

## Use of Plant Introductions in Peanut Cultivar Development

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### ABSTRACT

The genetic base of peanut (*Arachis hypogaea* L.) in the U.S.A. has at times been extremely narrow, particularly in specific production areas where a single cultivar may be grown in near-monoculture. Because peanut is not a native North American species, all U.S. cultivars necessarily trace their ancestry to plant introductions (PIs), but most of the genetic base of current cultivars rests on selections from farmer-stock peanuts of obscure origin. The objectives of this study were to (a) summarize and document the use of introduced genetic resources in cultivar development and (b) estimate the resulting economic impact. Different PIs were used as parents in early breeding programs. B.B. Higgins used Gambian line Basse as a parent of the GA 207 cross that gave rise to selections used in Georgia, Florida, and North Carolina as the basis for further improvement. PI 121067 was one of seven parents used by W.C. Gregory to initiate the program in North Carolina. A different set of PIs including PI 121070, PI 161317, PI 168661, and *A. monticola* Krapov. & Rigoni were used in the Texas and Oklahoma programs. Recycling of lines as parents and exchange of germplasm among breeding programs proliferated these PIs in the pedigrees of cultivars released since 1960. Over the past 20 yr, there have been concerted efforts to incorporate additional germplasm into U.S. breeding populations, usually with the purpose of improving resistance to diseases or pests, but also with the objective of broadening the genetic base. These efforts have had a significant economic impact on U.S. peanut farmers, the largest from the development of cultivars with resistance to Sclerotinia blight (*Sclerotinia minor* Jagger), root-knot nematodes (*Meloidogyne* spp.), and tomato spotted wilt virus. Use of these resistant cultivars has an economic impact of more than \$200 million annually for U.S. peanut producers. In the runner and virginia market types, the average PI ancestry of all cultivars was 17.9%. There are several examples of successful cultivars with up to 25% ancestry from a single PI, including Georgia Green and NC-V11. In the spanish market type, most successful cultivars have derived 50% or more of their ancestry from PIs. Several recent or impending releases

incorporate PI germplasm but have not yet been proven in the U.S. seed market.

Key Words: *Arachis hypogaea* L., coancestry, disease resistance, genetic vulnerability, genetic resources.

As of July 2000, 119 peanut (*Arachis hypogaea* L.) cultivars had been released in the U.S. (Table 1), 53 released prior to 1961 when the Crop Science Society of America (CSSA) began to register crop cultivars and germplasm (Isleib and Wynne, 1992) and 66 registered with the CSSA after 1960 (registered cultivars, germplasm lines, parental lines, and genetic stocks can be obtained from the National Plant Germplasm System web site at [www.ars-grin.gov/npgs](http://www.ars-grin.gov/npgs)) or with the Plant Variety Protection Office. Forty-eight were protected under Plant Variety Protection certificates, 24 of them current, eight expired, and 16 pending. One hundred twenty-eight additional lines have been released as germplasm, parental lines, or genetic stocks (Table 2).

In spite of the large number of cultivars available to growers, the U.S. peanut crop has been characterized as being genetically vulnerable to diseases and insect pests (Hammons, 1972, 1976; Knauff and Gorbet, 1989). This has been due to the commercial success of specific cultivars grown in particular production areas. For example, the runner-type cultivar Florunner dominated production in the southeastern U.S. (Georgia, Florida, and Alabama) from 1972 to 1993, occupying over 60% of the acreage in those years, over 80% from 1974 through 1987, and over 95% from 1976 through 1984. Currently, southeastern production is dominated by the cultivar Georgia Green. The Virginia-Carolina (VC) production area has had less tendency toward monoculture. The most dominant cultivar of the past 40 yr in the VC area was Florigiant which averaged over 60% of the area's acreage from 1974 through 1985, and with a maximum of over 80% in 1979. Since the decline of Florigiant, no cultivar in the VC area has occupied more than 50% of the acreage in a given year. In the southwestern (SW) production area (Texas and Oklahoma) where spanish-type cultivars were the main type produced through the 1970s, there was near-monoculture of Starr in the late 1960s and early 1970s. Over the past 20 yr, there has

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Table 1. Cultivars released or registered in the United States.

Name	Market type	CSSA no.	NPGS no.	PVP cert. no.	Yr of release	Releasing agency	Reference
AD-1	virginia			8600030	1986	J. Ashley Darden	
AgraTech 1-1	runner			20000134*	2000	AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
AgraTech AT 108	runner			9600322	1997	AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
AgraTech AT 120	runner			9700275	1997	AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
AgraTech AT 127	runner			9200066	1996	AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
AgraTech AT 201	runner			200000135*	2000	AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
AgraTech AT 225	runner			9700010	1997	AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
High Oleic							
AgraTech GK 3	virginia			7300094**	1976	Gold Kist Peanut Co.	E. Harvey <sup>a</sup> , 1990/D.A. Knauff <sup>a</sup> , 1990
AgraTech GK 7	runner			8200141*	1982	Gold Kist Peanut Co.	E. Harvey <sup>a</sup> , 1990
AgraTech GK 7	runner			9800019*	1997	AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
High Oleic							
AgraTech GK 19	runner			7300005**	1973	Gold Kist Peanut Co.	E. Harvey <sup>a</sup> , 1990
AgraTech VC-1	virginia			9200014		AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
AgraTech VC-2	virginia			200000136*	2000	AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
AgraTech ViruCard <sup>®</sup>	runner			9800022*	1997	AgraTech Seeds Inc.	K. Moore <sup>a</sup> , 2000
ALR 2	spanish	CV-61	PI 599975		1994	ICRISAT; Tamil Nadu Agric. Univ.	Varman <i>et al.</i> , 1998
Altika	runner	CV-18	PI 565453		1972	Guyana Min. Agric., FLAES	Norden & Gorbet, 1974
Andru 93	runner	CV-53	PI 566905	9300153	1993	FLAES	Gorbet & Knauff, 1995
Argentine	spanish		PI 300585		1951	OKAES	McGill, 1963
Avoca 11	virginia			7100110**	1976	R.J. Reynolds Co.	Mozingo <i>et al.</i> , 1987
C-99R	runner			200000182*	1999	FLAES	Gorbet <i>et al.</i> , 1999
COAN	runner			9900338*	1999	TXAES	C.E. Simpson <sup>a</sup> , 2000
Comet	spanish				1970	OK & GA AES	J.S. Kirby <sup>a</sup> , 1990
David	virginia			9500120	1996	Israel Agric. Res. Org.	I. Wallerstein <sup>a</sup> , 1999
Dixie Runner	runner				1943	FLAES	Carver & Hull, 1950
Dixie Spanish	spanish				1950	GAAES	Higgins & Bailey, 1955
Early Bunch	virginia	CV-21	PI 565458		1977	FL & GAAES, USDA-ARS	Norden <i>et al.</i> , 1978
Early Runner	runner				1952	FLAES	Carver <i>et al.</i> , 1952
Flavorrunner 458	runner			9600242	1997	Mycogen	
Florida MDR 98	runner			9900212*	1998	FLAES	D.W. Gorbet <sup>a</sup> , 2000
Florigiant	virginia	CV-01	PI 565445		1961	FLAES	Carver, 1969
Florispant Runner	spanish				1953	GAAES	Carver, 1953
Florunner	runner	CV-02	PI 565448		1969	FLAES	Norden <i>et al.</i> , 1969
Georgia 119-20	virginia	CV-12	PI 565440		1954	GAAES & USDA-ARS	Hammons, 1971
Georgia Bold	runner	CV-60	PI 601980	9800041*	1997	GAAES	Branch, 1998
Georgia Browne	runner	CV-52	PI 574450	9400043	1993	GAAES	Branch, 1994
Georgia Green	runner	CV-55	PI 587093	9500165	1995	GAAES	Branch, 1996
Georgia Red	valencia	CV-33	PI 508278		1986	GAAES & USDA-ARS	Branch & Hammons, 1987
Georgia Runner	runner	CV-41	PI 542960	9200029	1990	GAAES	Branch, 1991
GFA Spanish	spanish				1941	GAAES	Higgins & Bailey, 1955
Goldin I	spanish			7100035**	1976	Wilson Co. Peanut Co.	R.O. Hammons <sup>a</sup> , n.d.
Gregory	virginia	CV-62	PI 608666	9900337*	1997	NCARS	Isleib <i>et al.</i> , 1999
H&W Valencia 101	valencia			9700337*		H&W GENETEX, Inc.	
H&W Valencia 102	valencia			9700336*		H&W GENETEX, Inc.	
Holland Jumbo	virginia				1945	VAAES	Batten, 1945
Holland Va. Runner	virginia				1943	VAAES	Batten, 1945
Hughes Runner	runner			200000203*	2000	Kenneth E. Hughes	
ICGS 1	spanish	CV-43	PI 478780		1990	ICRISAT	Nigam <i>et al.</i> , 1991b
ICGS 11	spanish	CV-38	PI 478788		1986	ICRISAT	Nigam <i>et al.</i> , 1990a
ICGV 86325	spanish <sup>b</sup>	CV-54	PI 590879		1994	ICRISAT	Dwivedi <i>et al.</i> , 1996a
ICGV 86590	valencia	CV-49	PI 562530		1991	ICRISAT	Reddy <i>et al.</i> , 1993
ICGV 87128	spanish	CV-37	PI 537112		1988	ICRISAT	Nigam <i>et al.</i> , 1990b
ICGV 87141	spanish <sup>b</sup>	CV-42	PI 546372		1989	ICRISAT	Nigam <i>et al.</i> , 1990c
ICGV 87160	spanish	CV-47	PI 478787		1990	ICRISAT	Reddy <i>et al.</i> , 1992
ICGV 87187	spanish	CV-45	PI 550930		1990	ICRISAT	Nigam <i>et al.</i> , 1992b
Improved Spanish 2B	spanish					GAAES	Beattie <i>et al.</i> , 1954
Jeokwangtangkong	spanish	CV-63	PI 607913		1997	ICRISAT	Oh <i>et al.</i> , 2000
Keel 29	virginia			8000155*	1974	Keel Peanut Seed Co.	Mozingo <i>et al.</i> , 1987
Keel 76	virginia					James T. Keel	
KH20	runner			8700094		Kenneth E. Hughes	
Langley	runner	CV-31	PI 506237		1986	TXAES	Simpson <i>et al.</i> , 1987
MARC 1	runner	CV-46	PI 552555	9200115	1990	FLAES	Gorbet <i>et al.</i> , 1992
McRan	valencia			7300066**	1973	Borden Peanut Seed Co.	R.O. Hammons <sup>a</sup> , n.d.
NC 1	virginia				1952	NCAES	W.C. Gregory <sup>a</sup> , n.d.

