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Authors

Ruiz-Macias, Omar Zarrouk, Pauline Cole, Shaun et al.

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Preliminary Target Selection for the DESI Bright Galaxy Survey (BGS)

Omar Ruiz-Macias, ^{1,2} Pauline Zarrouk, ¹ Shaun Cole, ¹ Peder Norberg, ^{1,3} Carlton Baugh, ^{1,2} David Brooks, ⁴ Arjun Dey, ⁵ Yutong Duan, ⁶ Sarah Eftekharzadeh, ⁷ Daniel J. Eisenstein, ⁸ Jaime E. Forero-Romero, ⁹ Enrique Gaztañaga, ^{10,11} ChangHoon Hahn, ^{12,13} Robert Kehoe, ¹⁴ Martin Landriau, ¹² Dustin Lang, ^{15, 16} Michael E. Levi, ¹² John Lucey, ³ Aaron M. Meisner, ⁵ John Moustakas, ¹⁷ Adam D. Myers, ¹⁸ Nathalie Palanque-Delabrouille, ¹⁹ Claire Poppett, ²⁰ Francisco Prada, ²¹ Anand Raichoor, ²² David J. Schlegel, ¹² Michael Schubnell, ²³ Gregory Tarlé, ²⁴ David H. Weinberg, ²⁵ M. J. Wilson, ^{12, 13} and Christophe Yèche ¹⁹

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<sup>1</sup>Institute for Computational Cosmology, Department of Physics, Durham University, South Road, Durham DH1 3LE, UK
                   <sup>2</sup>Institute for Data Science, Durham University, South Road, Durham DH1 3LE, UK
 <sup>3</sup>Centre for Extragalactic Astronomy, Department of Physics, Durham University, South Road, Durham DH1 3LE, UK
          ^4Department of Physics and Astromomy, University College London, Gower Street, London WC1e 6BT
    <sup>5</sup>NSF's National Optical-Infrared Astronomy Research Laboratory, 950 N. Cherry Ave., Tucson, AZ 85719, USA
                             <sup>6</sup>Physics Department, Boston University, Boston, MA 02215, MA
 Department of Physics and Astronomy, The University of Utah, 115 South 1400 East, Salt Lake City, UT 84112, USA
             <sup>8</sup> Harvard-Smithsonian Center for Astrophysics, 60 Garden Street, Cambridge, MA 02138, USA
        <sup>9</sup>Departamento de Física, Universidad de los Andes, Cra. 1 No. 18A-10, CP 111711, Bogotá, Colombia
                            <sup>10</sup>Institute of Space Sciences (ICE, CSIC), 08193 Barcelona, Spain
                      11 Institut d' Estudis Espacials de Catalunya (IEEC), 08034 Barcelona, Spain
                 <sup>12</sup>Lawrence Berkeley National Laboratory, 1 Cyclotron Road, Berkeley, CA 94720, USA
                        <sup>13</sup>Berkeley Center for Cosmological Physics, UC Berkeley, CA 94720, USA
          <sup>14</sup>Department of Physics, Southern Methodist University, 3215 Daniel Avenue, Dallas, TX 75275, USA
        <sup>15</sup>Perimeter Institute for Theoretical Physics, 31 Caroline Street N, Waterloo, Ontario, N2L 2Y5, Canada
            <sup>16</sup>Department of Physics and Astronomy, University of Waterloo, Waterloo, ON N2L 3G1, Canada
            <sup>17</sup>Department of Physics and Astronomy, Siena College, 515 Loudon Road, Loudonville, NY 12211
                       <sup>18</sup>University of Wyoming, 1000 E. University Ave., Laramie, WY 82071, USA
                          <sup>19</sup>IRFU, CEA, Université Paris-Saclay, F-91191 Gif-sur-Yvette, France
               <sup>20</sup>Space Sciences Laboratory at University of California, 7 Gauss Way, Berkeley, CA 94720
           <sup>21</sup>Instituto de Astrofísica de Andalucía (CSIC), Glorieta de la Astronomía, E-18080 Granada, Spain
              <sup>23</sup>Department of Physics, University of Michigan, 450 Church St., Ann Arbor, MI 48109, USA
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²²Institute of Physics, Laboratory of Astrophysics, Ecole Polytechnique Fédérale de Lausanne (EPFL), Observatoire de Sauverny, 1290 Versoix, Switzerland ²⁴Department of Physics, University of Michigan, Ann Arbor, MI 48109, USA

²⁵Department of Astronomy and the Center for Cosmology and Astroparticle Physics, The Ohio State University, 140 West 18th Avenue, Columbus OH 43210,

ABSTRACT

The Dark Energy Spectroscopic Instrument (DESI) will execute a nearly magnitude-limited survey of low redshift galaxies (0.05 $\leq z \leq$ 0.4, median $z \approx$ 0.2). Clustering analyses of this Bright Galaxy Survey (BGS) will yield the most precise measurements to date of baryon acoustic oscillations and redshift-space distortions at low redshift. DESI BGS will comprise two target classes: (i) BRIGHT (r < 19.5 mag), and (ii) FAINT (19.5 < r < 20 mag). Here we present a summary of the star-galaxy separation, and different photometric and geometrical masks, used in BGS to reduce the number of spurious targets. The selection results in a total density of ~ 800 objects/deg² for the BRIGHT and ~ 600 objects/deg² for the FAINT selections. A full characterization of the BGS selection can be found in Ruiz-Macias et al. (2020).

