

Supplementary Table 1. List of identification keys used for each of the genera sampled in the San Bernardino Valley

Andrenidae

Anylandrena (Zavortink 1974)

Andrena – many identifications by John S. Ascher, morphospecies by J.S. Ascher and R.L. Minckley (Bouseman & LaBerge 1979; LaBerge 1969, 1971, 1973, 1977, 1985, 1986, 1987; LaBerge & Bouseman 1970; LaBerge & Ribble 1975; Ribble 1968; Thorp 1969)

Protoxaea (Hurd Jr. & Linsley 1976)

Calliopsis (Danforth 1994; Rozen Jr. 1958; Shinn 1967)

Macroterea (Danforth 1996; Snelling & Danforth 1992; Timberlake 1954)

Perdita – many identifications by Douglas A. Yanega (Portman et al. 2016; Timberlake 1954, 1956, 1958, 1960, 1962, 1964a, 1968, 1971, 1980a)

Protandrena - (Timberlake 1964b, 1967, 1975, 1976)

Pseudopanurgus (Timberlake 1973)

Apidae

Ancyloscelis (Michener 1942)

Anthophora- identifications of *Anthophora (Micranthophora)* by Michael C. Orr and some other *Anthophora* by Robert Brooks (Brooks 1988; Orr et al. 2018)

Centris(Snelling 1974, 1984; Vivallo 2020)

Diadasia - Sedonia Sipes, unpublished key

Eucera (Timberlake 1969)

Martinapis (Zavortink & LaBerge 1976)

Melissodes - many identifications by Karen Wright (LaBerge 1956a,b, 1961)

Svastra (LaBerge 1958)

Peponapis and *Xenoglossa* (Hurd Jr. & Linsley 1966, 1967a,b)

Tetraloniella (Laberge 2001)

Anthophorula and *Exomalopsis* (Silveira 1995; Timberlake 1980b)

Zacosmia (Hurd Jr. & Linsley 1951)

Oreopasites (Linsley 1941; Rozen Jr. 1992)

Holcopasites and *Neopasites* (Hurd Jr. & Linsley 1972; Linsley 1943a)

Brachynomada (Rozen Jr. 1994; Snelling & Rozen Jr. 1987)

Paranomada (Linsley 1943b, 1945)

Triopasites (Rozen Jr. 1997)

Epeolus (Onufko 2018)

Triepeolus - material identified by M. Rightmyer (Rightmyer 2008)

Melecta (Hurd Jr. & Linsley 1951; Linsley 1939)

Neolarra (Michener 1939a; Shanks 1978)

Nomada- identified to morphospecies by R.L. Minckley

Ceratina (Daly 1973)

Xylocopa (Hurd Jr. 1961)

Colletidae

Colletes (Stephen 1954)

Caupolicana (Vergara & Michener 2004)

Hylaeus (Snelling 1966a,b)

Halictidae

Augochlorella (Ordway 1966)

Agapostemon (Roberts 1972)

Halictus (Sandhouse 1924)

Lasioglossum- many identifications by Jason Gibbs and Joel Gardner (Gardner & Gibbs 2020;
McGinley 1986; Sandhouse 1924)

Sphecodes- morphospecies separated by R.L. Minckley

Dieunomia and *Nomia* (Blair 1935; Cross 1958; Ribble 1965)

Conanthalictus and *Sphecodosoma* (Timberlake 1961)

Dufourea – sorted to morphospecies by R.L. Minckley

Protodufourea (Bohart & Griswold 1997)

Megachilidae

Anthidium- identified by Victor Gonzalez-Betancourt (Gonzalez & Griswold 2013; Schwarz 1926)

Dianthidium (Grigarick & Stange 1968)

Stelis- most sorted to morphospecies by R.L. Minckley (Parker & Bohart 1979)

Dioxys (Hurd Jr. 1958)

Lithurgopsis (Mitchell 1937a; Snelling 1990)