Table 1 (Continued)

Name	Market type	CSSA no.	NPGS no.	PVP cert. no.	Yr of release	Releasing agency	Reference
NC 2	virginia	CV-05	PI 565446		1952	NCAES	Gregory, 1970
NC 4	virginia				1944	NCAES	W.C. Gregory <sup>a</sup> , n.d.
NC 4X	virginia				1959	NCAES	Gregory, 1960
NC 5	virginia	CV-06	PI 565447		1964	NCAES	Emery & Gregory, 1970
NC 6	virginia	CV-20	PI 565456	7605011 <sup>**</sup>	1976	NCARS	Campbell <i>et al.</i> , 1977
NC 7	virginia	CV-22	PI 565459	7900104 <sup>**</sup>	1978	NCARS	Wynne <i>et al.</i> , 1979
NC 8C	virginia	CV-27	PI 565476		1982	NCARS	Wynne & Beute, 1983
NC 9	virginia	CV-30	PI 565484	8500201	1985	NCARS	Wynne <i>et al.</i> , 1986
NC 10C	virginia	CV-39	PI 540460	8900116	1988	NCARS	Wynne <i>et al.</i> , 1991a
NC-V11	virginia	CV-40	PI 540461		1989	NCARS	Wynne <i>et al.</i> , 1991b
NC 12C	virginia	CV-57	PI 596406	9700074	1996	NCARS	Isleib <i>et al.</i> , 1997
NC-Fla 14	virginia	CV-17	PI 565466		1973	NCAES & FLAES	Emery <i>et al.</i> , 1974
NC 17	virginia	CV-07	PI 565451		1969	NCARS	Emery, 1970
New Mexico Valencia A	valencia	CV-14	PI 565452		1971	NMAES	Hsi & Finkner, 1972
New Mexico Valencia C	valencia	CV-24	PI 565461		1979	NMAES	Hsi, 1980
Okrun	runner	CV-36	PI 531499		1986	USDA-ARS & OKAES	Banks <i>et al.</i> , 1989
Perry	virginia			20000022 <sup>5</sup>	2000	NCARS	T.S. Isleib <sup>a</sup> , 2000
Pronto	spanish	CV-28	PI 565475		1980	USDA-ARS, OK & GAAES	Banks & Kirby, 1983
Shosh	virginia			9400123		Israel Agric. Res. Org.	I. Wallerstein <sup>a</sup> , 1997
Shulamith	virginia		PI 372572		1968	Israel Agric. Res. Org. & GAAES	Goldin, 1970
Sinkarzei (ICGS 114)	spanish	CV-48	PI 561673		1989	ICRISAT	Nigam <i>et al.</i> , 1993b
Southeastern Runner 56-15	runner	CV-09	PI 565439		1947	GAAES & USDA-ARS	Hammons, 1970a
Southern Runner	runner	CV-32	PI 506419	8700093	1984	FLAES	Gorbet <i>et al.</i> , 1987
Southwest Runner	runner	CV-58	PI 599178		1995	OKAES & USDA-ARS	Kirby <i>et al.</i> , 1998
Spanco	spanish	CV-35	PI 531500		1981	OKAES & USDA-ARS	Kirby <i>et al.</i> , 1989
Spancross	spanish	CV-03	PI 565449		1970	GAAES, USDA-ARS, & OKAES	Hammons, 1970b
Spanette	spanish				1959	GAAES	NAS, 1972
Spanhoma	spanish				1969	OKAES & GAAES	NAS, 1972
Spanish 18-38	spanish					GAAES	Gregory <i>et al.</i> , 1951
Spanish No. 146	spanish						Higgins & Bailey, 1955
Spantex	spanish	CV-15	PI 565442		1948	TXAES	Simpson, 1972a
Starr	spanish	CV-16	PI 565443		1961	TXAES	Simpson, 1972b
Sunbelt Runner	runner	CV-26	PI 565473		1982	USDA-ARS & GAAES	Mixon, 1982
SunOleic 95R	runner	CV-56	PI 578304	9400148	1995	FLAES	Gorbet & Knauff, 1997
SunOleic 97R	runner	CV-65	PI 596800	9700182	1997	FLAES	Gorbet & Knauff, 2000
Sunrunner	runner	CV-29	PI 565433		1982	FLAES	Norden <i>et al.</i> , 1985
Tamnut 74	spanish	CV-19	PI 564855		1974	TXAES, OKAES, & GAAES	Simpson & Smith, 1975
Tamrun 88	runner	CV-34	PI 520600		1988	TXAES	Smith & Simpson, 1989
Tamrun 96	runner	CV-59	PI 601819	9800338 <sup>*</sup>	1996	TXAES	Smith <i>et al.</i> , 1998
Tamrun 98	runner	CV-64	PI 608737	9900189 <sup>*</sup>	1998	TXAES	Simpson <i>et al.</i> , 2000
Tamspan 90	spanish	CV-44	PI 550721		1990	TXAES & USDA-ARS	Smith <i>et al.</i> , 1991
Tennessee Red	valencia						Handy, 1896
Tifrun	runner				1977	GAAES & USDA-ARS	R.O. Hammons <sup>a</sup> , 1977; D.A. Knauff <sup>a</sup> , 1990
Tifspan	spanish	CV-04	PI 565450		1964	GAAES, USDA-ARS, & OKAES	Hammons, 1970c
Toalson	spanish	CV-23	NSL 102066		1979	TXAES	Simpson <i>et al.</i> , 1979
VA-C 92R	virginia	CV-50	PI 561566	9200252	1992	VAAES, NCARS, & USDA-ARS	Mozingo <i>et al.</i> , 1994
VA 93B	virginia	CV-51	PI 561568		1993	VAAES & USDA-ARS	Coffelt <i>et al.</i> , 1994b
VA 98R	virginia	CV-66	PI 607566	9900419 <sup>*</sup>	1998	VAAES & USDA-ARS	Mozingo <i>et al.</i> , 2000
Virginia 56R	virginia	CV-10	PI 565441		1956	VAAES	Alexander & Allison, 1970a
Virginia 61R	virginia	CV-11	PI 565444		1961	VAAES	Alexander & Allison, 1970b
Virginia 72R	virginia	CV-13	PI 565454		1972	VAAES	Alexander & Allison, 1972
Virginia 81 Bunch	virginia	CV-25	PI 565474		1981	VAAES & USDA-ARS	Coffelt <i>et al.</i> , 1982
Virginia Bunch 46-2	virginia				1952	VAAES & USDA	Beattie & Batten, 1953
Virginia Bunch 67	runner	CV-08	PI 565438		1945	GAAES	Hammons, 1970d
Virginia Bunch G2	virginia				1952	GAAES	Higgins & Bailey, 1955
Virginia Bunch G26	virginia				1952	GAAES	Higgins & Bailey, 1955

<sup>\*\*</sup>Plant variety protection certificate pending or expired, respectively.

<sup>a</sup>Denotes personal communication.

<sup>b</sup>Denotes lines with spanish-type seeds but botanically classified as var. *hypogaea*.

Table 2. Germplasm lines, parental lines, and genetic stocks registered with the Crop Science Society of America (CSSA).