Keywords: BGS, surveys, large-scale structure, star-galaxy separation

INTRODUCTION

The DESI BGS (DESI Collaboration et al. 2016) will be a flux-limited r-band selected sample of 10 million galaxies to z = 0.4. The DESI target selection will be done on the Legacy Surveys (LS) imaging (Dey et al. 2019). Results presented here are based on the DR8 release¹. The DESI BGS is expected to have a target density of about 800 galaxies/deg² in a primary selection with a magnitude limit in the r-band of 19.5 and 600 galaxies/deg² in an additional sample defined by the magnitude range 19.5 < r < 20(Smith et al. 2017). Henceforth, we will refer to these BGS samples as BRIGHT and FAINT respectively.

BGS TARGET SELECTION

The BGS targets are selected using the three optical fluxes in the Legacy Surveys (Ruiz-Macias et al. 2020):

$$r < 19.5 \text{ for BRIGHT}$$
, $19.5 < r < 20 \text{ for FAINT}$ (1)

$$(-1 < g - r < 4) \tag{2}$$

$$(-1 < r - z < 4) \tag{3}$$

where g, r, and z indicate the extinction-corrected AB magnitudes in the corresponding band (using LS extinction corrections). In addition, we require all targets to be covered by at least one image in each optical band. We remove sources near bright stars, large galaxies, or globular clusters by requiring that LS MASKBITS 1, 12 and 13 are not set as defined in the LS data model. The bright star mask (defined by bit=1) combines stars from Gaia DR2 (Gaia Collaboration et al. 2018) and the Tycho2 (Høg et al. 2000) catalog, corrected for epoch and proper motions. This mask consists of a circular exclusion region with a radius that depends on the magnitude of the star, m. The magnitude is either the Tycho2 MAG_VT or Gaia G-mag with Gaia G-mag taking precedence. Stars fainter than m = 13 are not masked. The large galaxies mask (bit=12) was defined by the Siena Galaxy Atlas - 2020 (Moustakas, J. in prep), an angular diameter-limited sample of galaxies with mostly HyperLeda objects(Makarov et al. 2014). To mask around large galaxies and elliptical mask is used defined by the diameter at the 25 mag/arcsec² (optical) surface brightness, the ratio of the semi-minor axis to the to semi-major axis and the position angle. The globular cluster (GC) mask (bit=13) consists of a circular exclusion zone around known GCs from the OpenNGC catalog².

BGS star-galaxy separation is based on Gaia DR2. Gaia is highly complete for BGS and has a better point spread function (PSF) than ground-based surveys. A galaxy in BGS is defined by (G - rr > 0.6) or (G = 0) where G is Gaia G-mag and rr is LS r-band magnitude (without any extinction correction). We apply Equation 2 and 3 to remove sources that are beyond the color range of BGS galaxies. To remove spurious large and low-surface-brightness galaxies, we implement a cut that compares the r-band magnitude (r) to the predicted r-band fiber magnitude (rfibmag), as in Equation 4. Finally, we impose a minimum quality for the data reduced by The Tractor³ (Lang et al. 2016), in FRACMASKED_i < 0.4, FRACIN_i > 0.3, FRACFLUX_i < 5, where $i \equiv g, r$ or z. FRACIN is used to select sources for which a large fraction of the model flux lies within the contiguous pixels to which the model was fitted, FRACFLUX is used to reject objects that are swamped by flux from adjacent sources, and FRACMASKED is used to veto objects with a high fraction of masked pixels.

CONCLUSION

We have presented the BGS target selection from LS DR8 divided into two samples, the BRIGHT and FAINT samples. The BRIGHT sample, which will have a higher fiber-allocation priority, comprises ~800 targets/deg². The FAINT sample, which will be assigned at lower priority, comprises ~600 objects/deg². Both samples undergo the spatial and photometric cuts outlined by the MASKBITS and the number of observations required, and by Equations 2-4 respectively, passed the Gaia based star-galaxy classification and our quality cuts. Fig. 1 shows the BGS target density as a function of r-band magnitude and the N(z) of BGS targets cross-matched to GAMA DR4 (Driver et al. 2012; Liske et al. 2015; Baldry et al. 2017). The preliminary selection described in this note is public⁴.

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¹ http://legacysurvey.org/dr8/

² https://github.com/mattiaverga/OpenNGC

³ https://github.com/dstndstn/tractor

⁴ Available at https://data.desi.lbl.gov/public/ets/target/catalogs/ and detailed at https://desidatamodel.readthedocs.io

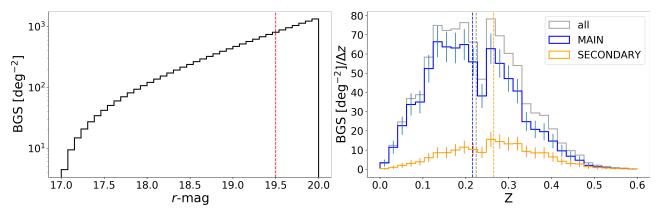


Figure 1. Left: Target density of BGS as a function of r-band magnitude in DECaLS DR8. The red dashed line at 19.5 marks the boundary between the BRIGHT and FAINT selections. **Right:** Redshift distribution of DECaLS DR8 BGS targets cross-matched with GAMA DR4 in redshift bins of $\Delta z = 0.02$. The solid blue, orange and gray lines show the BRIGHT, FAINT and combined samples respectively. The dashed lines show the means of the samples at 0.21 (BRIGHT), 0.26 (FAINT) and 0.22 (combined). Error bars for BRIGHT and FAINT distributions are Poisson. Because GAMA has a magnitude limit of approximately r = 19.8 in (SDSS DR7) r-petrosian (Abazajian et al. 2009; Driver et al. 2012), only 30 per cent of the FAINT sample appears in GAMA and the mean redshift is expected to be somewhat higher. While for the BRIGHT sample, the GAMA completeness is as high as 95 per cent.

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