Coelioxys (Mitchell 1973)

Megachile (Mitchell 1935, 1936, 1937b)

Ashmeadiella (Hurd Jr. & Michener 1955; Michener 1939b)

Atoposmia (Hurd Jr. & Michener 1955; Michener 1943)

Heriades (Hurd Jr. & Michener 1955; Michener 1938, 1954)

Hoplitis (Hurd Jr. & Michener 1955; Michener 1947)

Osmia- identified by Molly G. Rightmyer (Griswold & Rightmyer 2017; Rightmyer et al. 2010)

Melittidae

Hesperapis (Stage 1966)

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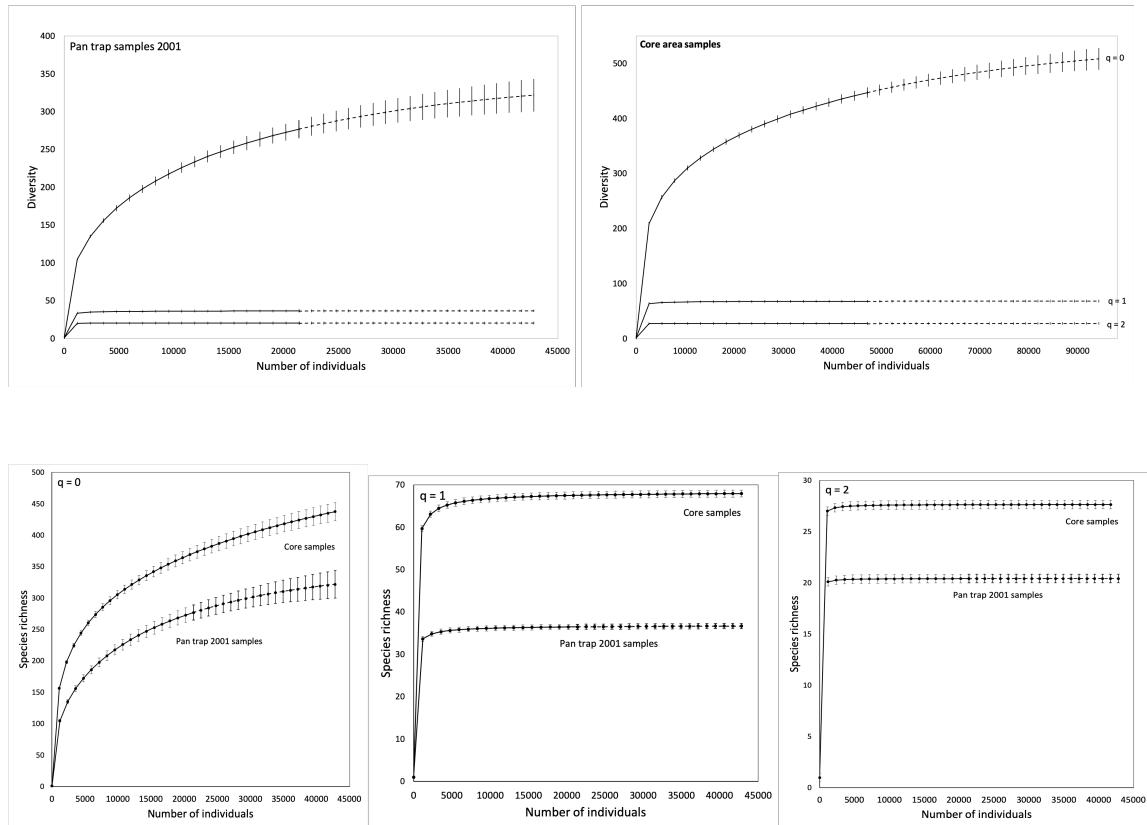
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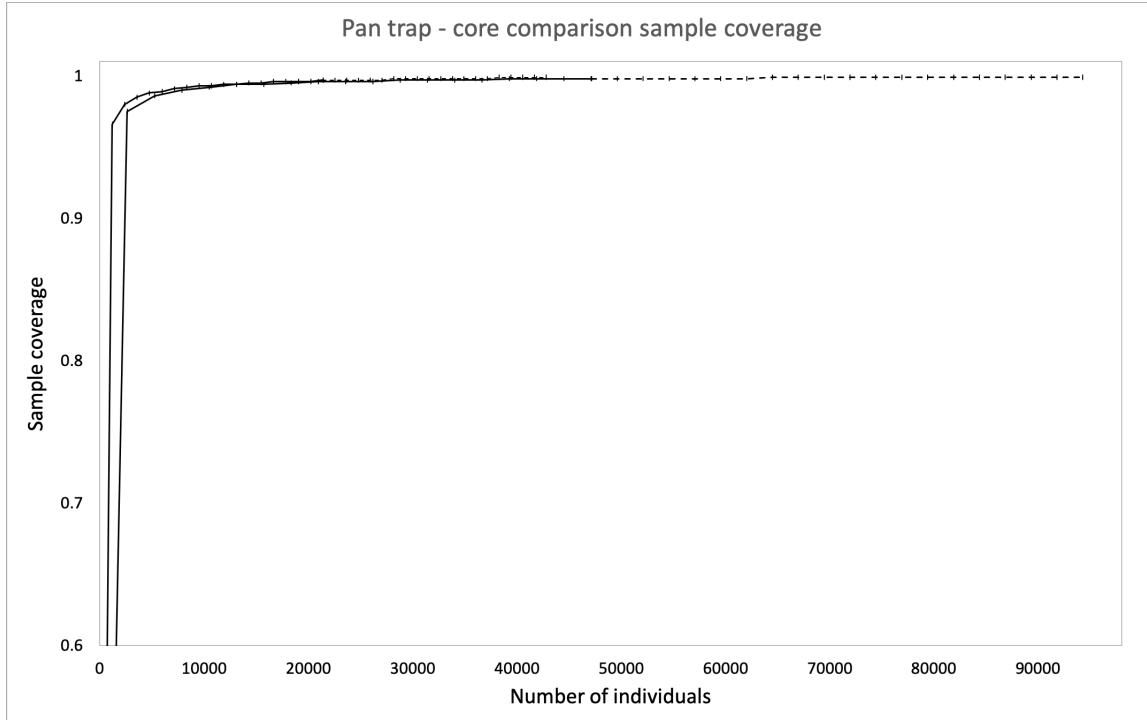
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Supplementary Table 2. Table 4 from Meiners et al (2019) of bee species density for native bee inventories with at least 100 species in natural or semi-natural areas across the United States. Information here is updated for the San Bernardino Valley (this study).