Line	CSSA reg. no.	NPGS no.	Year of release	Releasing agency	Reference
AR-1	GP-35	PI 565480	1983	USDA-ARS & GAAES	Mixon, 1983a
AR-2	GP-36	PI 565481	1983	USDA-ARS & GAAES	Mixon, 1983a
AR-3	GP-37	PI 565482	1983	USDA-ARS & GAAES	Mixon, 1983a
AR-4	GP-38	PI 565483	1983	USDA-ARS & GAAES	Mixon, 1983a
CBR-R1	GP-12	PI 565470	1981	USDA-ARS	Hammons, 1981
CBR-R2	GP-13	PI 565469	1981	USDA-ARS	Hammons, 1981
CBR-R3	GP-14	PI 565468	1981	USDA-ARS	Hammons, 1981
CBR-R4	GP-15	PI 565471	1981	USDA-ARS	Hammons, 1981
CBR-R5	GP-16	PI 565467	1981	USDA-ARS	Hammons, 1981
CBR-R6	GP-17	PI 565472	1981	USDA-ARS	Hammons, 1981
Chico	GP-02	PI 565455	1974	USDA-ARS, GA, VA, & OKAES	Bailey & Hammons, 1975
CPES population	GP-55	PI 542961	1990	GAAES & USDA-ARS	Branch & Holbrook, 1991
F334A-B-14	GP-32	PI 565477	1983	USDA-ARS, FL & GAAES	Hammons <i>et al.</i> , 1983
GFA-1	GP-33	PI 565478	1983	USDA-ARS & GAAES	Mixon, 1983b
GFA-2	GP-34	PI 565479	1983	USDA-ARS & GAAES	Mixon, 1983b
NC-GP343	GP-01	PI 565465	1952	NCARS	Campbell <i>et al.</i> , 1971
NC-GPWS 1	GP-59	PI 564844	1992	NCARS	Stalker & Beute, 1993
NC-GPWS 2	GP-60	PI 564845	1992	NCARS	Stalker & Beute, 1993
NC-GPWS 3	GP-61	PI 564846	1992	NCARS	Stalker & Beute, 1993
NC-GPWS 4	GP-62	PI 564847	1992	NCARS	Stalker & Beute, 1993
NC-GPWS 5	GP-103	PI 619169	1998	NCARS	Stalker <i>et al.</i> , 2002a
NC-GPWS 6	GP-104	PI 619170	1998	NCARS	Stalker <i>et al.</i> , 2002a
NC-GPWS 7	GP-105	PI 619171	1998	NCARS	Stalker & Lynch, 2002
NC-GPWS 8	GP-106	PI 619172	1998	NCARS	Stalker & Lynch, 2002
NC-GPWS 9	GP-107	PI 619173	1998	NCARS	Stalker & Lynch, 2002
NC-GPWS 10	GP-108	PI 619174	1998	NCARS	Stalker & Lynch, 2002
NC-GPWS 11	GP-109	PI 619175	1998	NCARS	Stalker <i>et al.</i> , 2002b
NC-GPWS 12	GP-110	PI 619176	1998	NCARS	Stalker <i>et al.</i> , 2002b
NC-GPWS 13	GP-111	PI 619177	1998	NCARS	Stalker <i>et al.</i> , 2002b
NC-GPWS 14	GP-112	PI 619178	1998	NCARS	Stalker <i>et al.</i> , 2002b
NC-GPWS 15	GP-113	PI 619179	1998	NCARS	Stalker <i>et al.</i> , 2002b
ICGL1	GP-50	PI 544348	1989	ICRISAT	Nigam <i>et al.</i> , 1991a
ICGL2	GP-51	PI 544349	1989	ICRISAT	Nigam <i>et al.</i> , 1991a
ICGL3	GP-52	PI 544350	1989	ICRISAT	Nigam <i>et al.</i> , 1991a
ICGL4	GP-53	PI 544351	1989	ICRISAT	Nigam <i>et al.</i> , 1991a
ICGL5	GP-54	PI 544352	1989	ICRISAT	Nigam <i>et al.</i> , 1991a
ICGS35	GP-67	PI 577819	1981/1986	ICRISAT	Nigam <i>et al.</i> , 1994
ICGV-SM 83305	GP-91	PI 598135	1995	ICRISAT	Nigam <i>et al.</i> , 1998b
ICGV-SM 83708	GP-68	PI 585000	1990	ICRISAT	Nigam <i>et al.</i> , 1995a
ICGV-SM 85048	GP-90	PI 598134	1992	ICRISAT	Nigam <i>et al.</i> , 1998a
ICGV-SM 86715	GP-89	PI 598133	1992	ICRISAT	Moss <i>et al.</i> , 1998
ICGV 86015	GP-73	PI 585005	1993	ICRISAT	Nigam <i>et al.</i> , 1995b
ICGV 86031	GP-58	PI 561917	1991	ICRISAT	Dwivedi <i>et al.</i> , 1993
ICGV 86143	GP-87	PI 596359	1994	ICRISAT	Upadhaya <i>et al.</i> , 1997b
ICGV 86155	GP-79	PI 594969	1995	ICRISAT	Upadhaya <i>et al.</i> , 1997a
ICGV 86156	GP-80	PI 594970	1995	ICRISAT	Upadhaya <i>et al.</i> , 1997a
ICGV 86158	GP-81	PI 594971	1995	ICRISAT	Upadhaya <i>et al.</i> , 1997a
ICGV 86252	GP-69	PI 585001	1993	ICRISAT	Dwivedi <i>et al.</i> , 1995
ICGV 86388	GP-77	PI 593239	1995	ICRISAT	Dwivedi <i>et al.</i> , 1996b
ICGV 86393	GP-70	PI 585002	1993	ICRISAT	Dwivedi <i>et al.</i> , 1995
ICGV 86455	PI-71	PI 585003	1993	ICRISAT	Dwivedi <i>et al.</i> , 1995
ICGV 86462	GP-72	PI 585004	1993	ICRISAT	Dwivedi <i>et al.</i> , 1995
ICGV 86564	GP-65	PI 573007	1992	ICRISAT	Dwivedi <i>et al.</i> , 1994
ICGV 86699	GP-76	PI 591815	1994	ICRISAT	Reddy <i>et al.</i> , 1996
ICGV 87121	GP-57	PI 478784	1989	ICRISAT	Nigam <i>et al.</i> , 1992a
ICGV 87157	GP-56	PI 556992	1990	ICRISAT	Nigam <i>et al.</i> , 1992b
ICGV 87165	GP-78	PI 594923	1994	ICRISAT	Moss <i>et al.</i> , 1997
ICGV 87378	GP-82	PI 594972	1995	ICRISAT	Upadhaya <i>et al.</i> , 1997a
ICGV 87921	GP-83	PI 594973	1995	ICRISAT	Upadhaya <i>et al.</i> , 1997a
ICGV 88145	GP-74	PI 585006	1993	ICRISAT	Rao <i>et al.</i> , 1995
ICGV 88438	GP-84	PI 596514	1995	ICRISAT	Hadjichristodoulou <i>et al.</i> , 1997
ICGV 89104	GP-75	PI 585007	1993	ICRISAT	Rao <i>et al.</i> , 1995
ICGV 89214	GP-85	PI 596515	1995	ICRISAT	Hadjichristodoulou <i>et al.</i> , 1997
ICGV 91098	GP-86	PI 596516	1995	ICRISAT	Hadjichristodoulou <i>et al.</i> , 1997
ICGV 92196	GP-92	PI 599344	1996	ICRISAT	Upadhaya <i>et al.</i> , 1998
ICGV 92206	GP-93	PI 599345	1996	ICRISAT	Upadhaya <i>et al.</i> , 1998
ICGV 92234	GP-94	PI 599346	1996	ICRISAT	Upadhaya <i>et al.</i> , 1998
ICGV 92243	GP-95	PI 599347	1996	ICRISAT	Upadhaya <i>et al.</i> , 1998

Table 2 (Continued)

