. The genus topology test, in particular, tests the data set for the standard inflationary prediction that the fluctuations should derive from a Gaussian random-phase distribution. All of our tests have confirmed this prediction dramatically.

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