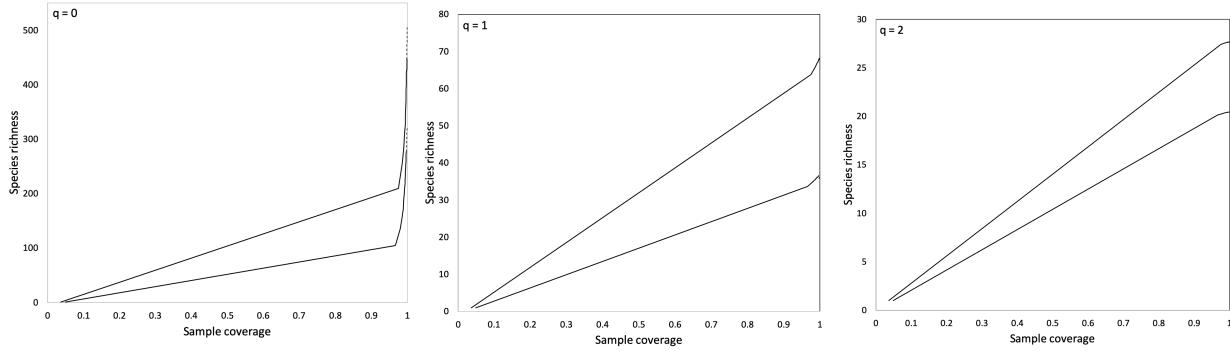
Study location	Study dates	Species	Approx. total area (km ²)	References
Grand Staircase Escalante National Monument, Utah	2000-2003	656	7,610	(Carril et al. 2018)
Clark County, Nevada	1998; 2005, 2006	598	20,487	(Griswold et al. 1999)
Yosemite National Park, California	2006-2009	554	3,028	pers. comm. T. Griswold in Meiners et al (2019)
Pinnacles National Park, California	1996-1999, 2002, 2011-2012	450	109	(Meiners et al. 2019)
San Bernardino Valley, Arizona/ Sonora	2000-2008	473	16	This study
Carlsbad Caverns National Park, New Mexico	2010-2011	364	189	pers. comm. T. Griswold in Meiners et al (2019)
Curlew Valley, Idaho	1969-1974	340	4,999	(Bohart & Knowlton 1973)
San Rafael Desert, Utah	1979-1992	333	5,180	(Griswold et al. 1998)
Mojave National Preserve California	1975-1995	305	6,475	pers. comm. T. Griswold in Meiners et al (2019)
Black Hills of South Dakota and Wyoming	2010-2011	290	12,950	(Drons 2012)
Carlinville, Illinois	1884-1916	288	256	(Minckley 2008)
MPG Ranch, Montana	2013-2015	229	39	(Kuhlman & Burrows 2017)
Indiana Dunes, Indiana	2003, 2004, 2010	204	60	(Grundel et al. 2011)
Albany County, Wyoming	1995-1996	200	11,160	(Tepedino & Stanton 1981)
Palouse Prairie, Idaho	2012-2013	174	2,122	(Rhoades et al. 2017)
Dugway Proving Ground, Utah	2003, 2005	163	3,243	(Wilson et al. 2009)
Channel Islands, California	Not specified	154	904	(Rust et al. 1985)
Black Rock Forest Reserve, New York	2003	144	15.5	(Giles & Ascher 2006)
Tonasket Ranger District, Washington	2004	140	1,678	(Wilson et al. 2010)
Black Belt Prairie, Mississippi	1991-2001	118	803	(Smith et al. 2012)
Archibald Biological Station, Florida		113	21	(Deyrup et al. 2002)
Hattiesburg, Mississippi	1943-1944	104	140	(Michener 1947)