Line	CSSA reg. no.	NPGS no.	Year of release	Releasing agency	Reference
NC 10247	GP-05	PI 565434	1975	NCARS	Campbell <i>et al.</i> , 1975
NC 10272	GP-06	PI 565435	1975	NCARS	Campbell <i>et al.</i> , 1975
NC 15729	GP-07	PI 565436	1975	NCARS	Campbell <i>et al.</i> , 1975
NC 15745	GP-08	PI 565437	1975	NCARS	Campbell <i>et al.</i> , 1975
NC 3033 (ICG 5816)	GP-09	PI 565460	1974	NCARS	Beute <i>et al.</i> , 1976
PI 109839 (Mani)	GP-10	PI 109839	1979	GAAES & USDA-ARS	Hammons <i>et al.</i> , 1980
PI 337394F	GP-03	PI 337394F	1974	USDA-ARS, AL, & GAAES	Mixon & Rogers, 1975
PI 337409 (Rosado)	GP-04	PI 337409	1974	USDA-ARS, AL, & GAAES	Mixon & Rogers, 1975
Tifrust-1	GP-18	PI 561676	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982d
Tifrust-10 (ICG 7890)	GP-27	PI 561685	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982c
Tifrust-11 (ICG 7893)	GP-28	PI 561686	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982c
Tifrust-12 (ICG 7891)	GP-29	PI 561687	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982c
Tifrust-13 (ICG 7883)	GP-30	PI 561688	1981	USDA-ARS, GAAES, ICRISAT & Israel Agric. Res. Org.	Hammons <i>et al.</i> , 1982b
Tifrust-14 (ICG 7882)	GP-31	PI 561689	1982	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982a
Tifrust-2 (ICG 7886)	GP-19	PI 561677	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982d
Tifrust-3 (ICG 7887)	GP-20	PI 561678	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982d
Tifrust-4 (ICG 7898)	GP-21	PI 561679	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982d
Tifrust-5 (ICG 7894)	GP-22	PI 561680	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982c
Tifrust-6 (ICG 7895)	GP-23	PI 561681	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982c
Tifrust-7 (ICG 7896)	GP-24	PI 561682	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982c
Tifrust-8 (ICG 7888)	GP-25	PI 561683	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982c
Tifrust-9 (ICG 7889)	GP-26	PI 561684	1981	USDA-ARS, GAAES, & ICRISAT	Hammons <i>et al.</i> , 1982c
Tifton-8	GP-39	PI 565463	1985	USDA-ARS, VAAES, & GAAES	Coffelt <i>et al.</i> , 1985
TxAG-1	GP-40	NSL 199440	1985	TXAES	Simpson & Smith, 1986
TxAG-2	GP-41	NSL 199441	1985	TXAES	Simpson & Smith, 1986
TxAG-4	GP-48	PI 535816	1990	TXAES, USDA-ARS, & OKAES	Smith <i>et al.</i> , 1990
TxAG-5	GP-49	PI 535817	1990	TXAES, USDA-ARS, & OKAES	Smith <i>et al.</i> , 1990
TxAG-6	GP-63	PI 565287	1992	TXAES	Simpson <i>et al.</i> , 1993
TxAG-7	GP-64	PI 565288	1992	TXAES	Simpson <i>et al.</i> , 1993
VGP 1	GP-11	PI 565462	1979	USDA-ARS & VAAES	Coffelt, 1980
VGP 10	GP-88	PI 584772	1997	USDA-ARS & VAAES	Coffelt & Mozingo, 1998
VGP 11	GP-96	PI 584773	1997	USDA-ARS & VAAES	Coffelt <i>et al.</i> , 1998
VGP 2	GP-42	PI 509536	1986	USDA-ARS & VAAES	Coffelt <i>et al.</i> , 1987
VGP 3	GP-43	PI 509537	1986	USDA-ARS & VAAES	Coffelt <i>et al.</i> , 1987
VGP 4	GP-44	PI 509538	1986	USDA-ARS & VAAES	Coffelt <i>et al.</i> , 1987
VGP 5	GP-45	PI 509539	1986	USDA-ARS & VAAES	Coffelt <i>et al.</i> , 1987
VGP 6	GP-46	PI 509540	1986	USDA-ARS & VAAES	Coffelt <i>et al.</i> , 1987
VGP 7	GP-47	PI 509541	1986	USDA-ARS & VAAES	Coffelt <i>et al.</i> , 1987
VGP 9	GP-66	PI 561567	1993	USDA-ARS & VAAES	Coffelt <i>et al.</i> , 1994a
GA 207-3-4	PL-02	PI 565464	1984	GAAES & USDA-ARS	Branch & Hammons, 1984
Jenkins Jumbo	PL-01	PI 565457	1978	GA & FLAES & USDA-ARS	Hammons & Norden, 1979
Curly-leaf	GS-03	PI 578012	1993	GAAES	Branch, 1995
Georgia Non-Nod	GS-06	PI 595385	1994	GAAES	Branch, 1997
ICGL 6	GS-01	PI 561916	1991	ICRISAT	Nigam <i>et al.</i> , 1993a
Krinkle-leaf <sup>a</sup>					USDA-ARS Hammons, 1964
Rusty-Leaf	GS-07	PI 608669	1998	GAAES	Branch, 1999a
Variegated-Leaf	GS-02	PI 561736	1992	GAAES	Branch, 1993
VGS 1	GS-04	PI 584770	1994	USDA-ARS & VAAES	Coffelt, 1995
VGS 2	GS-05	PI 584771	1994	USDA-ARS & VAAES	Coffelt, 1995
White-Spot Testa	GS-08	PI 608670	1998	GAAES	Branch, 1999b

<sup>a</sup>A genetic stock whose description was published in Crop Science, but not registered with the Crop Science Society of America.

been an influx of runner-type cultivars into the SW area, resulting in relatively great genetic diversity among the cultivars grown there.

One of the ways that plant breeders can increase the genetic diversity of a crop is to incorporate diverse germplasm into the breeding populations from which cultivars derive. *Arachis hypogaea* is a genetically diverse species with two subspecies and six botanical vari-

eties. The first peanut introduction of the modern era was PI 4253, collected by B. Lathrop and D.G. Fairchild in 1899 and identified as the prize-winning peanut from the 1898 exposition of the Khedival Agricultural Society of Cairo, Egypt (USDA, 1900, 1901). There have been thousands of accessions of *A. hypogaea* introduced and numbered by the USDA since that time. Many were donated by U.S. diplomats, missionaries, and travelers in

foreign countries. Others were provided by foreign governments and agricultural research institutions as part of germplasm exchanges with U.S. institutions. Still others were collected as part of a coordinated effort by the USDA and international agencies to collect and preserve natural genetic diversity before it erodes through the displacement of farmer-held seed stocks by improved cultivars. The current USDA collection contains over 8000 *A. hypogaea* accessions. Many of these are duplicated within the collection as a result of wholesale exchange of germplasm with collections in other countries. In the early years of the national plant germplasm system, breeding lines from U.S. programs were often cataloged, and accessions were sometimes renumbered each time seed was produced. Many exotic accessions have been lost from the U.S. collection because early introductions of minor crops were often increased only once or not at all, then distributed to the agricultural experiment stations in states where the crop was grown. Some accessions without obvious value were discarded by recipients after cursory observation. No doubt some lines were lost due to special requirements or characteristics that made them unadapted to the prevailing conditions in the southern U.S. Some accessions were used as parents in peanut breeding programs, and some have descendants among released cultivars and germplasm lines. The objectives of this study were to (a) summarize and document the use of genetic resources in cultivar development and (b) estimate the resulting economic impact.

## Materials and Methods

The importance of ancestors of current runner, virginia, and spanish cultivars was assessed by pedigree analysis of improved cultivars (Tables 3-5; Figs. 1-3). The parentage of each cultivar was traced back to ancestors for which no further information could be found. Pedigree information was gathered from published registration articles, release notices, internal documents of the North Carolina and Florida breeding projects, and by personal communication with breeders in other states.

Each cultivar was assigned to a breeding cycle based on the number of cycles of crossing and selection it was removed from the founding ancestors. In this system, ancestors were assigned to Cycle 0, selections from Cycle 0 ancestors or progeny from crosses between Cycle 0 lines were assigned to Cycle 1, and so on. Each line was assigned to a cycle one higher than the higher of its parents. Coefficients of coancestry (Malécot, 1948) of the cultivars with each of the ancestors were then calculated. Each such coancestry value represents the proportion of the genome of the cultivar that should trace to the ancestor. Coancestries were totaled across PI ancestors to indicate the proportion of the cultivar's genome attributable to PIs.

The coancestry between lines X and Y, ( $\theta_{XY} = \theta_{XC} + \theta_{XD}$ )/2 = ( $\theta_{AY} + \theta_{BY}$ )/2 = ( $\theta_{AC} + \theta_{AD} + \theta_{BC} + \theta_{BD}$ )/4. The expressions are expanded back until they intersect at a common ancestor, W, at which point they include a term  $\theta_{WW}$ , the coancestry of a line with itself which is  $\theta_{WW} = (1 + F_w)/2$ , where  $F_w$  is the coefficient of inbreeding or the probability that the two genes carried by W are identical by descent. For purposes of this study, all lines were assumed to be completely inbred ( $F = 1$ ). These rules were developed for outcrossing

species, but can be adapted for use in self-pollinated species. If one assumes that the probability of fixation under self-pollination of the gene derived from either parent of a cross is 0.5 (St. Martin, 1982), then the computational rules for self-pollinators are identical to those for cross-pollinators. Additional rules are required to determine the coancestry of lines derived by self-fertilization from the same cross (Cockerham, 1983). For lines derived by selection within an existing line, either with or without mutagenesis, it was assumed that the coancestry of the parental and selected lines was 0.99.

Averages and ranges were calculated for the coancestries of cultivars from different breeding cycles and market types to assess the contributions of the ancestors to particular groups. These averages were not weighted by the acreage planted to each cultivar. The total ancestry derived from plant introductions was summed for each cultivar, and distributions were calculated in increments of 12.5%. Distributions were compared through the use of contingency tables and chi-square testing.

The economic impact from the use of genetic resources in cultivars was assessed through surveys of peanut breeders, pathologists, and extension specialists. Estimates were based on the yield advantage of new cultivars in comparison to older ones, land area occupied by new cultivars, and any reduction in pesticide usage that may have occurred because of the release of new cultivars.

## Results and Discussion

**Use of Genetic Resources in Cultivar Development.** Because peanuts as a crop were introduced to what is now the U.S.A., all U.S. peanut cultivars necessarily trace back to plant introductions. However, much of the genetic base of current cultivars traces back to ancestors that were developed by mass selection from farmer stock peanuts in the various production areas (Isleib and Wynne, 1992). Much of the base of improved runner and virginia cultivars rests on four ancestors used as parents in the early years of peanut improvement in the U.S., including var. *hypogaea* lines Dixie Giant and Basse and var. *vulgaris* Harz lines Small White Spanish and Spanish 18-38. Of these, only Basse is known to have been introduced in the modern era of plant collection. The initial cross of Small White Spanish by Dixie Giant was made by W.A. Carver at the Univ. of Florida in the early to mid-1930s, and the cross, numbered F230 in the Florida system, eventually gave rise to the successful runner-type cultivar Early Runner and numerous sibling lines derived from  $F_2$  plant selection F230-118. Meanwhile, B.B. Higgins at the Univ. of Georgia crossed Basse by Spanish 18-38, giving rise to several selections in the GA 207 series. Most current runner- and virginia-type cultivars trace their ancestry back to these two crosses through Florispan and its close siblings, derived from a cross between GA 207-3 and F230-118-2-2, and their immediate descendants Florunner and Florigiant.

In addition to the four primary ancestors of runner-type cultivars, the early virginia market-type cultivars had additional infusion of ancestry from farmer stock selection Jenkins Jumbo, a large-seeded selection from farmer stock used as a parent in the Florida program, a group of five lines (NC 4, NC Bunch, White's Runner, Improved



Table 3 (Continued)

Cy- cle	Name	Origin	PI 203 Basse 396	PI 109 839	Goldin I	PI 331 334	TxA G- 6	PI 121 070	PI 161 317	PI 259 785	PI 475 871	PI 121 067	PI 221 057	Total
6	C-99R	PI 203396 / F427B-3-1-7-4 (Altika sib), UF81206-1 // 72x32B-13-1-3, PI 259785 / Florigiant	.047	.250	.000	.000	.000	.000	.000	.250	.000	.000	.000	.547
6	Georgia Browne	Southern Runner / Sunbelt Runner	.078	.250	.000	.000	.000	.000	.000	.000	.000	.000	.000	.328
6	Georgia Green	Southern Runner / Sunbelt Runner	.078	.250	.000	.000	.000	.000	.000	.000	.000	.000	.000	.328
6	GK 193	GK 3 / Early Bunch	.047	.000	.000	.000	.000	.000	.000	.000	.000	.000	.000	.047
6	MARC I	Early Runner / Florispan, F439-17-2-1-1 (Florunner sib) // F459B-3-2-4-6-2-2-1 (Early Bunch component)	.078	.000	.000	.000	.000	.000	.000	.000	.000	.000	.000	.078
6	SunOleic 97R	SunOleic 95R / Sunrunner	.121	.000	.000	.000	.000	.000	.000	.000	.000	.000	.000	.121
7	Georgia Bold	Southern Runner / Sunbelt Runner, Georgia Bold Parent 1 / Sunbelt	.086	.125	.000	.000	.000	.000	.000	.000	.000	.000	.000	.211
8	Florida MDR 98	Southern Runner / 3 / Florida MDR parent 2, F439-17-2-1-1 (Florunner sib) / F459B-3-2-4-6-2-2-1 (Early Bunch sib), UF79308 // UF81206-1, PI 203396 / F427B-3-1-7-4 (Altika sib)	.059	.375	.000	.000	.000	.000	.000	.000	.000	.000	.000	.434
	Number of descendants among 41 cultivars		32	6	3	1	1	1	1	1	1	1	1	33
	Mean coancestry with introduction		.076	.043	.024	.012	.012	.012	.008	.006	.006	.002	.002	.216
	Mean of nonzero coancestries		.097	.292	.333	.500	.500	.490	.155	.250	.250	.063	.063	.268

\*Current cultivar, i.e., cultivar in use in the 2000 season.





Spanish 2B, and PI 121067) among seven used by W.C. Gregory to initiate the breeding program at N. C. State Univ., and Atkins Runner, an ancestor used by the USDA breeding program in Virginia. Of these additional early ancestors of the virginia market type, only PI 121067 is a modern plant introduction. A different set of introductions including PI 121070, PI 161317, PI 268661, and *A. monticola* Krapov. & Rigoni were used as parents in the Texas and Oklahoma breeding programs.

Thirteen plant introductions occur in the pedigrees of runner-type cultivars (Table 3; Fig. 1):

- **Basse** is by far the most common introduction in the pedigrees of cultivars of the runner and virginia market types, appearing in the pedigrees of 32 of 41 improved runner-type cultivars. Basse has been identified as an introduction from Gambia, and the specimen in the collection in North Carolina is a runner-type line, but the exact source of the original introduction has become obscure. Several lines identified as Basse or selections from Basse have been reintroduced to the U.S.A. There were other early peanut introductions from Gambia [PI 5358 (USDA, 1902), PI 15785 (USDA, 1907), PI 21198 (“Nkate or Nbatie’,” received from the Gold Coast in 1907) (USDA, 1908)], but Basse is likely PI 24114 which was sent to the USDA in 1908 by R.P. Skinner, U.S. consul general at Marseille, France, at the request of W.R. Beattie and C.S. Scofield and was identified as having originated in Gambia. Beattie is quoted in the Plant Inventory (USDA, 1909) as regarding the line as “exceptionally valuable for use in the manufacture of candy and other products where shelled nuts are required.”
- **PI 203396**, A.A. Beetle Col. No. 2201, collected in a market at Porto Alegre, Brazil in 1952 (USDA, 1957), was used as a parent in the Univ. of Florida breeding program in the 1970s and 1980s. It was a source of resistance to late leaf spot [*Cercosporidium personatum* (Berk. & Curt.) Deighton], southern stem rot (*Sclerotium rolfsii* Sacc.), and tomato spotted wilt virus. PI 203396 appears in the pedigrees of six runner-type cultivars through its use as a parent of the multiple disease-resistant cultivar Southern Runner and sibling line UF81206-1. Through Southern Runner, PI 203396 is a grandparent of Georgia Browne and Georgia Green and a great-grandparent of Georgia Bold. Georgia Green is currently the most extensively grown runner cultivar in the U.S.A., particularly in the southeastern production area where tomato spotted wilt virus is the primary constraint to production. Through UF81206-1, PI 203396 is a grandparent of disease-resistant Florida releases Florida MDR 98 and C-99R.
- **PI 109839**, W.A. Archer Col. No. 2971, identified locally as “Mani,” (“peanut”) collected at a market in Caracas, Venezuela in 1935 (USDA, 1939), is a source of resistance to early leaf spot (*Cercospora arachidicola* Hori). It was used in the 1980s as a parent in the Texas A&M Univ. breeding program and appears in the pedigrees of the Texas cultivars

Langley, Tamrun 96, and Tamrun 98. PI 109839 has been used also as a parent in the N.C. State Univ. program but, due to its small seed size, its progeny have not been suited to the virginia-type market.

- **PI 259785**, “U/A 29-S 185” presented by the Agric. Res. Sta., Lilongwe, Nyasaland (Malawi) in 1959 (USDA, 1966), is a grandparent of C-99R, a disease-resistant release from the Univ. of Florida.
- **PI 331334**, R.O. Hammons & W.R. Langford Col. No. 169, known as “Criollo” (“local”) at its original site of collection in Bolivia, was obtained from the INTA collection at Manfredi, Argentina in 1968 (USDA, 1970) and is a parent of Georgia Runner.
- **PI 475871**, GKSPSc Col. No. 223, collected from a farmer at Guajara-Mirim, Bolivia in 1979 (USDA, 1984), is a grandparent of Tamrun 96.
- **F21** is an introduction with no known PI number but identified in Florida records as “Rasteiro,” obtained by W.A. Carver and used in the Florida breeding program in the early to mid-1930s. Two Brazilian lines named Rasteiro are listed in the USDA Plant Inventory as having been introduced in 1935 (USDA, 1940); PI 111885 donated by the Inst. Agronomico at Campinas, São Paulo, and PI 111965 presented by the Estação Exp. de Cana de Assucar e Plantas Oleaginosas, Piracicaba, São Paulo. F21 provided 6.25% of the ancestry of AgraTech release AT 120.
- **Goldin I**, obtained from E. Goldin (Faculty of Agric., Hebrew Univ. of Jerusalem, Rehovot, Israel) by the Wilson Peanut Co. of Pleasanton, TX is probably derived at least in part from U.S. breeding materials transported to Israel. No records are extant to permit determination of the relationship between Goldin I and U.S. germplasm. Goldin I is a parent of Tamrun 88.
- **TxAG-6** is a synthetic allotetraploid created by C.E. Simpson by making a three-way cross among diploid ( $2n = 2x = 20$ ) wild peanut species PI 298639 (*A. batizocoi* Krapov. and W.C. Gregory, K 9484) // PI 262141 (*A. cardenasii* Krapov. and W.C. Gregory, GKP 10017) / PI 276235 (*A. diogeni* Hoehne, GKP 10602) (USDA, 1966, 1967b, 1968) and treating the hybrid with colchicine to induce tetraploidy. TxAG-6 is a parent of COAN, a runner-type cultivar that derives its resistance to root-knot nematodes [*Meloidogyne arenaria* (Neal) Chitwood] from the interspecific hybrid.