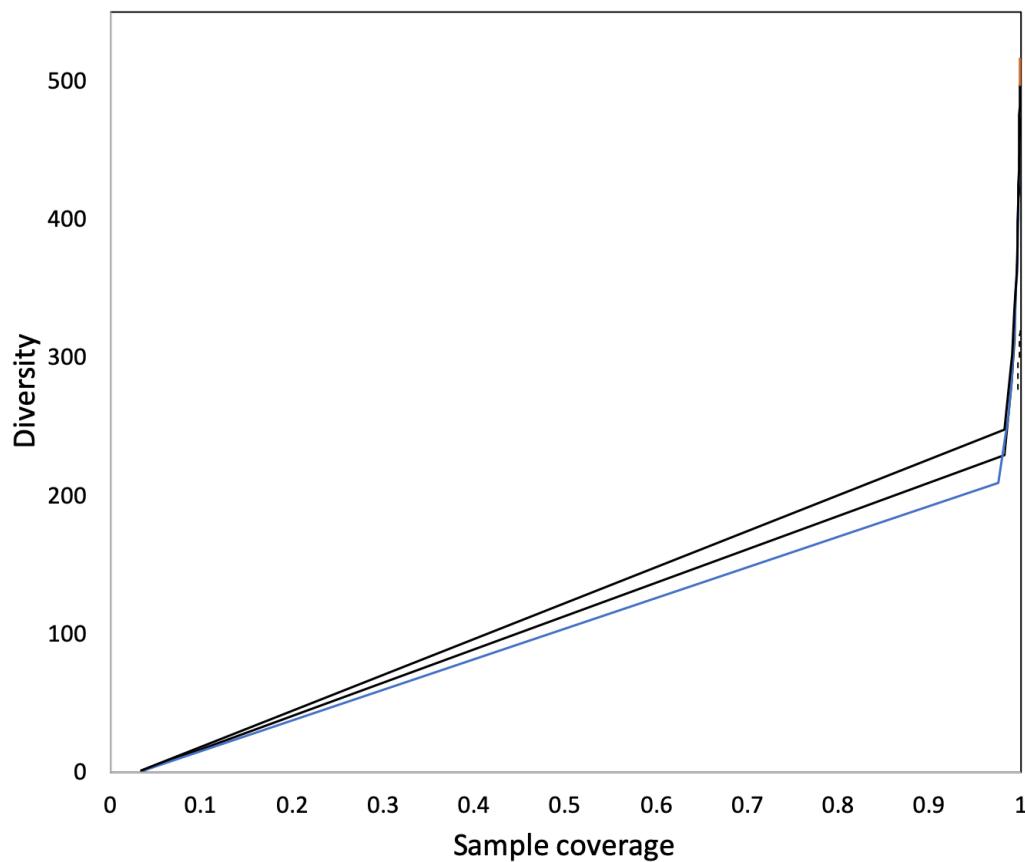
Supplemental Figure 1. Comparison of sample-sized-based rarefaction (solid lines) and extrapolation (dashed line) from the pan trap samples at permanent sites in 2001 and the samples made by netting and pan traps throughout the same area encompassed by the permanent sites over 9 years (the “core” area) following the approach of Chao et al (34). Bee species richness at the base sample size of 42,798 individuals (i.e., double the smaller reference size) is 406.0 for samples from the core area and 321.8 from the 2001 pan traps. The two top graphs are sample-size-based rarefaction and extrapolation curves for the pan trap samples from 2001 and the samples made in the core area from 2000-2008. The bottom three graphs are comparisons of samples from the pan trap samples from 2001 and the core area for each order of q . Although bees were sampled from the same area, differences in sampling and years sampled gave different species richness estimates for each order of q . The first graph in lower row is the same as Fig. 1 in the paper and presented here for completeness.