The remaining five introductions that appear in the pedigrees of runner-type cultivars (PI 121067, PI 121070, PI 616317, PI 259785, and PI 221057) do so through crosses of runner-type parents with virginia-type and spanish-type parents. Their descriptions are included below in the sections for those market types. Only three plant introductions appear in the pedigrees of improved virginia-type cultivars (Table 4; Fig. 2): Basse, PI 121067, and PI 337396.

- **Basse** was described earlier; its genetic contribution is present in 24 of 33 improved virginia-type cultivars, including all that are currently grown. Al-

Table 4. Improved virginia-type cultivars released in the U.S.A., their origin, breeding cycle, and coancestry with three introduced ancestors.

Cycle	Name	Origin	Basse	PI 121067	PI 337396	Total
1	NC 1	NC 4 / Improved Spanish 2B	.000	.000	.000	.000
1	NC 4X	Selection from irradiated 'NC 4'	.000	.000	.000	.000
1	VA 56R	Selection from 'Atkins Runner'	.000	.000	.000	.000
1	VA 61R	Selection from 'Atkins Runner'	.000	.000	.000	.000
1	Virginia Bunch 46-2	Selection from 'Virginia Bunch Large'	.000	.000	.000	.000
2	GA 119-20	Southeastern Runner / Dixie Giant, 210-4 // Virginia Runner	.000	.000	.000	.000
2	NC 2	Basse / Spanish 18-38, GA 207-2 // White's Runner	.000	.000	.000	.000
2	NC 5	NC 4 / Improved Spanish 2B, NC 1 // C12, PI 121067 / NC Bunch	.000	.250	.000	.250
2	NC 6 <sup>a</sup>	NC Bunch / PI 121067, C12 // C37 (same as C12), GP-NC 343 (selection from NC Ac 4508) // VA 61R	.000	.500	.000	.500
2	VA 72R	VA 61R / VA A89-15 (selection from farmers stocks, perhaps Atkins Runner)	.000	.000	.000	.000
3	Avoca 11	Selection from 'NC 2'	.248	.000	.000	.248
3	Florigiant	Basse / Spanish 18-38, GA 207-3 // F230-118-2-2 (same as F230), F334A-5-5-1 / 3 / F359-1-3-14, Jenkins Jumbo // F230-118-5-1, Dixie Giant / Small White Spanish 3x-2	.125	.000	.000	.125
3	NC-Fla 14	Jenkins Jumbo / F334A-3-5-5-1 (Florispans derivative)	.125	.000	.000	.125
3	NC 17	F334A-3-5-5-1 / Jenkins Jumbo	.125	.000	.000	.125
4	Altika	F393-7-1 (NC-Fla 14 sib) / 3 / GA 119-20, Southeastern Runner / Dixie Giant, 210-14 // Virginia Runner	.063	.000	.000	.063
4	GK 25	Florigiant / GA 186-28	.063	.000	.000	.063
4	Keel 29	Selection from 'Florigiant'	.124	.000	.000	.124
4	NC 7 <sup>a</sup>	NC 5 // F393, F334-3-5-5-1 (Florispans derivative) / Jenkins Jumbo	.063	.125	.000	.188
4	NC 8C	NC 2 // A48, NC 4 / Spanish 2B, NC Ac 3139 / 3 / Florigiant	.125	.000	.000	.125
4	NC 9 <sup>a</sup>	NC 2 // Florigiant	.188	.000	.000	.188
4	Shulamith	Florigiant / F334A-B-17-1 (Florispans derivative)	.188	.000	.000	.188
4	VA 81B	F392-8 (Florigiant sib) / 3 / GA 119-20, Southeastern Runner / Dixie Giant, 210-14 // Virginia Runner	.063	.000	.000	.063
5	Early Bunch	Virginia Station Jumbo / 4 / F385-1-7-4, Pearl (F228) // F68-74 S3-1-2, McSpan (F13, Small White Spanish) / Virginia Jumbo Runner (F14), F249-42-3-1 / 3 / Jenkins Jumbo, F406A / 5 / F420, F231-51 (Dixie Runner sib) / F392-12-1-7 (Florigiant sib)	.031	.000	.000	.031
5	Gregory <sup>a</sup>	NC 7 / NC 9	.125	.063	.000	.188
5	NC 10C <sup>a</sup>	NC 8C / Florigiant	.125	.000	.000	.125
5	NC-V11 <sup>a</sup>	Florigiant / NC 5 // Florigiant / PI 337396	.063	.063	.250	.375
5	NC 12C <sup>a</sup>	NC 7 / NC 9	.125	.063	.000	.188
5	VA-C 92R <sup>a</sup>	Florigiant / F393-7-47-1-7-1, NC Ac 17213 // NC 7	.094	.063	.000	.156
6	Perry <sup>a</sup>	VA 81B / 3 / VA 78039 (NC-V11 sib), Florigiant / NC 5, NC Ac 17257 // NC Ac 17922, Florigiant / PI 337396	.125	.000	.000	.125
6	VA 93B <sup>a</sup>	VA 81B / 3 / VA 78039 (NC-V11 sib), Florigiant / NC 5, NC Ac 17257 // NC Ac 17922, Florigiant / PI 337396	.063	.031	.125	.219
6	VA 98R <sup>a</sup>	GA 207-3 / F230-118-2-2, F334A-3-5-5-1 // Jenkins Jumbo, F393-2-1-2 / 4 / Florunner, GK 1A / 5 / AgraTech VC1 Parent 2, Florigiant / 3 / GA 186-28, Virginia Runner / Pearl // Dixie Giant / Carolina Runner, GK 25 / 4 / Florunner	.063	.031	.125	.219
6	AgraTech VC-1	Florigiant / 3 / GA 186-28, Virginia Runner / Pearl // Dixie Giant / Carolina Runner, GK 25 / 4 / Florunner	.109	.000	.000	.109
7	AgraTech VC-2 <sup>a</sup>	F435-2-3-B-2-1-b4-B-3-b3-1-B / 3 * GK 193	.086	.000	.000	.086
	Number of descendants among 33		24	9	3	26
	Mean coancestry with introduction		.084	.036	.015	.135
	Mean of nonzero coancestries		.115	.132	.167	.171

<sup>a</sup>Current cultivar, i.e., cultivar in use in the 2000 season.

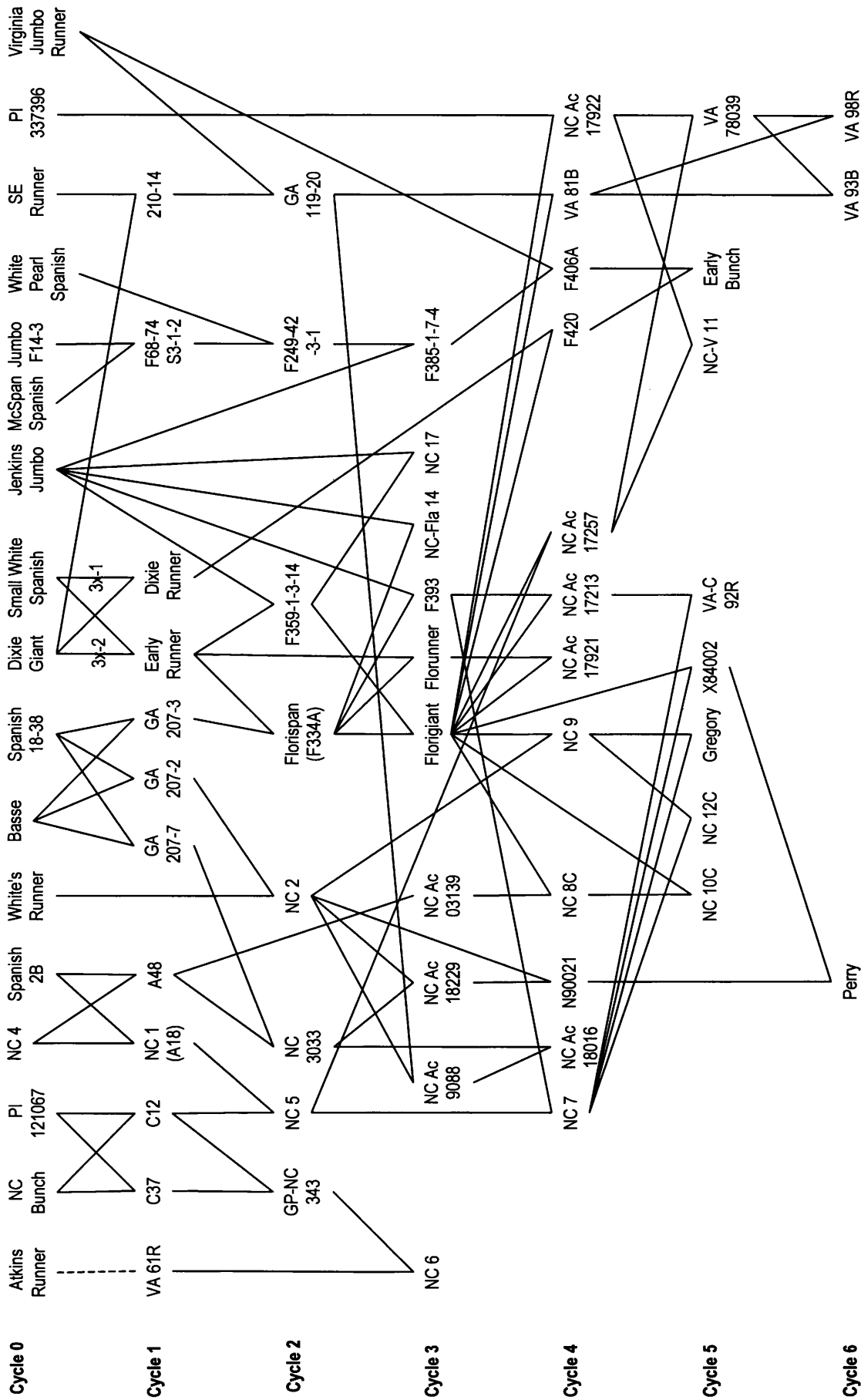


Fig. 2. Parentage of virginia-type cultivars. Solid lines indicate parents and progeny of crosses; dashed lines indicate selections from existing lines with or without mutation.

**Table 5. Spanish-type cultivars released in the USA, their origins, breeding cycle, and coancestry with seven introduced ancestors.**

Cycle	Name	Origin	PI 161317	PI 121070	Chico	PI 221057	Spanish No. 146	PI 405933	<i>A.</i> <i>monticola</i>	Total
0	Spantex	Selection from farmers □ Spanish stocks	.000	.000	.000	.000	.000	.000	.000	.000
1	Argentine	Selection from PI 121070	.000	.990	.000	.000	.000	.000	.000	.990
1	Dixie Spanish	Selection from Spanish introduction (Coll. No. 146)	.000	.000	.000	.000	.990	.000	.000	.990
1	Spanette	Selection from Spanish 18-38	.000	.000	.000	.000	.000	.000	.000	.000
1	Starr	Spantex / PI 161317	.500	.000	.000	.000	.000	.000	.000	.500
2	Comet	Selection from Starr	.495	.000	.000	.000	.000	.000	.000	.495
2	Spancross	Argentine (PI 121070-1) / PI 405933	.000	.495	.000	.000	.000	.500	.000	.995
2	Spanhoma	Selection from □ Argentine □	.000	.980	.000	.000	.000	.000	.000	.980
2	Tamnut 74	Starr // TPL 647-2-5, Spantex / <i>A. monticola</i>	.250	.000	.000	.000	.000	.000	.250	.500
2	Tifspan	Argentine (PI 121070-1) / Spanette	.000	.495	.000	.000	.000	.000	.000	.495
2	Toalson	PI 221057 / Selection 26 (Spantex sib), TPL 673-A // Starr	.250	.000	.000	.250	.000	.000	.000	.500
3	Pronto	Chico / Comet	.248	.000	.500	.000	.000	.000	.000	.748
3	Spanco	Chico / Comet	.248	.000	.500	.000	.000	.000	.000	.748
4	Tamspan 90*	Selection in TxAG-5 (Toalson / UF 734022)	.124	.000	.000	.124	.000	.000	.000	.248
Number of descendants among 14			7	4	2	2	1	1	1	12
Mean coancestry with introduction			.151	.211	.071	.027	.071	.036	.018	.585
Mean of nonzero coancestries			.302	.740	.500	.187	.990	.500	.250	.682

\*Current cultivar, *i.e.*, cultivar in use in the 2000 season.

though Basse ancestry is common across most virginia-type cultivars, it accounts for only about 10% of the total ancestry.

- **PI 121067**, W.A. Archer Col. No. 4998; “Indio” (A.H. 1119-A) from Misiones, Argentina; presented by Sr. Clos of the Min. de Agric., Buenos Aires, Argentina in 1937 (USDA, 1942), was one of the seven parents used by W.C. Gregory to initiate the peanut breeding program at N.C. State Univ. It has partial resistance to *C. arachidicola* and southern corn rootworm (*Diabrotica undecimpunctata* Howard Barber) and is an ancestor of many of the North Carolina breeding populations and releases including insect-resistant germplasm line GP-NC 343 and seven cultivars—NC 5, insect-resistant NC 6, NC 7, NC-V 11, *Cylindrocladium* black rot-resistant NC 12C, Gregory, and VA-C 92R. Through NC 5, PI 121067 is also an ancestor of Virginia releases VA 93B and VA 98R. Through NC 7, it is an ancestor of runner-type cultivar Virugard.
- **PI 337396**, R.O. Hammons & W.R. Langford Col. No. 250. “FAV 70,” a valencia-type line obtained from the INTA collection at Manfredi, Misiones, Argentina in 1968 (USDA, 1970), is a grandparent of NC-V 11 and a sibling line that was a parent of VA 93B and VA 98R.

Most runner- and virginia-type cultivars are characterized as having had some introgression of genes from subsp. *fastigiata* Waldron, mostly from var. *vulgaris* but to some extent from var. *fastigiata*. Spanish-type cultivars are more varietally pure than other market types for the most part. Seven introductions appear in the pedigrees of spanish-type cultivars; most are spanish-type (var. *vulgaris*) lines themselves. Two are accessions of *A. monticola*, an alternate-branching tetraploid ( $2n = 4x$

= 40) species that crosses readily with *A. hypogaea*.

- **PI 161317**, J.L. Stephens & W. Hartley Col. No. 31-A; a spanish-type line collected in Salto, Uruguay in 1947 (USDA, 1954), appears in the pedigrees of seven spanish-type cultivars—Starr, Comet, Tamnut 74, Toalson, Pronto, Spanco, and Tamspan 90.
- **PI 121070**, W.A. Archer Col. No. 5001; spanish-type Strain A.H. 1131-1 collected at Chiarí, Entre Rios, Argentina in 1937 (USDA, 1942), is ancestral to four spanish-type cultivars. It was the source from which Argentine was selected. Argentine was used in the Oklahoma breeding program and is a parent of Spanhoma and the source from which the cultivar Spanhoma was selected. Argentine is also a parent of Tifspan, developed by the USDA program at Tifton, GA.
- **Chico**, a selection from spanish-type PI 268661; “APAXUC 70” [“*Arachis* 70”] from the USSR; presented by the Mount Makulu Res. Sta., Chilanga, Rhodesia (Zambia) in 1960 (USDA, 1967a), is a very early maturing line. Chico has been used as a parent in several programs in attempts to shorten maturity; it is a parent of Pronto and Spanco.
- **PI 221057**, spanish type line “J.B.M. 19/3” presented by the Inst. Agronômico at Campinas, São Paulo, Brazil in 1954 (USDA, 1960), is a grandparent of Toalson and a great-grandparent of Tamspan 90.
- **Spanish No. 146**, a spanish-type introduction (Col. No. 146) obtained from India by Tom Huston Peanut Co., was used in the Georgia breeding program where it was the source from which Dixie Spanish was selected.
- **PI 405933**, *A. monticola* T-1119 collected in Yala,

Jujuy, Argentina in 1950; presented to the USDA by V.A. Rigoni, Estacion Exp. Agropecuaria, Manfredi, Misiones, Argentina (USDA, 1978), was a parent of Spancross in the USDA program at Stillwater, OK.

- *A. monticola*, an accession not identified by PI number, is a grandparent of Tamnut 74 in the Texas A&M Univ. program. As the number of *A. monticola* accessions in the national collection is limited, and common accessions are generally held by several breeding programs, it is possible that this is also PI 405933.

When the distribution of lines with different proportions of PI ancestry (Table 6) were compared for the runner, virginia, and spanish market types, there were marked differences overall ( $\chi^2 = 39.07$  with 14 df,  $P < 0.001$ ), but most of the difference was between the spanish type and the two alternate-branching types (runner and virginia). The distribution of PI ancestry in improved cultivars was similar for the runner and virginia market types ( $\chi^2 = 5.65$  with 5 df,  $P < 0.35$ ) while the

distribution of the pooled runner and virginia types were very different from the spanish group ( $\chi^2 = 33.53$  with 7 df,  $P < 0.001$ ). In the runner and virginia groups, the average PI ancestry of all improved cultivars was 17.9% with a range of 0 to 56.2%. Due to the proliferation of ancestry derived from early PI ancestors, later breeding cycles had no lines without PI ancestry, and after the fourth cycle there remained a slight trend toward increased PI ancestry in later cycles. Overall, 28 of 41 runner-type cultivars and 31 of 33 virginia-type cultivars derived 25% or less of their ancestry from plant introductions. Only seven of the 74 total cultivars derived more than half of their ancestry from PIs, and in all cases that total was due to the contributions of more than one PI in the pedigree of the cultivar. No single introduction contributed more than half the ancestry of any cultivar.