Supplemental Figure 2. Plot of sample coverage for rarified samples (solid line) and extrapolated samples (dashed line) as a function of sample size for the pan trap samples at permanent sites in 2001 and the samples made by netting and pan traps throughout the same area encompassed by the permanent sites over 9 years (the “core” area). 95% confidence intervals calculated from 50 bootstrap replications are not prominent because of their small size. Both samples have greater than 95% coverage and broadly overlap up to their reference sizes (21,403 individuals for 2001 pan trap samples, 47,120 for core area samples) and when extrapolated to twice the reference sample size.



Supplemental Figure 3. Comparison of the coverage-based rarefaction (solid line) and extrapolation (dashed line), for the pan trap samples at permanent sites in 2001 and the samples made by netting and pan traps throughout the same area encompassed by the permanent sites over 9 years (the “core” area) for all three orders of q . In each of the three graphs the lower line is the 2001 pan trap samples and the upper line is the samples from the core area. Note that the lines do not intersect at any sample coverage.



Supplemental Figure 4. Comparison of the coverage-based rarefaction (solid line) and extrapolation (dashed line), for the samples made in the core area (middle line), core area combined with the pan trap sampling at permanent sites (blue line on bottom), and all samples made in the San Bernardino Valley (top line) at $q = 0$. The confidence intervals are very small they do not show on the graph. Sample coverage is above 95% for diversity values below 200 species for the datasets and the extrapolations overlap broadly at higher sampling coverage indicating all three datasets predict very similar bee species richness in the San Bernardino Valley. Note that the analyses at other orders of q are not shown because the results match that shown here.