Many crosses involving plant introductions as parents are made with the objective of transferring some desirable trait from the PI into an adapted breeding population. Because most improved cultivars contain between

**Table 6. Distribution of ancestry derived from plant introductions for improved US peanut cultivars by market-types and breeding cycles.**

Market-type	No. of lines	$\theta_{total} = 0$	0 < $\theta_{total} \leq 0.125$	0.125 < $\theta_{total} \leq 0.25$	0.25 < $\theta_{total} \leq 0.375$	0.375 < $\theta_{total} \leq 0.5$	0.5 < $\theta_{total} \leq 0.625$	0.625 < $\theta_{total} \leq 0.75$	0.75 < $\theta_{total} \leq 0.875$	0.875 < $\theta_{total} \leq 1$	Mean $\theta_{total}$	Minimum $\theta_{total}$	Maximum $\theta_{total}$
<b>Runner</b>													
Cycle 0	1	1	—	—	—	—	—	—	—	—	.000	.000	.000
Cycle 1	7	6	—	—	—	1	—	—	—	—	.071	.000	.500
Cycle 2	1	—	—	1	—	—	—	—	—	—	.250	.250	.250
Cycle 3	1	—	1	—	—	—	—	—	—	—	.125	.125	.125
Cycle 4	11	1	3	1	1	—	5	—	—	—	.334	.000	.562
Cycle 5	8	—	4	2	—	1	1	—	—	—	.211	.062	.562
Cycle 6	10	—	5	2	2	—	1	—	—	—	.195	.047	.547
Cycle 7	1	—	—	1	—	—	—	—	—	—	.211	.211	.211
Cycle 8	1	—	—	—	—	1	—	—	—	—	.434	.434	.434
All cycles	41	8	13	7	3	3	7	—	—	—	.216	.000	.562
<b>Virginia</b>													
Cycle 1	5	5	—	—	—	—	—	—	—	—	.000	.000	.000
Cycle 2	5	2	—	2	—	1	—	—	—	—	.200	.000	.500
Cycle 3	4	—	3	1	—	—	—	—	—	—	.156	.125	.248
Cycle 4	8	—	5	3	—	—	—	—	—	—	.125	.062	.188
Cycle 5	6	—	2	3	1	—	—	—	—	—	.177	.031	.375
Cycle 6	4	—	2	2	—	—	—	—	—	—	.168	.109	.219
Cycle 7	1	—	1	—	—	—	—	—	—	—	.086	.086	.086
All cycles	33	7	13	11	1	1	—	—	—	—	.135	.000	.500
<b>Pooled runner and virginia</b>													
Cycle 0	1	1	0	0	0	0	0	0	0	0	.000	.000	.000
Cycle 1	12	11	0	0	0	1	0	0	0	0	.042	.000	.500
Cycle 2	6	2	0	3	0	1	0	0	0	0	.208	.000	.500
Cycle 3	5	0	4	1	0	0	0	0	0	0	.150	.125	.248
Cycle 4	19	1	8	4	1	0	5	0	0	0	.246	.000	.562
Cycle 5	14	0	6	5	1	1	1	0	0	0	.196	.031	.562
Cycle 6	14	0	7	4	2	0	1	0	0	0	.188	.047	.547
Cycle 7	2	0	1	1	0	0	0	0	0	0	.148	.086	.211
Cycle 8	1	0	0	0	0	1	0	0	0	0	.434	.434	.434
All cycles	74	15	26	18	4	4	7	0	0	0	.179	.000	.562
<b>Spanish</b>													
Cycle 0	1	1	—	—	—	—	—	—	—	—	.000	.000	.000
Cycle 1	4	1	—	—	—	1	—	—	—	2	.620	.000	.990
Cycle 2	6	—	—	—	—	4	—	—	—	2	.661	.495	.995
Cycle 3	2	—	—	—	—	—	—	2	—	—	.748	.748	.748
Cycle 4	1	—	—	1	—	—	—	—	—	—	.248	.248	.248
All cycles	14	2	—	1	—	5	—	2	—	4	.585	.000	.995

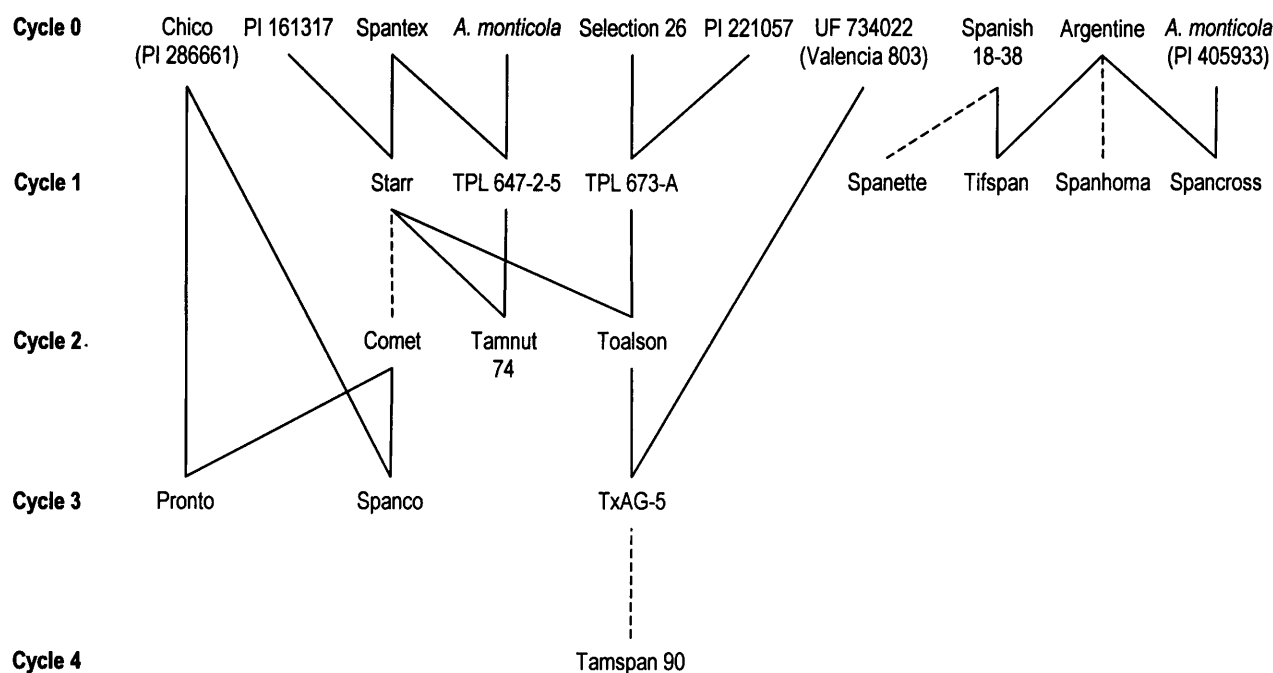


Fig. 3. Parentage of Spanish-type cultivars. Solid lines indicate parents and progeny of crosses; dashed lines indicate selections from existing lines with or without mutation.

12.5 and 25% PI ancestry, it would appear that successful incorporation of PI ancestry into adapted cultivars is best achieved through some form of backcrossing. This might be achieved in the classical way by testing for the trait under transfer from the exotic parent at the  $F_1$  or  $F_2$  generation following a given backcross, or it might be performed in a less conventional way with selection of lines from later generations of the initial cross to the PI followed by backcrossing to the adapted parent as was the case in the development of lines derived from PI 203396.

**Economic Impact of Genetic Resources.** The genetic variability contributed to breeding populations by the introgression of genes from introductions has no doubt contributed to genetic gain in the development of cultivars that possess no salient disease resistance or other introduction-derived trait. However, the economic impact of such introgression is difficult if not impossible to measure. Genetic resources have been particularly useful in adding disease resistance to peanut cultivars. This has had significant economic impact on U.S. peanut farmers, the largest from the development of cultivars with resistance to Sclerotinia blight, root-knot nematode, and tomato spotted wilt virus.

PI 221057 served as a source of resistance to Sclerotinia blight in the development of the Spanish-type cultivar Tamspan 90 and, more recently, in the runner-type cultivar Tamrun 98. There are approximately 16,200 ha (40,000 acres) of Sclerotinia-infested land in the southwestern peanut production area, and fungicide treatments for the control of this disease are costly. Use of Sclerotinia-resistant cultivars increases yield and reduces fungicide costs. The estimated impact from the use of these resistant cultivars is \$5 million annually.

COAN, a nematode-resistant runner-type cultivar, was developed by introgressing resistance from wild diploid peanut species. Approximately 29,000 ha (72,000 acres) of peanut land in the Southwest is infested with *M. arenaria* race 1. A nematode-resistant cultivar would save southwestern peanut growers an estimated \$6.5 million annually in increased yields and reduced nematocide costs. Unfortunately, COAN is susceptible to TSWV and is therefore not adapted to the southeastern peanut production area. Breeders are attempting to combine resistance to root-knot nematode with field resistance to TSWV.

The use of PIs has had its largest economic impact through the development of runner-type cultivars with resistance to TSWV. In the Southeastern U.S., TSWV was detected in peanut in 1987, and its incidence has increased substantially in the ensuing years (Culbreath *et al.*, 1992). PI 203396 was an ancestor to many breeding populations in the Southeast due to its resistance to late leaf spot. Fortunately, PI 203396 also has resistance to TSWV which was transmitted to resistant runner-type cultivars Southern Runner, Georgia Green, Florida MDR 98, and C-99R. Under severe TSWV pressure, the additional economic return from growing these cultivars in comparison to previous susceptible cultivars is in excess of \$500 per acre. Assuming that half of the peanut acreage in the Southeast has severe TSWV pressure, the economic impact of this resistance is more than \$200 million annually.

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